

Mechanisms and Consequences of Reduced Dendritic Growth of Cerebellar Purkinje Cells

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Olivia Sarah Gugger

aus Frauenfeld (TG)

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auf Antrag von**

**Prof. Josef Kapfhammer
Prof. Markus Rüegg
Prof. Peter Scheiffele**

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List of Abbreviations

AMPA	(RS)- α -Amino-3-hydroxy-5-methyl-4-isoxazolepropionic acid (AMPA)
BDNF	brain-derived neurotrophic factor
bHLH	basic helix-loop-helix
Ca	calcium
CB6	B6CF1, mouse strain
CF	climbing fiber
CGN	cerebellar granule cell
CNS	central nervous system
DCN	deep cerebellar neuron
DHPG	(RS)-3,5-Dihydroxyphenylglycine
DIV	days in vitro
E	embryonic day
EAAT	excitatory amino acid transporter
EGL	external granule layer
ES cell	external sensory cells
EtD	a 1:1 mixture of ethanol and dimethyl sulfoxide
FBP	fructose 1,6-bisphosphate
GluR δ 2	glutamate receptor δ 2
GNP	granule neuron precursor
IGL	internal granule layer
LTD	long term depression
MD cell	multidendritic cell
mGluR1	metabotropic glutamate receptor, type 1
ML	molecular layer
MW	molecular weight
mwk	moonwalker
Nasp	naphtyl-acetyl spermine
NGF	nerve growth factor
NT	neurotrophin
P	postnatal day
P/Q-block	combination of 100 nM ω -agatoxin IVA and 1 μ M ω -conotoxin MVIIC
PTN	pleiotrophin

PTP ζ	protein tyrosine phosphatase zeta
PB	phosphate buffer
PC	Purkinje cell
PCL	Purkinje cell layer
PCR	polymerase chain reaction
PF	parallel fiber
PKC	protein kinase C
PLC	phospholipase C
PM	preparation medium
PMA	phorbol-12-myristate-13-acetate
PNS	peripheral nervous system
RGC	retinal ganglion cell
RL	rhombic lip
ROR α	retinoid-related orphan receptor α
SCA	spinocerebellar ataxia
sEPSP	slow excitatory postsynaptic potential
SGC	soluble guanylate cyclase
Shh	sonic hedgehog
SKF	SKF&F 96365
SS	stock solution
TBI	traumatic brain injury
TF	transcription factor
TRPC3	transient receptor potential cation channel, type 3
UBN	unipolar brush neuron
VDCC	voltage dependent calcium channel
vGluT	vesicular glutamate transporter
VZ	ventricular zone

Summary

The cerebellum is a brain region crucial for coordination and motor learning. Being the principal output cell of the cerebellar cortex, Purkinje cell loss and degeneration play an important role in many cerebellar diseases. The most striking feature of cerebellar Purkinje cells is their large and extensively branched dendritic tree, which is almost flat and strictly arranged in the sagittal plane. The factors and molecules which control the growth and patterning of neuronal dendrites are still poorly understood. Previous research in our lab has shown that chronic activation of metabotropic glutamate receptor 1 (mGluR1) (Sirzen-Zelenskaya et al., 2006) or Protein Kinase C (PKC) (Metzger & Kapfhammer, 2000; Schrenk et al., 2002) in organotypic cerebellar slice cultures of postnatal mice severely inhibits the growth and development of the Purkinje cell dendritic tree.

Although we found that the similar effect induced by mGluR1 or PKC is mediated by independent mechanisms (Sirzen-Zelenskaya et al., 2006), the signaling events leading to inhibition of dendritic growth after both mGluR1 and PKC activation remain largely unknown. Another intriguing question is that of the physiological relevance of limiting dendritic size after chronic activation of a metabotropic glutamate receptor.

We addressed both aspects using organotypic cerebellar slice cultures. In this culture model, a 300 μm thick slice of cerebellar tissue is kept in culture, where the natural microenvironment of a cell with neighboring cell-cell interactions and local neuronal networks are preserved. In order to study Purkinje cell dendritic development, cerebella from mice were cultured shortly before Purkinje cells enter the developmental stage of rapid dendritic growth and expansion, and were maintained for 10-12 days.

Purkinje cells are especially affected in various diseases involving excitotoxicity. We have tested the hypothesis that it is the size of the dendritic tree which determines Purkinje cell sensitivity to excitatory overload. Therefore, we have grown Purkinje cells under conditions which result in a strong reduction of dendritic tree size. Then we have exposed the cultures to (RS)- α -Amino-3-hydroxy-5-methyl-4-isoxazolepropionic acid (AMPA), which is an appropriate model system to study excitotoxic neurodegeneration in the cerebellum as it selectively kills Purkinje cells but leaves cerebellar granule cells unharmed. Our results indicate that a reduction of the size of the dendritic tree in Purkinje cells does not offer any protection from glutamate-induced excitotoxicity. Staining for the vesicular glutamate transporter

vGluT1 revealed a high density of glutamatergic synapses on the stunted dendritic trees of pre-treated Purkinje cells. This suggests that receptor density rather than total receptor load is important for determining the sensitivity of Purkinje cells to AMPA-mediated neurotoxicity.

In a second study we have searched for potential mechanisms limiting Purkinje cell dendritic growth and have concentrated on channels allowing the entry of Ca^{2+} ions, especially the TRPC3, P/Q-type and T-type Ca^{2+} channels.

To analyze the roles of these channels in mediating dendritic reduction induced by mGluR1 or PKC we have treated Purkinje cells in cerebellar slice cultures with the mGluR1 activator DHPG or the PKC activator PMA and simultaneously with different combinations of Ca^{2+} channel inhibitors. After the culture period, the size of the dendritic trees was evaluated.

Co-treatment with a combination of P/Q- and T-type inhibitors partially rescued the dendrite-reducing effects induced by DHPG or PMA treatment. In contrast, no improvement of dendritic growth was found in mice lacking TRPC3 function or using an acute pharmacological TRPC3 inhibitor. Furthermore, the partial rescue obtained by P/Q- and T-type inhibition was not further increased by additional co-treatments with inhibitors of GluR δ 2 - or R-type Ca^{2+} channels, or of Ca^{2+} release from internal stores.

In conclusion, our results suggest that T-type and P/Q-type Ca^{2+} channels are part of the signaling pathways induced after chronic mGluR1 and PKC stimulation resulting in the inhibition of dendritic growth, while no involvement of TRPC3-, GluR δ 2-, R-type Ca^{2+} channels or Ca^{2+} release from intracellular stores was found.

1. Introduction

1.1. General Dendritic Development

1.1.1. Overview

There are a large number of different neurons in the nervous system, and each type develops a specific set of dendrites which is precisely adapted to a specific neuron's function and role in neuronal circuits. Demonstrating the essential significance of dendrites in cognition, aberrant dendritic morphologies have been found in several conditions of mental retardation (reviewed in (Kaufmann & Moser, 2000)) as well as in diseases such as schizophrenia (Broadbelt *et al.*, 2002).

Dendritic structures vary in size and complexity and, as a consequence, in the number of synapses they bear (Purves & Hume, 1981). Dendrites carry some characteristic features that are responsible for the way a neuron integrates various synaptic inputs and reacts to them with action potentials and/or synaptic plasticity (Hausser *et al.*, 2000). Given this highly specialized structure of individual dendrites it is not surprising that dendritic development is a complicated process that involves genetic programs running in concert with reactions to environmental extracellular signals.

In general, dendritic development starts after the axon has emerged from the cell body (Gao *et al.*, 1999; Wu *et al.*, 1999). It can be subdivided into three major stages (Wu *et al.*, 1999) (see Fig. 1):

1. (See Fig. 1 A). During the outgrowth of the primary dendrites overall dendritic growth is dynamic but slow and few branches with growth cones are present (Dailey & Smith, 1996; Wu *et al.*, 1999).
2. (See Fig. 1 B). Concurrent with the beginning of synaptogenesis, filopodia emerge from the dendritic shafts (Dailey & Smith, 1996). Synapses can be found on filopodia as well as on dendritic shafts. The dendritic trees gain complexity in a highly dynamic manner with branches and filopodia rapidly rearranging by addition, retraction, resorption or elongation and stabilization to become dendritic branches or spines (Papa *et al.*, 1995; Dailey & Smith, 1996; Saito *et al.*, 1997; Fiala *et al.*, 1998; Cline, 2001). In fact, turnover of filopodia

occurs within less than a minute (Dailey & Smith, 1996; Williams & Truman, 2004).

Besides interstitial branching, where a filopodium emerges from the dendritic trunk, an alternative mechanism for branching is the splitting of growth cones, which happens under some circumstances in cultured neurons (Bray, 1973).

The rapid dynamic motility of dendrites has been proposed to be intrinsically regulated (Dunaevsky *et al.*, 1999), while other studies have found dendritic motility to be responsive to external stimulation (Maletic-Savatic *et al.*, 1999) as well as to application of glutamate and NMDA (Segal, 1995; Korkotian & Segal, 1999). The differences between findings in different studies are to be expected because the effects of activity on dendrites depend on the age of the dendrites as well as on the exact way, intensity and even location of the applied activity (Bravin *et al.*, 1999).

3. (See Fig. 1 C). In the end-phase of synaptogenesis, filopodia are more and more stabilized to become spines and overall motility of terminal branches is slowing down as the dendritic tree takes its terminal shape (Dailey & Smith, 1996; Dunaevsky *et al.*, 1999). Limitation of dendritic growth plays an important role at this stage of development. A striking example is a phenomenon called tiling, which has been described in the vertebrate retina, where it has first been observed (Wassle *et al.*, 1981), as well as in several types of neurons in *Drosophila* (Grueber *et al.*, 2002; Millard *et al.*, 2007; Soba *et al.*, 2007; Ting *et al.*, 2007). One speaks of tiling when dendrites of the same type of neurons completely cover a receptive area, but their dendritic fields do not overlap. They do overlap, however, with dendrites of other classes of neurons. To establish this situation, there must be a repulsive signal upon interaction with neighboring dendrites of the same type. Dscam2, a transmembrane adhesion molecule of the immunoglobulin superfamily, mediates tiling behavior in L1 lamina neurons in *Drosophila* by homophilic binding and repulsion (Millard *et al.*, 2007). Also a very important role in tiling and branching in *Drosophila* sensory neurons play the products of the genes tricornered (*trc*) and furry (*fry*) (Emoto *et al.*, 2004). The serine/threonine kinase Trc (Tamaskovic *et al.*, 2003) is activated by the large protein Fry to mediate a repulsive response after contact with like dendrites (Emoto *et al.*, 2004). Appropriately, mammalian Fry has been shown to bind to microtubules

(Tamaskovic *et al.*, 2003), thus having a possibility to influence dendritic growth patterns.

A related phenomenon to tiling is self-avoidance, which is crucial for segregation of dendritic branches of the same cell (Kramer & Stent, 1985). The molecular basis of self-avoidance was identified in *Drosophila* as Dscam1, another protein of the immunoglobulin superfamily, which can be spliced in over 38'000 of different ways (Schmucker *et al.*, 2000). Each neuron expresses on its surface a specific set of Dscam1 splice variants (Neves *et al.*, 2004). If Dscam1 molecules of the same splice variant bind to each other, as it statistically only happens when dendritic branches of the same cell meet, they generate a repulsive signal that causes the branches to turn and grow in a different direction (Wojtowicz *et al.*, 2004; Hattori *et al.*, 2007a; Matthews *et al.*, 2007).

Vertebrate Dscam was recently shown to carry out a similar function in the mouse retina. Mutant mice lacking functional Dscam had severe defects in arborization and mosaic-like distribution of cell bodies in only those retinal amacrine cell types which naturally express Dscam (Fuerst *et al.*, 2008). Unlike Dscam1 in *Drosophila*, vertebrate Dscam appeared to be responsible for both self-avoidance (arborization) and tiling. In about 60 % of retinal ganglion cells and retinal interneurons, either Dscam or a closely related adhesion molecule of the immunoglobulin superfamily (DscamL, Sidekick-1 or Sidekick-2) has been found to be expressed and play a role in arborization of the neurons in a specific sublamina of the inner plexiform layer (Yamagata & Sanes, 2008).

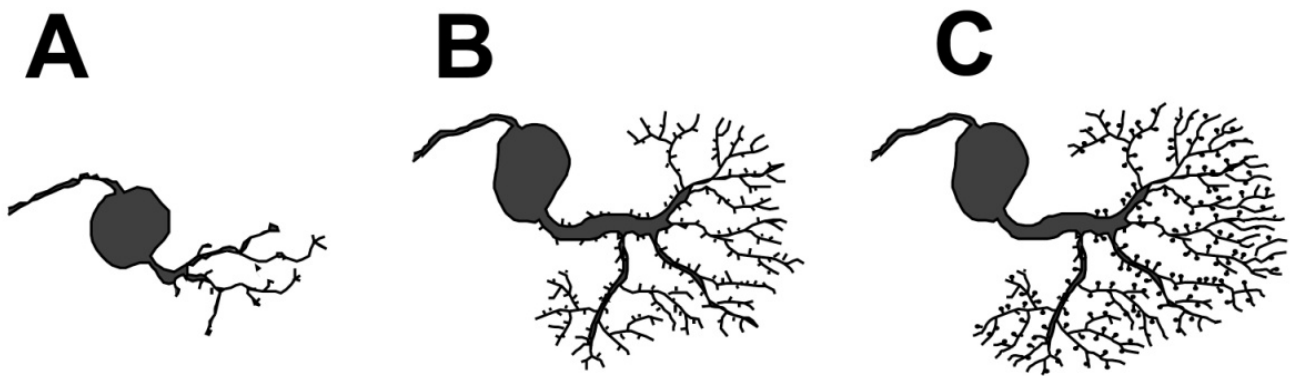


Figure 1: Stages of dendritic development.

A) Outgrowth of primary dendrites, slow overall dendritic growth.

B) Fast and highly dynamic dendritic growth and synaptogenesis

C) Slow dendritic growth, refinement of dendritic tree, stabilization of synapses and spines, tiling. Picture adapted from (Wong & Wong, 2000).

Regarding dendritic development in a larger context than motility of single branches, directed neuronal activity has been shown to be highly influential in various cell types and species. For example, mice which were raised in a so-called enriched environment, promoting learning and social interaction, developed larger dendritic trees selectively in the Hippocampus, a brain region essential for learning and memory (Faherty *et al.*, 2003). Similarly, in *Xenopus laevis* tadpoles, 4 hours of light stimulation proved sufficient to enhance dendritic growth in optical tectal neurons even during a following 4 hour's period in the dark (Sin *et al.*, 2002). Similarly, pharmacological inhibition of synaptic activity in the CA1 region of early postnatal mice resulted in decreased branching of pyramidal cells (Groc *et al.*, 2002).

1.1.2. Dendritic growth regulation by extracellular molecules

Apart from neurotransmitter-conveyed signals there are other extrinsic cues that have been found to be influential on dendritic development and mediate interneuronal communication. I would like to introduce a few interesting examples.

Several **hormones** have been found to have specific effects on dendritic growth in certain brain regions. For example, Pyramidal cells in the CA3 region of the Hippocampus reacted to hyperthyroidism with increased dendritic growth and branching (Gould *et al.*, 1990b) and, consistently, to hypothyroidism with reduced

dendritic trees (Rami *et al.*, 1986). Pyramidal cells in the CA1 region, on the other hand, responded to thyroid treatment with a decrease in spine density, but no changes in dendritic growth and branching were observed (Gould *et al.*, 1990a; Gould *et al.*, 1990b).

The **neurotrophins** comprise at least four proteins: nerve growth factor (NGF), which was first discovered in the 1950's (Cohen *et al.*, 1954), neurotrophin-3 (NT-3), neurotrophin-4 (NT-4) and brain-derived neurotrophic factor (BDNF). Their receptors are the low-affinity receptor p75 and the high-affinity receptor-kinases TrkA, TrkB and TrkC. Trks are tyrosine kinases and differentially expressed in the developing CNS (Martin-Zanca *et al.*, 1986; Ringstedt *et al.*, 1993; Segal, 2003).

In one of the first studies to investigate the role of a neurotrophins in dendritic development, neonatal rats were treated with NGF for 1 or 2 weeks, which resulted in highly significant increase of dendritic length, branching and number of primary dendrites in sympathetic ganglion neurons (Snider, 1988).

The effect of neurotrophins on cortical pyramidal neurons in the CNS has been addressed in several studies. For example, in organotypic slice cultures of the developing visual cortex of ferrets, treatment with NGF, BDNF, NT-3 or NT-4 over 3 days showed that basal dendrites of pyramidal neurons in each cortical layer responded predominantly to a single neurotrophin, while the apical dendrites responded to a wider range of neurotrophins (McAllister *et al.*, 1995). Interestingly, the effects of neurotrophins on different cells, and even on different dendrites on the same cell, varied widely. BDNF, for example, enhanced dendritic growth of basal dendrites in layers 4 and 5, but decreased it in layer 6. Since the neurons in one specific layer of the mammalian cortex build a functionally and morphologically related community, neurotrophins do not generally promote dendritic growth but act specifically in the development of typical growth patterns (McAllister *et al.*, 1995; McAllister *et al.*, 1997). Additionally, responsiveness of pyramidal cells to neurotrophins has been shown to require synaptic activity (McAllister *et al.*, 1996), thus providing a possible mechanism of how bioelectrical activity may influence dendritic growth.

Expression of the small membrane-bound protein **CPG15** is induced by neuronal activity (injection of kainate or light-exposure) in adult rats, and it is also expressed during development (Nedivi *et al.*, 1996). When CPG15 was transfected into the optic tectum of *Xenopus* tadpoles, the dendritic complexity and branch length of projection

neurons, but not of interneurons, was increased (Nedivi *et al.*, 1998). Similar results were obtained in cats (Corriveau *et al.*, 1999).

CPG15 protein is concentrated in axons of retinal ganglion cells during the period when the dendritic development of their target cells was sensitive to CPG15 (Nedivi *et al.*, 1998). This may provide another mechanism through which neuronal activity is transduced into dendritic growth. Recently, a paralogue of CPG15, CPG15-2, was discovered, which has a different expression pattern but has many functional similarities with CPG15 (Fujino *et al.*, 2008).

Semaphorins are a large family of guidance molecules for axons and dendrites. **Semaphorin3A** (Sema3A) is produced and secreted by the cortical plate during development from late embryonic stages (Giger *et al.*, 1996; Skalióra *et al.*, 1998) and has been shown to act as a chemorepellent on growing axons of cortical pyramidal cells (Polleux *et al.*, 1998). Conversely, it acts as a chemoattractant on apical dendrites of the same cells (Polleux *et al.*, 2000), as well as a regulator of spine maturation (Morita *et al.*, 2006). At least one part of the Sema3A receptor, Neuropilin-1, is apparently involved in both repelling axons and attracting dendrites, as it is expressed in pyramidal neurons in axons as well as in dendrites (He & Tessier-Lavigne, 1997; Kolodkin *et al.*, 1997; Polleux *et al.*, 2000). Whether Sema3A acts as a repellent or an attractant has been shown to be dependent on the presence of cyclic GMP (Song *et al.*, 1998). Indeed, in cortical pyramidal neurons, the distribution of soluble guanylate cyclase (SGC) has been found to be asymmetric with a high concentration of SGC in the apical pole already before the outgrowth of the apical dendrite, and SGC was shown to specifically mediate dendritic orientation via regulation of cGMP-dependent protein kinase (PKG) (Polleux *et al.*, 2000). Besides several effector-molecules which have been supposed to be involved, the two kinases Fyn and CDK5 have been shown to take part in mediation of the Sema3A effect on apical dendrites (Sasaki *et al.*, 2002; Morita *et al.*, 2006).

Glial cells promote dendritic growth even after they are fully matured (Le Roux & Reh, 1995), in contrast, they lose their ability to promote axon growth after the initial phase of development (Smith *et al.*, 1990). It has been found that rat embryonic RGCs require direct contact to neighboring astrocytes to become able to form dendritic synapses (Barker *et al.*, 2008). Direct contact with astrocytes caused dislocation of the protein Neurexin out of the dendrites, and overexpression of Neurexin in dendrites abolished the astrocytic effect on synaptogenesis, thus

proposing a possible mechanism (Barker *et al.*, 2008). In postnatal RGCs, on the other hand, astrocytes promote synaptogenesis without requiring direct cell-cell interaction; instead astrocytes secrete signals that promote synaptogenesis (Nagler *et al.*, 2001). Two proteins of the thrombospondin family (TSPs), TSP-1 and TSP-2, have been identified to be a necessary part of this astrocyte-secreted signal to induce synaptogenesis in cultured RGCs and *in vivo* (Christopherson *et al.*, 2005). Another mechanism for how neuron-glia-interaction may be mediated has been suggested for pyramidal cells in the CA1 region of the Hippocampus in mice, where contact with astrocytes regulates spine morphology. Adult hippocampal spines express the EphA4 receptor on their surface, and its ligand, Ephrin-A3, is expressed on the surface of astrocytes (Murai *et al.*, 2003). EphA4-activation causes spines to retract, which is important for proper spine organization and regulation of spine morphology (Murai *et al.*, 2003).

1.1.3. Internal pathways to integrate external signals

In Chick retinal ganglion cells it has been found that neurotransmitter-mediated input leading to Ca^{2+} induced Ca^{2+} release from internal stores and subsequently to local increases of internal Ca^{2+} concentration is required for maintaining dendritic structures, while global increases of Ca^{2+} concentration (concerning the whole neuron) were not required (Lohmann *et al.*, 2002).

In other studies, Ca^{2+} entry through L-type VDCCs has been found to play an important role in activity-induced dendritic changes (Wu *et al.*, 2001; Redmond *et al.*, 2002).

All movement of cellular processes eventually involves changes in the cytoskeleton. While the cytoskeleton in filopodia and spines fundamentally consists of actin, dendritic branches are filled with microtubules (Fischer *et al.*, 1998; Harada *et al.*, 2002). How are external signals transferred to affect actin and microtubules?

In cultured rat sympathetic neurons, neuronal activity also caused dendrite formation, and this effect was enhanced by addition of NGF. Activation of a key regulator of dendritic stability (Harada *et al.*, 2002), the microtubule-associated protein MAP2 by phosphorylation through two types of kinases, CamKII and Erks was necessary to stabilize microtubules and thus mediate the effect of activity (Vaillant *et al.*, 2002).

The necessity of CamKs and the MEK-ERK-pathway to work together in activating MAPK2 after stimulation was also found in a study in cultured hippocampal neurons (Wu *et al.*, 2001). Interestingly, the pattern of activation had a profound effect: only spaced, repetitive impulses resulted in a prolonged phosphorylation of dendritic MAPK, which correlated with stable morphological changes in spines. Activity of CaMKs was only required for short-term activation of MAPK.

In another study in hippocampal neurons, the effect of neuronal activity on actin has been found to require **β -catenin** (Yu & Malenka, 2003). β -catenin is an important part of the Wnt signaling pathway, which regulates gene transcription in cancer and embryonic development (Cadigan & Nusse, 1997; Bienz & Clevers, 2000; Polakis, 2007). Additionally, it functions in the cadherin/catenin complex to mediate cell adhesion and ties the actin-cytoskeleton to cadherins in the cell membrane (Gumbiner, 1996). In rat hippocampal neurons β -catenin is present during development (Benson & Tanaka, 1998) and high levels of β -catenin (and other components of the cadherin/catenin complex) have been shown to enhance dendritic branching (Yu & Malenka, 2003). Furthermore, membrane depolarization (thus mimicking electrical activity) also increased dendritic branching and this effect required β -catenin to be mediated. β -catenin influenced dendritic branching independently of its role in gene transcription, instead it has been shown that the extracellular domain of N-cadherin as well as actin-binding was necessary (Yu & Malenka, 2003).

Similarly, **Rho**, **Rac** and **CdC42**, the best-studied subfamilies of the Rho family of small GTPases, which are of major importance for regulating the actin cytoskeleton (Hall, 1998; Hall & Nobes, 2000), have been shown to enhance dendritic elaboration in cultured cortical neurons of E18 rats (Threadgill *et al.*, 1997).

Concerning the dynamics of dendritic remodeling during synaptogenesis in chick retinal ganglion cells at E12-13, Rho and Rac have been found to have a reciprocal influence (Wong *et al.*, 2000). While Rac activation increased glutamatergic transmission-dependent motility and number of fine terminal processes, Rho decreased it.

Rho, Rac and CdC42 have also been shown to mediate the effects of activity on dendritic growth in the optic tectum of *Xenopus laevis* tadpoles (Li *et al.*, 2002; Sin *et al.*, 2002). Consistent with the results obtained by Wong *et al.* (Wong *et al.*, 2000),

Rho activity was found to inhibit activity-induced dendritic growth, while Rac and Cdc42 enhanced it (Sin *et al.*, 2002).

On the other hand, in cortical cultures from embryonic rats, electrical stimulation and following Ca^{2+} entry via L-VSCCs exerted their growth-promoting effect on dendrites through induction of protein synthesis (Redmond *et al.*, 2002). CaMKIV activated the transcription factor CREB and its co-activator CBP which resulted in changes of gene transcription.

1.1.4. Dendritic growth regulation by transcriptional programs

Although synaptic input and environmental interactions are undoubtedly necessary for dendritic development, a study by Verhage and colleagues provided evidence that intrinsic programs make up a substantial part of brain development (Verhage *et al.*, 2000). Deletion of the *munc-18-1* gene in mice abolished all neurotransmitter release and resulted in completely paralyzed mice. Although neurons eventually underwent apoptosis, all brain regions initially developed in an apparently normal way including circuit assembly and synaptogenesis (Verhage *et al.*, 2000).

The importance of a complicated transcriptional control for morphological development is undeniable. Transcription factors are the core components of every cell-intrinsic growth pattern.

The transcription factor **hamlet**, for example, is expressed only in the neuronal precursor cell IIB in the *Drosophila* PNS and is inherited by both daughter cells, an external sensory (ES) neuron and a glial cell (Moore *et al.*, 2002). Hamlet expression persists in ES neurons throughout morphogenesis of their single dendrite, which does not develop branches. Without hamlet expression, however, ES cells become multidendritic (MD) neurons, which have highly branched dendritic trees and are derived from the IIB precursor cell, which gives also rise to the IIB precursor. Hamlet has been found to determine both cell fate specification and dendritic morphology (Moore *et al.*, 2002). Similarly in mammals, the characteristic unipolar dendritic morphology of pyramidal neurons in the neocortex and hippocampus has been found to be determined by expression of the transcription factor Neurogenin2 (**Ngn2**) in their progenitors (Hand *et al.*, 2005).

Interesting examples for how specific expression of transcription factors endows neurons with their characteristic dendritic arbors are the *Drosophila* dendritic

arborization (da) neurons, a subgroup of the md neurons and an extensively studied model. According to their dendritic arbor complexity, they have been categorized into four sub-classes (Grueber *et al.*, 2002) (see Fig. 2). Each sub-class expresses distinct levels of the three transcription factors Cut, Abrupt and Knot (Grueber *et al.*, 2003; Sugimura *et al.*, 2004; Jinushi-Nakao *et al.*, 2007). In classes I-III, increasing expression of Cut correlates with increased dendritic branching (Grueber *et al.*, 2003). Class I da neurons have the least complex branching pattern and are the only da subclass with no detectable **Cut** expression (Grueber *et al.*, 2003), instead they express **Abrupt** (Sugimura *et al.*, 2004). Abrupt has been shown to be responsible for the class-I-specific dendritic comb-like morphology (Sugimura *et al.*, 2004). Although Cut and Abrupt are expressed complementary, they do not influence each other at the level of gene expression (Sugimura *et al.*, 2004). When they are both overexpressed, however, they functionally antagonize each other with Abrupt limiting dendritic growth and branching (Sugimura *et al.*, 2004).

Class III neurons have on their long primary and secondary dendrites many so-called spikes, which contain actin but no microtubules, whereas class IV neurons have microtubule-filled higher-order branches (Grueber *et al.*, 2002). Class IV da neurons express lower levels of Cut than class III neurons (Grueber *et al.*, 2003). Additionally, they express **Knot** which causes branching and represses Cut-driven spike-formation (Hattori *et al.*, 2007b; Jinushi-Nakao *et al.*, 2007; Crozatier & Vincent, 2008). The Knot-induced effect on branching has been shown to be essentially mediated by upregulation of spastin, a protein influencing microtubule organization. Both Cut and Knot are required for class IV-specific dendritic morphology, since they regulate different features of the dendritic arbor, by targeting different aspects of the cytoskeleton (Jinushi-Nakao *et al.*, 2007). While Knot has no influence on Cut expression, Cut influences the amount of Knot that is expressed, although not its initial expression (Jinushi-Nakao *et al.*, 2007; Crozatier & Vincent, 2008).

Furthermore, all four classes of neurons, like all sensory neurons in the *Drosophila* embryonic PNS, express the transcription factor Spineless independently of Cut and Abrupt (Kim *et al.*, 2006). In classes I and II, Spineless limits dendritic branching, while in classes III and IV, Spineless promotes branching (Kim *et al.*, 2006).

To underline that these explanations are only a rough simplification of the complicated interplay between transcription factors in dendritic morphogenesis, I would like to mention a study by Parrish and colleagues (Parrish *et al.*, 2006), where

RNA interference was used to identify over 70 transcription factors involved in dendritic development of class I da neurons alone.

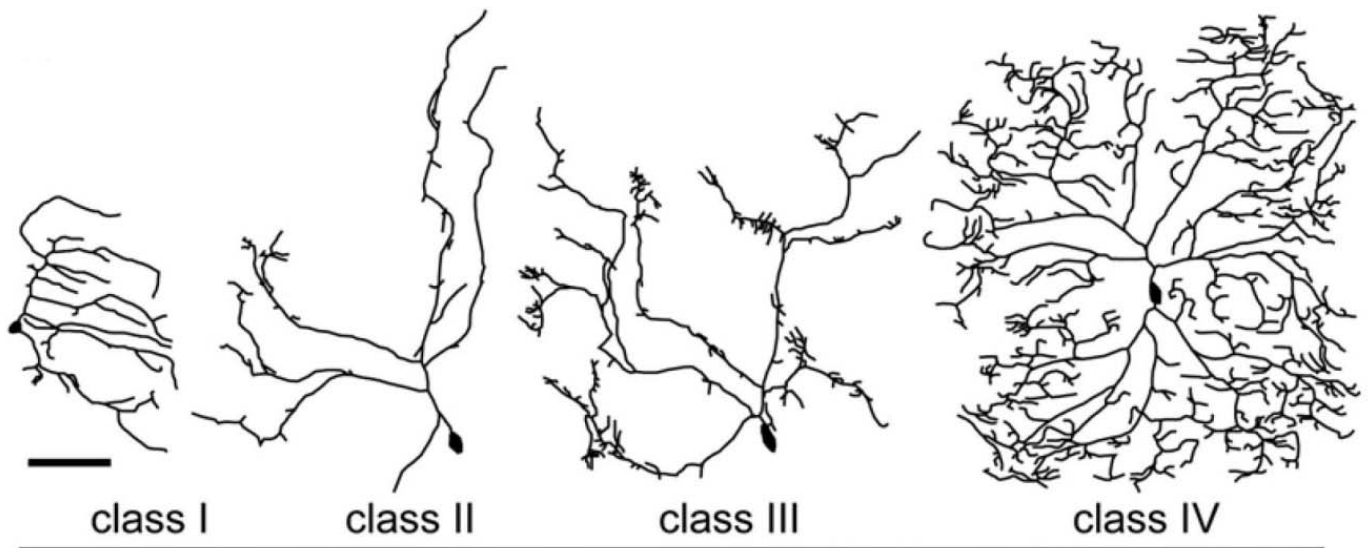


Figure 2: Dendritic branching pattern of the four subclasses of *Drosophila* dendritic arborization neurons according to (Grueber *et al.*, 2002). Picture from (Jinushi-Nakao *et al.*, 2007)

1.2. Cerebellum

Anatomically, the cerebellum fundamentally consists of the cerebellar cortex, the deep cerebellar nuclei and the cerebellar peduncles.

The cerebellar cortex is furled into parallel folds running transversely. While there are numerous anatomical subdivisions of the cerebellum which have little or no functional meaning, its hemispheres can be functionally divided into three main sections (see Fig. 3):

- I. The **cerebrocerebellum** is the largest part in humans. It is concerned with intricate movements such as speech or playing a musical instrument, as well as with visual guidance of ongoing movements. Damage to the cerebrocerebellum consequently results in impairment of such learned skills which depend on fine coordination (Dale Purves, 2008).
- II. The **spinocerebellum** occupies the median parts of the hemispheres, including the midline, which is called vermis. It is primarily concerned with movements of the limbs, as well as with some eye movements (e.g.

saccades). The somatic sensory input from the whole body surface is represented as somatotopic maps in the spinocerebellum. The lower limbs, for example, are mapped in the anterior part of the spinocerebellum. Chronic alcohol abuse damages specifically the anterior portion of the cerebellum and consequently, patients suffer from a staggering, shuffling gait but can still fluently move their arms and hands. Damage in larger parts of the spinocerebellum, however, result in intention tremor and general impairment of well-coordinated movements (Dale Purves, 2008). The vermis region has been found to be involved in fear-related involuntary movements such as freezing in rats (Supple *et al.*, 1987), while the more lateral parts of the spinocerebellum are crucial for learning conditioned eyeblink reflexes (Bracha *et al.*, 1997).

- III. The **vestibulocerebellum** projects to Lower motor neurons in the spinal cord and brainstem and is responsible for posture and balance, as well as for the vestibulo-ocular reflex. Difficulties with a balanced stance or following a moving object with one's eyes as well as loss of muscle tone are typically related with lesions of the vestibulocerebellum (Dale Purves, 2008).

This classification is mainly based on the source of the afferents and the destination of efferents connecting these functional divisions with the rest of the brain. These three divisions receive input from different regions of the CNS and project through different deep cerebellar nuclei and peduncles, and finally, to distinct parts in the CNS (Dale Purves, 2008).

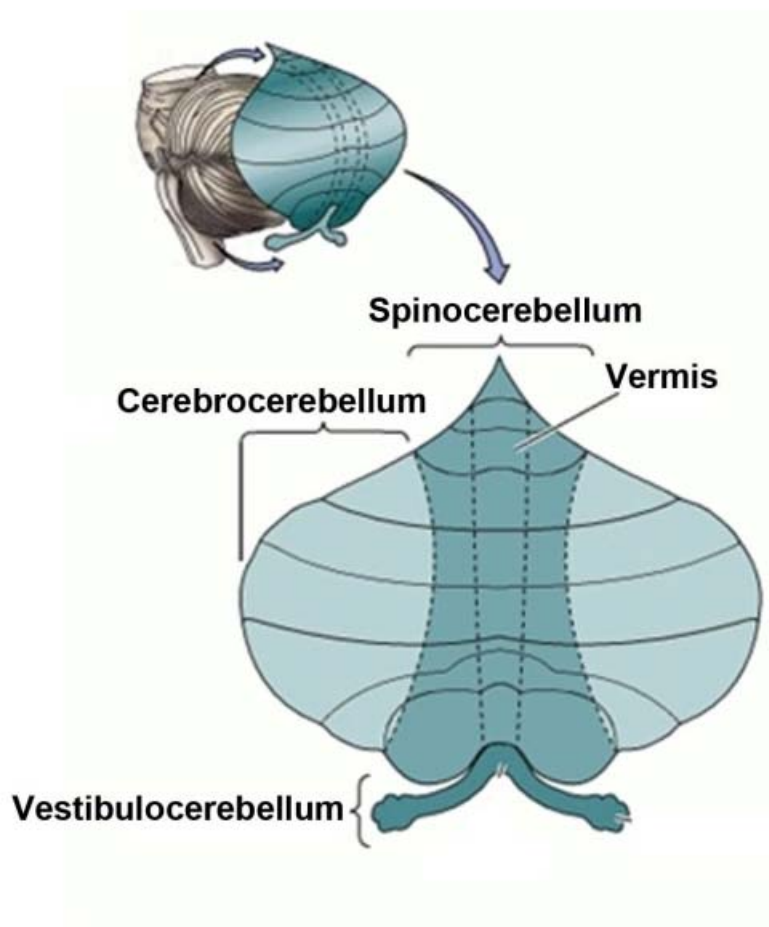


Figure 3: Flat projection of the cerebellar surface showing the three main areas.
Adapted from (Dale Purves, 2008).

1.2.1. Cerebellar Cytology

The density of neurons in the cerebellum is very high compared to other parts of the brain, and the cerebellar cortex is highly organized. It consists of three layers (see Fig. 4):

1. The molecular layer (ML) is the outermost layer of the cerebellar cortex, and it principally consists of the Purkinje cell dendritic trees, which are strictly arranged in the sagittal plane, and the T-shaped axons of the cerebellar granule neurons (CGN) called parallel fibers (PF). They run perpendicular to the Purkinje cell dendritic arbors and one PF builds synapses on the dendritic

spines of tens of thousands of Purkinje cells, while, on the other hand, one Purkinje cell gets contacted by up to 200 000 PFs (Napper & Harvey, 1988; Dale Purves, 2008). The relatively few cell bodies present in the molecular layer belong to two types of inhibitory interneurons, which are modulating the excitatory input from PFs on PCs:

- The basket cells synapse on Purkinje cell bodies
- The stellate cells synapse on Purkinje cell dendrites

The second excitatory input Purkinje cells receive is the one from the climbing fibers (CF), which are axons from neurons in the inferior olive. One single CF makes numerous synapses on the dendritic shaft of one single PC.

2. Below the ML, the Purkinje cell bodies are aligned in a monolayer, which is called Purkinje cell layer (PCL). Purkinje cells are GABAergic neurons and provide the principal output of the cerebellar cortex.
3. Below the PCL and right above the white matter lies the granular layer (GL). It contains the most numerous cell type in the whole brain, the cerebellar granule neurons (CGNs), as well as the Golgi interneurons, which synapse on CGN dendrites and whose dendrites get input from the PFs in the ML, thus providing a negative feedback loop to the CGNs. The CGNs receive excitatory input from the mossy fibers which are the major afferents of the cerebellum coming from various brain regions (see above). A remarkable type of excitatory interneuron, the unipolar brush neurons (UBN), intensify this afferent input of mossy fibers on CGNs, especially in the vestibulocerebellum (Kalinichenko & Okhotin, 2005). Another cell type in the GL, situated just below the PCL, are the Lugaro cells, an inhibitory interneuron which contacts Basket and stellate cells in the ML, and Golgi cells in the GL, while Lugaro cells themselves receive input from Purkinje cell collaterals (reviewed in (Geurts et al., 2003))

Through the white matter underneath the cortex, the Purkinje cell axons project to neurons in the deep cerebellar nuclei (DCN). Their axons leave the cerebellum through the cerebellar peduncles.

All input to the cerebrocerebellum is relayed onto neurons in the pontine nuclei and the inferior olive. From the pontine nuclei, the mossy fibers enter the cerebellum through the cerebellar peduncles, split off collaterals to neurons in the deep cerebellar nuclei and ascend further to excite the CGNs. The CGNs convey

the excitatory signal to the Purkinje cells, where the integration of information is controlled by stellate and basket cells, as well as by CF input. The CFs from the inferior olive also project collaterals to DCNs before they contact Purkinje cells. CF input on Purkinje cells reduces their excitability and thus the effectiveness of the PF-PC synapses. Immediate motor error corrections as well as long-term adaptations like motor learning have been shown to depend on CF activity (Dale Purves, 2008).

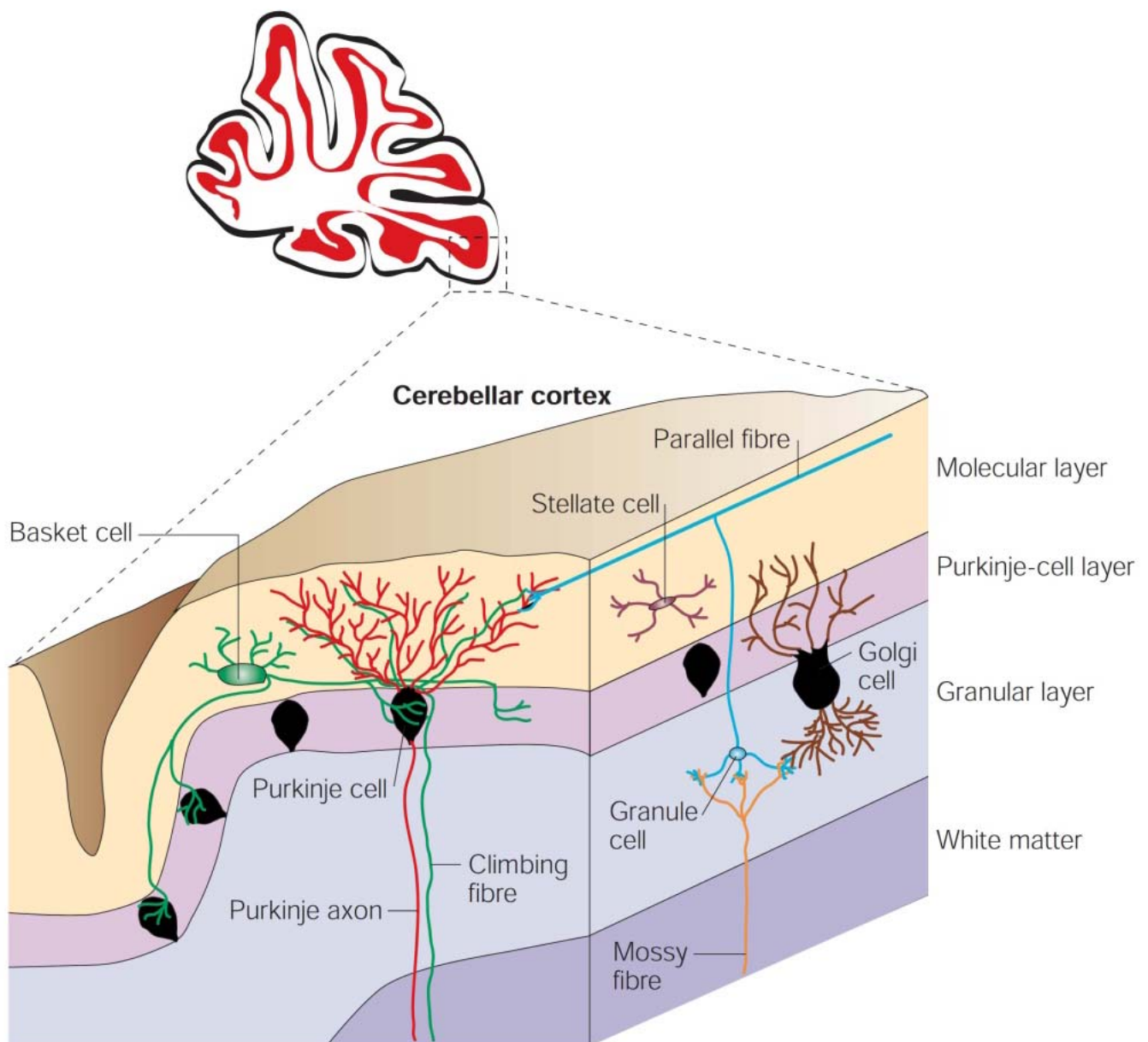


Figure 4: Schematic enlargement from a sagittal section through the cerebellum. Composition of the cerebellar cortex with the most important cell types is illustrated. Adapted from (Kobielak & Fuchs, 2004).

1.2.2. Functional aspects

The cerebellum is a brain region crucial for coordination and motor learning, and is involved in planning and execution of movements as well as in ocular reflexes and maintaining balance and posture. According to more recent findings, the cerebellum also plays a role in spatial orientation and modulating cognitive programs.

Cerebellar diseases and lesions are characterized by difficulties with fluid and precise movements rather than paralysis. In cerebellar ataxia the coordination of muscle groups is disturbed, which leads to clumsiness. Typical signs are intention tremor, applying a wrong amount of muscle force to a task, reaching too high or too low (dysmetria), problems with keeping an upright and balanced stance, unsteady muscle tone or carrying out movements at the wrong speed. These symptoms can affect all skeletal muscle groups, involving speech, eye movements, movements of limbs and trunk. There are a group of congenital diseases called spinocerebellar ataxias (SCA), which are all caused by different genetic disorders concerning the cerebellum. SCAs are usually slowly progressing disorders caused by mutation in a single gene and include autosomal-dominant as well as recessive patterns of inheritance (reviewed in (Perlman, 2011)). Dominant SCAs are generally caused by a “gain-of-function” gene mutation, which most often includes an expansion of CAG triplet repeats. These polyglutamine gene products become toxic and often lead to degeneration of cerebellar structures (reviewed in (Perlman, 2011)). The detailed proceedings and extent of a cerebellar ataxia strongly depend on the exact location of the lesion or damage in the cerebellum (see above).

Altogether, the cerebellum receives input from various areas of the cerebral cortex, spinal cord and brainstem, and it mainly projects back to the Premotor and Motor cortices, Thalamus, spinal cord and brainstem. This constitutes a feedback-loop which allows the cerebellum to carry out one of its most important functions, which is the correction of motor errors. During the execution of a movement, the cerebellum receives constant sensory information on position and motion of the body, which allows detection of differences in the originally planned movement and the actual situation. The cerebellum then sends a corrective signal to motor neurons, in order to adapt the movement immediately, while the correction can also be stored for motor learning (Dale Purves, 2008).

1.2.3. Basics of cerebellar development

Early in development, the neural tube is subdivided along the rostral-caudal axis into four areas. The prosencephalon at the front of the neural tube will later bring about the forebrain, the mesencephalon will develop mainly into the midbrain, and the rhombencephalon, after dividing into the metencephalon and the myelencephalon, will form the hindbrain. The cerebellum principally evolves from the metencephalon, with parts of the mesencephalon contributing (Wang & Zoghbi, 2001; Dale Purves, 2008). By embryonic day 9.5 (E9.5) in the **mouse**, the neural tube is virtually closed, but at the border between the mesencephalon and the metencephalon, there is a gap which forms a specialized region of the dorsal neuroepithelium, the rhombic lip (RL) (see Fig. 5) (Morales & Hatten, 2006). In the developing cerebellum, there are basically two different germinal matrices where neurons are produced: the RL and the ventricular zone (VZ) (Wingate, 2001; Hoshino et al., 2005; Butts et al., 2011).

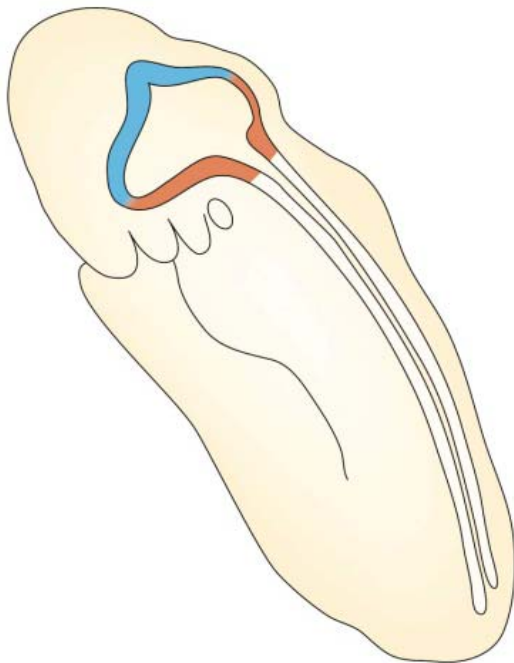
All GABAergic neuron populations derive from the VZ from progenitors expressing the basic helix-loop-helix (bHLH) transcription factor Ptf1a (Hoshino et al., 2005). Between E10 and E17, the GABAergic fraction of DCNs is born, Purkinje cells are produced from E11-E13, and GABAergic interneurons subsequently (Miale & Sidman, 1961; Wang & Zoghbi, 2001).

In the RL, TGF β -signaling from the neighboring roof plate induces the expression of a different bHLH transcription factor, Math1 (often also referred to as Atoh1). All glutamatergic neuron populations in the cerebellum arise from the RL, and they are born during a particular time frame (Gilthorpe et al., 2002; Hoshino et al., 2005; Butts et al., 2011):

At E10.5 in the mouse, the glutamatergic fraction of deep cerebellar neurons is born. They switch off expression of Math1 after migrating away from the RL towards the cerebellar nuclei.

2 days later, the RL starts to produce unipolar brush neurons and progenitors of cerebellar granule neurons (GNP). GNPs migrate first along the surface of the neural tube and form the external granule Layer (EGL) (see Fig. 6). This process lasts until after birth. In the EGL, GNP transit amplification is induced by sonic hedgehog (Shh) signaling from underlying Purkinje cells and is dependent of maintained Math1 expression (Flora et al., 2009). The extent and duration of Shh signaling (and thus GNP proliferation in the EGL) is an essential determinant for the development of

cerebellar lobules and complexity of foliation (Corrales et al., 2006). From 3 days until 3 weeks after birth, postmitotic GNPs switch off *Math1* expression and start to migrate radially away from the surface past the PCs along glial extensions to finish differentiation in the internal granule layer (IGL) (Hatten, 1999; Machold & Fishell, 2005; Butts *et al.*, 2011).



E9-E10

Figure 5: The rhombic lip at E9. Blue and orange regions highlight the rostral and caudal parts of the RL, respectively. The cerebellar granule neurons derive from the rostral area (blue region). Adapted from (Wang & Zoghbi, 2001)

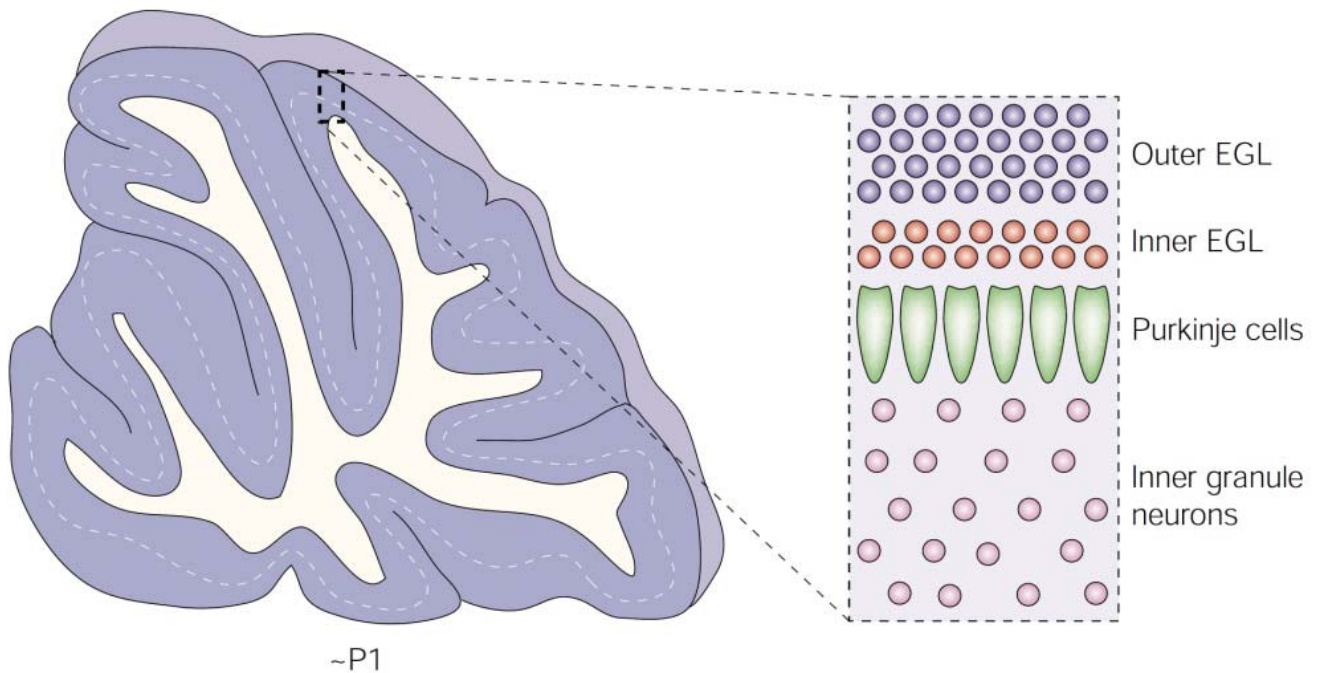


Figure 6: Schematic enlargement from a sagittal section through the cerebellum. Composition of the cerebellar cortex around 1 day before birth is illustrated. Postmitotic granule neuron precursors migrate from the EGL into the IGL. Adapted from (Wang & Zoghbi, 2001).

1.3. Purkinje cell development

1.3.1. Overview

The following studies have been carried out either on rats or mice. To compare the data acquired from both species, it can be assumed that the development of the mouse is generally one day ahead of the development of the rat (<http://embryology.med.unsw.edu.au/OtherEmb/Rat.htm>).

As mentioned above, Purkinje cells are produced in the mouse from E11-E13 in the ventricular zone (Miale & Sidman, 1961; Wang & Zoghbi, 2001). From E16 in the rat, the Purkinje cell marker calbindin is expressed (Armengol & Sotelo, 1991), and from E19 in the mouse, climbing fibers build synapses on Purkinje cells (Morara *et al.*, 2001). Purkinje cell axons have been found to project to cerebellar nuclei at E18 (Eisenman *et al.*, 1991). The Purkinje cell monolayer in the cortex is established

within the first week after birth (Armengol & Sotelo, 1991; Kapfhammer, 2004). Dendritic development starts around birth and is closely paralleled by development of electrophysiological properties (Woodward *et al.*, 1969; Gruol *et al.*, 1992; McKay & Turner, 2005).

P0-P9 (see Fig. 7):

At birth, Purkinje cells are polarized and have one smooth dendrite (Armengol & Sotelo, 1991). During the first 6 days after birth, this simple appearance becomes transiently more complex before dendrites retract and then rapidly regrow again from the soma in an unpolarized manner, a stage termed “stellate cell with disoriented dendrites” (Armengol & Sotelo, 1991). Since Purkinje cells pass the stages at an individual rate, all different stages are present at all times in these first 6 days (Armengol & Sotelo, 1991). Dendritic development at this early stage of dendritic remodeling has been found to be controlled by expression of the transcription factor retinoid-related orphan receptor α (ROR α) (Boukhtouche *et al.*, 2006). *Staggerer* mutant mice are lacking ROR α function in Purkinje cells and suffer from cerebellar atrophy after cell death of Purkinje- and, subsequently, granule cells (Hamilton *et al.*, 1996; Steinmayr *et al.*, 1998). The remaining Purkinje cells fail to proceed in their development beyond this early stage (Shirley & Messer, 2004).

Despite the rapid dendritic rearrangements in the first postnatal week, there is little total dendritic expansion (McKay & Turner, 2005). Cell bodies, on the other hand, grow considerably to reach their adult size by P9 in the rat (McKay & Turner, 2005). Electrophysiologically, Purkinje cells reacted to current injections with an immature pattern of Na⁺ spikes of low amplitude (McKay & Turner, 2005). The big majority of synaptic input comes from climbing fibers, which synapse still perisomatically on Purkinje cells and are not yet confined to a one CF-PC relation with only one CF innervating one Purkinje cell ((Mason *et al.*, 1990), abstract). Activation of mGluR1, and subsequently PLC β and PKC γ , has been shown to be crucial for this process of multiple CF elimination, which starts around P5 in rodents and is finished by 3-4 weeks postnatally (Kano *et al.*, 1995; Kano *et al.*, 1997; Kano *et al.*, 1998).

P9- week 4 after birth (see Fig. 7):

Until P12, Purkinje cell dendrites rearrange again to develop a single primary dendrite (McKay & Turner, 2005). Purkinje cells then enter the phase of rapid

dendritic expansion, with the dendritic trees increasing dramatically in length and complexity. Dendritic growth at this stage is now restricted to the sagittal plane of the cerebellum (Ramon y Cajal, 1911). As granule neurons migrate past the Purkinje cells into the IGL during this time, Parallel Fibers extend and form synapses on Purkinje cells dendrites. Also synapses between Purkinje cells and inhibitory interneurons are formed. Synapses of CFs dislocate from the somata to the proximal dendrites and are restricted to one CF per Purkinje cell ((Mason *et al.*, 1990), abstract). The phase of rapid dendritic growth lasts until 3 weeks after birth in mice and until 4 weeks in rats ((Berry & Bradley, 1976; Weiss & Pysh, 1978), abstracts).

Electrophysiological input resistance decreased remarkably concurrent with the appearance of the prominent stem dendrite (McKay & Turner, 2005), and while the dendritic trees grow, the cells first become able to produce so-called Ca^{2+} - Na^{+} -bursts (a burst of Na^{+} spikes terminated by a Ca^{2+} spike) and later develop a trimodal pattern of intrinsic activity (Womack & Khodakhah, 2004; McKay & Turner, 2005). This trimodal pattern is independent of synaptic input and consists of three modes of spontaneous activity; Purkinje cells fire tonically, then accelerate to burst (Ca^{2+} - Na^{+} -bursts) and then go silent, before they restart the pattern with firing tonically again (Womack & Khodakhah, 2002). The duration of one cycle has been found to vary widely between Purkinje cells of the same age (20 seconds to 20 minutes), but did not change within the same cell (Womack & Khodakhah, 2002). After P18 in rats, only minor changes in electrophysiological output, as well as in dendritic growth were detected (McKay & Turner, 2005).

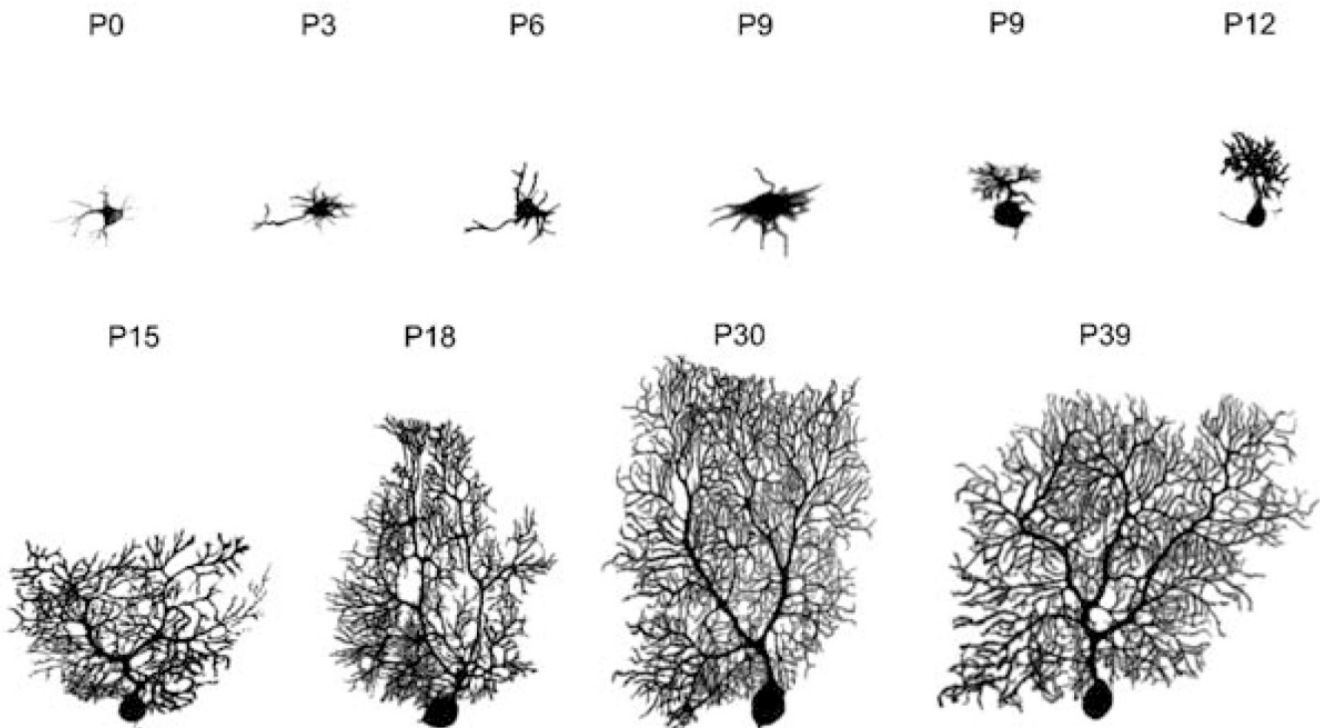


Figure 7: morphological stages of Purkinje cell dendritic development from P0-P39 in rats. Adapted from (McKay & Turner, 2005).

Given these dynamic morphological and electrophysiological processes, it is not surprising that also the expression of ion channels in Purkinje cells is differentially regulated during the first 3 weeks of development. For example, the voltage-independent, small conductance Ca^{2+} -activated K^+ channel type 2 (SK2) is already expressed at P1 in Purkinje cells and is continuously downregulated within 3 weeks (Cingolani *et al.*, 2002). Exclusively in young Purkinje cells, SK2 channels have been found to be very important for the regularity and patterning of spontaneous activity (Cingolani *et al.*, 2002). The expression of large conductance voltage- and Ca^{2+} -activated K^+ (BK) channels, on the other hand, increases during the first 2 weeks postnatally and is high in adult Purkinje cells (Muller *et al.*, 1998).

While in Purkinje cells of 2 weeks and older animals P/Q-type Ca^{2+} channels have been shown to mediate the main Ca^{2+} influx (Regan, 1991; Mintz *et al.*, 1992a), in P4-P7 rats L-type Ca^{2+} channels have been found to be required for spontaneous intracellular Ca^{2+} oscillations, a striking feature of developing Purkinje cells which may play a role in regulation of gene expression (Liljelund *et al.*, 2000). Intriguingly, immunostaining of P/Q-type Ca^{2+} channels on P5 was equally intense as of L-type channels, thus P/Q type channels were also expressed in Purkinje cells, but had a

very small influence on Ca^{2+} oscillations (Liljelund *et al.*, 2000). In a later study by this group, they found that both P/Q- and L-type channels contributed to the Ca^{2+} signal observed after K^{+} -induced membrane depolarization, but only L-type channels were found to lead to activation of the transcription factor CREB and subsequent gene transcription (Gruol *et al.*, 2005).

1.3.2. External aspects influencing Purkinje cell dendritic development

Synaptic input

How much of this late dendritic development in Purkinje cells is dependent on parallel fiber input has been the content of several studies. In dissociated Purkinje cell cultures, dendritic development without neuronal activity such as glutamate treatment, co-culture with granule cells or under TTX treatment was rather poor (Schilling *et al.*, 1991; Baptista *et al.*, 1994; Tanaka *et al.*, 2006). In cerebellar organotypic slice cultures, on the other hand, only slightly smaller dendritic trees were found after inhibition of all glutamate receptors (Adcock *et al.*, 2004). Additionally, extinction of granule cells by X-irradiation (Altman & Anderson, 1972) or absence of granule cell contacts in the cerebellum of weaver mutant mice (Rakic & Sidman, 1973) revealed that although Purkinje cells lacked third and higher order branches and spiny branchlets, an overall normal looking dendritic tree, including spines, developed.

Hormones

Purkinje cells of neonatal rats have been found to produce the sex hormones progesterone and estrogen in especially high levels during the phase of dendritic development, at the same time when they also have been found to express progesterone- and estrogen receptors (Ukena *et al.*, 1999; Jakab *et al.*, 2001; Sakamoto *et al.*, 2003a; Sakamoto *et al.*, 2003b), and both hormones have been shown to increase dendritic growth and spine formation (Sakamoto *et al.*, 2002; Sakamoto *et al.*, 2003a). In addition, Purkinje cells have also been found to respond to the environmental chemicals octylphenol (OP) and bisphenol A (BPA), which act on estrogen receptors (Khurana *et al.*, 2000; Paris *et al.*, 2002). When one week old rats were injected with OP or BPA for 4 days into the cerebrospinal fluid around the cerebellum, the outgrowth of dendrites was promoted (Shikimi *et al.*, 2004).

While treatment with thyroid hormones has been shown to enhance dendritic growth of Purkinje cells (Lindholm *et al.*, 1993), their absence in hypothyroid rats resulted in a significant reduction and delay of dendritic development ((Vincent *et al.*, 1982), abstract). Interestingly, this effect has been found to be caused by adverse actions of thyroid receptors that do not have thyroid hormones bound rather than by the absence of thyroid receptor signal transduction itself (Gothe *et al.*, 1999; Hashimoto *et al.*, 2001).

Neurotrophins

The roles of neurotrophins have been extensively studied in dissociated cultures of Purkinje cells. While NT-3 and NT-4/5 have been shown to increase Purkinje cell survival when applied to the cultures ((Mount *et al.*, 1994b), abstract) (Larkfors *et al.*, 1996), NGF has been found to require direct or indirect activation of mGluRs to mediate cell survival, cell size and neurite development in a p75 neurotrophin receptor mediated manner (Cohen-Cory *et al.*, 1991; Mount *et al.*, 1994a; Mount *et al.*, 1998).

For BDNF, however, results have been controversial, even among studies using dissociated Purkinje cell cultures. BDNF did not influence Purkinje cell survival in a study by Mount and colleagues ((Mount *et al.*, 1994b), abstract), while it did so in another study (Larkfors *et al.*, 1996). In Purkinje-granule cell co-cultures, Shimada and colleagues found an influence of BDNF on Purkinje cell spine development, but not on dendritic complexity (Shimada *et al.*, 1998), whereas Hirai and Launey found that inhibition of the BDNF receptor Trk-B reduced Purkinje cell dendritic growth, most likely an indirect effect mediated by less promoted granule cell survival (Hirai & Launey, 2000). It was suggested that the culture medium used might have an impact on the expression of Trk receptors in the culture, thus influencing the need for BDNF (Hirai & Launey, 2000).

In BDNF^{-/-} mice, Purkinje cell dendrites have been shown to be severely reduced during the first three weeks after birth (Schwartz *et al.*, 1997), but at P24 they did not look very different from *wt* mice (Carter *et al.*, 2002). However, BDNF^{-/-} mice were reported to have general growth retardation and reduced weight (Ernfors *et al.*, 1994; Korte *et al.*, 1995). In cerebellar slice cultures from BDNF-deficient mice, no differences in several Purkinje cell dendritic growth parameters were found compared to *wt* mice (Adcock *et al.*, 2004). Furthermore, Purkinje cell dendrites had a normal

appearance in mice lacking a functional BDNF receptor, TrkB (Rico *et al.*, 2002). In summary, these results certainly reflect the *in vivo* situation closer than the dissociated culture system and thus support the concept that BDNF does not play a significant role in Purkinje cell dendritic development. Instead, TrkB signaling, and thus BDNF and NT-4, have been found to be essential to the development and activity-dependent synaptogenesis of inhibitory interneurons in the cerebellum (Rico *et al.*, 2002; Seil, 2003; Spatkowski & Schilling, 2003) .

Bergmann glia

Purkinje cell dendritic processes have been shown to grow preferentially in the vertical direction (up towards the pia) when they are in close contact with fibers of the cerebellar astrocytes called Bergmann glia (Lordkipanidze & Dunaevsky, 2005). Thus, the radial processes of the Bergmann glia supposedly build a substrate which helps shape the Purkinje cell dendritic tree, although the dendritic growth rate was unchanged by these interactions (Lordkipanidze & Dunaevsky, 2005).

Interactions with Bergmann glia have also been found to be important in proceeding Purkinje cell morphology from multiple disoriented dendrites to a single primary dendrite with directed growth in cerebellar slice cultures (Tanaka *et al.*, 2003). Signal transduction of the receptor protein tyrosine phosphatase zeta (PTP ζ) after binding of one of its ligands, the heparin-binding growth factor Pleiotrophin (PTN), has been shown to be required for this process, as well as activity of the glutamate transporter GLAST (Tanaka *et al.*, 2003). PTP ζ is a chondroitin sulfate proteoglycan, which is expressed by Bergmann glia until adulthood and by Purkinje cells during the time of dendritic development (Maeda *et al.*, 1992; Canoll *et al.*, 1993; Tanaka *et al.*, 2003). GLAST is expressed on the Bergmann glia lamellate processes which have been found to closely surround differentiating Purkinje cell dendritic trees (Yamada *et al.*, 2000). Double immunostaining for PTP ζ and GLAST revealed an overlap of the two proteins and treatments affecting PTN-PTP ζ signaling reduced GLAST immunoreactivity, suggesting that GLAST activity or expression is regulated downstream of PTN-PTP ζ signaling (Tanaka *et al.*, 2003). Interestingly, comparable morphological defects have been observed in Purkinje cells of the mutant mouse ducky, where a defect in the $\alpha 2\delta$ -2 protein, a subunit of voltage-gated Ca²⁺ channels strongly expressed in Purkinje cells, leads to absence epilepsy and ataxia (Barclay *et*

al., 2001; Brodbeck *et al.*, 2002). Thus, there might be a connection between PTN-PTP ζ signalling, GLAST and voltage-gated Ca²⁺ channels.

Moreover, mice lacking GLAST function had defects in motor coordination, persistent Purkinje cell multiple climbing fiber innervation, and were more susceptible to brain injury (Watase *et al.*, 1998), thus underlining the importance of Bergmann glia for correct synaptogenesis as well as limiting excitotoxicity after cerebellar damage.

1.3.3. *The role of PKC in Purkinje cell dendritic development*

PKC γ is the PKC isoform which is in the cerebellum specifically expressed in Purkinje cells, where it is also the most abundant isoform (Kose *et al.*, 1988; Barmack *et al.*, 2000). Like PKC α and β , it belongs to the “classical” subgroup of PKC molecules, which are characterized by being activated in the presence of phosphatidylserine by diacylglycerol (DAG) and Ca²⁺ (for review see (Saito & Shirai, 2002)). Besides PKC γ , PKC α is also expressed in Purkinje cells, as well as in all cells in the cerebellum (Huang *et al.*, 1988). Expression of PKC γ is developmentally regulated, with low levels around birth and subsequent upregulation until 2-3 weeks postnatally (Hashimoto *et al.*, 1988; Yoshida *et al.*, 1988). PKC γ plays a major role in cerebellar Long term depression (for review see (Ito, 2001)). Missense mutations in PKC γ have been found in patients suffering from spinocerebellar ataxia type 14 (SCA14), an autosomal dominant disease which comes along with degenerated Purkinje cells (Chen *et al.*, 2003). Moreover, PKC γ is an important regulator of Purkinje cell dendritic growth under natural conditions; chronic pharmacological inhibition of PKC in rat cerebellar slice cultures during the time of rapid dendritic expansion lead to excessive growth and branching (Metzger & Kapfhammer, 2000). These findings were confirmed in PKC γ deficient mice, although a decreased PKC γ activity in mice resulted in increased branching rather than in larger dendritic trees of Purkinje cells (Schrenk *et al.*, 2002). Chronic pharmacological activation of PKC, on the other hand, limited dendritic growth in slice cultures from *wild type* rats and mice, probably by retraction of dendrites (Metzger & Kapfhammer, 2000; Schrenk *et al.*, 2002). It was excluded that the negative effect of PKC activation on dendritic growth was due to downregulation of the enzyme (Metzger & Kapfhammer, 2000). Interestingly, PKC γ deficient mice still reacted to pharmacological PKC activation with limited dendritic growth, while mice lacking the PKC α isoform were protected (Gundlfinger *et al.*,

2003). In conclusion, PKC γ naturally regulates dendritic growth in Purkinje cells, while a reserve pool of PKC α is activated or upregulated upon stronger stimulation (Schrenk *et al.*, 2002; Gundlfinger *et al.*, 2003).

1.3.4. The role of mGluR1 in Purkinje cell dendritic development

The 8 genes of metabotropic glutamate receptors (mGluR1-8) identified so far have been put into 3 groups according to their signal transduction pathways, sequence similarities and pharmacological features. mGluR1 and mGluR5 both couple to phospholipase C, thus inducing phosphoinositide hydrolysis upon activation and constitute group I of the mGluRs (for reviews see (Nakanishi, 1992; Conn & Pin, 1997; Knopfel & Grandes, 2002)). The most abundant mGluR subtype in Purkinje cells is mGluR1 (Lein *et al.*, 2007), with lower levels of mGluR7 found (Kinzie *et al.*, 1995). mGluR1 is expressed in Purkinje cells already before birth (Masu *et al.*, 1991; Lopez-Bendito *et al.*, 2001) and during the period of rapid dendritic expansion it spreads into the dendrites to take its final localization in spines, perisynaptically of excitatory synapses (Nusser *et al.*, 1994; Lopez-Bendito *et al.*, 2001). mGluR1a is the mGluR1 splice variant with the strongest expression in the cerebellum, with increasing expression levels during development until adulthood (Casabona *et al.*, 1997; Lopez-Bendito *et al.*, 2001). mGluR1 activation occurs after repeated discharges by either climbing fibers or parallel fibers, the two excitatory synaptic inputs to Purkinje cells (Batchelor & Garthwaite, 1993; Batchelor *et al.*, 1994; Dzubay & Otis, 2002). Two responses have been found to be initiated:

- An increase in IP₃ due to PLC activation leads to Ca²⁺ release from the endoplasmic reticulum (ER) in dendrites and spines (Finch & Augustine, 1998; Takechi *et al.*, 1998; Okubo *et al.*, 2004), followed by activation of PKC. This pathway has been shown to be crucial for elimination of multiple climbing fiber innervation and establishment of LTD (Kano *et al.*, 1995; Kano *et al.*, 1997; Offermanns *et al.*, 1997; De Zeeuw *et al.*, 1998; Kano *et al.*, 1998; Ichise *et al.*, 2000; Ito, 2001).
- A slow excitatory postsynaptic potential (sEPSP) is induced (Batchelor & Garthwaite, 1993; Batchelor *et al.*, 1994), which consists of a considerable Na⁺ - and, to a lesser extent, Ca²⁺ influx (Knopfel *et al.*, 2000; Tempia *et al.*, 2001).

The nonselective, Ca²⁺ permeable TRPC3 channel has been shown to mediate the sEPSC (Hartmann *et al.*, 2008) and its activation depends on tyrosine phosphatase activity (Canepari & Ogden, 2003).

Both mGluR1-induced responses are mediated by the G-protein Gq (Hartmann 2004), but diverge upstream of PLC, since PLC activity is not required for sEPSC induction (Hirono *et al.*, 1998; Canepari *et al.*, 2001; Canepari & Ogden, 2003).

Mutant mice lacking mGluR1 function suffer from an ataxic gait, intention tremor, impaired LTD and associative learning (Aiba *et al.*, 1994). In addition, elimination of multiple climbing fiber innervation is incomplete (Kano *et al.*, 1997). All cerebellar phenotypic symptoms were rescued by transgenic induction of the mGluR1a splice variant in Purkinje cells of mGluR1 k.o. mice (Ichise *et al.*, 2000; Kishimoto *et al.*, 2002). No significant aberrations in cerebellar anatomy, including Purkinje cell morphology, have been observed in mGluR1 k.o. mice, however (Aiba *et al.*, 1994; Ichise *et al.*, 2000). Inhibition of mGluR1 in mixed rat Purkinje cell/CGN cultures and *in vivo* as early as P3-9, on the other hand, has been shown to adversely affect Purkinje cell morphology and survival (Catania *et al.*, 2001). Pharmacological mGluR1 inhibition over 12 days in cerebellar slice cultures at a later time in development (starting at P8 in mice) had very little effect on Purkinje cell morphology (Adcock *et al.*, 2004). On the contrary, treatment of cerebellar slice cultures during the time of rapid dendritic expansion with the class I mGluR activator DHPG resulted in severely reduced Purkinje cell dendritic tree sizes and numbers of branch points in a concentration dependent manner (Sirzen-Zelenskaya *et al.*, 2006). DHPG co-treatments with inhibitors of either PKC, PLC, mGluR5 or glutamatergic synaptic transmission (TTX combined with inhibitors of AMPA and NMDA receptors) were ineffective in modulating the inhibitory effect of DHPG, thus indicating that a direct mGluR1 receptor activation is responsible and that the inhibitory effect is mediated independent of bioelectrical activity and the classical pathway involving PKC and PLC.

1.4. The cerebellar slice culture model

In dissociated cell cultures, cells of one type - or a small selection of co-cultured cell types - are studied in isolation. In contrast, in organotypic slice cultures, a thick slice of the whole tissue is kept in culture, where the natural microenvironment of a cell with neighboring cell-cell interactions and, as in the case of nervous tissue, local neuronal networks are preserved ((Dupont *et al.*, 2006), abstract). This provides the possibility for various kinds of studies, like cell development, synaptogenesis, regeneration or electrophysiology.

Slice cultures from nervous tissue, however, can only be obtained from immature tissue, as survival of neurons critically depends on their stage of development. Purkinje cells, for example, are extremely sensitive to axotomy at P1-P5 in rats (P4-P6 in mice) and do not survive in slice cultures prepared during this period (Dusart *et al.*, 1997). In general, slice cultures can be taken from developing nervous tissue as long as there is no myelination (Kapfhammer, 2010).

Some special measures have to be taken to ensure a thorough supply with nutrients and oxygen to all cells in the typically 0.3 – 0.5 mm thick cultures. In the technically simplest method, the static slice culture, which allows cerebellar slices to be cultured for several weeks, the slices are placed on top of a membrane which is permeable to all ingredients of the culture medium (Yamamoto *et al.*, 1989; Stoppini *et al.*, 1991). The membrane is held by a culture insert and placed over the medium, which is drawn through the membrane into the slice culture by capillary forces. Thus, the top of the culture is at the interface between culture medium and air, which allows both nutrients and oxygen to reach the whole culture (see Fig. 8).

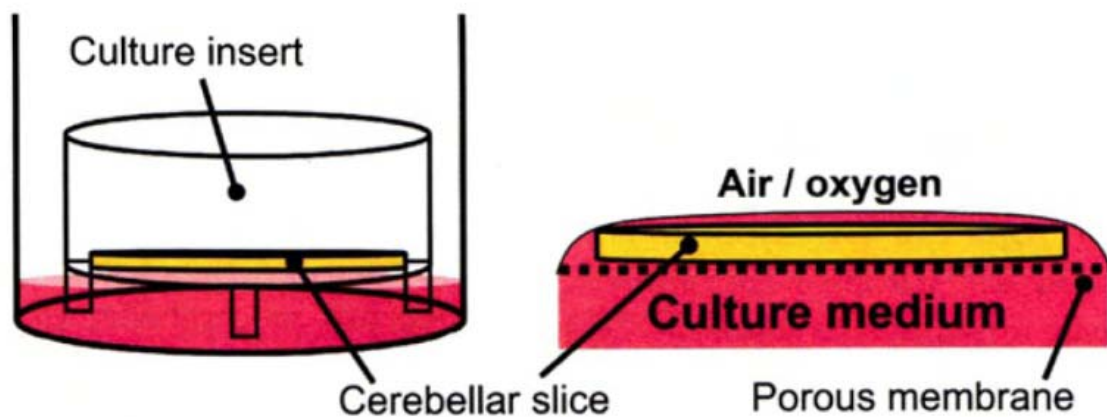


Figure 8: Illustration of the experimental device for static slice culture. The bottom of the culture insert is a porous membrane which is permeable to the culture medium below. The slice culture is lying on this membrane at the interface between culture medium and air. By capillary forces, the medium is drawn through the membrane into the tissue slice. Picture from (Kapfhammer, 2010).

Obtaining slice cultures from mouse cerebella is possible until 2 weeks postnatal. They are usually cut in the sagittal plane, since this is the natural orientation of the Purkinje cell dendritic tree, therefore allowing morphological analyses of Purkinje cells. Besides Purkinje cells, granule neurons, Golgi cells, basket cells, stellate cells, unipolar brush cells, astrocytes and Bergmann glia, as well as an extracellular matrix have been found in slice cultures (Jaeger *et al.*, 1988; Nunzi & Mugnaini, 2000; Bruckner & Grosche, 2001). Since the axons of granule cells run at a right angle to the Purkinje cell dendritic trees, they are being cut during the culturing process. Due to their immaturity in the early days after birth, they regenerate and reorganize to form synapses on Purkinje cells (Tanaka *et al.*, 1994). In the median zone of the cerebellum Purkinje cell axons are cut as well, and although the cells survive the axotomy well after P6 in the mice, they do not regrow (Dusart *et al.*, 1997). In more lateral areas of the cerebellum, the deep cerebellar nuclei are part of the slice culture, and Purkinje cells project to their original target, the deep cerebellar neurons (Mouginot & Gahwiler, 1995). Climbing and Mossy fibers, which originate from outside of the cerebellum, are absent. During the culture period, the slices become thinner and although the cytological architecture of the cerebellar cortex is generally preserved, Purkinje cells are not strictly arranged in a monolayer anymore. The slice culture model is also well suitable for electrophysiological studies, as electrical

activity develops spontaneously in the cultures (Leiman & Seil, 1973; Kapoor *et al.*, 1988; Mougnot & Gahwiler, 1995; Dupont *et al.*, 2006).

After the culture period, the slices can be fixed and immunostained for various targets. To study Purkinje cells, an antibody against calbindin D-28K, which stains no other cells in the cerebellum (Celio, 1990), is conveniently used to visualize whole Purkinje cells including dendrites and axons. Granule cell survival can be simultaneously shown with NeuN-staining as a control (Weyer & Schilling, 2003).

Slices taken from P7-P9 mice and kept in culture for approximately 10 days provide good survival of Purkinje and granule cells and cover the period of rapid Purkinje cell dendritic growth. All things considered, the cerebellar slice culture model is a very suitable system to study Purkinje cell morphological development.

2. Specific Aims of the thesis

2.1. The role of Purkinje cell dendritic tree size in excitotoxicity

The aim of this project was to test the hypothesis that it is the extraordinary size of their dendritic tree, with an accordingly large amount of AMPA receptors, which makes Purkinje cells especially susceptible to excitotoxic death.

Chronic activation of either mGluR1 or PKC in organotypic slice cultures strongly inhibits Purkinje cell dendritic growth (Metzger & Kapfhammer, 2000; Schrenk *et al.*, 2002; Sirzen-Zelenskaya *et al.*, 2006). The question arises what the physiological relevance of this effect might be. One possibility is that growth of the Purkinje cell dendritic tree is inhibited as soon as a certain amount of excitatory input reaches the cell, thus creating a feedback mechanism which protects the cell from excitatory overload. In fact, Purkinje cells appear to be highly sensitive to excitotoxic death (Garthwaite & Garthwaite, 1984). In several models of traumatic brain injury (TBI), delayed Purkinje cell loss has been found even if the injury was inflicted to other parts of the brain than the cerebellum (Mauter *et al.*, 1996; Park *et al.*, 2006). Evidently, this Purkinje cell loss does not occur by mechanical damage or haemorrhage, but involves excitotoxic mechanisms (Ai & Baker, 2002; Park *et al.*, 2006; Igarashi *et al.*, 2007).

Treatment with (RS)-a-Amino-3-hydroxy-5-methyl-4-isoxazolepropionic acid (AMPA) was used to induce excitotoxic neurodegeneration in cerebellar slice cultures. AMPA-exposure does not affect cerebellar granule cells, but kills Purkinje cells already in low doses by a mechanism called dark-cell degeneration, which includes aspects of both, apoptosis and necrosis (Garthwaite & Garthwaite, 1990; Meldrum & Garthwaite, 1990; Garthwaite & Garthwaite, 1991b; Leist & Jaattela, 2001; Strahlendorf *et al.*, 2003). All things considered, AMPA-treatment is a suitable model to study sensitivity of Purkinje cells to excitotoxicity.

Purkinje cells with either normal or reduced dendritic tree sizes after chronic activation of mGluR1 or PKC were exposed to AMPA, and cell survival was compared.

Purkinje cells with a reduced dendritic tree were equally sensitive to AMPA treatment as were control cells with a norm-sized dendritic tree. Thus, our results indicate that the expansion of the dendritic tree is no major determinant for Purkinje cell susceptibility to excitatory overload.

2.2. Analysis of the signaling pathways mediating dendritic growth inhibition after chronic mGluR1 or PKC activation

The aim of this project was to identify components of the signaling pathways which lead to dendritic growth inhibition in Purkinje cells after chronic mGluR1 or PKC activation.

Activation of either mGluR1 or PKC result in a severely reduced Purkinje cell dendritic tree, but intriguingly, the pathways mediating this effect appear to be independent of each other (Metzger & Kapfhammer, 2000; Schrenk *et al.*, 2002; Sirzen-Zelenskaya *et al.*, 2006) and remain, in both cases, largely unknown.

A similar reduction of Purkinje cell dendritic trees has been found in two independent mouse models, the *moonwalker* (*mwk*) and the *lurcher* mouse (Becker *et al.*, 2009; Zanjani *et al.*, 2009). In both mouse models, abnormal activity of a nonselective, Ca²⁺ permeable cation channel has been found. In *lurcher* mice, a mutation of the $\delta 2$ glutamate receptor (GluR $\delta 2$) results in a constitutively open cation channel and leads to death of Purkinje cells and stunted dendrites in surviving ones (Zuo *et al.*, 1997; Zanjani *et al.*, 2009). In *mwk* mice, a gain-of-function mutation in the *Trpc3* gene

results in facilitated TRPC3 channel opening and leads to reduced Purkinje cell dendritic trees and an ataxic phenotype (Becker *et al.*, 2009).

Based on these findings, we focused our research on Ca^{2+} permeable cation channels in the Purkinje cell membrane, especially on TRPC3, P/Q-type and T-type Ca^{2+} channels, as they are all strongly expressed in Purkinje cells and can be associated with mGluR1. TRPC3 channels have been shown to mediate the sEPSC occurring after mGluR1 stimulation (Hartmann *et al.*, 2008). P/Q-type and T-type Ca^{2+} channels account for up to 95% of all Ca^{2+} currents in Purkinje cells (Usowicz *et al.*, 1992; Watanabe *et al.*, 1998; Swensen & Bean, 2003; Isope & Murphy, 2005; Isope *et al.*, 2010). P/Q-type channels have been found to directly interact with mGluR1 in Purkinje cell spines (Kitano *et al.*, 2003), and T-type channels of the Cav3.1 isoform have been shown to co-localize with mGluR1 in Purkinje cell spines and to be potentiated upon mGluR1 activation (Hildebrand *et al.*, 2009).

We have treated cerebellar slice cultures of P8 mice with several pharmacological inhibitors of Ca^{2+} channels simultaneously with activators of mGluR1 or PKC. Treatment with inhibitors of R-type Ca^{2+} channels, GluR δ or TRPC3 channels did not reveal an involvement of these channels. Concerning the TRPC3 channels, the results were in line with experiments using mice lacking TRPC3 function. Inhibitors of T-type or P/Q-type Ca^{2+} channels partially rescued the dendrite reducing effect seen after chronic PKC activation. The combination of T-type and P/Q-type Ca^{2+} channel inhibitors conferred a partial rescue from the dendrite reducing effect seen after chronic mGluR1 activation.

Our results strongly suggest an involvement of T-type and P/Q-type Ca^{2+} channels in mediating dendritic growth inhibition after both, mGluR1 and PKC activation.

3. Dendritic tree size in excitotoxicity

Reduced size of the dendritic tree does not protect Purkinje cells from excitotoxic death

Olivia S. Gugger and Josef P. Kapfhammer

Anatomical Institute, Department of Biomedicine Basel, University of Basel, Pestalozzistr. 20, CH - 4056 Basel, Switzerland.

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3.1. Abstract

Purkinje cell loss by excitotoxic damage is a typical finding in many cerebellar diseases. One important aspect of this high sensitivity of Purkinje cells to excitotoxic death might be the enormous size of their dendritic tree with a high load of excitatory glutamate receptors. We have studied whether the reduction of the size of the dendritic tree might confer resistance against excitotoxic death to Purkinje cells.

We have grown Purkinje cells in organotypic cerebellar slice cultures under chronic activation of metabotropic glutamate receptors or of Protein kinase C. Both treatments strongly reduced dendritic tree size. After this treatment, cells were exposed to the glutamate receptor agonist AMPA which has a strong excitotoxic effect on Purkinje cells. We found that Purkinje cells with small dendritic trees were equally sensitive to AMPA exposure as untreated control cells with large dendritic trees. Immunostaining against vesicular glutamate transporter 1 revealed that the small dendritic trees were densely covered by glutamatergic terminals.

Our results indicate that the expansion of the dendritic tree and the total number of AMPA receptors per neuron do not play a major role in determining the susceptibility of Purkinje cells to excitotoxic death.

3.2. Introduction

The loss of Purkinje cells (PCs) is a hallmark of many cerebellar diseases causing functional impairment of motor performance. An important factor for the involvement of Purkinje cells in cerebellar disease is their high sensitivity to excitotoxic death (Garthwaite & Garthwaite, 1984). In addition, Purkinje cells are often affected in traumatic brain injuries (TBI). Loss of motor control and coordination, symptoms typically associated with cerebellar dysfunction, play an important role within the pathophysiology of TBI. It has been shown that Purkinje cell loss is a common finding in TBI in experimental animals, even if the cerebellum is remote from the site of impact, e.g. a focal injury in the forebrain (Mautes *et al.*, 1996; Park *et al.*, 2006). Delayed neurodegeneration in the cerebellum is a generalized response to TBI and the specific local pattern of cell loss (with a special vulnerability of the vermis region

of the cerebellum) suggests involvement of mechanisms other than mechanical damage and haemorrhage (Park *et al.*, 2006; Igarashi *et al.*, 2007). There is evidence that the PC loss in TBI does occur by excitotoxic mechanisms (Ai & Baker, 2002; Park *et al.*, 2006). In experimental animals as well as in the cerebrospinal fluid of patients with TBI, glutamate levels were significantly elevated after injury (Baker *et al.*, 1993; Palmer *et al.*, 1993). Purkinje cell degeneration induced by (RS)- α -Amino-3-hydroxy-5-methyl-4-isoxazolepropionic acid (AMPA) is an appropriate model system to study excitotoxic neurodegeneration. The cytological features of Purkinje cell degeneration after AMPA exposure are characterized by a process called dark-cell degeneration (Meldrum & Garthwaite, 1990; Garthwaite & Garthwaite, 1991a). This is a form of programmed cell death with features from both, apoptosis and necrosis (Leist & Jaattela, 2001; Strahlendorf *et al.*, 2003). Degenerating Purkinje cells become positive for TUNEL staining and caspase 3 immunoreactivity (Strahlendorf *et al.*, 1999; Strahlendorf *et al.*, 2003). Since AMPA-exposure kills Purkinje cells but leaves cerebellar granule cells unharmed (Garthwaite & Garthwaite, 1990; Garthwaite & Garthwaite, 1991a), it is well suited to investigate susceptibility of Purkinje cells to excitotoxic death.

We have previously shown that chronic activation of metabotropic glutamate receptor 1 (mGluR1) (Sirzen-Zelenskaya *et al.*, 2006) or Protein kinase C (PKC) (Metzger & Kapfhammer, 2000; Schrenk *et al.*, 2002) in organotypic cerebellar slice cultures of postnatal mice severely inhibits the growth and development of the Purkinje cell dendritic tree. A potential physiological function of growth inhibition by mGluR activation might be to limit the size of the Purkinje cell dendritic tree in a kind of feedback mechanism which would limit the number of excitatory inputs and the total AMPA receptor load to make the cell less sensitive to excitotoxic death (depicted in Fig. 9 A). In this study, we have tested this hypothesis and have grown Purkinje cells in organotypic slice cultures under conditions which result in a severe reduction of dendritic tree size. We have then analyzed the effects of AMPA exposure on Purkinje cells with different dendritic tree sizes. Our results indicate that a reduction of the size of the dendritic tree in Purkinje cells does not offer any protection from glutamate-induced excitotoxicity.

3.3. Materials and Methods

3.3.1. Organotypic slice cultures

Animal experiments were carried out in accordance with the European Communities Council Directive of 24 November 1986 (86/609/EEC) and were reviewed and permitted by Swiss authorities. Cultures were prepared from B6CF1 mice as described previously (Adcock *et al.*, 2004; Kapfhammer, 2005). Briefly, mice were decapitated at postnatal day 8 (P8), their brains were aseptically removed and the cerebellum was dissected in ice-cold preparation medium (minimal essential medium (MEM), 1% glutamax (Gibco, Invitrogen), pH 7.3). Sagittal sections 350 μm thick were cut on a McIlwain tissue chopper under aseptic conditions. Slices were separated, transferred onto permeable membranes (Millicell-CM, Millipore) and incubated on a layer of incubation medium (50% MEM, 25% Basal Medium Eagle, 25% horse serum, 1% glutamax, 0.65% glucose) in a humidified atmosphere with 5% CO_2 at 37°C. The medium was changed every 2-3 days for a total of 13-14 days. Pharmacological compounds were added to the medium at each change, according to the treatment schedule (see Fig. 1 B): (RS)-3,5-Dihydroxyphenylglycine (DHPG, Tocris) or Phorbol 12-myristate 13-acetate (PMA, Tocris) were added starting at 3-4 days in vitro (DIV) and washed out at DIV 10-11 (after 7 days of treatment), (RS)- α -Amino-3-hydroxy-5-methyl-4-isoxazolepropionic acid (AMPA, Tocris) was added 24 hours after end of the pre-treatment (at DIV 11-12) for 4 hours. The following concentrations were used: 15 μM DHPG, 100 nM PMA, 30 μM AMPA. Slices were kept in culture for two additional days before fixation and immunohistochemical staining.

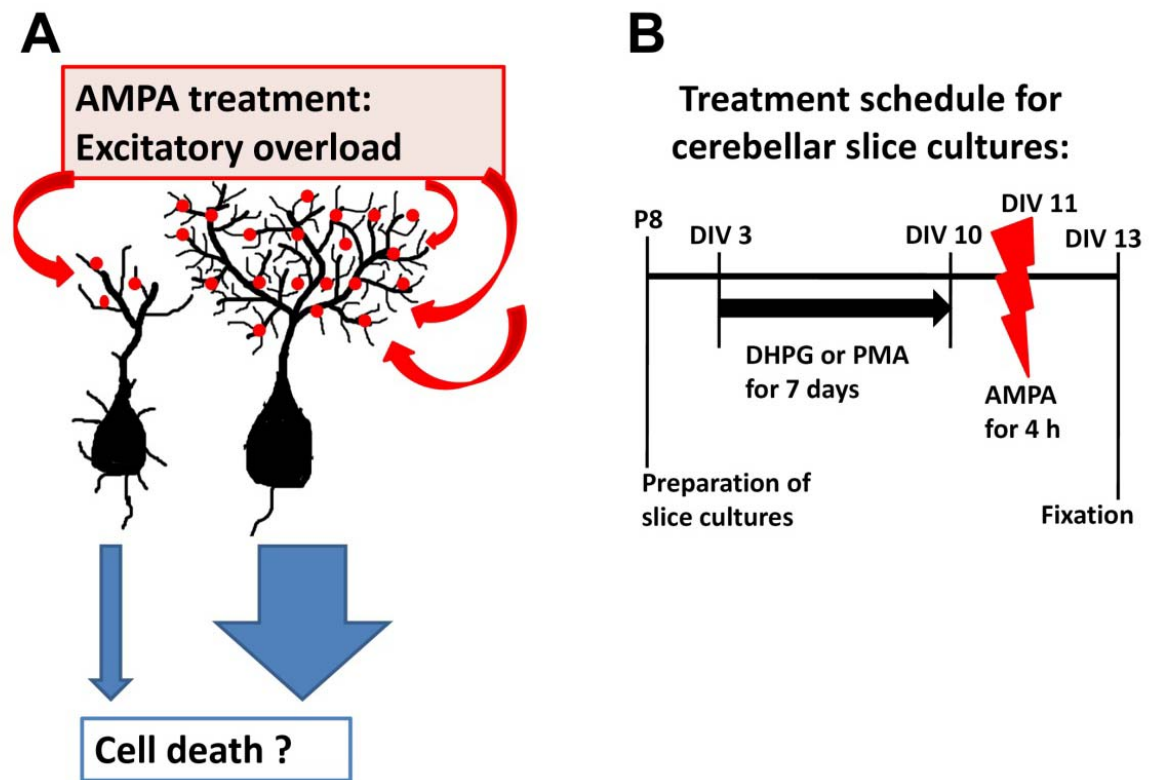


Figure 9: Schematic drawings of starting hypothesis and treatment schedule.

A) Starting hypothesis of this study: On an extensive dendritic tree, a greater number of glutamatergic synapses provide more entry points for excitatory input. This could make a large Purkinje cell more susceptible to excitatory overload through AMPA compared to a cell with a reduced dendritic tree.

B) Treatment schedule for cerebellar slice cultures. At postnatal day 8 (P8), slice cultures were prepared. The pharmacological treatments with either DHPG or PMA were started at day in vitro 3 (DIV 3) and stopped at DIV10. After 24 hours of recovery time, the cultures were challenged with AMPA for 4 hours and kept in culture for two more days without treatment.

3.3.2. Immunohistochemistry

At DIV 13-14 cultures were fixed in 4% paraformaldehyde overnight at 4°C. All reagents were diluted in 100 mM phosphate buffer (PB), pH 7.3. Slices were incubated in blocking solution (0.5% Triton X-100, 3% normal goat serum) for a minimum of 30 minutes in order to permeabilize the tissue and block non-specific antigen binding. Two different primary antibodies were simultaneously added to the slices in fresh blocking solution and incubated overnight at 4°C. After washing in PB, secondary antibodies were added to the slices in PB containing 0.1% Triton X-100 in order to prevent non-specific antigen binding for at least 1 hour at room temperature.

For the analysis of Purkinje and granule cell survival and Purkinje cell dendritic size, rabbit anti-Calbindin D-28K (Swant, 1:1000) and monoclonal anti-NeuN (Chemicon, Millipore, 1:500) were used as primary antibodies and goat anti-rabbit Alexa 546 (Molecular Probes, Invitrogen, 1:500) and goat anti-mouse Alexa 488 (Molecular Probes, Invitrogen, 1:500) were used as secondary antibodies to visualize Purkinje cells and cerebellar granule cells, respectively. For the synaptic analysis, mouse anti-Calbindin D-28K (Swant, 1:1000) and rabbit anti-vGluT1 (Synaptic Systems, 1:3000) were used as primary antibodies and goat anti-mouse Alexa 546 (1:500) and goat anti-rabbit Alexa 488 (1:500) were used as secondary antibodies to visualize Purkinje cells and glutamatergic synapses, respectively. Stained slices were mounted on cover slips with Mowiol. Cultures were viewed on an Olympus AX-70 microscope equipped with a Spot digital camera. Recorded images were adjusted for brightness and contrast with Photoshop image processing software. For Fig. 2, multiple images of cerebellar slice cultures photographed with the 10x lens were stitched together to show the entire slice using the Photoshop panorama function.

3.3.3. Quantitative analysis of Purkinje cell survival and dendritic tree size

The quantification of the effects of the pharmacological treatments on Purkinje cell dendritic tree size was done as previously described (Adcock *et al.*, 2004; Sirzen-Zelenskaya *et al.*, 2006). Purkinje cells which had an axon and dendritic tree which could be isolated from its surroundings were selected for analysis. As described earlier (Adcock *et al.*, 2004), the measurement of dendritic tree size was only possible in Purkinje cells in which the dendritic field did not overlap with the one of neighbouring cells. Cells were viewed on a with a 20x lens and photographed with a digital camera (Coolsnap-Pro, Photometrics, Canada). An image analysis program (Image Pro Plus) was used to trace the outline of the Purkinje cell dendritic trees yielding the area covered by the dendritic tree. Cells were acquired from four independent experiments with a minimum number of 20 cells per experiment. The total number of analyzed individual cells ranged from 101 for control to 134 for DHPG treatment. The data were analyzed using GraphPad Prism software. The mean value of the dendritic tree area of untreated control cells was set to 100% and the results

were expressed as percentage of controls +/- standard deviation with 95% confidence intervals. The statistical significance of differences in parameters was assessed by non-parametric analysis of variance (Kruskal-Wallis test) followed by Dunn's post test. For comparisons of single data columns, Mann-Whitney's non-parametric test was used.

For the evaluation of Purkinje cell survival, we used a semi-quantitative rating similar to a one described earlier (Dusart *et al.*, 1997; Schrenk *et al.*, 2002). Briefly, five classes of slices were defined according to the number and arrangement of Calbindin D-28K stained Purkinje cells. Class I included slices with very few dispersed cells, i.e. with no compact group containing more than 20 cells and a total Purkinje cell number < 50. Class II included slices with at least one cluster with more than 20 cells or a total Purkinje cell number >50. In the third class (III), slices with 4 or more clusters of over 20 cells or with one intact folium were included. In class IV, slices with 2 or more intact folia but at least two missing or incomplete folia were included. Class V contained all slices with not more than one missing or incomplete folium. The total number of analyzed slices ranged from 45 for DHPG treatment to 72 for PMA treatment and slices for each treatment came from at least 4 independent experiments. In order to compare results from different treatment groups, the arithmetic mean of the survival class values for all slices from one group was calculated and named the "survival score" for this treatment group. A possible protective effect on Purkinje cell survival would be detected with high sensitivity because additional survival of only few Purkinje cells in a slice with no Purkinje cell survival (as typically seen with AMPA treatment) would lead to classification in class II instead of class I.

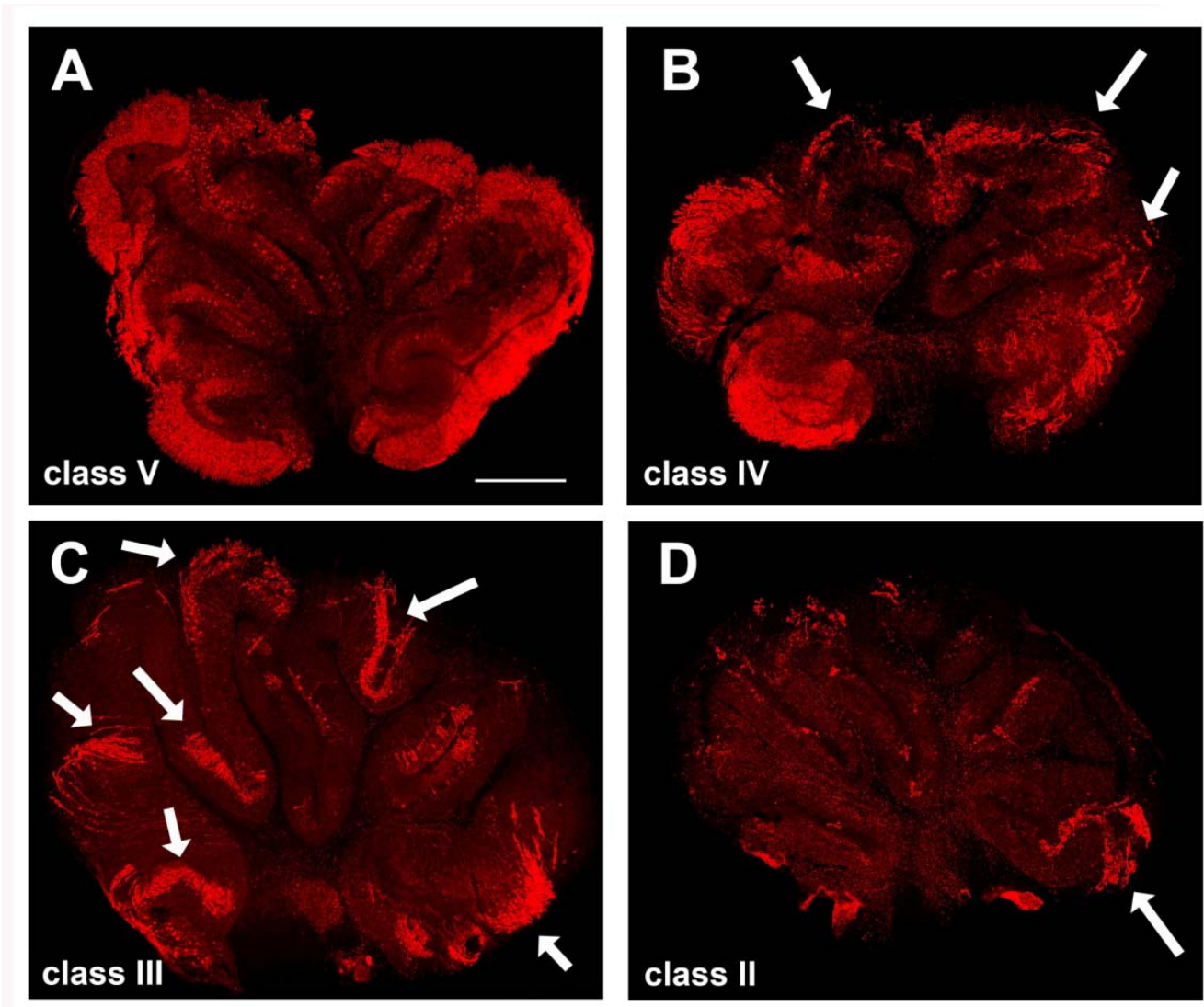


Figure 10: Representative slices for survival classes V (A), IV (B), III (C) and II (D). Calbindin D-28K staining shows Purkinje cells.

A) Class V, virtually complete survival of Purkinje cells.

B) Class IV, Purkinje cell growth is thinned out in 3 folia (arrows).

C) Class III, moderate Purkinje cell survival, more than 4 clusters with compact growth (arrows).

D) Class II, few dispersed surviving Purkinje cells, only one cluster with more than 20 Purkinje cells (arrow).

Scale bar 500 μ m.

3.4. Results

3.4.1. AMPA kills Purkinje cells in a dose dependent manner

In order to evaluate cell survival, we used a semi-quantitative rating of Purkinje cell survival in the slice cultures similar to the one described in (Dusart *et al.*, 1997). Five groups of slices were defined according to the number and arrangement of Calbindin D-28K stained Purkinje cells. Class V would represent slices with virtually complete survival of Purkinje cells throughout the slice and class I would represent almost complete death of Purkinje cells (details in Methods). Figure 10 shows representative examples of three of the five groups. For the different treatment conditions, a mean survival score was calculated which represents the mean of the survival scores of a particular treatment group. A mean survival score of 5 would mean that 100% of the slices were in class V and a score of 1 that all slices were in class I.

In order to assess the neurotoxic potential of AMPA treatment for Purkinje cells, slices grown under control conditions for 11 days (see treatment schedule in Methods, Fig. 9 B) were treated with different concentrations of AMPA for 4 hrs. After the treatment, the slices were kept in culture for 2 additional days. Figure 11 A shows the survival scores of Purkinje cells exposed to different AMPA concentrations. Without AMPA treatment the mean survival score was 4.69, i.e. most slices belonged to class V, demonstrating good survival of Purkinje cells under control conditions. Survival of Purkinje cells in slices treated with 2 μ M AMPA for 4 h was similar to control slices with a mean survival score of 4.66. After treatment with 5 μ M AMPA, the slice cultures displayed several regions with remarkable cell loss and the mean survival score dropped to 3.88. After 10 μ M AMPA there was a considerable cell loss throughout the slices with the mean survival score dropping to 2.25. After 20 μ M AMPA 89% of slices was in class I and the mean survival score was down to 1.13. After treatment with 30 μ M AMPA virtually all Purkinje cells were lost with 98% of the slices belonging to class I and a mean survival score of 1.02. AMPA toxicity affected Purkinje cells in an all-or-nothing kind of reaction: within one slice some Purkinje cells were seen disintegrating, while neighbouring cells looked healthy without any signs of impairment. Figure 11 B shows a dying Purkinje cell after AMPA treatment. Note that the remnants of the former dendritic tree are still visible.

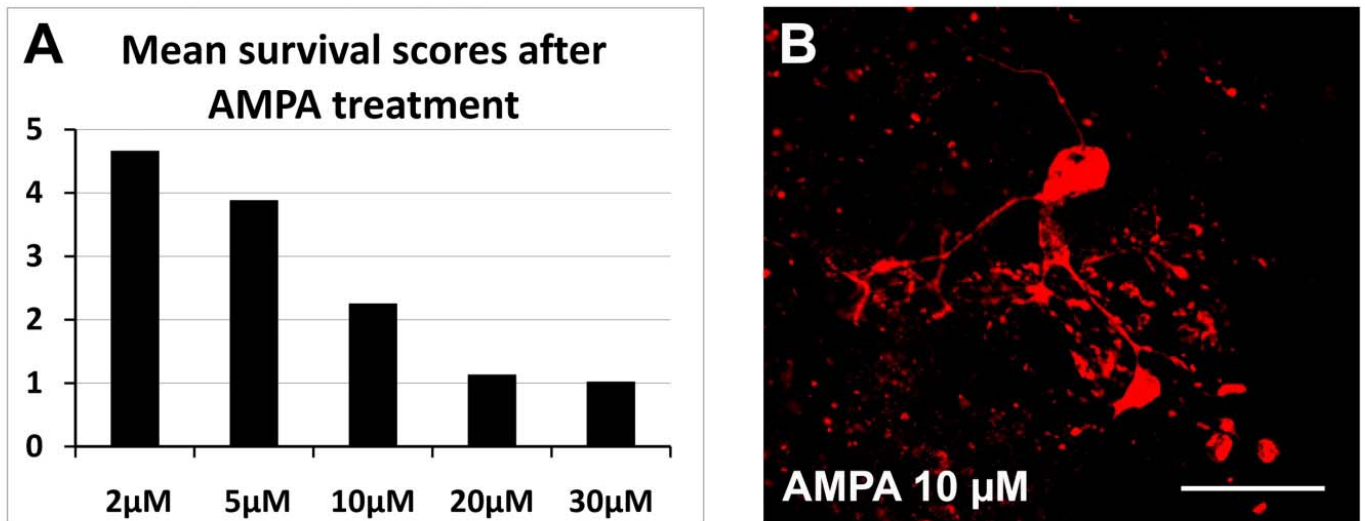


Figure 11: AMPA kills Purkinje cells in a dose dependent manner.

A) Mean survival score of the slice cultures treated with increasing AMPA concentrations. Each slice was given a value according to the survival class it was categorized into and the arithmetic mean was calculated. A value of 5 would mean complete Purkinje cell survival, a value of 1 complete Purkinje cell death. The slices in the control group remained untreated. With increasing AMPA concentrations a dose-dependent loss of Purkinje cells is evident which became virtually complete at 20 μM and 30 μM AMPA.

B) Calbindin D-28K staining of a Purkinje cell after treatment with 10 μM AMPA. The dendritic tree and the cell body are disintegrating. Scale bar 50 μm .

3.4.2. Pre-treatment with DHPG or PMA reduces dendritic tree size without a strong impact on cell survival

A strong reduction of the dendritic tree size in cerebellar Purkinje cells in slice cultures can be achieved by chronic activation of either mGluR1 (Sirzen-Zelenskaya *et al.*, 2006) or PKC (Metzger & Kapfhammer, 2000; Schrenk *et al.*, 2002). We have treated organotypic slice cultures according to these studies for 7 days with either the PKC activator PMA or the group I mGluR activator DHPG, starting at DIV 3-4. After completion of these treatments, slices were kept in culture for 4 more days in parallel with the cultures from the experimental groups which were exposed to AMPA (see below). Measurement of the dendritic tree area showed that the PMA treatment reduced the size of the dendritic tree to 43% of that of control cells, and the DHPG treatment to 49%, meaning that dendritic tree area was reduced to less than half of its original size (see Fig. 12 A and B). This reduction was clearly evident in the cell morphology (Fig. 12 C-E). It should be noted that the size reduction probably was

even more pronounced at the beginning of the AMPA treatment (1 day after the 7 day treatment period) because the dendritic trees had 4 days for recovery in control medium after ending the DHPG or PMA treatment. Recovery of the dendritic tree size has been shown to occur after washout of PMA (Metzger & Kapfhammer, 2000).

In order to verify that none of the pharmacological agents used severely affected Purkinje cell survival by itself, we measured cell survival at DIV 13-14 in DHPG-treated, PMA-treated and control cultures. PMA-treatment had no adverse effect on cell survival, and DHPG treatment resulted in a mild reduction of Purkinje cell survival with more slices falling into classes 4 and 3. The mean survival scores were 4.52 for control cultures, 4.56 for PMA-treated cultures and 3.8 for DHPG-treated cultures. The distribution of the slices in the different classes is shown in Figure 13.

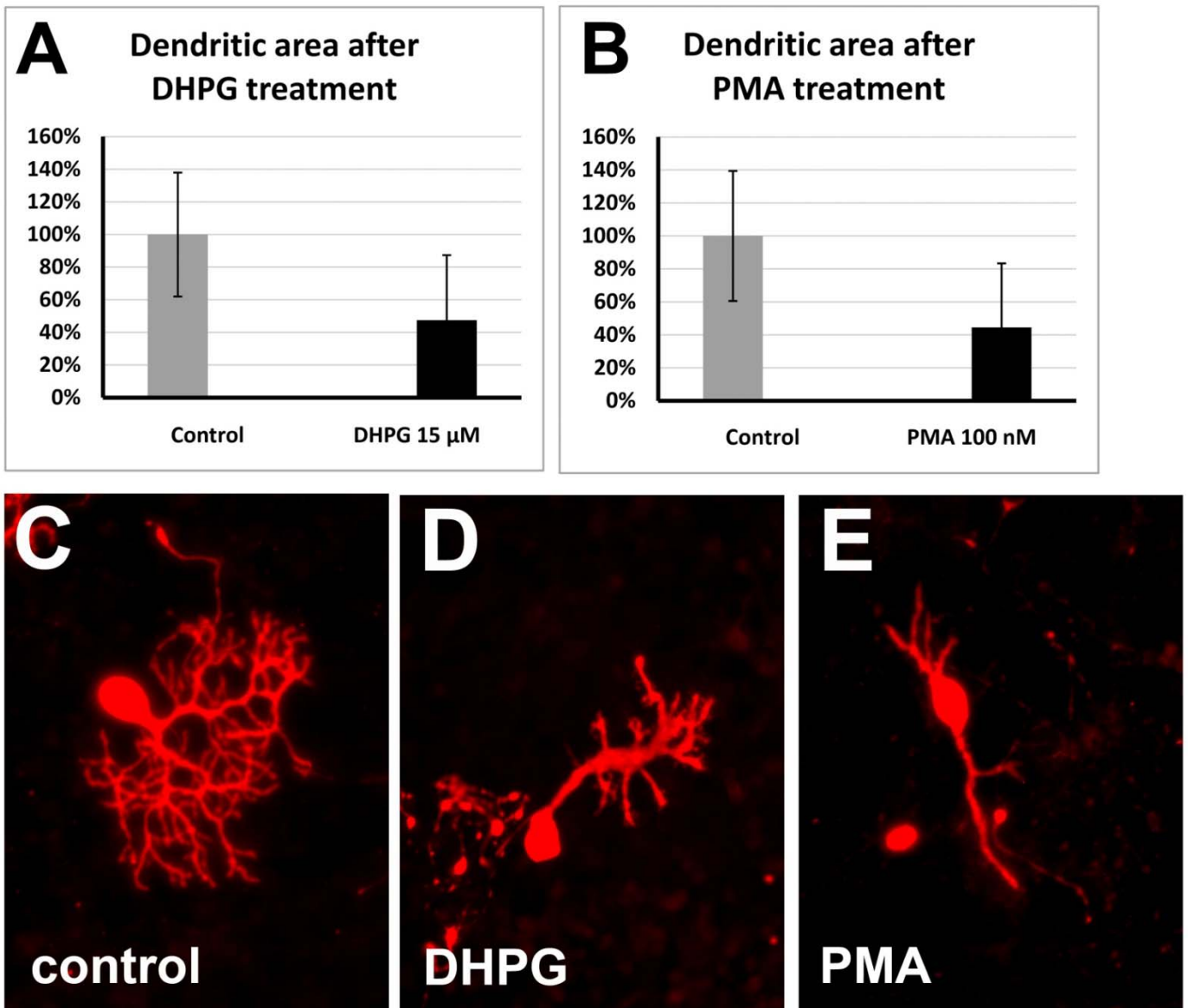


Figure 12: Size of the area covered by a Purkinje cell dendritic tree. Standard size of control cells was rated 100%.

A) Cells were treated with DHPG (15 μ M) without AMPA exposure. The average dendritic tree size was reduced to 43% of that of control cells (100%).

B) After treatment with PMA (100 nM) without AMPA exposure the mean size of the dendritic tree was reduced to 49% of that of control cells (100%).

Differences were significant with $p < 0.01$, Mann-Whitney test. Error bars represent the mean \pm standard deviation; standard deviation was 39% for PMA-treated cells (A), 40% for DHPG-treated cells (B) and 39% for control cells.

C – E) Examples of the morphology of dendritic trees of Purkinje cells grown under control conditions (C) or treated with DHPG (D) or PMA (E) for 7 days, without the AMPA exposure. Dendritic trees were visualized by immunostaining for Calbindin D-28K. The strong reduction of dendritic tree size after the treatments is evident. Scale bar 50 μ m.

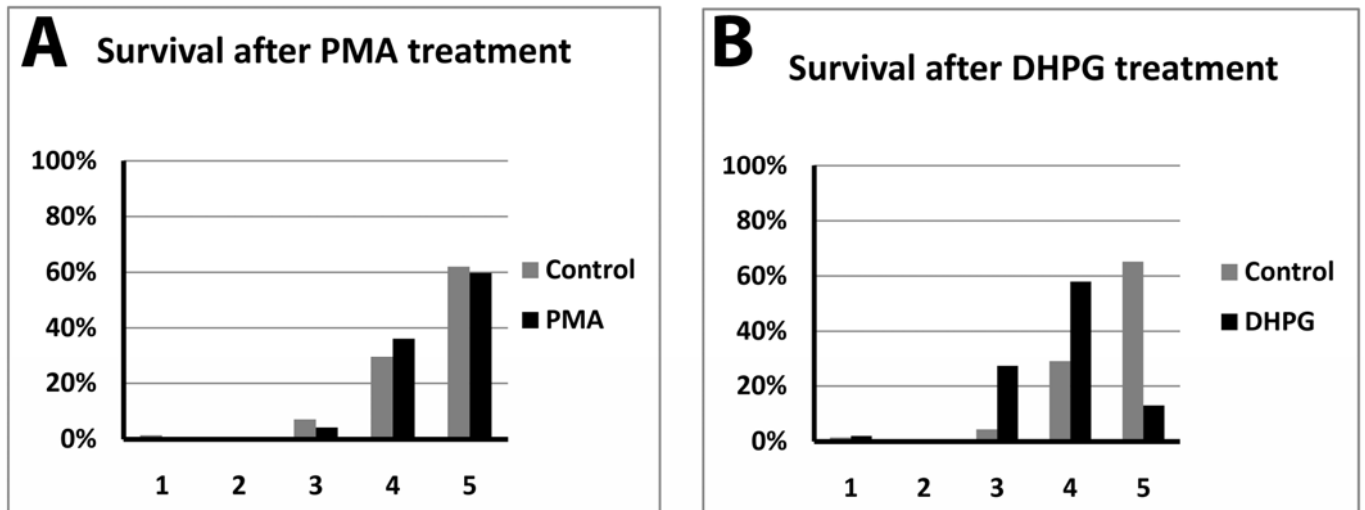


Figure 13: Distribution of the survival classes in PMA- and DHPG-treated slice cultures. The total number of classified slices of each treatment was set as 100%. The slices in the control group remained untreated.

A) Survival of slices receiving PMA 100 nM for 7 days was identical to control slices.

B) Survival of slices treated with DHPG 15 μ M for 7 days was moderately reduced compared to control slices with more slices falling into classes III and IV.

3.4.3. Normal sensitivity to AMPA induced neurotoxic death of Purkinje cells with small dendritic trees after PMA-treatment or DHPG-treatment

To test Purkinje cell susceptibility to AMPA treatment, we chose an AMPA concentration of 30 μ M, the lowest concentration which caused an explicit, almost complete Purkinje cell loss in control cultures (Fig. 11 A). At this point of the AMPA dose-response curve, a higher resistance to AMPA would result in a clearly detectable improvement of Purkinje cell survival.

When we exposed Purkinje cells from either DHPG-treated or PMA-treated cultures with small dendritic trees to 30 μ M AMPA for 4 hours we found no difference in survival compared to Purkinje cells in control cultures with normal sized dendritic trees. After the AMPA treatment, the mean survival score of control cultures was down to 1.15, with 85% of the slices grown under control conditions were in class I and 15% in class II. In slices pretreated with DHPG, the survival score after AMPA exposure was 1.16 with 84% falling into class I and to 16% into class II. With PMA pretreatment, the AMPA-mediated Purkinje cell loss was similar with a mean survival

score of 1.19. Of these slices, 85% were in class I, 10.5% in class II and 4.5% reached class III. These results indicate that the reduction of the dendritic tree size by pretreatment with either DHPG or PMA had no beneficial effect on AMPA mediated toxicity, i.e. the Purkinje cells retained the full sensitivity to AMPA neurotoxicity despite the strong reduction of dendritic tree size. The distribution of the slices in the different classes is shown in Figure 14.

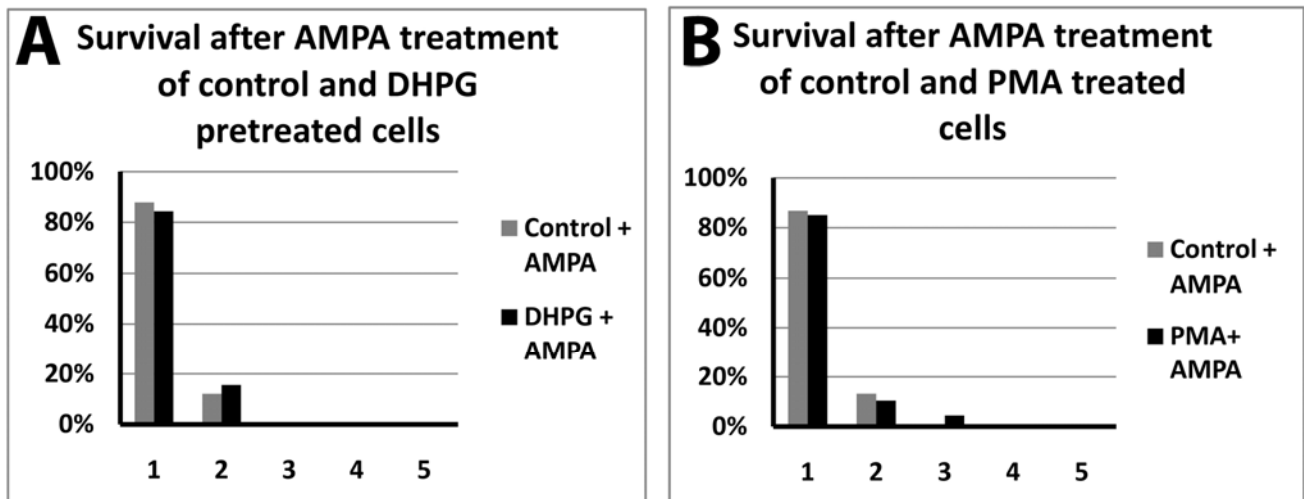


Figure 14: Survival of Purkinje cells with size-reduced dendritic trees after AMPA exposure.

A) Slice cultures were pre-treated with 100 nM PMA for 7 days and then exposed to 30 μ M AMPA for four hours (at DIV 11). There was no improved survival of the Purkinje cells in the pre-treated cultures; survival classes were similar to control cultures.

B) Slice cultures were pre-treated with 15 μ M DHPG for 7 days and then exposed to 30 μ M AMPA for four hours (DIV 11). There was no improved survival of the Purkinje cells in the pre-treated cultures; survival classes were similar to control cultures.

3.4.4. Glutamatergic synapses are abundantly available even on severely reduced dendritic trees

Since Purkinje cells with reduced dendritic trees were equally sensitive to excitatory overload by AMPA as those in control cultures, we have studied the presence of glutamatergic synapses on these Purkinje cells by immunohistochemistry. In order to visualize the presynaptic parts of the parallel fiber/Purkinje cell synapses, we stained for the vesicular glutamate transporter (vGluT) 1 which is expressed on the terminals of parallel fibers (Fremeau *et al.*, 2001; Takamori, 2006). Immunostaining with vGluT2 which specifically labels mossy fibers and climbing fibers (Hioki *et al.*, 2003)

was negative throughout the culture confirming the absence of these extrinsic fiber populations (data not shown). As shown in Figure 15, vGluT1 immunoreactivity appeared to be equally intense on the normal dendritic tree of control neurons and on the size-reduced Purkinje cell dendritic trees from PMA- or DHPG-treated cultures, revealing the presence of tightly clustered glutamatergic terminals. Interestingly, many of the somata of Purkinje cells in PMA-treated, but not in DHPG- treated or control cultures, were also covered by vGluT1-positive terminals. Despite the absence of inferior olive-derived climbing fibers in the cultures the PC soma was not covered by vGluT1-positive terminals under control conditions indicating a specific affinity of parallel fiber terminals for the more distal parts of the PC dendritic tree even in the absence of climbing fibers. This specificity appears to be lost after PKC activation, but not after mGluR activation.

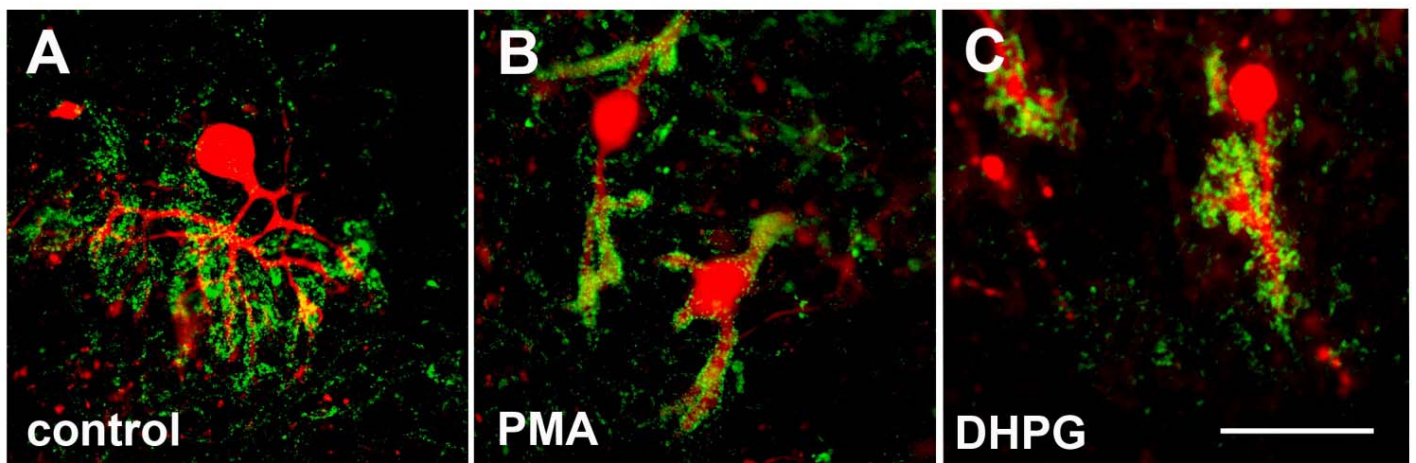


Figure 15: vGluT1 staining on Purkinje cells to visualize glutamatergic terminals. Double staining with Calbindin D-28K showing Purkinje cells in red and vGluT1 showing glutamatergic afferent terminals in green clustered around dendrites of Purkinje cells.

A) A Purkinje cell grown under control conditions shows dense clusters of vGluT1-positive afferent terminals along the dendrites. The cell soma and the proximal part of the dendritic tree are devoid of vGluT1-positive afferent terminals.

B) Two Purkinje cells after treatment with 100 nM PMA. The small dendrites are completely and densely covered by vGluT1-positive afferent terminals. In the lower cell vGluT1- positive afferent terminals are also present on the cell soma and the proximal dendrites.

C) A Purkinje cell after treatment with 15 μ M DHPG. The small dendrites are completely and densely covered by vGluT1-positive afferent terminals. The cell soma and the proximal part of the dendrite are devoid of vGluT1-positive afferent terminals.

Scale bar 50 μ m.

3.4.5. Cerebellar granule cells survive high doses of AMPA and start to express Calbindin D-28K immunoreactivity

While we observed an almost complete loss of Purkinje cells after 4 hrs of AMPA exposure, we did not observe any noticeable loss of granule cells in these experiments. Granule cell survival was assessed by staining with NeuN which in the cerebellum selectively labels granule cells (Weyer & Schilling, 2003). Neither DHPG nor PMA treatment changed the overall appearance of granule cells (data not shown). After exposure to AMPA, granule cells survived with apparently normal cell density. In some experiments we used concentrations of up to 1 mM AMPA for several days without observing a considerable loss of granule cells (data not shown) demonstrating the remarkable resistance of these cells to AMPA exposure. Interestingly, after AMPA treatment, granule cells started to express Calbindin D-28K at the end of the culture period (see Fig. 16). Calbindin D-28K immunoreactivity on granule cells could be detected as early as 24h after AMPA exposure (data not shown).

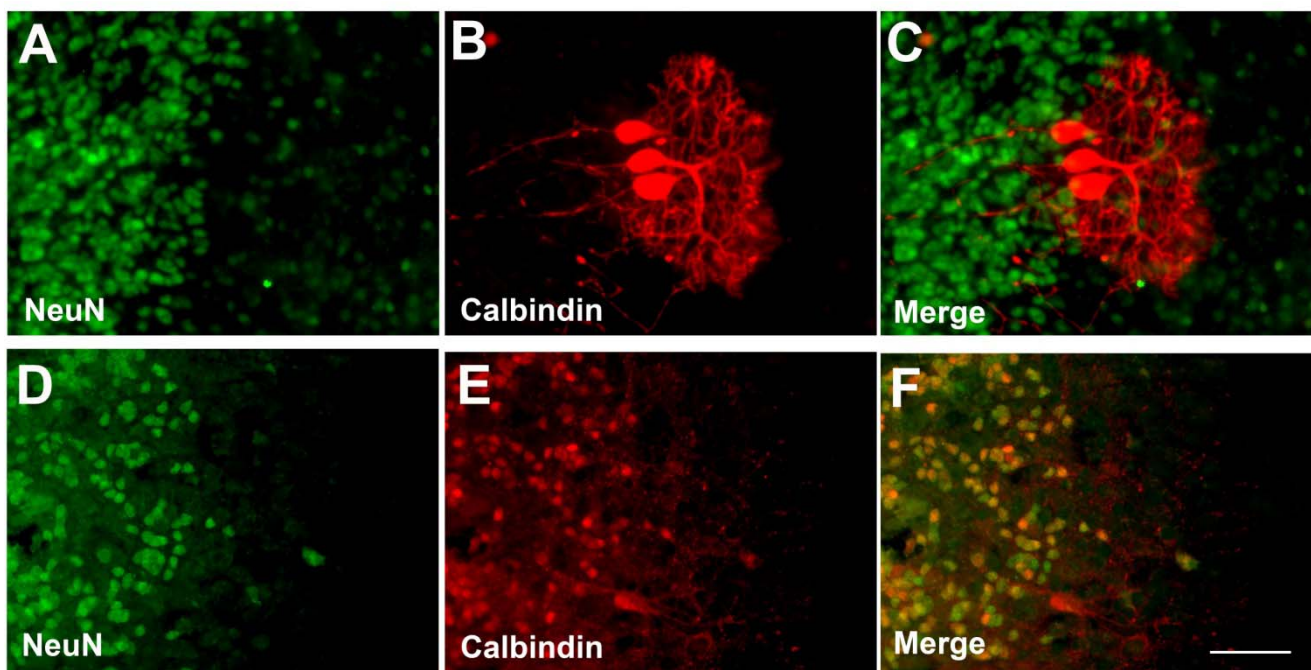


Figure 16: Double staining of Calbindin D-28K (red) and NeuN (green) reveals induction of Calbindin D-28K expression in AMPA challenged granule cells.

A) - C) In control cultures granule cells are positive for NeuN (A) and are negative for Calbindin D-28K, which labels specifically Purkinje cells (B). Merged image in (C).

D) – F) After AMPA exposure many NeuN-positive granule cells (D) became positive for Calbindin D-28K immunoreactivity (E). Most Purkinje have been lost due to the neurotoxic action of AMPA (E). Merged image in (F). Scale bar 50 μ m.

3.5. Discussion

Neurons display a large variety in their sensitivity to excitotoxic activation. In the cerebellum, Purkinje cells are highly susceptible to AMPA-mediated neurotoxic injury and death. In contrast, cerebellar granule cells, despite the presence of AMPA receptors, are almost insensitive to AMPA neurotoxic effects. While many of the differences in sensitivity of different neurons to neurotoxic damage probably result from different intracellular defence and compensatory mechanisms, one possible determinant for this sensitivity also might be the total receptor load of the neuron and, related to this, the size of the dendritic tree. We have tested whether a reduction of the dendritic tree size of Purkinje cells, induced by two different independent pharmacological treatments, may confer a certain protection from AMPA mediated neurotoxicity. Our results show that the reduction of the dendritic tree size offers no protection from AMPA-mediated neurotoxicity. Immunohistochemical staining for the vesicular glutamate transporter vGluT1 revealed that glutamatergic synapses were present in high density on the small dendritic trees of pre-treated Purkinje cells suggesting that receptor density rather than total receptor load is important for determining the sensitivity of Purkinje cells to AMPA-mediated neurotoxic death.

3.5.1. Purkinje cell excitotoxic death

Purkinje cells are very sensitive to AMPA-mediated excitotoxic death. A brief 30 min long exposure of Purkinje cells to 30 μ M AMPA is sufficient to cause death of most Purkinje cells by dark degeneration (Garthwaite & Garthwaite, 1991a). Neurotoxic death is mediated by a rise in intracellular calcium (Garthwaite & Garthwaite, 1991b; Strahlendorf *et al.*, 1998). Our finding that a 4 hour exposure to 20-30 μ M AMPA essentially wipes out the Purkinje cell population in cerebellar slice cultures is in good agreement with these findings. Furthermore, we show that even lower concentrations of AMPA beginning with 5 μ M AMPA for 4 hours lead to a considerable loss of Purkinje cells. This is in agreement with our earlier finding that chronic exposure of cerebellar slice cultures to 5 μ M AMPA resulted in an almost complete loss of Purkinje cells in cerebellar slice cultures (Sirzen-Zelenskaya *et al.*, 2006). The high sensitivity of Purkinje cells to AMPA-mediated excitotoxic death is also reflected by

the finding that lesions in the cerebral cortex can induce Purkinje cell damage (Park *et al.*, 2006; Slemmer *et al.*, 2007). In this paradigm, a strong activation of the cortico-ponto-cerebellar projection by the cortical lesion is sufficient to kill a fraction (up to 50%) of the Purkinje cells. These experiments show that even a strong activation of the regular endogenous excitatory projection pathway is sufficient to cause Purkinje cell excitotoxic death. Similarly, a hypoxic challenge of acute cerebellar slices induces Purkinje cell death which can be prevented by glutamate receptor antagonists demonstrating that Purkinje cells in this experimental paradigm are killed not by the hypoxia directly, but by the increased release of glutamate from afferent terminals (Barenberg *et al.*, 2001).

3.5.2. The role of the dendritic compartment for excitotoxic neuronal damage

Dendritic calcium signals play an important role in the development of an excitotoxic lesion. In C57/Bl6 mouse strains with different sensitivities to excitotoxic neuronal death there is a remarkable difference in the handling of dendritic calcium signals by hippocampal neurons. While these signals remain small and rather local in the resistant strain they are stronger and are able to spread over the entire cell in the sensitive strain (Shuttleworth & Connor, 2001). Purkinje cells have a dendritic tree of an enormous size which is tightly covered by AMPA-type glutamate receptors (Bergmann *et al.*, 1996; Douyard *et al.*, 2007). The large size of the dendritic tree could be an important determinant of Purkinje cell sensitivity to AMPA-mediated excitotoxic death. We have previously shown that chronic activation of metabotropic glutamate receptors or of protein kinase C leads to a strong reduction of the size of the Purkinje cell dendritic tree. One potential developmental function of this growth limiting effect obviously could be the control and limitation of Purkinje cell dendritic tree size, which in turn might be protective with respect to excitotoxic injury. We have experimentally tested this hypothesis in these studies. Our results clearly demonstrate that Purkinje cells with small dendritic trees are equally sensitive to AMPA exposure as their counterparts with large dendritic trees. Since pharmacological treatments were suspended 24h prior to the AMPA exposure it is unlikely that the used drugs could have enhanced AMPA neurotoxicity and influenced

Purkinje cell survival. Furthermore, pharmacological treatments alone had no or only a minor effect on Purkinje cell survival. These findings clearly indicate that the high sensitivity of Purkinje cells to AMPA-mediated excitotoxic death is unrelated to the large size of the Purkinje cell dendritic tree and to total AMPA receptor load. They rather suggest that important factors for the sensitivity to neurotoxic death are glutamate receptor density and the handling of calcium signals by the neuron. This interpretation is supported by studies showing that Purkinje cell neurotoxic death is influenced by the presence and activity of neuronal and glial glutamate transporters (Yamashita *et al.*, 2006; Slemmer *et al.*, 2007) or by the activation of calpain (Mansouri *et al.*, 2007). It should be noted that the experiments in this study were done in the organotypic slice culture model system, where the afferent innervation of Purkinje cells is severely altered compared to the *in vivo* situation. The climbing fiber innervation is missing in these cultures, and innervation by parallel fibers is not by parallel fiber beams running orthogonal to the Purkinje cell dendrite, but rather by fibers originating from local granule cells. Nevertheless, expression of glutamate receptors in Purkinje cells is very similar to the *in vivo* situation, and most electrophysiological characteristics of Purkinje cells are nicely preserved in the slice cultures (Dupont *et al.*, 2006). We are therefore convinced that our findings are not specific to the culture situation, but also apply to Purkinje cells *in vivo*.

3.5.3. Resistance of granule cells to AMPA treatment

In our experimental paradigm granule cells in the cerebellar slice cultures proved to be remarkably resistant to AMPA treatment. Even after chronic exposure to 1mM AMPA granule cells survived without any apparent cell loss in our cultures. This is in agreement with earlier studies showing that granule cells are insensitive to AMPA mediated excitotoxicity (Garthwaite & Garthwaite, 1991a; Resink *et al.*, 1994) despite the presence of Calcium-permeable GluR2 receptor subunits and AMPA-mediated Calcium influx (Hack *et al.*, 1995). Apparently granule cells are able to handle a certain Calcium load evoked by AMPA treatment, but AMPA induces granule cell loss when receptor desensitization is blocked (Hack *et al.*, 1995) or energy metabolism is compromised (Cebers *et al.*, 1998). In this study we observed that granule cells exposed to AMPA became distinctly immunoreactive for Calbindin D-28K. Typically,

granule cells do not express Calbindin D-28K immunoreactivity, which in the cerebellum is a specific marker for Purkinje cells (Bastianelli, 2003). Our finding that AMPA-challenged cerebellar granule cells rapidly become immunoreactive for Calbindin D-28K shows that the specificity of Calbindin D-28K immunoreactivity for Purkinje cells in the cerebellum is not absolute, and that also granule cells can express this protein. This finding suggests that the upregulation of Calcium-buffering proteins may be an important mechanism by which granule cells protect themselves from AMPA-mediated neurotoxic death.

3.6. Conclusion

Our results show that the reduction of the dendritic tree size offers no protection from AMPA-mediated neurotoxicity. Vesicular glutamate transporter vGluT1 staining revealed that glutamatergic synapses were present in high density on the small dendritic trees of pre-treated Purkinje cells suggesting that receptor density rather than total receptor load is important for determining the sensitivity of Purkinje cells to AMPA-mediated neurotoxic death. Our finding that AMPA-challenged cerebellar granule cells rapidly become immunoreactive for Calbindin D-28K shows that the specificity of Calbindin D-28K immunoreactivity for Purkinje cells in the cerebellum is not absolute, and that also granule cells can express this protein. This finding suggests that the upregulation of Calcium-buffering proteins may be an important mechanism by which granule cells protect themselves from AMPA-mediated neurotoxic death.

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4. Mechanisms of dendritic growth inhibition mediated by chronic mGluR1 or PKC activation

P/Q-type and T-type calcium channels, but not TRPC3 channels, are involved in inhibition of dendritic growth after chronic mGluR1 and PKC activation in cerebellar Purkinje cells

Olivia S. Gugger¹, Jana Hartmann², Lutz Birnbaumer³ and Josef P. Kapfhammer¹

¹ Anatomical Institute, Department of Biomedicine Basel, University of Basel, Pestalozzistr. 20, CH - 4056 Basel, Switzerland

² Institute for Neuroscience, TU Munich, Biedersteiner Str. 29, D - 80802 Munich, Germany

³ National Institute of Environmental Health Sciences, Research Triangle Park, NC27709, USA

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4.1. Abstract

The development of a neuronal dendritic tree is modulated by signals from afferent fibers as well as by an intrinsic program. We have previously shown that chronic activation of either class I metabotropic glutamate receptors or Protein Kinase C in organotypic cerebellar slice cultures of mice and rats severely inhibits the growth and development of the Purkinje cell dendritic tree. The signaling events linking receptor activation to the regulation of dendritic growth remain largely unknown. We have studied whether channels allowing the entry of Ca^{2+} ions into Purkinje cells, in particular the TRPC3 channels, P/Q-type and T-type Ca^{2+} channels, might be involved in the signaling after mGluR1 or PKC stimulation. We show that the inhibition of dendritic growth seen after mGluR1 or PKC stimulation is partially rescued by pharmacological blockade of P/Q-type and T-type Ca^{2+} channels indicating that activation of these channels mediating Ca^{2+} influx contributes to the inhibition of dendritic growth. In contrast, absence of Ca^{2+} -permeable TRPC3 channels in TRPC3-deficient mice or pharmacological blockade had no effect on mGluR1- and PKC-mediated inhibition of Purkinje cell dendritic growth. Similarly, blockade of Ca^{2+} influx through GluRdelta2 or R-type Ca^{2+} channels or inhibition of release from intracellular stores did not influence mGluR1- and PKC-mediated inhibition of Purkinje cell dendritic growth. These findings suggest that both T- and P/Q-type Ca^{2+} channels, but not TRPC3 or other Ca^{2+} -permeable channels, are involved in mGluR1 and PKC signaling leading to the inhibition of dendritic growth in cerebellar Purkinje cells.

4.2. Introduction

A large and highly elaborate dendritic tree is a characteristic feature of cerebellar Purkinje cells. The factors and molecules controlling the growth and patterning of neuronal dendrites are not yet well understood. We have previously shown that chronic activation of metabotropic glutamate receptor 1 (mGluR1) (Sirzen-Zelenskaya et al., 2006) or Protein Kinase C (PKC) (Metzger & Kapfhammer, 2000; Schrenk *et al.*, 2002) in organotypic cerebellar slice cultures of postnatal mice severely inhibits the growth and development of the Purkinje cell dendritic tree.

The signaling events leading to inhibition of dendritic growth after both mGluR1 and PKC activation remain largely unknown, but PKC activation is not required for the inhibitory effect of mGluR1 suggesting that both use independent signaling pathways (Sirzen-Zelenskaya et al., 2006). In this study we have searched for potential mechanisms limiting Purkinje cell dendritic growth and have concentrated on channels allowing the entry of Ca^{2+} ions. Increased influx of Ca^{2+} into Purkinje cells occurs in two independent mouse models which are characterized by Purkinje cells with greatly reduced dendritic arbors similar to the ones seen after mGluR or PKC stimulation: the *lurcher* (Zanjani et al., 2009) and the *moonwalker* (*mwk*) mice (Becker et al., 2009). In the *mwk* mouse model, a gain-of-function mutation in the *Trpc3* gene leads to altered TRPC3 channel gating resulting in an ataxic phenotype with impaired growth and development of Purkinje cells with a drastically reduced dendritic tree reminiscent of that seen after mGluR or PKC activation (Becker et al., 2009). The TRPC3 channel is a nonselective cation channel which mediates the slow excitatory postsynaptic potential (sEPSP) seen after stimulation of mGluR1 (Hartmann et al., 2008). *Trpc3* is strongly expressed in Purkinje cells during dendritogenesis (Huang et al., 2007).

Purkinje cell dendrites have a high density of voltage dependent Ca^{2+} channels (VDCC), which consist mostly of the high-threshold P/Q-type (Usowicz et al., 1992). P/Q-type Ca^{2+} channels in dendrites have been shown to be essential for the trimodal pattern of Purkinje cell spontaneous activity (Womack & Khodakhah, 2002; Womack & Khodakhah, 2004). Contributing to the low-threshold fraction of Ca^{2+} currents and to dendritic burst firing of Purkinje cells are the T-type Ca^{2+} channels (Pouille *et al.*, 2000; Swensen & Bean, 2003; Womack & Khodakhah, 2004). The Cav3.1 and Cav3.3 T-type channel isoforms are expressed in Purkinje cells and the Cav3.1

isoform has been shown to co-localize with mGluR1 in spines and to be potentiated upon mGluR1 activation (Hildebrand et al., 2009).

In this study we have analyzed the role of P/Q-type, T- type and TRPC3 channels in mGluR1- and PKC-mediated dendritic growth inhibition. Pharmacological inhibition of the channels in cerebellar slice cultures revealed that both T- and P/Q-type Ca^{2+} channels are involved in mediating growth inhibition after chronic mGluR1 and PKC activation. In contrast, using TRPC3 deficient mice (Hartmann et al., 2008) or pharmacological blockade of TRPC3, release from intracellular stores, GluRdelta2 or R-type Ca^{2+} channels, no involvement of TRPC3 or other sites of Ca^{2+} entry in the mGluR1 and PKC-mediated inhibition of dendritic growth was found.

4.3. Materials and Methods

4.3.1. Organotypic slice cultures

Animal experiments were carried out in accordance with the European Communities Council Directive of 24 November 1986 (86/609/EEC) and were reviewed and permitted by Swiss authorities. Cultures were prepared from B6CF1 mice (CB6) and TRPC3^{-/-} mice (Hartmann et al., 2008) as described previously (Adcock et al., 2004; Kapfhammer, 2005). Briefly, mice were decapitated at postnatal day 8 (P8), their brains were aseptically removed and the cerebellum was dissected in ice-cold preparation medium (minimal essential medium (MEM), 1% glutamax (Gibco, Invitrogen), pH 7.3). Sagittal sections 350 μ m thick were cut on a McIlwain tissue chopper under aseptic conditions. Slices were separated, transferred onto permeable membranes (Millicell-CM, Millipore) and incubated on a layer of incubation medium (50% MEM, 25% Basal Medium Eagle, 25% horse serum, 1% glutamax, 0.65% glucose) in a humidified atmosphere with 5% CO₂ at 37°C. The medium was changed every 2-3 days. Pharmacological compounds were added to the medium at each change for a total of 7 days, starting at 2-4 days in vitro (DIV). Slices were kept in culture for a total of 9-11 days before fixation and immunohistochemical staining. The following compounds were used: (RS)-3,5-Dihydroxyphenylglycine (DHPG, Tocris, Bristol, UK), Phorbol 12-myristate 13-acetate (PMA, Tocris), ω -Agatoxin IVA (Bachem, Bubendorf, Switzerland), ω -Conotoxin MVIIIC (Bachem), mibefradil dihydrochloride (Tocris), NNC 55-0396 dihydrochloride (Tocris), SNX-482 (Biotrend,

Köln, Germany), 1-naphthyl acetyl spermine (Nasp, Sigma), U73122 (Tocris), Ethyl-1-(4-(2,3,3-trichloroacrylamide)phenyl)-5-(trifluoromethyl)-1H-pyrazole-4-carboxylate (Pyr3, Tocris). The following concentrations were used: 10 μ M DHPG (Sirzen-Zelenskaya *et al.*, 2006), 50 nM PMA (Metzger & Kapfhammer, 2000; Schrenk *et al.*, 2002), 100 nM ω -Agatoxin IVA (shown to be effective and selective in (Mintz *et al.*, 1992a; Mintz *et al.*, 1992b)), 1 μ M ω -Conotoxin MVIIC (McDonough *et al.*, 1996), 500nM SNX-482 (Newcomb *et al.*, 1998), 100 μ M Nasp (as used in (Zanjani *et al.*, 2009), 3 μ M Pyr3 (Kiyonaka *et al.*, 2009), 10 μ M U73122 (as used in (Canepari & Ogden, 2006)), 2 μ M mibefradil, 1 μ M NNC 55-0396. When organotypic slice cultures were treated for 7 days with the specific T-type Ca^{2+} channel inhibitor NNC 55-0396, we found that the compound was toxic to Purkinje cell axons in concentrations above 1 μ M. Purkinje cell survival and dendritic development, however, was not affected up to concentrations of 6 μ M (Fig. 17). The same observation was made with the T-type Ca^{2+} channel inhibitor mibefradil in higher concentrations (Gugger and Kapfhammer, unpublished observations). Since mibefradil has been reported to inhibit T-type currents on Purkinje cells at 2 μ M (McDonough & Bean, 1998) and to have an IC_{50} value about 1.5 times higher than NNC 55-0396 (Huang *et al.*, 2004), we decided to apply 1 μ M NNC 55-0396 to study dendritic development.

4.3.2. Immunocytochemistry

At DIV 9-11 cultures were fixed in 4% paraformaldehyde overnight at 4°C. All reagents were diluted in 100 mM phosphate buffer (PB), pH 7.3. Slices were incubated in blocking solution (0.3% Triton X-100, 3% normal goat serum) for a minimum of 30 minutes in order to permeabilize the tissue and block non-specific antigen binding. The primary antibody was added to the slices in fresh blocking solution and incubated overnight at 4°C. After washing in PB, the secondary antibody was added to the slices in PB containing 0.1% Triton X-100 in order to prevent non-specific antigen binding for at least 1 hour at room temperature. For the analysis of Purkinje cell survival and Purkinje cell dendritic size, rabbit anti-Calbindin D-28K (Swant, Marly, Switzerland, 1:1000) was used as a primary antibody to visualize Purkinje cells. Stained slices were mounted on glass slides and coverslipped with Mowiol. Cultures and sections were viewed on an Olympus AX-70 microscope

equipped with a Spot digital camera. Recorded images were adjusted for brightness and contrast with Photoshop image processing software.

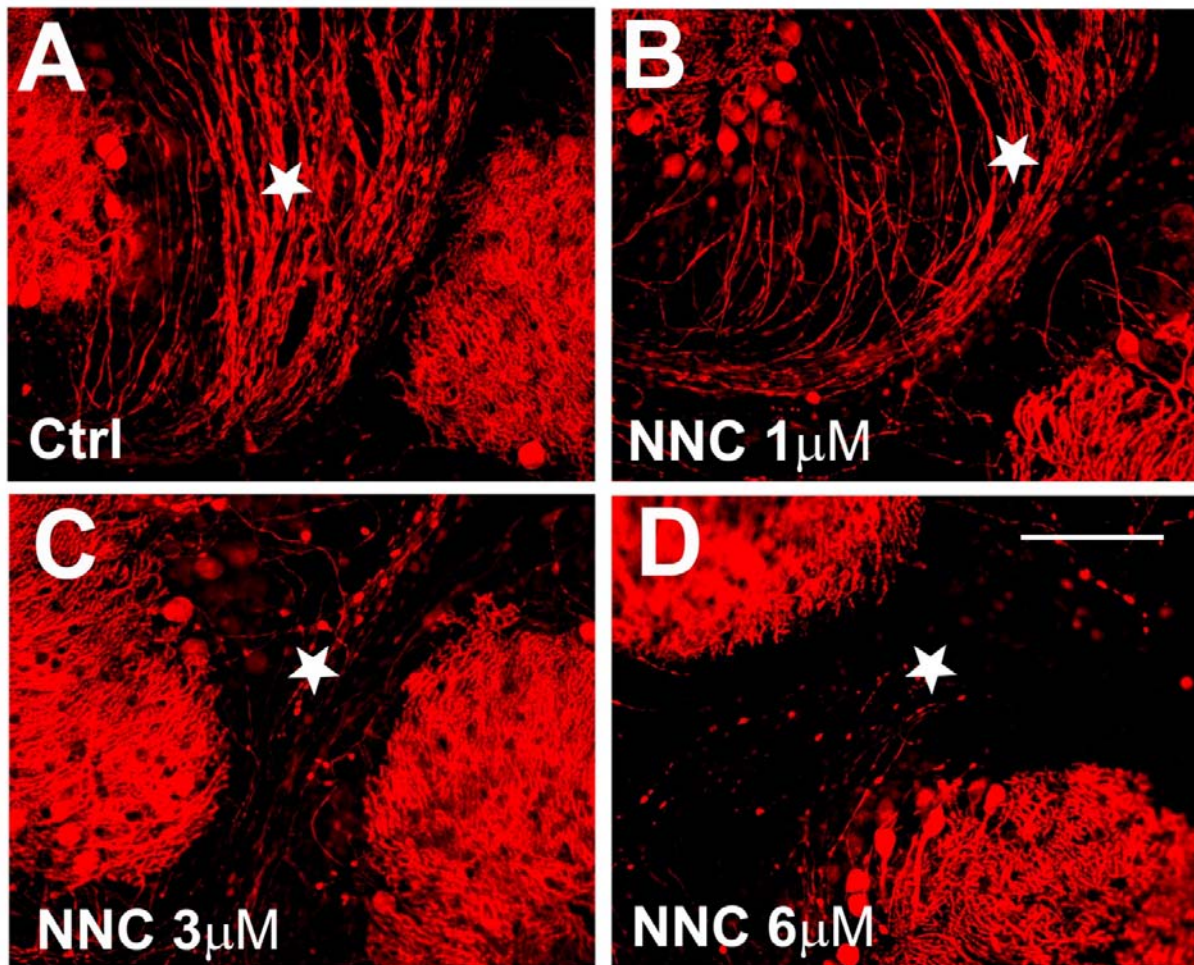


Figure 17: Toxicity of the T-type channel inhibitor NNC 55-0396 on Purkinje cell axons. Staining of Calbindin D-28K shows Purkinje cells. Cells were treated with NNC 55-0396 for 7 days. Scale bar: 100 μm

- A)** In untreated control cultures the Purkinje cell axons are clearly visible (asterisk in A).
- B)** Treatment with NNC 1 μM does not harm Purkinje cell axons (asterisk in B)
- C)** Treatment with NNC 3 μM causes damage of Purkinje cell axons (asterisk in C)
- D)** After treatment with NNC 6 μM most Purkinje cell axons are missing (asterisk in D)

4.3.3. Quantitative analysis of cultured Purkinje cells

The quantification of the effects of the pharmacological treatments on Purkinje cell dendritic tree size was done as previously described (Adcock et al., 2004; Sirzen-Zelenskaya et al., 2006). Purkinje cells which had a dendritic tree which could be isolated from its surroundings were selected for analysis. Cells were viewed with a

25x lens and photographed with a digital camera (Coolsnap-Pro, Photometrics, Canada). Dendritic branch points were counted and an image analysis program (Image Pro Plus) was used to trace the outline of the Purkinje cell dendritic trees yielding the area covered by the dendritic tree. Cells were acquired from three independent experiments with an average number of 20 cells per experiment and per growth condition. For detailed information on the number of cells measured for each diagram see supporting Table 1. The data were analyzed using GraphPad Prism software. The mean value of the dendritic tree area and number of branch points of untreated control cells were set to 100 % and the results were expressed as percentage of controls. Error bars represent the standard error of the mean (SEM). The statistical significance of differences in parameters was assessed by non-parametric analysis of variance (Kruskal-Wallis test) followed by Dunn's post test. For comparisons of single data columns, Mann-Whitney's non-parametric test was used. Confidence intervals were 95 %, statistical significance when $p < 0.05$. The number of measured cells is indicated as n value in the figure legends.

4.4. Results

4.4.1. Inhibition of T-, N- and P/Q-type Ca^{2+} channels in cerebellar slice cultures does not affect dendritic tree size in Purkinje cells

T- and P/Q-type Ca^{2+} channels are abundantly expressed in Purkinje cell dendrites (Usowicz *et al.*, 1992; Hildebrand *et al.*, 2009) and are one of the major sources of Ca^{2+} influx into Purkinje cells (Usowicz *et al.*, 1992; Watanabe *et al.*, 1998; Isope & Murphy, 2005; Isope *et al.*, 2010). Furthermore, Ca^{2+} influx through these channels has been shown to be potentiated by mGluR1 activation (Kitano *et al.*, 2003; Hildebrand *et al.*, 2009; Johnston & Delaney, 2010). In order to exclude the possibility that inhibition of P/Q- and T-type Ca^{2+} channels by itself has a specific effect on Purkinje cell dendritic trees independent of mGluR1 or PKC stimulation we tested the effect of channel antagonists on Purkinje cell dendritic trees. For P/Q-type Ca^{2+} channel inhibition, a combination of ω -Agatoxin IVA and ω -Conotoxin MVIIC was used (further referred to as P/Q-block) (Mintz *et al.*, 1992a; McDonough *et al.*, 2002). ω -Conotoxin MVIIC is also a strong inhibitor of N-type Ca^{2+} channels

(McDonough *et al.*, 1996), which contribute about 5 % to the total high-threshold Ca^{2+} currents in Purkinje cells (Regan, 1991; Mintz *et al.*, 1992a; Watanabe *et al.*, 1998; McKay & Turner, 2005). We further refer to the combination of ω -Agatoxin IVA and ω -Conotoxin MVIIC merely as P/Q-block although it is in fact a P/Q/N block.

T-type Ca^{2+} channels were blocked with the inhibitors mibefradil or NNC 55-0396 (McDonough & Bean, 1998; Huang *et al.*, 2004).

Organotypic slice cultures were treated for 7 days with one of the T-type Ca^{2+} channel inhibitors mibefradil or NNC 55-0396 and the P/Q-block starting at DIV 2-4. The average size of dendritic tree areas in cultures treated with either combination of a T-type Ca^{2+} channel inhibitor and the P/Q block was minimally reduced compared to untreated control cultures. Average dendritic branch point number in cultures treated with NNC 55-0396 and the P/Q-block was slightly increased. With both combinations, the effects on dendritic tree size and branch points were not statistically significant and no changes in Purkinje cell morphology were visible. Inhibition of T- and P/Q-type Ca^{2+} channels alone thus had no effect on Purkinje cell dendritic development (Fig. 18, A and B).

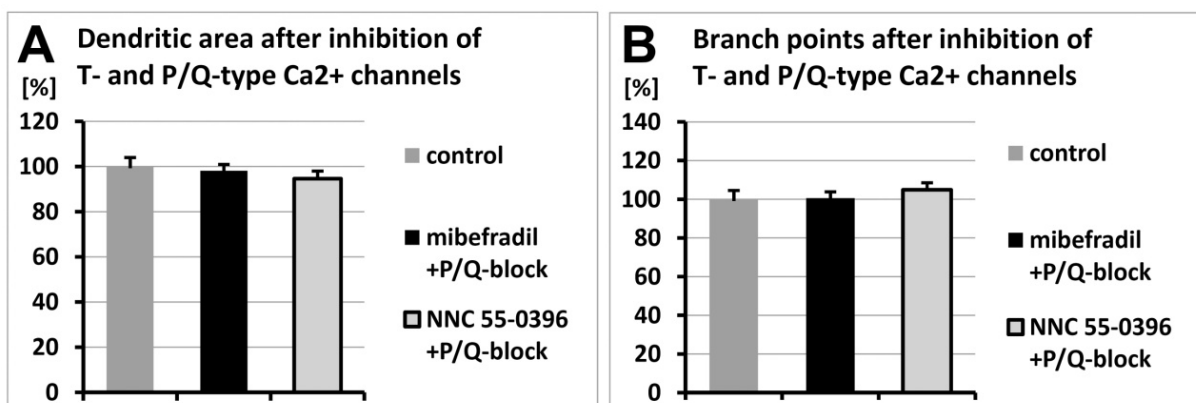


Figure 18. Dendritic tree size after treatment with P/Q-type and T-type channel blockers. A) and B): Size of the area covered by a single dendritic tree (A) and number of branch points (B). Cells were treated with the P/Q-block and one of the T-type Ca^{2+} channel inhibitors mibefradil or NNC 55-0396 combined. Standard growth of control cells was rated 100%. No statistical significance. Number of cells analyzed is given as n-value. Error bars represent the SEM.

A) Treatment with mibefradil and the P/Q-block reduced average area to 98% (n=60). Treatment with NNC 55-0396 and the P/Q-block reduced average area to 95% (n=86). No statistical significant difference to control (n=58).

B) Treatment with Mibefradil and the P/Q-block left average branch point number at 100% (n=50). Treatment with NNC 55-0396 and the P/Q-block increased average branch point number to 105% (n=66). No statistical significant difference to control (n=50).

4.4.2. Inhibition of either T- or P/Q-type calcium channels alone could not rescue dendritic growth after mGluR1 stimulation

In order to find out whether T- and P/Q type Ca^{2+} channels are involved in mGluR1 mediated dendritic growth inhibition, we treated the cultures with the group I mGluR activator DHPG and simultaneously with inhibitors of T- or P/Q-type Ca^{2+} channels. Organotypic slice cultures were treated for 7 days with DHPG (10 μM), starting at DIV 2-4. Alternatively, cultures were treated in the same way with DHPG and the P/Q-block or one of the T-type Ca^{2+} channel inhibitors mibefradil or NNC 55-0396 (Fig. 19, A and B).

The treatment with DHPG resulted in reduced dendritic areas of 48% and dendritic branch point numbers of 37% of untreated control cultures. Co-treatment with mibefradil resulted in an average dendritic area of 53% and an average branch point number of 40%, co-treatment with NNC 55-0396 resulted in an average dendritic area of 53% and an average branch point number of 37% of control cultures. Co-treatment with the P/Q-block resulted in an average dendritic area of 51% and an average branch point number of 43% of control cultures. Thus, all co-treatments showed a trend towards an improvement of dendritic growth which did not reach statistical significance. No difference in effectiveness between the two T-type Ca^{2+} channel inhibitors mibefradil and NNC 55-0396 was found.

4.4.3. Combined inhibition of T- and P/Q-type calcium channels rescued Purkinje cell dendritic growth after mGluR1 stimulation

Because both types of channels mediate Ca^{2+} entry into Purkinje cells, the inhibition of one type of channel might have been compensated by the other. Therefore, we did experiments with simultaneous inhibition of T- and P/Q-type Ca^{2+} channels along with DHPG treatment. Organotypic slice cultures were treated for 7 days with DHPG (10 μM), starting at DIV 2-4. Alternatively, cultures were treated in the same way with DHPG, the P/Q-block and one of the T-type Ca^{2+} channel inhibitors mibefradil or NNC 55-0396.

The treatment with DHPG resulted in reduced dendritic areas of 48% and dendritic branch point numbers of 35% of untreated control cultures. Both combinations (including mibefradil or NNC 55-0396) were able to increase the average area and

branch point number of the dendritic trees significantly compared to cultures treated with DHPG alone. The P/Q/T inhibition including mibefradil increased the dendritic areas to 64% and the number of branch points to 50%, and the P/Q/T inhibition including NNC 55-0396 increased the dendritic areas to 65% and the number of branch points to 48% of control cultures (Fig. 19, C and D). Thus, the T-type Ca^{2+} channel inhibitors mibefradil and NNC 55-0396 were equally effective. The beneficial effect of P/Q/T inhibition on DHPG-treated cells was also evident qualitatively in cell morphology, with dendritic trees of a clearly higher degree of elaboration in rescued cultures (Fig. 19, E-G).

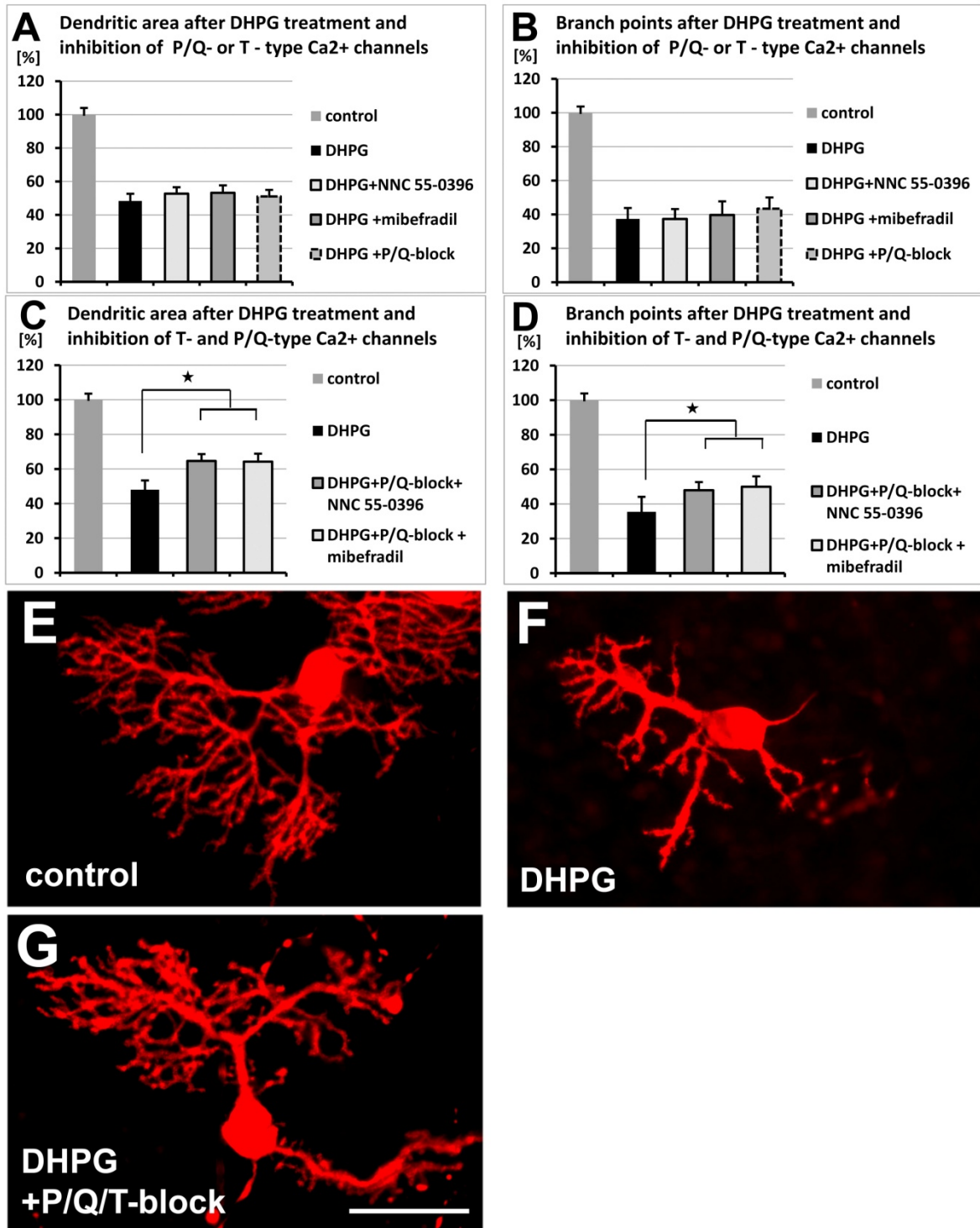


Figure 19: Dendritic trees after DHPG treatment and co-treatment with T- and/or P/Q-type channel blockers. Size of the area covered by a single dendritic tree (A, C) and number of branch points (B, D). Cells were treated with DHPG (10 μ M) for 7 days either alone or in the presence of the P/Q-block or one of the T-type Ca²⁺ channel inhibitors Mibefradil or NNC 55-0396 (A, B), or in the presence of the P/Q-block and one of the T-type Ca²⁺ channel inhibitors Mibefradil or NNC 55-0396 combined (C, D). Standard growth of control cells was rated 100%. Number of cells analyzed is given as n-value. Error bars represent the SEM. All pharmacologically treated growth conditions were significantly smaller than control ($P < 0.001$).

A) Treatment with DHPG reduced average area to 48% (n=98). Treatment with DHPG and NNC 55-0396 resulted in average areas of 53% (n=110, not significant), with DHPG and mibefradil of 53% (n=75, not significant) and with DHPG and the P/Q-block of 51% (n=91, not significant) of controls (n=85).

B) Treatment with DHPG reduced average branch point number to 37% (n=92). Treatment with DHPG and NNC 55-0396 resulted in average branch point numbers of 37% (n=92, not significant), with DHPG and mibefradil of 40% (n=62, not significant) and with DHPG and the P/Q-block of 43% (n=79, not significant) of controls (n=74).

C) Treatment with DHPG for 7 days (n=70) reduced average dendritic area to 48% of untreated controls (n=64). Treatment with P/Q-block and mibefradil simultaneously with DHPG increased the average area to 64% (n=77, significant *** $P < 0.001$). Treatment with P/Q-block and NNC 55-0396 simultaneously with DHPG increased average area to 65% (n=86, significant *** $P < 0.001$).

D) Treatment with DHPG for 7 days (n=67) reduced average branch point number to 35% of untreated controls (n=58). Treatment with P/Q-block and mibefradil simultaneously with DHPG increased average branch point number to 50% (n=69, significant $P < 0.05$). Treatment with P/Q-block and NNC 55-0396 simultaneously with DHPG increased average branch point number to 48% (n=78, significant $P < 0.05$).

E)-G) Examples of the morphology of Purkinje cell dendritic trees. Staining of Calbindin D-28K shows Purkinje cells. DHPG-treatment (F, 10 μM for 7 days) caused stunted dendrites compared to the control cell (E). Inhibition of P/Q- and T-type Ca^{2+} channels (NNC 55-0396 1 μM) along with DHPG treatment (G) resulted in visibly more dendritic branches and a general morphology close to the control cell.

Scale bar 40 μm .

4.4.4. T- and P/Q-type calcium channels mediate inhibition of Purkinje cell dendritic growth after PKC activation

A strong dendritic growth inhibition of cerebellar Purkinje cells in slice cultures can be achieved by chronic activation of either mGluR1 (Sirzen-Zelenskaya et al., 2006) or PKC (Metzger & Kapfhammer, 2000; Schrenk *et al.*, 2002). In order to find out whether T- and P/Q-type Ca^{2+} channels are also involved in mediating this effect after chronic PKC activation the same rescue experiment was carried out using the PKC stimulator PMA instead of DHPG. Organotypic slice cultures were treated for 7 days with PMA (50 nM), starting at DIV 2-4. Alternatively, cultures were treated in the same way with PMA and the P/Q-block, PMA and one of the T-type Ca^{2+} channel inhibitors mibefradil or NNC 55-0396, or PMA, P/Q-block and a T-type Ca^{2+} channel inhibitor combined.

Application of the P/Q-block or one of the T-type Ca^{2+} channel inhibitors as well as their combinations along with the PMA treatment was sufficient to increase the area and branch point number of the dendritic trees significantly (Fig. 20, A and B). The

rescue effect obtained by the combination of T- and P/Q-type Ca^{2+} channel inhibition was stronger than inhibition of either T- or P/Q-type Ca^{2+} channels. The size of the dendritic areas after PMA treatment was reduced to 42%, the average branch point number to 31% of untreated control cultures. Together with mibefradil, the dendritic areas reached 55% and the average branch point number 39%. PMA and NNC 55-0396 resulted in dendritic areas of 51% and a branch point number of 42%. Together with the P/Q-block the area was 56%, branch point number 44%. The combination of the P/Q-block and mibefradil resulted in an average dendritic area of 61% and a branch point number of 49%. In the combination with P/Q-block and NNC 55-0396 the areas were 64% and branch point numbers were 53%. In rescued cultures, the morphology of the dendritic trees was also visibly improved compared to cells treated with PMA only (Fig. 20, C-E).

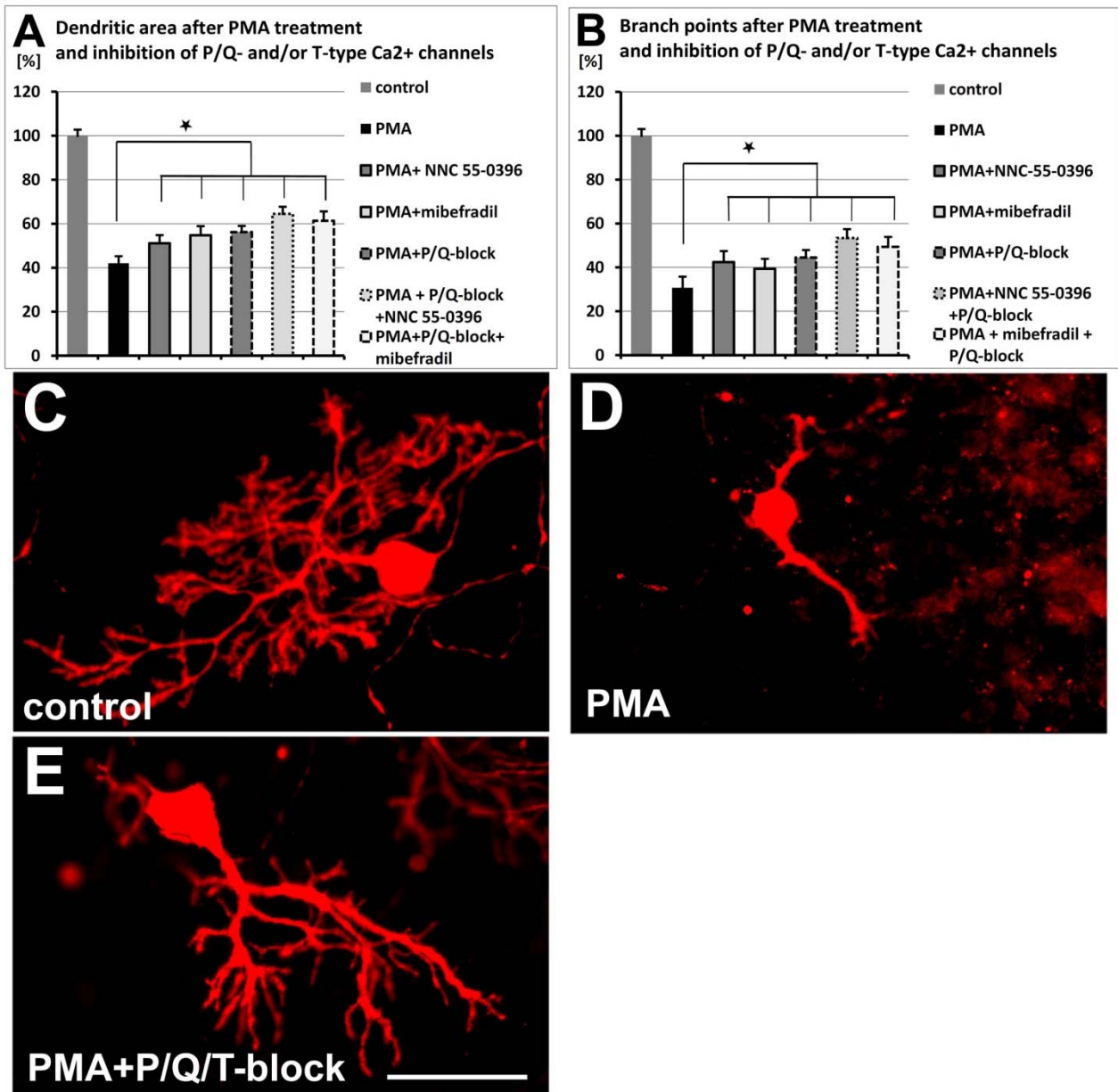


Figure 20: Dendritic trees after PMA treatment and co-treatment with T- and/or P/Q-type channel blockers Size of the area covered by a single dendritic tree (A) and number of branch points (B). Cells were treated with PMA (50 nM) for 7 days either alone or in the presence of the P/Q-block and/or one of the T-type Ca²⁺ channel inhibitors mibefradil or NNC 55-0396. Standard growth of control cells was rated 100%. Number of cells analyzed is given as n-value. Stars indicate statistical significance compared to PMA-only-treated cultures. All pharmacologically treated growth conditions were significantly smaller than control (P<0.001). Error bars represent the SEM.

A) PMA treatment (n=137) reduced average dendritic area to 42% of untreated controls (n=139). After co-treatment with NNC-55-0396, average area was 51% (n=119, significant P<0.05), after co-treatment with mibefradil 55% (n=80, significant P<0.001) and after co-treatment with the P/Q-block 56% (n=131, significant P<0.001).

Treatment with P/Q-block and mibefradil simultaneously with PMA increased average size to 61% (n=65, significant P<0.001). Treatment with P/Q-block and NNC 55-0396 simultaneously with PMA increased average size to 64% (n=100, significant P<0.001).

B) PMA treatment (n=143) reduced average dendritic branch point number to 31% of untreated controls (n=113). After co-treatment with NNC-55-0396, average branch point number was 42% (n=114, significant $P < 0.001$), after co-treatment with mibefradil 39% (n=69, significant $P < 0.05$) and after co-treatment with the P/Q-block 44% (n=117, significant $P < 0.001$).

Treatment with P/Q-block and mibefradil simultaneously with PMA increased average size to 49% (n=58, significant $P < 0.001$). Treatment with P/Q-block and NNC 55-0396 simultaneously with PMA increased average size to 53% (n=88, significant $P < 0.001$).

C-E) Examples of the morphology of Purkinje cell dendritic trees. Purkinje cells were visualized by immunostaining for Calbindin D-28K. PMA-treatment (**C**, 50 nM for 7 days) caused stunted dendrites compared to the control cell (**C**). Inhibition of P/Q- and T-type Ca^{2+} channels along with PMA treatment (**E**) resulted in visibly more dendritic branches and a general morphology close to the control cell. Scale bar 50 μ m.

4.4.5. TRPC3^{-/-} and wild-type cultures were equally sensitive to dendritic growth inhibition mediated by DHPG or PMA

In order to study TRPC3 involvement in mediating dendritic growth inhibition after chronic activation of mGluR1 or PKC, we used mice lacking TRPC3 function (TRPC3^{-/-} mice). CB6 mice which were born within two days before or after the TRPC3^{-/-} mice were used as a wild type control because TRPC3^{-/-} mice were bred in a C57Bl6 background (Hartmann *et al.*, 2008). Organotypic slice cultures were treated for 7 days with either the PKC activator PMA or the group I mGluR activator DHPG, starting at DIV 2-4.

No difference in Purkinje cell morphology could be detected between CB6 and TRPC3^{-/-} mice in control cultures as well as in cultures treated with DHPG or PMA (Fig. 21, A-F). The average size of the dendritic tree areas in TRPC3^{-/-} cultures did not vary significantly from CB6 cultures. The DHPG treatment reduced the size of the dendritic tree areas to 46% in CB6 as well as in TRPC3^{-/-} cells compared to CB6 controls. The PMA treatment reduced the size of the dendritic tree areas to 30% in CB6, and to 28% in TRPC3^{-/-}, compared to CB6 control cells (Fig. 21, G and H). No difference was found in branch point counts and in experiments with increasing concentrations of either DHPG or PMA the minimal dose resulting in a dendritic growth inhibition was identical in wildtype and in TRPC3^{-/-} mice (Gugger and Kapfhammer, unpublished observations).

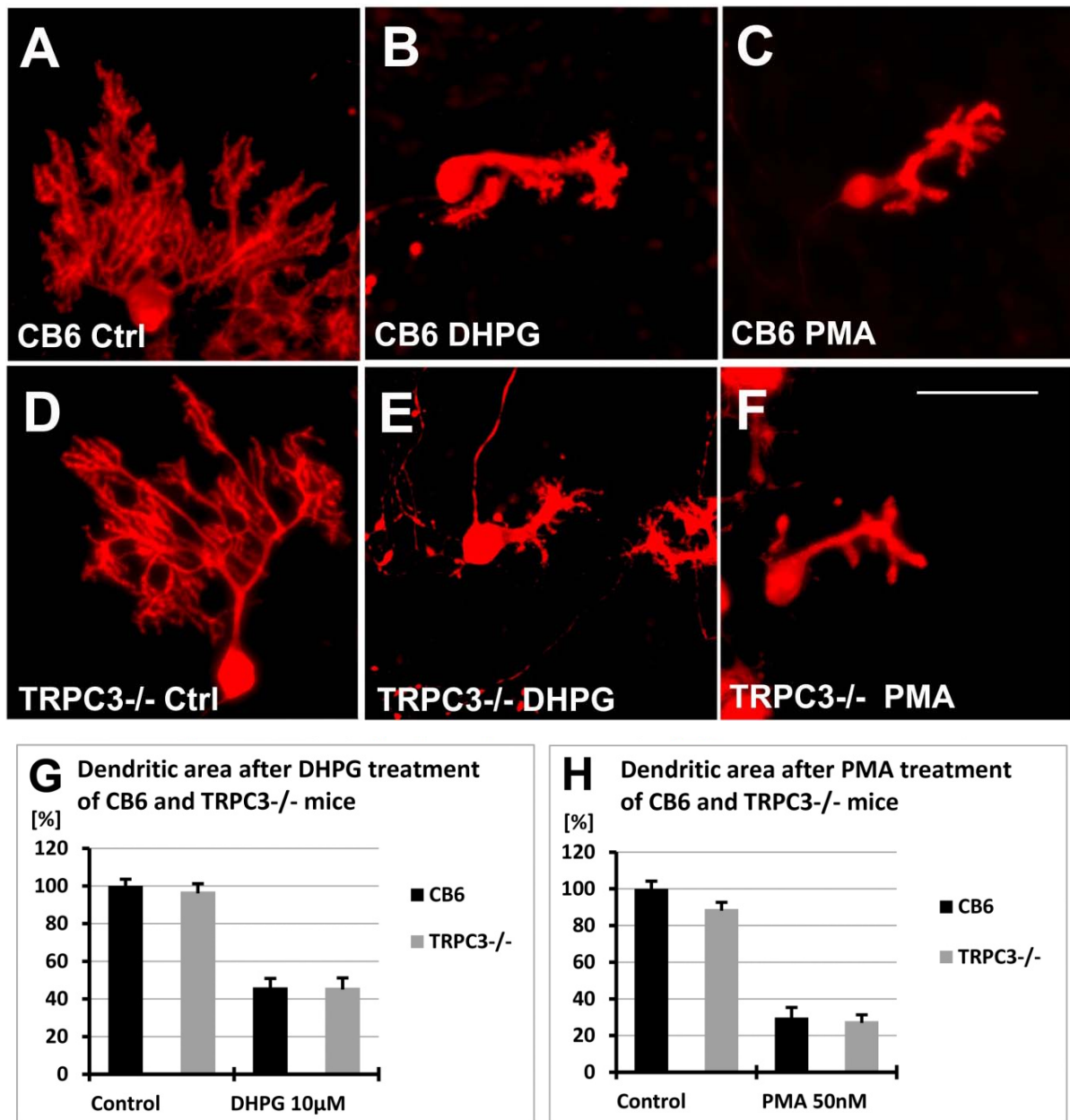


Figure 21: Dendritic trees after DHPG or PMA treatment in CB6 and TRPC3^{-/-} mice. A) – F) Staining of Calbindin D-28K shows Purkinje cells

A) – C) In cultures from CB6 mice, treatment with 10 μ M DHPG (B) or 50 nM PMA (C) limits dendritic growth compared to untreated cells (A).

D) – F) Cultures from TRPC3^{-/-} mice. No morphological difference to Purkinje cells from CB6 mice can be found. Untreated control culture (D). Treatment with 10 μ M DHPG (E) or 50 nM PMA (F) limits dendritic growth. Scale bar 50 μ m.

G) –H) Size of the area covered by a single dendritic tree. Standard size of CB6 control cells was rated 100%. Number of cells analyzed is given as n-value. All pharmacologically treated growth conditions were significantly smaller than control ($P < 0.001$). Error bars represent the SEM.

G) Cells were treated with DHPG (10 μ M) for 7 days. Average of dendritic tree size of TRPC3^{-/-} control Purkinje cells (n=57) reached 97% of CB6 controls (n=77, not significant). In cultures from CB6 mice, DHPG treatment reduced dendritic tree size to 46 % (n=91, significant $P < 0.001$) compared to CB6 control cells. In cultures from TRPC3^{-/-} mice, DHPG

reduced dendritic tree size to 46% (n=57, significant $P < 0.001$) compared to TRPC3^{-/-} control cells. In cultures from CB6 mice, PMA treatment reduced dendritic tree size to 30% (n=91, significant $P < 0.001$) compared to CB6 control cells. In cultures from TRPC3^{-/-} mice, PMA treatment reduced dendritic tree size to 28% (n=57, significant $P < 0.001$) compared to TRPC3^{-/-} control cells.

treatment reduced dendritic tree size to 46 % (n=92, significant P<0.001) compared to CB6 control cells.

H) Cells were treated with PMA (50 nM) for 7 days. Average of dendritic tree size of TRPC3^{-/-} control Purkinje cells (n=68) reached 89% of CB6 controls (n=62, not significant). In cultures from CB6 mice, PMA treatment reduced dendritic tree size to 30% (n=87, significant P<0.001) compared to CB6 control cells. In cultures from TRPC3^{-/-} mice, PMA treatment reduced dendritic tree size to 28% (n=76, significant P<0.001) compared to CB6 control cells.

4.4.6. Rescue of dendritic growth by the combined inhibition of P/Q- and T-type Ca²⁺ channels is not altered in the absence of TRPC3

Because TRPC3 forms Ca²⁺-permeable channels (Zitt et al., 1997; Kamouchi et al., 1999) an even increased rescue effect of T- and P/Q-type Ca²⁺ channel inhibition in cultures from TRPC3^{-/-} mice after mGluR1 or PKC activation was conceivable if TRPC3 channels were involved in the mGluR or PKC mediated signaling pathway. The same rescue experiments were carried out in cultures from TRPC3^{-/-} mice. Each single experiment was paralleled with cultures from CB6 mice which were born within 2 days. Organotypic slice cultures were treated for 7 days with DHPG (10 μM), starting at DIV 2-4. Alternatively, cultures were treated in the same way with DHPG and the P/Q-block, DHPG and the T-type Ca²⁺ channel inhibitor NNC 55-0396, or DHPG, P/Q-block and NNC 55-0396 combined.

As in cultures from CB6 mice, only the combination of P/Q- and T-type inhibition was able to increase the areas and branch point numbers of the dendritic trees significantly compared to cultures treated with DHPG alone. In TRPC3^{-/-} mice, the P/Q/T inhibition simultaneously with DHPG increased the dendritic areas to 65% of control cultures, compared to 47% in cultures which were treated only with DHPG. This constitutes a net increase of areas by 18% (Fig. 22 A). The number of branch points after DHPG treatment was 43% and after co-treatment with P/Q/T inhibitors 58%, which is an increase of 15% (Fig. 22 B). The results from the parallel experiments in CB6 mice were: DHPG only treated cultures reached dendritic areas of 50% of control, in combination with the P/Q/T-block they were increased to 66%. This constitutes a net increase of areas by 16% (Fig. 22 C). The number of branch points after DHPG treatment was 39% and after co-treatment with P/Q/T inhibitors 59%, which is an increase of 20% (Fig. 22 D). The dendritic areas of cells which were treated with either the P/Q-block or the T-type Ca²⁺ channel inhibitor NNC 55-0396

along with DHPG showed in both mouse strains only a slight trend of a rescue (Fig. 22, A-D). Taken together, there was no additional rescue effect by the absence of TRPC3 channels compared to treatment with the Ca²⁺ channel inhibitors alone.

Organotypic slice cultures were treated for 7 days with PMA (50 nM), starting at DIV 2-4. Alternatively, cultures were treated in the same way with PMA and the P/Q-block, PMA and the T-type Ca²⁺ channel inhibitor NNC 55-0396, or PMA, P/Q-block and NNC 55-0396 combined.

As in cultures from CB6 mice, all co-treatments were able to increase the areas and branch point numbers of the dendritic trees significantly compared to cultures treated with PMA alone, while the rescue effect in cultures co-treated with P/Q- and T-type-inhibitors combined was slightly stronger. In TRPC3^{-/-} mice, the PMA treatment resulted in reduced dendritic areas of 38% of controls. After co-treatment with NNC 55-0396 the areas were increased to 50%, after co-treatment with the P/Q-block to 47% and after co-treatment with P/Q- and T-type inhibition combined to 51% (Fig. 22 E). The number of branch points after PMA treatment was 25%, after co-treatment with NNC 55-0396 40%, after co-treatment with the P/Q-block 44% and after co-treatment with P/Q/T inhibitors combined 47% (Fig. 22 F). The results from the parallel experiments in CB6 mice were: PMA only treated cultures reached dendritic areas of 41% of control, after co-treatment with NNC 55-0396 49%, after co-treatment with the P/Q-block 52% and after co-treatment with P/Q/T inhibitors combined 62% (Fig. 22 G). The number of branch points after PMA treatment was 27%, after co-treatment with NNC 55-0396 39%, after co-treatment with the P/Q-block 43% and after co-treatment with P/Q/T inhibitors combined 50% (Fig. 22 H). Taken together, there was no additional rescue effect by the absence of TRPC3 channels compared to treatment with the Ca²⁺ channel inhibitors alone indicating that TRPC3 channels are not involved in the dendritic growth inhibition seen after mGluR1 or PMA stimulation.

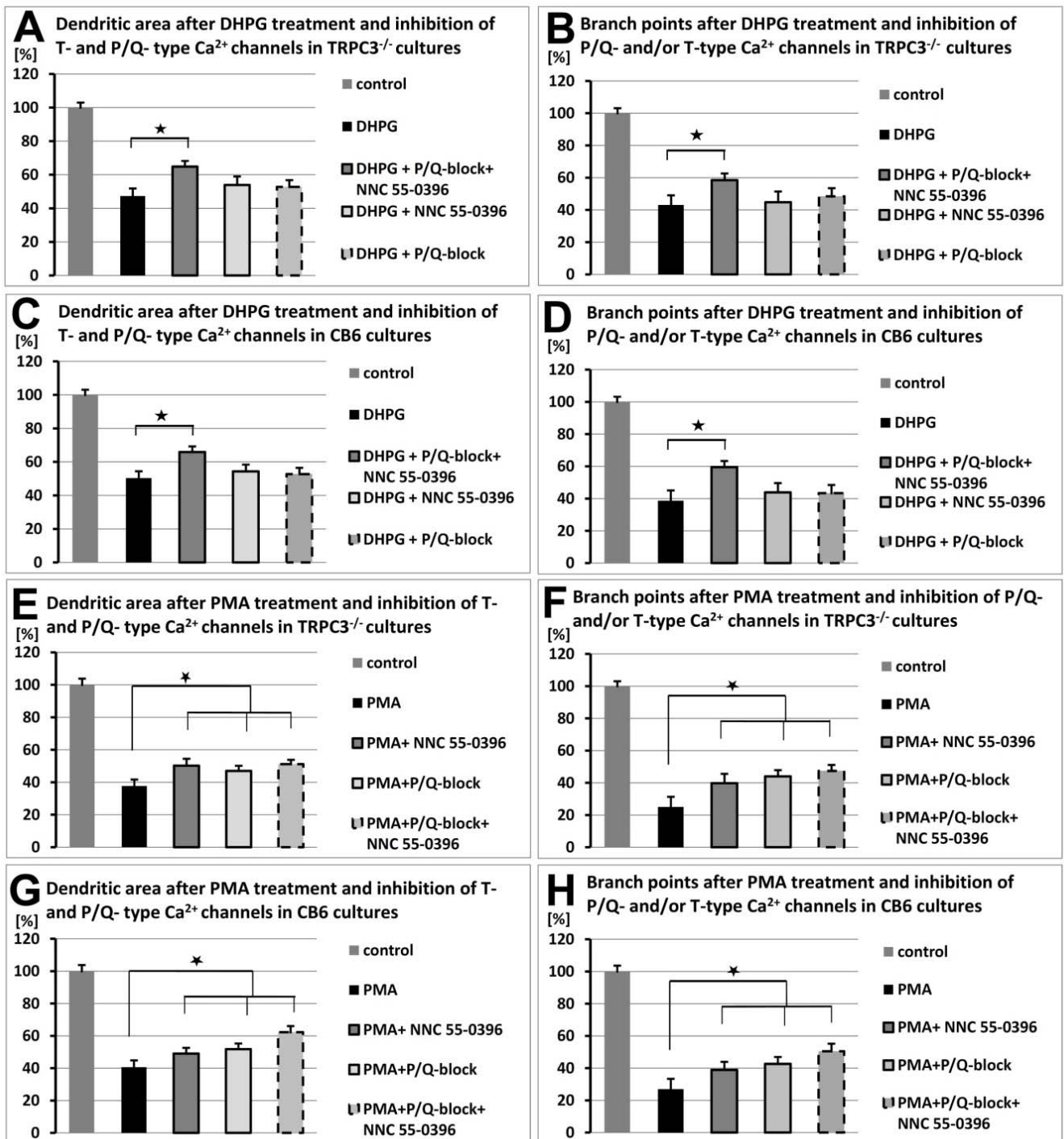


Figure 22: Dendritic trees after DHPG or PMA treatment and co-treatment with T-type and/or P/Q-type channel blockers in CB6 and $TRPC3^{-/-}$ mice. Size of the area covered by a single dendritic tree (A, C, E, G) and number of branch points (B, D, F, H). Cells were treated with DHPG (10 μ M) (A-D) or PMA (50 nM) (E-H) for 7 days either alone or in the presence of the P/Q-block and/or the T-type Ca^{2+} channel inhibitor NNC 55-0396. Standard growth of control cells was rated 100%. Star indicates statistical significance compared to DHPG- or PMA-only-treated cultures. Number of cells analyzed is given as n-value. All pharmacologically treated growth conditions were significantly smaller than control ($P < 0.001$). Error bars represent the SEM.

A) TRPC3^{-/-} cultures. DHPG treatment (n=99) reduced average dendritic area to 47% of untreated controls (n=77). After co-treatment with NNC-55-0396, average area was 54% (n=84, not significant) and after co-treatment with the P/Q-block 53% (n=96, not significant). Treatment with P/Q-block and NNC 55-0396 simultaneously with DHPG increased average size to 65% (n=93, significant P<0.001).

B) TRPC3^{-/-} cultures. DHPG treatment (n=93) reduced average dendritic branch point number to 43% of untreated controls (n=61). After co-treatment with NNC-55-0396, average branch point number was 45% (n=79, not significant) and after co-treatment with the P/Q-block 48% (n=88, not significant).

Treatment with P/Q-block and NNC 55-0396 simultaneously with DHPG increased average branch point number to 58% (n=88, significant P<0.01).

C) CB6 cultures. DHPG treatment (n=99) reduced average dendritic area to 50% of untreated controls (n=76). After co-treatment with NNC-55-0396, average area was 54% (n=93, not significant) and after co-treatment with the P/Q-block 53% (n=103, not significant). Treatment with P/Q-block and NNC 55-0396 simultaneously with DHPG increased average size to 66% (n=103, significant P<0.001).

D) CB6 cultures. DHPG treatment (n=91) reduced average dendritic branch point number to 39% of untreated controls (n=63). After co-treatment with NNC-55-0396, average branch point number was 44% (n=87, not significant) and after co-treatment with the P/Q-block 43% (n=87, not significant).

Treatment with P/Q-block and NNC 55-0396 simultaneously with DHPG increased average branch point number to 59% (n=89, significant P<0.001).

E) TRPC3^{-/-} cultures. PMA treatment (n=85) reduced average dendritic area to 38% of untreated controls (n=78). After co-treatment with NNC-55-0396, average area was 50% (n=70, significant P<0.01) and after co-treatment with the P/Q-block 47% (n=97, significant P<0.05).

Treatment with P/Q-block and NNC 55-0396 simultaneously with PMA increased average size to 51% (n=92, significant P<0.001).

F) TRPC3^{-/-} cultures. PMA treatment (n=86) reduced average dendritic branch point number to 25% of untreated controls (n=68). After co-treatment with NNC-55-0396, average branch point number was 40% (n=92, significant P<0.01) and after co-treatment with the P/Q-block 44% (n=97, significant P<0.001).

Treatment with P/Q-block and NNC 55-0396 simultaneously with PMA increased average branch point number to 47% (n=89, significant P<0.001).

G) CB6 cultures. PMA treatment (n=89) reduced average dendritic area to 41% of untreated controls (n=80). After co-treatment with NNC-55-0396, average area was 49% (n=93, significant P<0.05) and after co-treatment with the P/Q-block 52% (n=68, significant P<0.01). Treatment with P/Q-block and NNC 55-0396 simultaneously with PMA increased average size to 62% (n=78, significant P<0.001).

H) CB6 cultures. PMA treatment (n=87) reduced average dendritic branch point number to 27% of untreated controls (n=75). After co-treatment with NNC-55-0396, average branch point number was 39% (n=93, significant P<0.01) and after co-treatment with the P/Q-block 43% (n=66, significant P<0.001).

Treatment with P/Q-block and NNC 55-0396 simultaneously with PMA increased average branch point number to 50% (n=74, significant P<0.001).

4.4.7. Selective pharmacological inhibition of TRPC3 channels did not alter DHPG- or PMA-mediated dendritic growth inhibition or the rescue effects obtained by combined blockade of P/Q- and T-type Ca²⁺ channels

Chronic loss of TRPC3 function in TRPC3^{-/-} mice may be compensated by overexpression of other TRPC channel types. Therefore, we carried out the same rescue experiments in CB6 mice with an acute TRPC3 blockade by application of Pyr3, a selective pharmacological inhibitor of TRPC3 channels (Kiyonaka *et al.*, 2009).

Organotypic slice cultures were treated for 7 days with DHPG (10 μM) or PMA (50 nM), starting at DIV 2-4. Alternatively, cultures were treated in the same way with DHPG or PMA and the TRPC3 inhibitor Pyr3 (3 μM), DHPG or PMA and the combination of P/Q-block and T-type inhibitor NNC 55-0396 (1 μM), or DHPG or PMA, the P/Q- and T-type inhibitory block and Pyr3 combined.

Application of Pyr3 alone slightly reduced dendritic areas to 97 % of untreated control cultures and raised dendritic branch points to 101 %, which was statistically not significant (Fig. 23 A, B). Thus, TRPC3 inhibition had no major influence on dendritic tree size.

DHPG treatment resulted in reduced dendritic areas of 48 % and average branch point numbers of 42 %. Co-treatment with Pyr3 further reduced dendritic areas to 44 % and branch point number to 39 %. Co-treatment with the rescue-combination of P/Q-block and T-type inhibitor increased dendritic areas to 63 % and branch point number to 58 %. Additional co-treatment with Pyr3 to the P/Q- and T-type inhibitory block resulted in dendritic areas of 54% and dendritic branch points of 55% of control, which was somewhat less than with P/Q- and T-type inhibitory block alone (Fig. 23).

PMA treatment resulted in reduced dendritic areas of 36 % and average branch point numbers of 25 %. Co-treatment with Pyr3 further reduced dendritic areas to 29 % and branch point number to 17 %. Co-treatment with the rescue-combination of P/Q-block and T-type inhibitor increased dendritic areas to 49 % and branch point number to 43 %. Additional co-treatment with Pyr3 to the P/Q- and T-type inhibitory block resulted in dendritic areas and branch points of 38 %, again a reduced rescuing effect compared to P/Q- and T-type inhibitory block alone (Fig. 23).

Taken together, selective pharmacological inhibition of TRPC3 channels in CB6 mice had no significant influence on the effects of DHPG or PMA treatment and did not

convey an additional rescue effect to the combination of P/Q- and T-type inhibitors. This is in line with our results obtained with TRPC3^{-/-} mice.

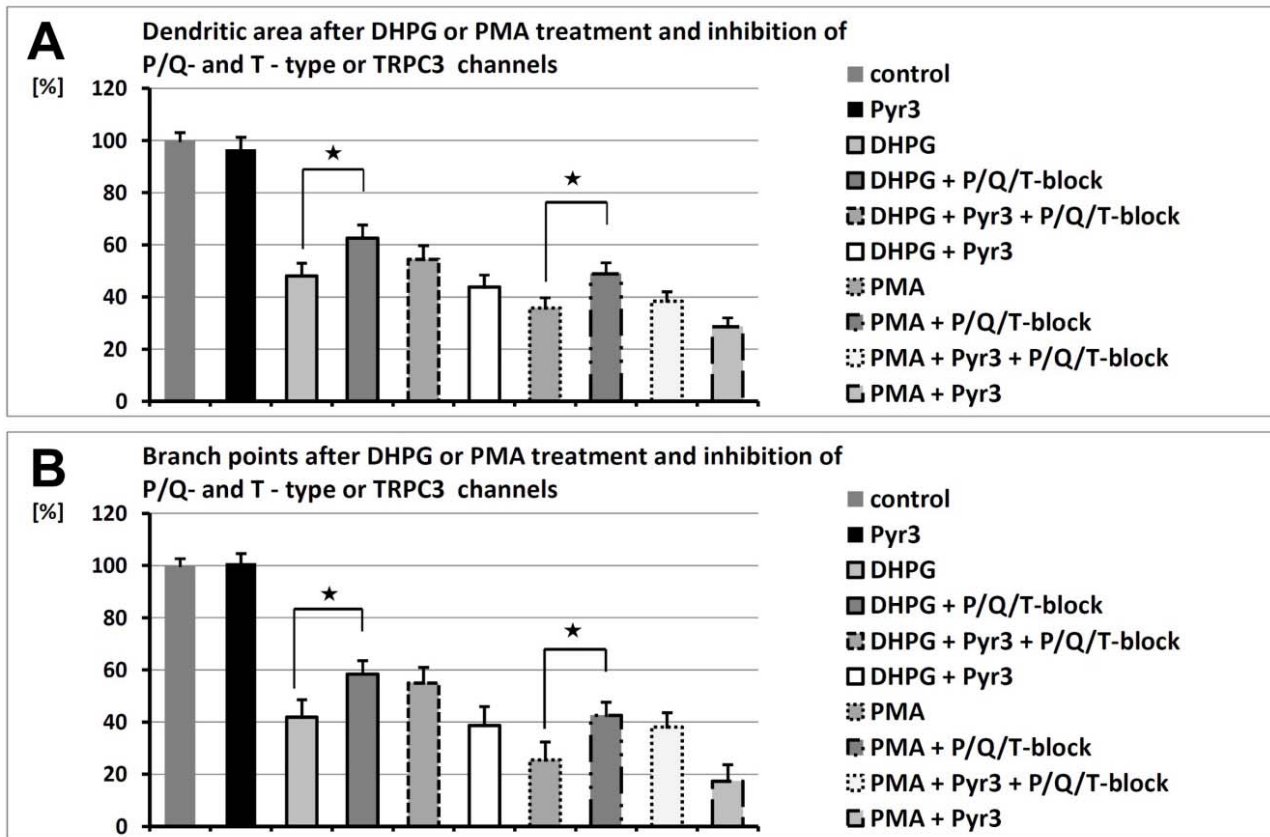


Figure 23: Dendritic trees after DHPG or PMA treatment and co-treatment with T/P/Q-type channel blockers and/or TRPC3 inhibitor pyr3. Size of the area covered by a single dendritic tree (A) and number of branch points (B). Cells were treated with the TRPC3 inhibitor Pyr3 (3 μ M) alone or with DHPG (10 μ M) or PMA (50 nM) for 7 days alone or in the presence of Pyr3 and/or the P/Q/T-block (including the T-type Ca²⁺ channel inhibitor NNC 55-0396). Standard growth of control cells was rated 100%. Star indicates statistical significance compared to DHPG- or PMA-only-treated cultures. Number of cells analyzed is given as n-value. All pharmacologically treated growth conditions were significantly smaller than control ($P < 0.001$), except for the condition Pyr3 alone, which was not significantly different from control. Error bars represent the SEM.

A) Treatment with Pyr3 (n=58) slightly reduced dendritic areas to 97% of untreated controls (n=107).

DHPG treatment reduced average dendritic area to 48% (n=91). After co-treatment with Pyr3 average area was 44% (n=70, not significant to DHPG), after co-treatment with the P/Q/T-block 63% (n=74, significant to DHPG $P < 0.05$) and after co-treatment with the P/Q/T-block and Pyr3 combined 54% (n=72, not significant to DHPG).

PMA treatment reduced average dendritic area to 36% (n=91). After co-treatment with Pyr3 average area was 29% (n=73, not significant to PMA), after co-treatment with the P/Q/T-block 49% (n=73, significant to PMA $P < 0.01$) and after co-treatment with the P/Q/T-block and Pyr3 combined 38% (n=70, not significant to PMA).

B) Treatment with Pyr3 (n=55) slightly increased dendritic branch point number to 101% of untreated controls (n=103).

DHPG treatment reduced average branch point number to 42% (n=91). After co-treatment with Pyr3 average branch point number was 39% (n=69, not significant to DHPG), after co-treatment with the P/Q/T-block 58% (n=74, significant to DHPG $P < 0.05$) and after co-treatment with the P/Q/T-block and Pyr3 combined 55% (n=71, not significant to DHPG). PMA treatment reduced average branch point number to 25% (n=92). After co-treatment with Pyr3 average branch point number was 17% (n=74, not significant to PMA), after co-treatment with the P/Q/T-block 43% (n=73, significant to PMA $P < 0.001$) and after co-treatment with the P/Q/T-block and Pyr3 combined 38% (n=70, not significant to PMA).

4.4.8. Inhibition of Phospholipase C, GluR δ 2 - or R-type Ca $^{2+}$ channels did not increase the rescue effects obtained by combined inhibition of P/Q- and T-type Ca $^{2+}$ channels

Co-treatment with a combination of P/Q- and T-type inhibitors partially rescued the dendrite-limiting effects of both, chronic mGluR1 or PKC stimulation. Yet, the rescued Purkinje cell dendritic trees did not reach the average size of untreated control cells. After mGluR1- or PKC stimulation with DHPG or PMA, respectively, calcium can enter the cytoplasm through GluR δ 2 - or R-type Ca $^{2+}$ channels (Meacham *et al.*, 2003; Cavalier *et al.*, 2008; Zanjani *et al.*, 2009) or could be released from internal stores (Canepari & Ogden, 2006) in addition to influx through P/Q-, T-, and N-type channels, which are effectively blocked by our combination of ω -Agatoxin IVA, ω -Conotoxin MVIIC and NNC 55-0396 (Mintz *et al.*, 1992a; McDonough *et al.*, 2002; Huang *et al.*, 2004). We investigated whether additional inhibition of these other sources of calcium entry into the cytoplasm would increase the rescue effect.

Organotypic slice cultures were treated for 7 days with DHPG (10 μ M) or PMA (50 nM), either with co-treatment of the P/Q/T-inhibitory combination or without, starting at DIV 2-4. Alternatively, cultures were co-treated with the R-type Ca $^{2+}$ channel inhibitor SNX-482 (500 nM) (Newcomb *et al.*, 1998), the open channel blocker 1-naphthyl-acetyl spermine (Nasp) which blocks influx through GluR δ 2 channels (Zanjani *et al.*, 2009) or the phospholipase C (PLC) inhibitor U73122 (Cruzblanca *et al.*, 1998). PLC activation leads to release of Ca $^{2+}$ from internal stores (Canepari & Ogden, 2006) and sole PLC inhibition by U73122 has been shown to be ineffective in rescuing the dendrite limiting effect mediated by DHPG (Sirzen-Zelenskaya *et al.*, 2006). A contribution of Ca $^{2+}$ from internal stores, however, might increase the rescue effect obtained by combined P/Q- and T-type inhibition.

Treatment of slice cultures for 7 days with SNX-482, Nasp or U73122 alone did not significantly influence dendritic size of Purkinje cells. SNX-482 (500 nM) increased dendritic areas to 101 % and reduced average branch point number to 98% compared to untreated controls. Nasp (100 μ M) reduced dendritic areas to 98% and average branch point number to 90 % and U73122 reduced dendritic areas to 98% and average branch point number to 88 % (Fig. 24 A, B).

DHPG treatment resulted in reduced dendritic areas of 52 % and an average branch point number of 43 % of controls. Co-treatment with SNX-482 further reduced average areas to 43 % and branch point number to 36 %. Co-treatment with Nasp resulted with an average area of 60 % and an average branch point number of 43 % in slightly increased dendritic trees, but the effect was not significant. The combination of P/Q-and T-type inhibitors significantly increased average dendritic areas to 74 % and branch point number to 62 %. Co-treatment with the P/Q/T-block and SNX-482 resulted in average dendritic areas of 64 % and branch point number of 63 %. Co-treatment with the P/Q/T-block and Nasp yielded dendritic areas of 68 % and a branch point number of 57 %. Co-treatment with the P/Q/T-block and U73122 resulted in dendritic areas of 62 % and branch point number of 53 % (Fig. 24 A, B). Thus, DHPG-mediated dendritic growth inhibition was not rescued by co-treatments with SNX-482 or Nasp, and the partial rescue obtained by P/Q- and T-type inhibition was not further increased by co-treatments with SNX-482, Nasp or U73122.

PMA treatment resulted in dendritic areas of 35 % and average branch point number of 25 % of controls. Co-treatment with SNX-482 reduced average areas to 36 % and branch point number to 23 %. Co-treatment with Nasp reduced average areas to 36 % and branch point number to 22 %. The combination of P/Q-and T-type inhibitors significantly increased average dendritic areas to 48 % and branch point number to 42 %. Co-treatment with the P/Q/T-block and Nasp significantly increased average dendritic areas to 45 % and branch point number to 37 %. Co-treatment with the P/Q/T-block and U73122 gave dendritic areas of 37 % and branch point number of 36 %. Dendritic areas were significantly smaller than in the co-treatment with the P/Q/T-block only. Co-treatment with the P/Q/T-block and SNX-482 yielded dendritic areas of 42 %, and branch point number of 37 % (Fig. 24 C, D). Thus, PMA-mediated dendritic growth inhibition was not rescued by co-treatments with SNX-482 or Nasp, and the partial rescue obtained by P/Q- and T-type inhibition was not further increased by co-treatments with SNX-482, Nasp or U73122.

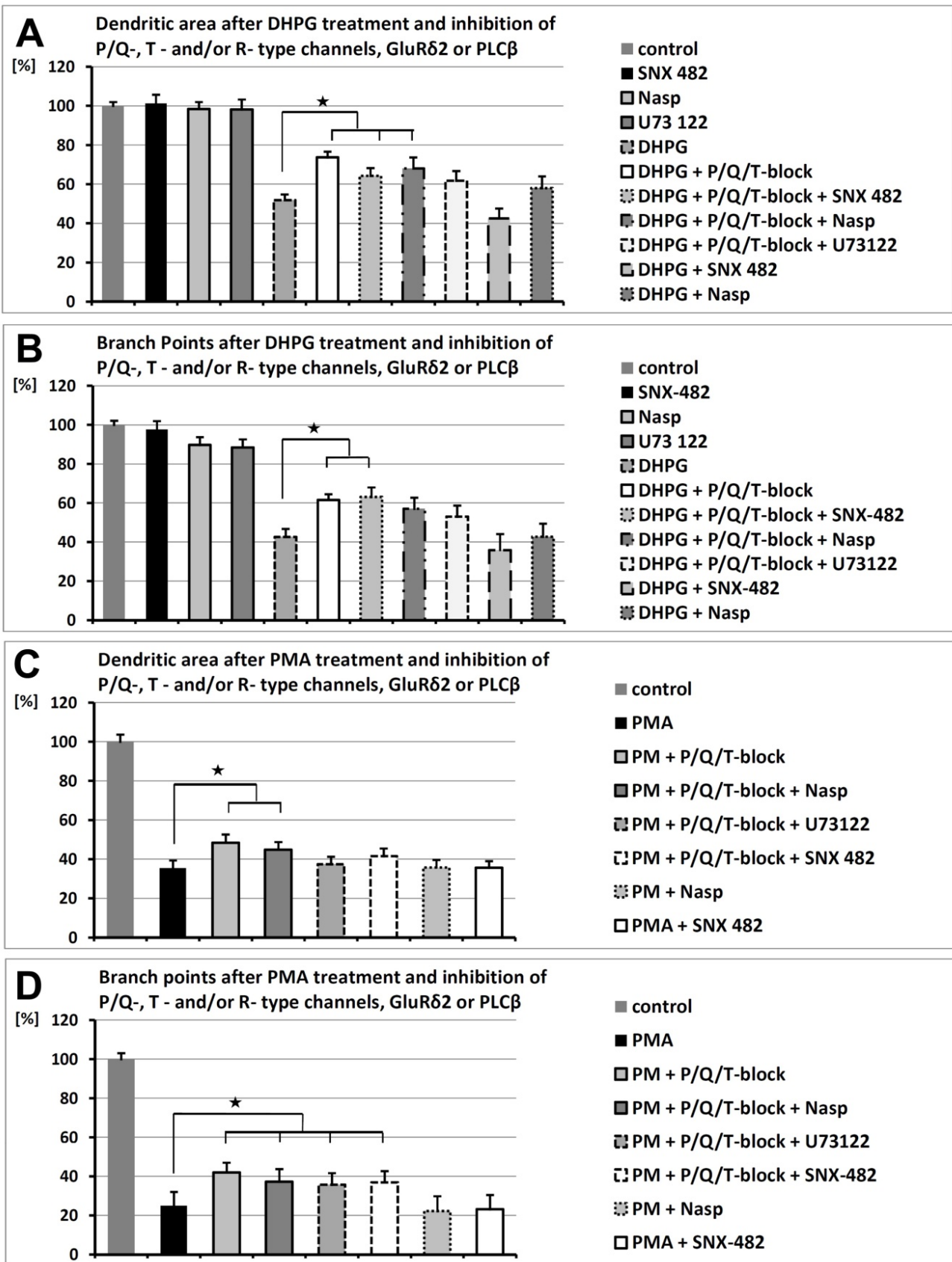


Figure 24: Dendritic trees after DHPG or PMA treatment and co-treatment with T/P/Q-type channel blockers and/or inhibitors of R-type channels, GluR δ 2 or PLC. Size of the average area covered by a single dendritic tree (A, C) and average number of branch points (B, D). Cells were treated for 7 days with SNX-482 (500 nM), Nasp (100 μ M), U73122 (10

μM) or DHPG (10 μM) alone, or with DHPG in the presence of SNX-482, Nasp or the P/Q/T-block (including the T-type Ca²⁺ channel inhibitor NNC 55-0396), or alternatively with DHPG, the P/Q/T-block and SNX-482, Nasp or U73122 combined (A, B). Cells were treated for 7 days PMA (50 nM) alone, or with PMA in the presence of SNX-482, Nasp or the P/Q/T-block, or alternatively with PMA, the P/Q/T-block and SNX-482, Nasp or U73122 combined (C, D) Standard growth of control cells was rated 100%. Star indicates statistical significance compared to DHPG- or PMA-only-treated cultures. Number of cells analyzed is given as n-value. All pharmacologically treated growth conditions were significantly smaller than control (P<0.001), except for the conditions SNX-482, Nasp and U73122 alone, which were not significantly different from control. Error bars represent the SEM.

A) Treatment with SNX-482 (n=66) slightly increased dendritic areas to 101% of untreated controls (n=219), Nasp (n=65) and U73122 (n=61) both reduced areas to 98%.

DHPG treatment reduced area to 52% (n=225). After co-treatment with SNX-482 area was 43% (n=67, not significant to DHPG), after co-treatment with Nasp 60% (n=75, not significant to DHPG), after co-treatment with the P/Q/T-block 74% (n=177, significant to DHPG P<0.001), after co-treatment with the P/Q/T-block and SNX-482 combined 64% (n=73, significant to DHPG P<0.05), after co-treatment with the P/Q/T-block and Nasp combined 68% (n=67, significant to DHPG P<0.01) and after co-treatment with the P/Q/T-block and U73122 combined 62% (n=75, not significant to DHPG).

B) Treatment with SNX-482 (n=63) slightly reduced dendritic branch point number to 98% of untreated controls (n=208), Nasp (n=64) to 90% and U73122 (n=59) to 88%.

DHPG treatment reduced branch points to 43% (n=216). After co-treatment with SNX-482 branch point number was 36% (n=67, not significant to DHPG), after co-treatment with Nasp 43% (n=71, not significant to DHPG), after co-treatment with the P/Q/T-block 61% (n=176, significant to DHPG P<0.001), after co-treatment with the P/Q/T-block and SNX-482 combined 63% (n=71, significant to DHPG P<0.001, not significant to DHPG with P/Q/T-inhibition alone), after co-treatment with the P/Q/T-block and Nasp combined 57% (n=66, not significant to DHPG) and after co-treatment with the P/Q/T-block and U73122 combined 53% (n=74, not significant to DHPG).

C) PMA treatment reduced area to 36% (n=91). After co-treatment with SNX-482 (n=71) or Nasp (n=65) area was unchanged 36%, after co-treatment with the P/Q/T-block 48% (n=73, significant to PMA P<0.001), after co-treatment with the P/Q/T-block and Nasp combined 45% (n=66, significant to PMA P<0.01), after co-treatment with the P/Q/T-block and U73122 combined 37% (n=66, not significant to PMA) and after co-treatment with the P/Q/T-block and SNX-482 combined 42% (n=64, not significant to PMA).

D) PMA treatment reduced branch points to 25% (n=92). After co-treatment with SNX-482 (n=71) branch point number was 23% and after co-treatment with Nasp (n=65) 22%. After co-treatment with the P/Q/T-block branch point number was 42% (n=73, significant to PMA P<0.001), after co-treatment with the P/Q/T-block and Nasp combined 37% (n=65, significant to PMA P<0.01), after co-treatment with the P/Q/T-block and U73122 combined 36% (n=66, significant to PMA P<0.05) and after co-treatment with the P/Q/T-block and SNX-482 combined 37% (n=63, significant to PMA P<0.01).

4.5. Discussion

In this study, we have analyzed the contribution of Ca^{2+} permeable channels for the inhibition of Purkinje cell dendritic growth after mGluR1 and after PKC stimulation. We have found that both, P/Q-type and T-type Ca^{2+} channels are involved in the inhibition of dendritic growth, because blockade of P/Q-type and T-type Ca^{2+} channels simultaneous with mGluR1 stimulation resulted in a partial rescue of the dendritic morphology. For PKC-stimulation, blockade of either P/Q-type or T-type Ca^{2+} channels was sufficient to achieve a rescue effect which was maximal after blockade of both types of channels. These findings imply that Ca^{2+} entry through voltage-gated Ca^{2+} channels is required for the inhibitory effects on dendritic growth. In contrast, absence or acute blockade of TRPC3 channels had no rescue effect indicating that Ca^{2+} entry through TRPC3 channels is not involved in the inhibition of Purkinje cell dendritic growth. Similarly, blockade of Ca^{2+} influx through GluRdelta2 channels or inhibition of release from intracellular stores did not influence mGluR1- and PKC-mediated inhibition of Purkinje cell dendritic growth. Thus, both P/Q-type and T-type Ca^{2+} channels, but not TRPC3 or other Ca^{2+} -permeable channels, are involved in the regulation Purkinje cell dendritic growth after chronic mGluR1 or PKC activation.

4.5.1. TRPC3 channels are not required for inhibition of dendritic growth of Purkinje cells

TRPC3 channels have an important role in mGluR1 signaling in Purkinje cells by mediating the so-called slow EPSP (Batchelor & Garthwaite, 1993; Canepari *et al.*, 2001). In mice which are deficient for the TRPC3 channels in Purkinje cells, the slow EPSC is completely absent after mGluR1 stimulation, whereas the mGluR1-mediated Ca^{2+} release from internal stores is not affected (Hartmann *et al.*, 2008). Because in previous experiments we have shown that the mGluR1-mediated inhibition of Purkinje cell dendritic growth is unlikely to be caused by Ca^{2+} release from internal stores (Sirzen-Zelenskaya *et al.*, 2006) the TRPC3 channel would be an attractive candidate for mediating this growth inhibitory effect. It is highly expressed on Purkinje cell dendrites during dendritic growth and it remains strongly expressed into adulthood (Huang *et al.*, 2007). Furthermore, in a recent publication it was shown that in the *moonwalker (mwk)* mouse mutant a facilitated opening of the TRPC3 channel

after sub-threshold mGluR1 stimulation results in a robust cerebellar phenotype with some loss of Purkinje cells and a pronounced reduction of the Purkinje cell dendritic tree (Becker et al., 2009). Interestingly, TRPC3 channels are inhibited by phosphorylation by Protein Kinase C, creating a possible link to the effects of PKC γ activation of Purkinje cell dendritic growth (Metzger & Kapfhammer, 2000; Schrenk et al., 2002). We have tested for an involvement of TRPC3 in mGluR1- and PKC-mediated dendritic growth inhibition by experiments with mice lacking TRPC3 channels (TRPC3^{-/-}) (Hartmann et al., 2008). Purkinje cells in cultures derived from TRPC3^{-/-}-mice reacted to mGluR1 or PKC-stimulation in exactly the same way as Purkinje cells in control cultures. Furthermore, experiments with an acute pharmacological blockade of TRPC3 channels yielded very similar results making functional compensation by other types of TRPC channels rather unlikely. Further evidence against compensation by other TRPC receptors comes from experiments with the unspecific TRPC inhibitor SKF 96365 which did not alter the dendritic growth inhibition by mGluR1 or PKC γ activation (see supporting information, Fig. 25). These experiments exclude that activation of the TRPC3 channels is required for the inhibitory effect on dendritic growth of Purkinje cells.

4.5.2. Inhibition of Purkinje cell dendritic growth involves Ca²⁺ entry through voltage gated channels

It is well known that Ca²⁺ signals play an important role in neuronal development by modulating dendritic growth and branching (Konur & Ghosh, 2005). There is also evidence that suggests the mGluR and PKC mediated inhibition of dendritic growth could be mediated by changes of the intracellular Ca²⁺ signaling. In two independent mouse models with increased Ca²⁺ entry into Purkinje cells also strongly reduced dendritic trees are found, similar to the dendritic trees seen after chronic mGluR1- or PKC-activation. In the *lurcher* mouse, a constitutively open Ca²⁺ permeable cation channel, the glutamate receptor $\delta 2$ (GluR $\delta 2$) channel, leads to disrupted Purkinje cell development followed by cell death (Zuo et al., 1997). Purkinje cells in organotypic cultures derived from *lurcher* mice have a much reduced dendritic tree. The growth of the dendritic tree can be rescued by blocking of cation channels with Naphtyl-acetyl spermine (Nasp, (Zanjani et al., 2009). Interestingly, Nasp was ineffective in mediating the growth inhibitory effects of mGluR1 or PKC-stimulation excluding that

they are mediated by the GluR δ 2 channel. Taken together with the similar phenotype of the *mwk* mice (Becker et al., 2009), see above) with a leaky TRPC3 channel it is very likely that a chronic Ca²⁺ overload of Purkinje cells during the period of dendritic development will result in inhibition of dendritic growth.

While we have excluded that Ca²⁺ influx either through TRPC3 channels or GluR δ 2 channels are involved in mGluR1 or PKC mediated dendritic growth inhibition of Purkinje cells we here provide evidence for an involvement of voltage dependent Ca²⁺ channels (VDCC) of the P/Q-type and T-type, but not of the R-type. The P/Q-type and T-type channels are responsible for up to 95% of the total Ca²⁺ currents in Purkinje cells (Usowicz et al., 1992; Watanabe et al., 1998; Swensen & Bean, 2003; Isope & Murphy, 2005; Isope et al., 2010). A direct interaction and functional coupling between mGluR1 and P/Q-type Ca²⁺ channels in Purkinje cell dendritic spines has been shown to result in a potentiation of Ca²⁺ influx (Kitano et al., 2003).

The T-type Ca²⁺ channels of the Ca_v3.1 isoform have been shown to be highly enriched in Purkinje cell spines where they co-localize with mGluR1 (Hildebrand et al., 2009). Furthermore, they are potentiated upon mGluR1 activation (Hildebrand et al., 2009), an effect which is mediated by a phospholipase C-independent, tyrosine-phosphatase-dependent pathway (Hirono et al., 1998; Canepari et al., 2004; Hildebrand et al., 2009). We have shown here that pharmacological blockade of these channels results in a rescue of the Purkinje cell dendritic tree after mGluR1 or PKC stimulation indicating that Ca²⁺ influx through these channels is critical for the inhibitory effect on dendritic growth. The observed rescue effect did not restore the full dendritic tree size. This may indicate that additional pathways are involved, but could also simply reflect an incomplete blocking activity of the channel blockers. In our experience the chronic activation of either mGluR1 receptors by DHPG or PKC activity by PMA over several days is an extremely powerful stimulant and will be very difficult to block by any drug directed at the downstream signaling pathways. Interestingly, there are subtle differences between mGluR1 and PKC-mediated inhibition of dendritic growth. For PKC activation, blockade of either P/Q-type channels or T-type channels was effective in providing a rescue of Purkinje cell dendrites, though combined blockade was the most effective. For mGluR1-activation the combined blockade of both channels was necessary to obtain a significant rescue effect. This indicates that, although in both cases P/Q-type and T-type channels are

involved, there are also differences in the signaling pathways following mGluR1 and PKC activation.

4.5.3. Pharmacological compounds blocking T-type Ca²⁺ channels are toxic to Purkinje cell axons

Unexpectedly, we found a consistent toxic effect on Purkinje cell axons with the use of the T-type channel blockers mibefradil and NNC 55-0396 when applied in concentrations higher than used in this study (Fig. 18). An even stronger toxic effect on Purkinje cell axons was found with the compound SK&F 96365 (SKF, see chapter 5, section 5.1., Fig. 26). Recently, SKF has been shown to be a potent blocker of low-voltage-activated (LVA) and high-voltage-activated (HVA) Ca²⁺ channels, with an especially high affinity for LVA T-type Ca²⁺ channels (Singh et al., 2010). Interestingly, there is evidence that T-type Ca²⁺ channels might indeed be present in the axon initial segment of Purkinje cells (Bender & Trussell, 2009). In cultured hippocampal neurons, activation of T-type Ca²⁺ channels induced relocation of the axon initial segment (Grubb & Burrone, 2010). It is not clear at the moment whether the axonotoxic activity of high concentrations of mibefradil and NNC55-0396 or SKF is due to blockade of T-type channels or rather due to an unknown side effect of these drugs. Further investigations on the role of voltage-gated Ca²⁺ channels in axons will be required to clarify this point.

4.5.4. The role of Ca²⁺ homeostasis for Purkinje cell dendritic growth

The observation that Purkinje cell dendritic growth is severely compromised after chronic activation of mGluR1 or PKC links electrical activity to the regulation of dendritic growth and might constitute a stop mechanism for the expansion of the Purkinje cell dendritic tree (Sirzen-Zelenskaya et al., 2006). The signaling pathways involved in this effect are of interest because they provide a link between the functional status of the cell and dendritic growth processes. Furthermore, pharmacological manipulation of several classical pathways (IP3/DAG pathway, PI3K/AKT/mTOR pathway, MAPK/ERK pathway) was not effective to rescue Purkinje cell dendritic morphology after either mGluR1 or PKC activation (Sirzen-Zelenskaya et al., 2006). While the initial steps of the signaling chain after mGluR or PKC

activation are still not clear, the findings presented in this study demonstrate that they will eventually result in an increased influx of Ca^{2+} ions into Purkinje cell dendrites through voltage-dependent Ca^{2+} channels. Whether there is also a contribution from intracellular Ca^{2+} stores is not known, but pharmacological blockade of IP3-R sensitive stores showed no rescue effect. The inhibition of Purkinje cell dendritic growth seen after chronic stimulation of mGluR1 or PKC shares many similarities with that observed in *lurcher* (Zanjani et al., 2009) or in *mwk* mutant mice (Becker et al., 2009). The *lurcher* mutation in the GluR δ 2 channel, the *mwk* mutation in the TRPC3 channel and the activation of mGluR1 or PKC clearly have in common a chronic long-lasting overload of Purkinje cells with Ca^{2+} ions entering through channels in the dendritic plasma membrane eventually compromising dendritic expansion. This mechanism is likely to be an important regulatory component of dendritic growth of cerebellar Purkinje cells.

Acknowledgements:

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4.6. Supporting information

To exclude the possibility that overexpression of other TRPC channel types might compensate for the chronic loss of TRPC3 function in TRPC3^{-/-} mice, we treated TRPC3^{-/-} mice with the unspecific TRPC channel inhibitor SKF 96365 (Kress *et al.*, 2008) in the presence of either DHPG or PMA. As visible in the diagrams in Figure 25, SKF treatment (25µM) had no significant influence on Purkinje cell dendritic growth and co-treatment with SKF (25µM) of cultures challenged with DHPG or PMA did not provide a rescue effect in cultures from both, CB6 and TRPC3^{-/-} mice. These results corroborate our findings with Pyr3, a selective pharmacological inhibitor of TRPC3 channels (Kiyonaka *et al.*, 2009).

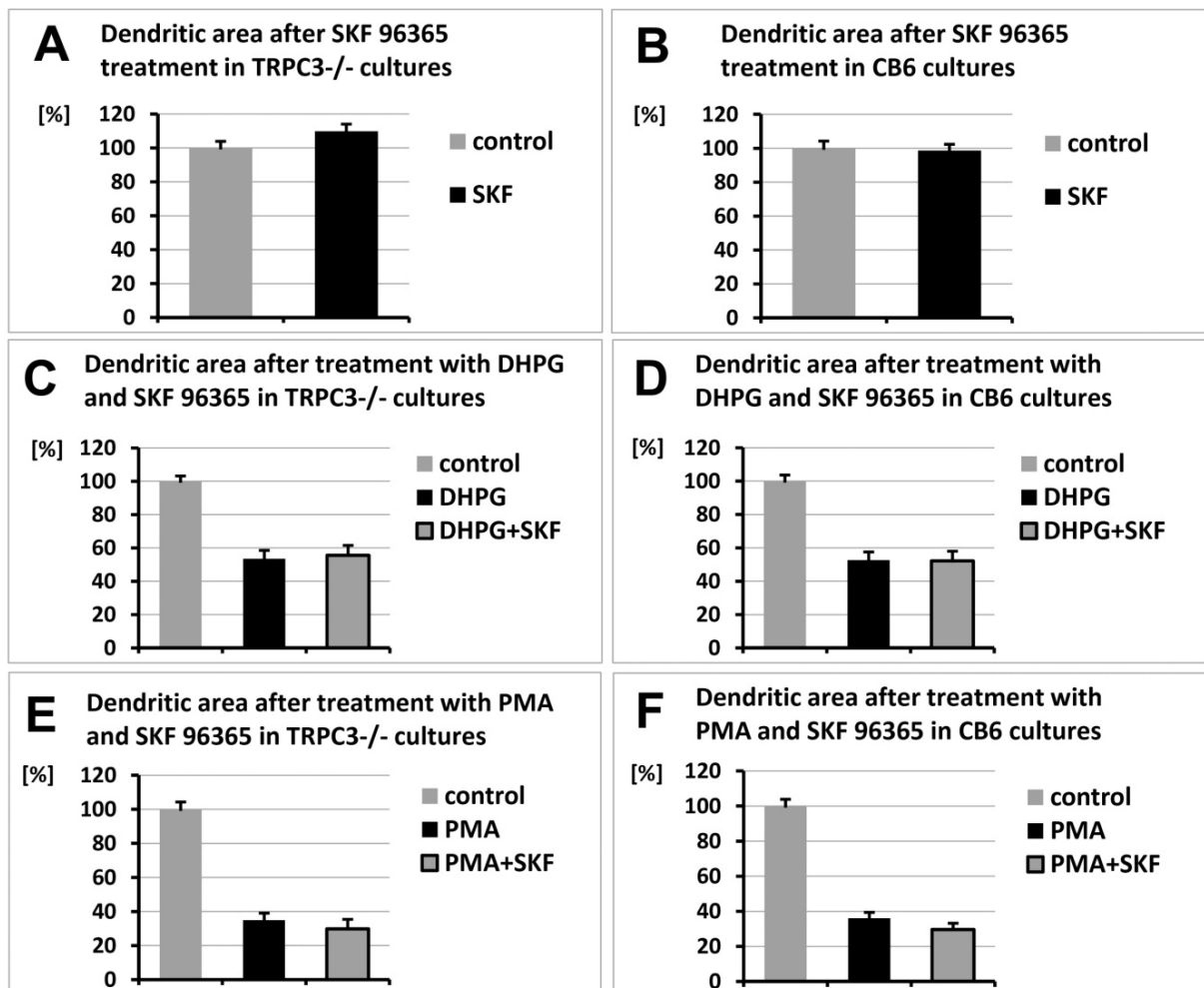


Figure 25: Dendritic trees after DHPG or PMA treatment and co-treatment with the TRPC inhibitor SKF 96365. Size of the area covered by a single dendritic tree. Cells were treated with the TRPC channel inhibitor SKF 96365 (25 µM) for 7 days either alone (A, B), in the presence of DHPG (10 µM) (C, D) or in the presence of PMA (100 nM) (E, F). The experiments performed in cultures from TRPC3^{-/-} mice (A, C, E) were paralleled with cultures from CB6 mice (B, D, F) born within two days before or after the TRPC3^{-/-} mice, as

a wild-type control. Standard growth of control cells was rated 100%. Error bars represent the SEM. Confidence intervals were 95%. Mann-Whitney's non-parametric test (A, B), Kruskal-Wallis test followed by Dunn's post test (C - F). Statistical significance when $p < 0.05$.

A) TRPC3^{-/-} mice. Treatment with SKF (n=56) increased average area to 110% of untreated controls (n=84).

B) CB6 mice. Treatment with SKF (n=65) reduced average area to 99% of controls (n=76, no significant difference).

C) TRPC3^{-/-} mice. Treatment with DHPG reduced average area to 53% (n=66, significant $p < 0.001$) of controls (n=83). Treatment with DHPG and SKF resulted in average areas of 56% (n=58, not significant to DHPG).

D) CB6 mice. Treatment with DHPG reduced average area to 53% (n=68, significant $p < 0.001$) of controls (n=74). Treatment with DHPG and SKF resulted in average areas of 52% (n=67, not significant to DHPG).

E) TRPC3^{-/-} mice. Treatment with PMA reduced average area to 35% (n=70, significant $p < 0.001$) of controls (n=64). Treatment with PMA and SKF resulted in average areas of 30 % (n=62, not significant to PMA).

F) CB6 mice. Treatment with PMA reduced average area to 36% (n=74, significant $p < 0.001$) compared to controls (n=87). Treatment with PMA and SKF resulted in average areas of 30% (n=66, not significant to PMA).

5. Additional data

The following section contains unpublished results.

5.1. The common TRPC inhibitor SK&F 96365 was in high concentrations toxic to Purkinje cell axons independent of TRPC3

The widely used TRPC3-inhibitor SK&F 96365 (SKF) (Kress *et al.*, 2008) caused toxic effects on Purkinje cells from CB6 mice in a concentration dependent manner (Fig. 26). At SKF 10 μ M, after a survival of 10 days the cultures displayed a moderate axonal damage (see Fig. 26 B) with the rest of the culture remaining unharmed. SKF 25 μ M caused strong axonal damage to the point of complete loss of axons (see Fig. 26 C). Survival of Purkinje cells was still equal to control cultures, but their dendritic trees showed clear signs of degeneration. After treatment with SKF 50 μ M, Purkinje cell density was remarkably reduced with the remaining cells displaying stunted dendrites and no axons (Fig. 26 D). In order to find out whether these effects were due to TRPC3 inhibition, we treated TRPC3^{-/-} cultures in the same way over 10 days. No difference between the two mouse strains could be found (Fig. 26 E-H).

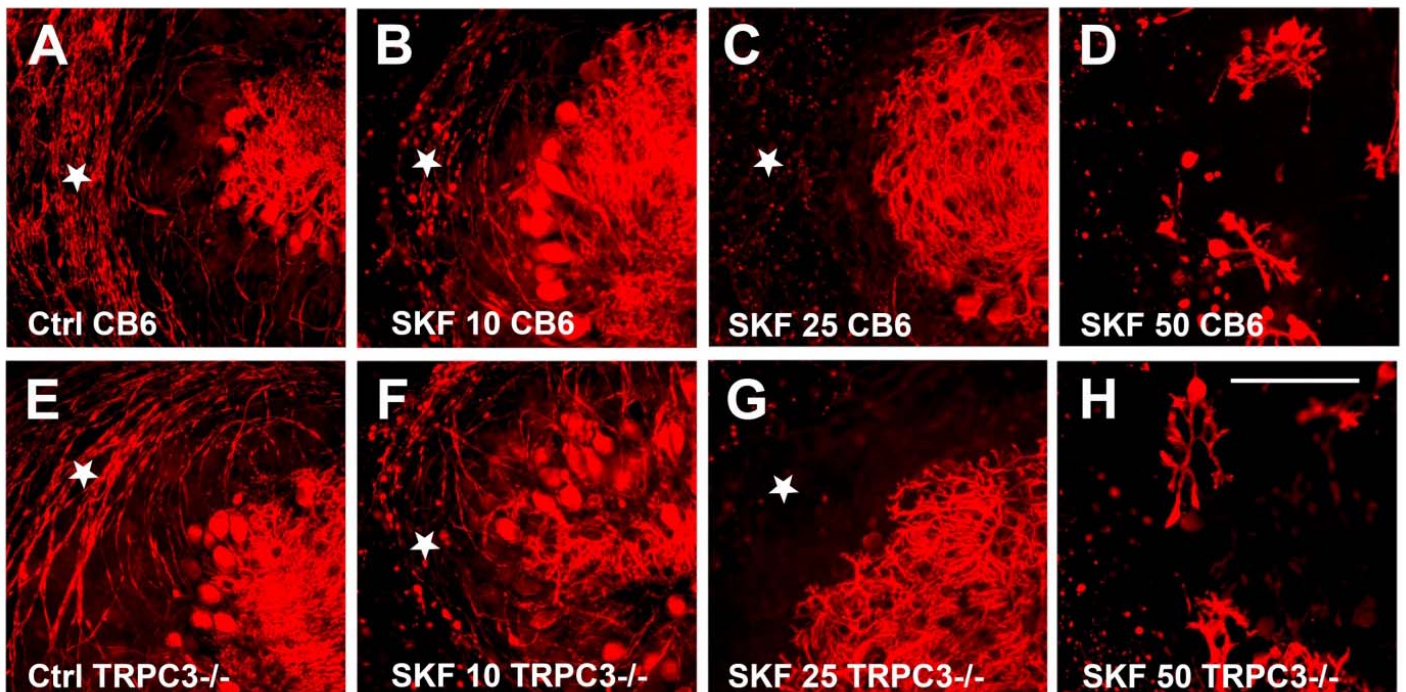


Figure 26: Toxicity of the TRPC inhibitor SKF 96365 on Purkinje cell axons. Staining of Calbindin D-28K shows Purkinje cells.

A) - D) Cultures from CB6 mice. In untreated control cultures the Purkinje cell axons are clearly visible (asterisk in **A**). Treatment with SKF 10 μM (**B**) and 25 μM (**C**) causes damage of Purkinje cell axons (asterisks in B, C). Treatment with SKF 50 μM additionally causes Purkinje cell death (**D**).

E) - H) Cultures from TRPC3^{-/-} mice. Untreated control cultures with clearly visible axons (asterisk in **E**). The toxic effects of treatment with SKF 10 μM (**F**), 25 μM (**G**) and 50 μM (**D**) on Purkinje cells are comparable to those in CB6 cultures.

5.2. Compounds which had no effect on dendritic growth after mGluR1 or PKC stimulation

5.2.1. Nimodipine: inhibitor of L-type calcium channels

P/Q-type and T-type Ca²⁺ are responsible for up to 95% of the total Ca²⁺ currents in Purkinje cells (Usowicz *et al.*, 1992; Watanabe *et al.*, 1998; Swensen & Bean, 2003; Isope & Murphy, 2005; Isope *et al.*, 2010). L-type Ca²⁺ channels are also expressed in Purkinje cells, making a small contribution to Ca²⁺ currents, however (Regan, 1991; Mintz *et al.*, 1992a; Watanabe *et al.*, 1998; McKay & Turner, 2005). We inhibited L-type channels additionally to T-type and P/Q-type Ca²⁺ channels simultaneous with mGluR1 activation to find out if the rescue effect on dendritic growth can be increased.

Treatment with the L-type channel inhibitor nimodipine had even a negative effect on the rescue achieved with T/P/Q-type inhibition in two experiments (see Fig. 27).

Due to these results, no further research or analysis was done with L-type channel inhibitors.

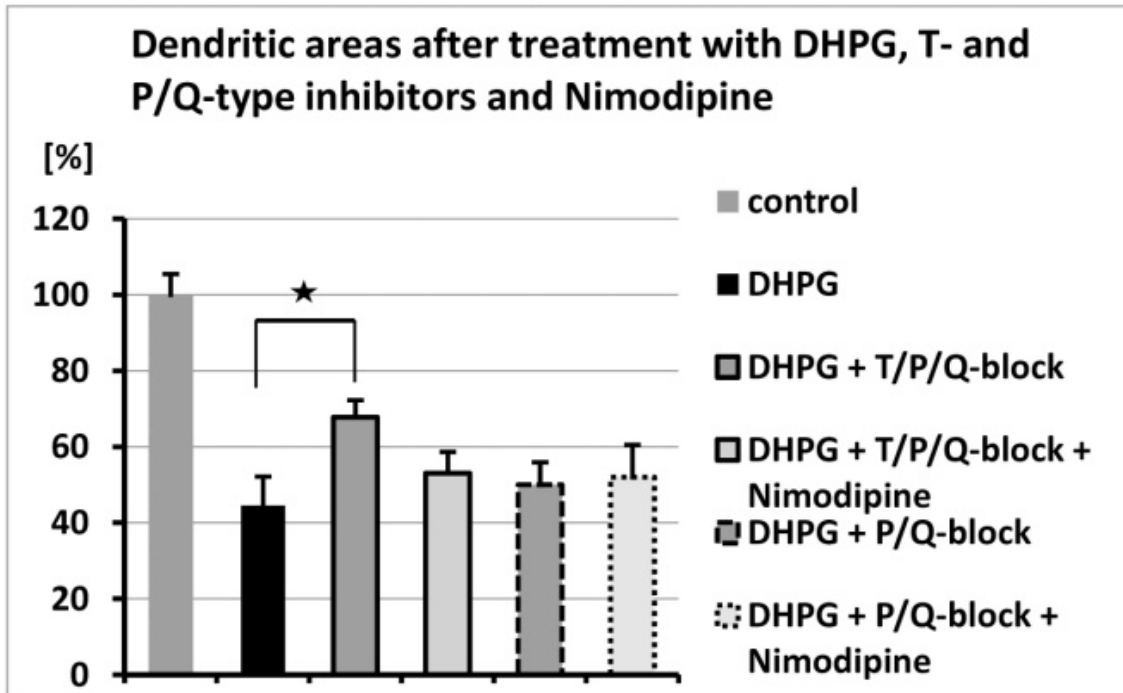


Figure 27: Dendritic tree areas after treatment with DHPG, T/P/Q-inhibitors and co-treatment with the L-type channel blocker nimodipine.

Size of the area covered by a single dendritic tree. Cells were treated with DHPG (10 μ M) for 7 days either alone or in the presence of the P/Q-block, the P/Q-block and the T-type Ca²⁺ channel inhibitor NNC 55-0396, or both combinations together with Nimodipine (2 μ M). Standard growth of control cells was rated 100 %, error bars represent the SEM. A minimum number of 23 cells per condition were measured. DHPG treatment reduced average dendritic area to 44 % of untreated controls. Treatment with P/Q-block and NNC 55-0396 simultaneously with DHPG increased the average area to 68 %. After addition of Nimodipine to T/PQ-inhibition, dendritic area was 53 %. After treatment with P/Q-block simultaneously with DHPG, the average area was 50 %. After addition of Nimodipine to P/Q-inhibition, dendritic area was 52 %. Star indicates statistical significance compared to DHPG-only-treated cultures.

5.2.2. BAPTA-AM: intracellular calcium chelation

BAPTA-AM is the membrane permeable form of the calcium chelator BAPTA. Since calcium entry through T-type and P/Q-type channels appears to mediate dendritic growth inhibition in Purkinje cells after mGluR1 and PKC activation, binding of intracellular calcium by a chelator can be expected to have similar effects as T-type and P/Q-type channel inhibition.

Organotypic slice cultures treated with 50 or 100 μ M BAPTA-AM simultaneous with DHPG or PMA for 7 days showed no substantial increase of dendritic areas or morphology in one experiment (see Fig. 28). In further experiments, BAPTA-AM was used in combination with P/Q-type and T-type inhibitors (see Fig. 29).

After dendritic growth inhibition with DHPG and inhibition of P/Q-type channels, the addition of BAPTA-AM 50 μ M slightly increased dendritic branch point number. In all other combinations, BAPTA-AM had no influence on dendritic areas or branch point numbers (see Fig. 29, A and B).

After dendritic growth inhibition with PMA and inhibition of T-type channels, the addition of BAPTA-AM 50 μ M caused a trend towards increased dendritic areas and branch point numbers (see Fig. 29, C and D). Inhibition of only T-type or P/Q-type channels simultaneous with PMA treatment caused a trend towards improved areas and branch point numbers compared to PMA alone, but only two experiments were done under these conditions and the number of cells measured (<75 cells) was too low to reach statistical significance. Due to the moderate effect of BAPTA-AM on dendritic development found in these pilot experiments, no further research or analysis was done.

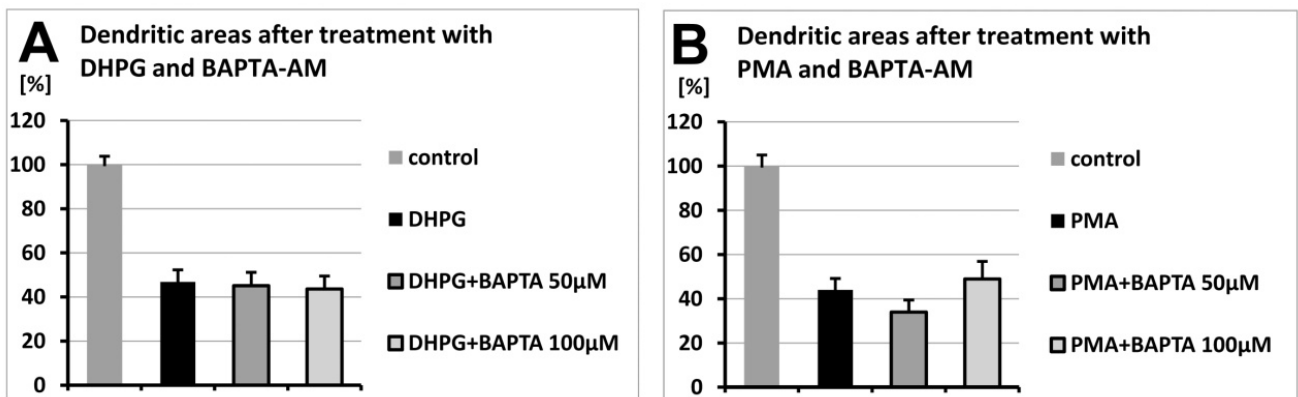


Figure 28: Dendritic tree areas after treatment with DHPG or PMA and co-treatment with the intracellular calcium chelator BAPTA-AM. Size of the area covered by a single dendritic tree. Cells were treated with DHPG (10 μ M) (A) or with PMA (50 nM) (B) for 7 days either alone or in the presence of BAPTA-AM (50 or 100 μ M). Standard growth of control cells was rated 100 %, error bars represent the SEM. A minimum number of 25 cells per condition were measured.

A) DHPG treatment reduced average dendritic area to 47 % of untreated controls. After co-treatment with BAPTA-AM 50 μ M, average area was 45 %, and after co-treatment with BAPTA-AM 100 μ M 44 %.

B) PMA treatment reduced average dendritic area to 44 % of untreated controls. After co-treatment with BAPTA-AM 50 μ M, average area was 34 %, and after co-treatment with BAPTA-AM 100 μ M 49 %.

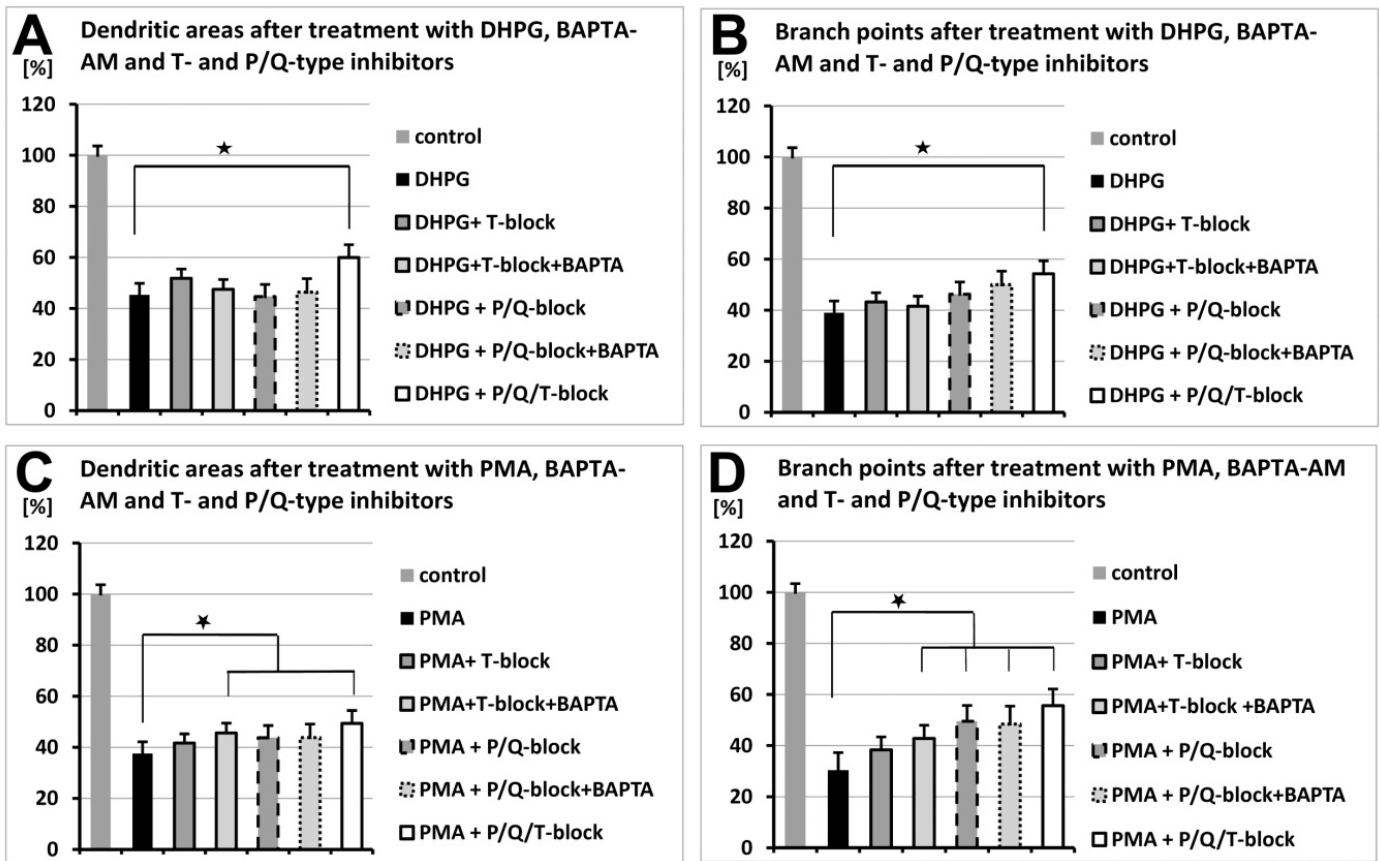


Figure 29: Dendritic tree areas after treatment with DHPG or PMA, T/P/Q-inhibitors and co-treatment with the intracellular calcium chelator BAPTA-AM.

Size of the area covered by a single dendritic tree (A, C) and number of branch points (B, D). Cells were treated with DHPG (10 μ M) (A, B) or with PMA (50 nM) (C, D) for 7 days either alone or in the presence of the P/Q-block and/or the T-type Ca^{2+} channel inhibitor NNC 55-0396 (1 μ M) and/or BAPTA-AM (50 μ M). Standard growth of control cells was rated 100 %, error bars represent the SEM. A minimum number of 42 cells per condition were measured.

A) DHPG treatment reduced average dendritic area to 45 % of untreated controls. After co-treatment with NNC-55-0396, average area was 52 %, after co-treatment with NNC-55-0396 and BAPTA-AM 47 %, after co-treatment with the P/Q-block 45 % and after co-treatment with the P/Q-block and BAPTA-AM 46 %.

Treatment with P/Q-block and NNC 55-0396 simultaneously with DHPG increased average size to 60 %.

B) DHPG treatment reduced average dendritic branch point number to 39 % of untreated controls. After co-treatment with NNC-55-0396, average branch point number was 43 %, after co-treatment with NNC-55-0396 and BAPTA-AM 42 %, after co-treatment with the P/Q-block 46 % and after co-treatment with the P/Q-block and BAPTA-AM 50 %.

Treatment with P/Q-block and NNC 55-0396 simultaneously with PMA increased average branch point number to 54 %.

C) PMA treatment reduced average dendritic area to 38 % of untreated controls. After co-treatment with NNC-55-0396, average area was 42 %, after co-treatment with NNC-55-0396 and BAPTA-AM 46 %, after co-treatment with the P/Q-block 44 % and after co-treatment with the P/Q-block and BAPTA-AM 44 %.

Treatment with P/Q-block and NNC 55-0396 simultaneously with PMA increased average size to 49 %.

D) PMA treatment reduced average dendritic branch point number to 30 % of untreated controls. After co-treatment with NNC-55-0396, average branch point number was 38 %, after co-treatment with NNC-55-0396 and BAPTA-AM 43 %, after co-treatment with the P/Q-block 50 % and after co-treatment with the P/Q-block and BAPTA-AM 48 %.

Treatment with P/Q-block and NNC 55-0396 simultaneously with PMA increased average size to 56 %.

Star indicates statistical significance compared to DHPG- or PMA-only-treated cultures, respectively.

5.3. Morphological analysis of Purkinje cell spines using confocal microscopy

To characterize the differences between the effects on Purkinje cell dendrites mediated by mGluR1 and PKC, we used confocal microscopy in a piloting experiment. Cultures grown in the presence of PMA (50nM) or DHPG (10 μ M) for 7 days were compared to untreated control cultures. Qualitative picture analysis revealed profound differences in spine morphology (see Fig. 30 A-C). Another observation was that in untreated control cultures, spines were only found on dendrites (see Fig. 31 A), while in PMA treated cultures, spines were also found on Purkinje cell somata (see Fig. 31 B) No definition of parameters describing spine morphology or quantification has been done.

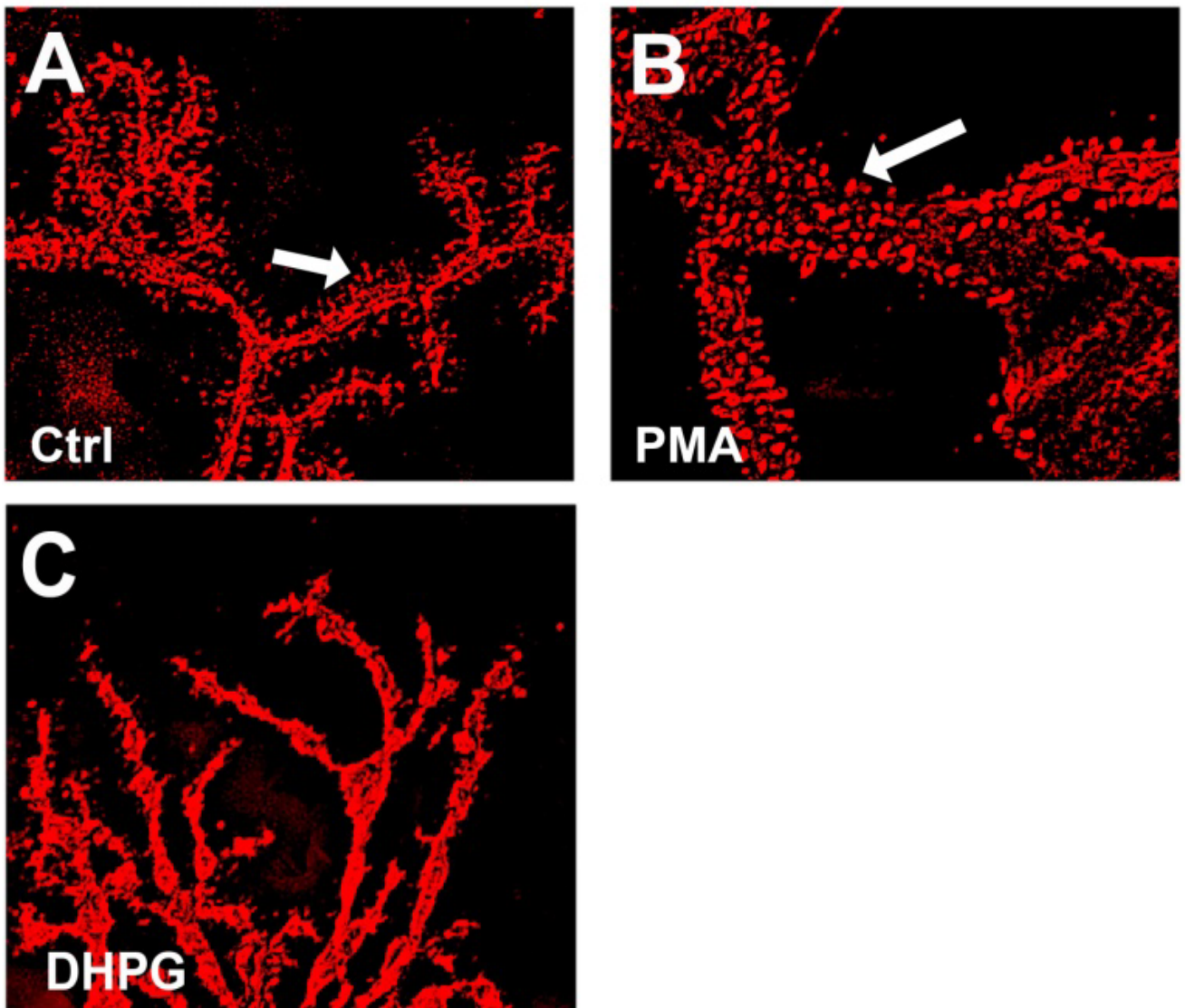


Figure 30: Examples for Purkinje cell dendrites viewed with confocal microscopy.

Pictures were taken with a 100 x lens and all digitally magnified by the same factor.

A) Untreated control culture, spines clearly visible (arrow in A points to a spine).

B) Culture treated with PMA (50nM) for 7 days, spines look different from control (arrow in B points to a spine).

C) Culture treated with DHPG (10 μ M) for 7 days, spines are virtually gone.

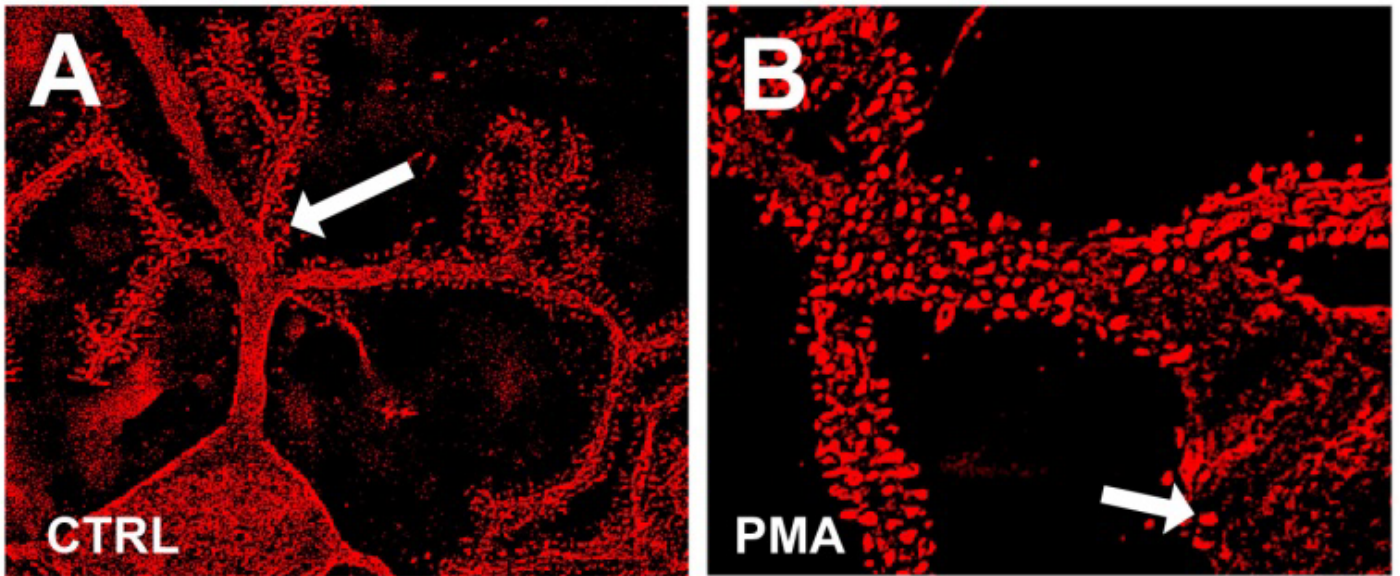


Figure 31: Examples for Purkinje cell dendrites viewed with confocal microscopy.

Pictures were taken with a 100 x lens and all digitally magnified by the same factor.

A) Untreated control culture, spines clearly visible on dendrites, no spines visible on soma (arrow in A points to a spine).

B) Culture treated with PMA (50nM) for 7 days, spines present on soma (arrow in B points to a spine).

6. General Discussion

6.1. What might be the cause of Purkinje cell vulnerability to excitotoxic death?

Cerebellar Purkinje cells have been found to be especially affected in various diseases with an excitotoxic aspect, such as ischemia, trauma or neurodegenerative disorders (reviewed in (Sarna & Hawkes, 2003)). When we treated cerebellar slice cultures with 30 μ M AMPA for 4 hours, virtually all Purkinje cells were killed independent of the size of their dendritic trees, which had been reduced by pre-treatment with either PMA or DHPG for 7 days. Staining for vGluT1 revealed that Purkinje cell dendrites were densely covered by glutamatergic terminals. This suggests that it is not the size, or total number of glutamatergic synapses on the dendritic tree that confers susceptibility to excitotoxic overload. In Purkinje cells, AMPA induces a process called dark cell degeneration, which resembles the cellular pathology seen after conditions such as hypoglycemia, stroke or trauma (Garthwaite & Garthwaite, 1991a). Excitotoxicity is mediated by an increase in intracellular calcium concentration (Garthwaite & Garthwaite, 1991b; Strahlendorf *et al.*, 1998). Although they do express functional AMPA receptors (Hack *et al.*, 1995), cerebellar granule cells in the same cultures did not show any impairment of survival after the AMPA treatment, instead they started to express calbindin D-28K, which normally is Purkinje cell specific in the cerebellum (Bastianelli, 2003). In several studies, the expression of calbindin has been found to be an important determinant for a cell's susceptibility to excitotoxic conditions. For example, in hippocampal granule cells of epileptic patients, loss of calbindin resulted in a faster Ca^{2+} -dependent inactivation of calcium entry through voltage-dependent calcium channels (VDCC) during seizure-like frequent neuronal discharges (Nagerl *et al.*, 2000). Probably by either direct interaction with VDCCs or accumulation around the Ca^{2+} entry points to buffer incoming Ca^{2+} , calbindin is thought to prevent an important negative feedback mechanism in controlling Ca^{2+} influx into neurons. Similarly, hippocampal CA1 pyramidal neurons of calbindin D-28k^{-/-} mice were remarkably less susceptible to experimental ischemia *in vivo* and *in vitro* than wild types (Klapstein *et al.*, 1998). It is conceivable that CGNs under excitotoxic stress benefit from faster Ca^{2+} dependent

Ca²⁺ inactivation due to absence of calbindin, while additional upregulation of calbindin buffers excess calcium ions.

The ability to cope with metabolic stress apparently also plays a role in excitotoxicity, as cerebellar granule cells have been shown to be more sensitive to AMPA when their energy metabolism was impaired ((Cebers *et al.*, 1998), abstract). The metabolic enzyme aldolase C has indeed been found to be a determinant for Purkinje cell excitotoxic susceptibility (Slemmer *et al.*, 2007). Aldolase C is an important enzyme in glycolysis, as it converts fructose 1,6-bisphosphate (FBP) to glyceraldehyde 3-phosphate and dihydroxyacetone phosphate, or reversibly catalyzes the reaction in the other direction. FBP has been found to protect neurons from excitotoxicity (Rogido *et al.*, 2003; Park *et al.*, 2004). Aldolase C is expressed in a subset of cerebellar Purkinje cells in a pattern of parasagittal stripes (Slemmer *et al.*, 2007) (see Fig. 32). Interestingly, Aldolase C expressing Purkinje cells were resistant to experimental TBI compared to Purkinje cells without the enzyme, within the same animal. Moreover, in dissociated cultures, Aldolase C-positive Purkinje cells survived a treatment with AMPA (30µM, 20 min), while Aldolase C-negative PCs did not. By transfecting the cells with aldolase C siRNA, which strongly exacerbated vulnerability to AMPA, an involvement of the enzyme was confirmed (Slemmer *et al.*, 2007). A small contribution of the glutamate transporter EAAT4, which co-localizes with Aldolase C expression in Purkinje cells, is possible, however. In fact, mice lacking the function of the glial glutamate transporter GLAST have been found to lose considerably more Purkinje cells after induced ischemia, with a special vulnerability of those Purkinje cells with low expression of EAAT4 (Yamashita *et al.*, 2006). Glutamate transporters remove glutamate from the extracellular space, which is a very important function when glutamate levels in the cerebrospinal fluid are elevated after injury (Baker *et al.*, 1993; Palmer *et al.*, 1993).

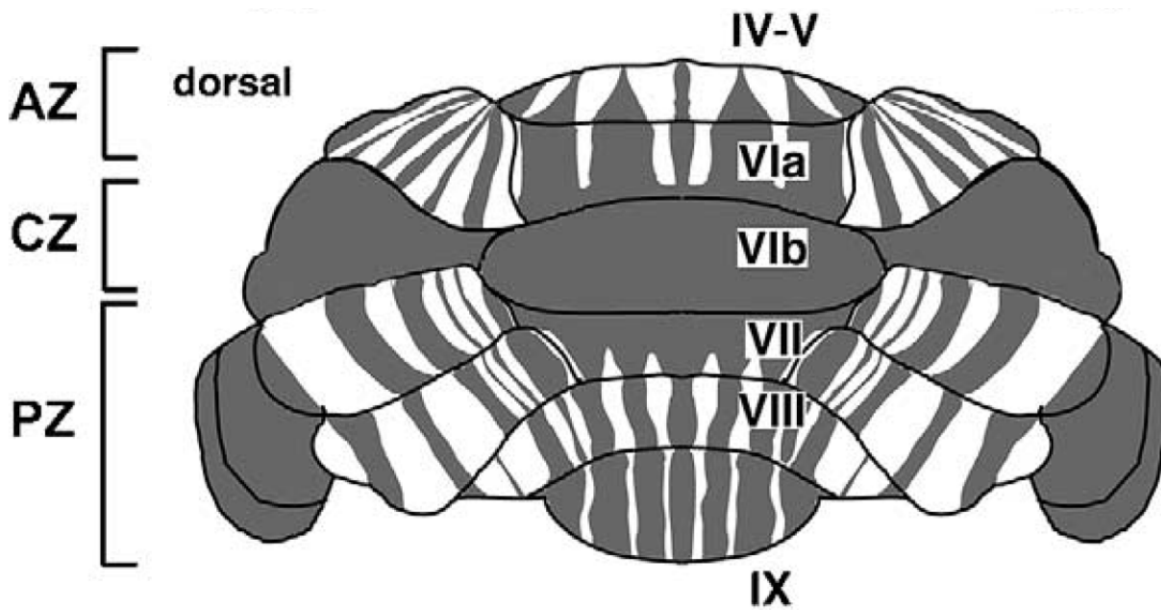


Figure 32: Schematic drawing of the pattern of aldolase C expression in the mouse cerebellum, dorsal view, according to (Sillitoe & Hawkes, 2002). Purkinje cells in the gray shaded areas express aldolase C (=zebrin II). Roman numbers I-X refer to the lobules of the cerebellar vermis region. In the anterior and posterior zones (AZ, PZ), aldolase C is expressed in a pattern resembling zebra stripes, while in the central zone (CZ), all Purkinje cells express aldolase C. picture adapted from (Sillitoe & Hawkes, 2002).

6.2. What are the differences between the similar effects induced by chronic mGluR1 and PKC activation?

Chronic activation of either mGluR1 or PKC results in a strong dendritic growth inhibition of the Purkinje cell dendritic tree in cerebellar slice cultures (Sirzen, Metzger & Kapf, Schrenk). Because PKC is a downstream effector in mGluR1 signalling mediating long term depression (Daniel *et al.*, 1998; Ito, 2001), it is interesting that PKC and mGluR1 appear to mediate dendritic growth inhibition via independent pathways (Sirzen-Zelenskaya *et al.*, 2006). The dendritic effects conveyed by chronic mGluR1 or PKC activation are similar, but they do differ in some aspects, however.

When we stained cerebellar slice cultures for vGluT1, glutamatergic terminals were revealed to be located at the more distal parts of the Purkinje cell dendritic trees (see Fig. 15 A). This was still the case after 7 days of treatment with DHPG (Fig. 15 C). After 7 days of treatment with PMA, on the other hand, vGluT1-positive terminals were also located on many Purkinje cell somata (Fig. 15 B). Somehow, the Purkinje

cell somata must have retained the characteristics of the dendritic compartments to build glutamatergic synapses. Preliminary results using confocal microscopy to visualize Purkinje cell spines revealed striking morphological differences between mGluR1- and PKC-induced dendritic growth inhibition (see Fig. 30). After chronic DHPG treatment, Purkinje cell dendrites were virtually devoid of spines, while after chronic PMA treatment, Purkinje cell dendrites, and often also somata, were densely covered by spines, which were clearly different in shape than the spines found in untreated controls. Moreover, after PMA treatment spines were found on Purkinje cell dendrites and somata, while in controls the somata were devoid of spines (see Fig. 31). This is in line with our findings on vGluT1-positive terminals on Purkinje cell somata after PMA treatment.

Combined inhibition of T-type and P/Q-type calcium channels provided a partial rescue from the dendritic growth inhibition mediated by PKC or mGluR1 activation. In PMA treated cultures, however, either inhibition of T-type *or* of P/Q-type channels was sufficient to increase dendritic growth parameters significantly, while in DHPG treated cultures, simultaneous inhibition of both channel types was required. This points toward additional, compensatory mechanisms being involved in mediating dendritic growth inhibition particularly after chronic mGluR1 activation.

6.3. Inhibition of Purkinje cell dendritic growth involves Ca²⁺ entry through voltage gated channels of the T- and the P/Q-type

We have shown here that pharmacological blockade of T-type and P/Q-type Ca²⁺ channels results in a rescue of the Purkinje cell dendritic tree after mGluR1 or PKC stimulation indicating that Ca²⁺ influx through these channels is important in mediating dendritic growth inhibition. In many aspects of neuronal development, Ca²⁺ signals play a critical role (Konur & Ghosh, 2005). Increased Ca²⁺ entry into Purkinje cells severely reduced dendritic trees in two independent mouse models. A constitutively open GluR δ 2 channel impairs Purkinje cell development and survival in the *lurcher* mouse (Zuo et al., 1997), while in the *mwk* mouse a leaky TRPC3 channel causes dendritic growth inhibition in Purkinje cells. Pharmacological inhibition of the GluR δ 2 channel with Nasp, which is able to rescue the effects in *lurcher* mice (Zanjani et al., 2009), did not convey any protection from dendritic growth inhibition after mGluR1 or PKC activation. Similarly, pharmacological

inhibition as well as experiments with mice lacking TRPC3 function strongly suggest that TRPC3 does not play a role in mediating mGluR1 or PKC induced dendritic growth inhibition. This was surprising, because a gain-of-function in the *trpc3* gene has a similar effect on Purkinje cell dendrites as DHPG- or PKC treatment, and, moreover, TRPC3 has been shown to be involved in both, mGluR1- and PKC signaling. TRPC3 mediates the sEPSC after mGluR1 activation (Hartman) and TRPC3 activity is modulated by PKC (Becker *et al.*, 2009).

Furthermore, no involvement or contribution of Ca²⁺ release from internal stores, from R-type and, in piloting experiments, L-type channels have been found, while the role of N-type Ca²⁺ channels cannot be fully excluded, as they were inhibited together with T- and P/Q- type channels with ω -conotoxin MVIIC (McDonough *et al.*, 1996). Since T- and P/Q-type channels together mediate up to 95% of the total Ca²⁺ currents in Purkinje cells (Usowicz *et al.*, 1992; Watanabe *et al.*, 1998; Swensen & Bean, 2003; Isope & Murphy, 2005; Isope *et al.*, 2010), and both have been shown to be involved in mGluR1 signaling (Kitano *et al.*, 2003; Hildebrand *et al.*, 2009), it is conceivable that they are also responsible for the major part of Ca²⁺ entry after mGluR1 or PKC activation.

In an attempt to corroborate our findings by supporting the theory that increased Ca²⁺ influx mediates the growth inhibiting effects after chronic DHPG or PKC treatment, we used BAPTA-AM as a chelator of intracellular calcium. In concentrations up to 100 μ M, BAPTA-AM did not rescue Purkinje cell dendrites which were simultaneously treated with DHPG or PMA (see Fig. 28), and BAPTA-AM at 50 μ M did not increase the rescue effect obtained by T/P/Q-inhibition (see Fig. 29). A possible explanation for this lack of effect could be that BAPTA-AM is required to traverse the cell membrane in order to perform its action. This can be problematic in organotypic slice cultures.

7. Conclusion and Outlook

7.1. Physiological relevance of limiting dendritic size after chronic activation of mGluR1 or PKC

Our results show that a reduced size of the dendritic tree does not protect Purkinje cells from AMPA-mediated neurotoxicity. Staining for the vesicular glutamate transporter vGluT1 revealed a high density of glutamatergic synapses on the stunted dendritic trees of pre-treated Purkinje cells. This suggests that receptor density rather than total receptor load is important for determining the sensitivity of Purkinje cells to AMPA-mediated neurotoxicity. Our finding that AMPA-challenged cerebellar granule cells rapidly become immunoreactive for Calbindin D-28K shows that the specificity of Calbindin D-28K immunoreactivity for Purkinje cells in the cerebellum is not absolute, and that also granule cells can express this protein. Intrinsic features like calbindin expression under normal conditions, coping with metabolic stress and the presence of glutamate transporters are possible reasons for Purkinje cell susceptibility to excitotoxic overload. Carrying out our experiments with calbindin^{-/-} mice (Klapstein *et al.*, 1998) might reveal interesting results.

It remains unclear what the physiological relevance of limiting dendritic size after chronic activation of a metabotropic glutamate receptor or PKC might be. The generation of transgenic mice expressing a constitutively active PKC γ , which is currently a project in our group, may not only clarify this point, but may also serve as a useful animal model to study spinocerebellar ataxia type 14 (Chen *et al.*, 2003). Similarly, experiments with a constitutively active mGluR1 receptor (Yamashita *et al.*, 2004), either *in vivo* or *in vitro* using a hand-held gene gun for transfection of cerebellar slice cultures, can be interesting projects for the future.

7.2. Further studies on dendritic growth inhibition induced by chronic mGluR1 or PKC activation

Chronic activation over several days of either mGluR1 or PKC results in a strong dendritic growth inhibition of the Purkinje cell dendritic tree in cerebellar slice culture (Metzger & Kapfhammer, 2000; Schrenk *et al.*, 2002; Sirzen-Zelenskaya *et al.*, 2006). Interestingly, PKC and mGluR1 appear to mediate dendritic growth inhibition via

independent pathways (Sirzen-Zelenskaya *et al.*, 2006). The effects on dendritic morphology and characteristics conveyed by chronic mGluR1 or PKC activation are similar, but differ in some aspects when they come under scrutiny. Qualitative piloting experiments with confocal microscopy revealed interesting differences in spine morphology. Further experiments and establishment of a method for quantification of spine density and definition of morphological parameters will be necessary to gain further insights.

7.3. Blockade of T-type and P/Q-type Ca²⁺ channels partially rescues dendritic trees from mGluR1 and PKC induced dendritic growth inhibition

Our results show that T-type and P/Q-type Ca²⁺ channels are involved in mediating dendritic growth inhibition after chronic mGluR1 and PKC activation. The rescued dendritic trees did not reach the size of untreated control cells. Although it is possible that additional pathways are involved, the reason for this partial rescue could also be an incomplete blocking of the channel blockers. We have experienced the permanent presence of DHPG or PMA in the culture medium over 7 days as a very potent activator of mGluR1 or PKC, respectively. Therefore, it might be problematic to thoroughly block any downstream target by pharmacological means. An alternative method which could help us clarify this point may be the introduction of siRNA against the calcium channels using a hand-held gene gun.

8. Materials and Methods

8.1. Cerebellar slice cultures

8.1.1. Media and prearrangements for cerebellar slice cultures

All ingredients were sterile and mixed in a laminar flow workbench.

Ingredients for 200 ml preparation medium (PM):

- 100 ml minimal essential medium (MEM) (Gibco 11012), in twofold concentration
- 98 ml Aqua bidest
- 1 ml Glutamax (Gibco 35050)
- 1N NaOH or 1N HCl to adjust pH to 7.2-7.4

Ingredients for 100 ml incubation medium:

- 25 ml MEM (Gibco 11012), in twofold concentration
- 23,5 ml Aqua bidest
- 25 ml Basal Medium Eagle, with Earl's salt, without glutamine (Gibco 21400)
- 25 ml horse serum, heat-inactivated (Gibco 26050)
- 1 ml Glutamax (Gibco 35050)
- 700 µl of a 10% glucose solution
- 1N NaOH or 1N HCl to adjust pH to 7.2-7.4

Pre-arrangements:

- A sterilized razorblade was installed on a tissue chopper (McIlwain).
- Petri dishes (greiner bio-one 627102, 35mm) were half-filled with ice-cold preparation medium and stored at 4°C until used. 2-3 dishes per mouse were required.
- 750µl of incubation medium was pipeted in each well of a tissue culture plate (6 wells, Falcon 353046). The plate was placed in an incubator providing a humidified atmosphere with 5% CO₂ at 37°C. One plate holds cultures from 3 mice.
- Surgical instruments were sterilized and placed ready.

8.1.2. Procedure

Animal experiments were carried out in accordance with the European Communities Council Directive of 24 November 1986 (86/609/EEC) and were reviewed and permitted by Swiss authorities. Cultures were prepared from B6CF1 mice at postnatal day 8 (P8) as described previously (Adcock *et al.*, 2004; Kapfhammer, 2005; 2010) and incubated according to the static method (Stoppini *et al.*, 1991). All steps were carried out in the aseptic environment of a horizontal laminar flow workbench (BDK, Germany).

- A P8 mouse pup was decapitated and the head was sprayed with 70% ethanol
- The skull was carefully opened in the sterile workbench.
- The cerebellum was removed together with surrounding brain structures (*colliculi inferiores, pons, medulla oblongata*) and placed immediately in a dish filled with ice-cold preparation medium. All further steps were carried out under a stereomicroscope (Zeiss, stemi2000).
- The cerebellum was isolated by cutting the cerebellar peduncles and removing most of the meninges from the surface.
- The cerebellum was placed on the tissue chopper and sagittal slices (350 µm) were cut.
- The sliced cerebellum was placed in a fresh dish with cold preparation medium and the slices were separated from each other. One cerebellum usually yielded 15-18 slices.
- The bottom membranes of two tissue culture inserts (Millicell CM, Millipore PICM 03050) were moistened with preparation medium.
- The slices were carefully laid on the membrane. Both culture inserts should contain about the same amount of slices from different regions of the cerebellum.
- The culture inserts were placed in the pre-arranged tissue culture plate and incubated immediately.

8.2. Experiments and maintenance of the cultures

- The incubation medium was changed every 2-3 days. Fresh incubation medium was pre-warmed to 37°C and the pH was adjusted to 7.2-7.4, if necessary.
- Drugs were added to the culture medium after every medium change.

8.2.1. Pharmacological agents

Solvents for stock solutions (SS, sterile):

- Preparation medium (PM)
- Aqua bidest (H₂O)
- 1:1 mixture of ethanol and dimethyl sulfoxide (EtD)

All Stock solutions 1 (see Table 1) were kept at -20°C. Stock solutions 2 were kept at -20°C if the solvent was EtD, and at -4°C for the duration of the experiment if the solvent was PM or H₂O. Concentrations of stock solutions 2 were chosen according to the desired end-concentration for the experiment, as to make sure that the concentration of solvent in the culture medium would not exceed 1 % (Kapfhammer, 2010).

Table 1: Substances and stock solutions.

Substance	MW [g/mol]	Stock solution 1	Stock solution 2	Supplier
ω-Agatoxin IVA	5202.3	0.1mM (0.1mg in 192μl PM)	50μM (dilute SS1 1:1 in EtD)	Bachem H- 1544.0100
AMPA (RS)-α-Amino-3-hydroxy-5-methyl-4-isoxazolepropionic acid	186.2	10mM (1.86mg/ml in H ₂ O)	5mM (dilute SS1 1:1 in EtD)	Tocris 169
BAPTA-AM (light sensitive)	764.7	10mM (7.65mg/ml in EtD)		Calbiochem 196419
ω-Conotoxin MVIIC	2639.1	1mM (0.5mg in 182μl PM)	500μM (dilute SS1 1:1 in EtD)	Bachem H- 8835.0500
DHPG (RS)-3,5-Dihydroxyphenylglycine	183.05	30mM (10mg in 1763μl PM)	7.5mM (dilute SS1 1:3 in EtD)	Tocris 0342

Substance	MW [g/mol]	Stock solution 1	Stock solution 2	Supplier
Mibefradil	568.55	30mM (10mg in 568 μ l H ₂ O)	1mM (dilute SS1 1:29 in H ₂ O)	Tocris 2198
Naspm 1-Naphthylacetyl spermine	479.9	50mM (5mg in 208 μ l PM),	25 mM (dilute SS1 1:3 in EtD)	Sigma N193
NNC 50-0396	564.6	25mM (10mg in 676 μ l H ₂ O)	0.5mM (dilute SS1 1:49 in H ₂ O)	Tocris 2268
Nimodipin (light sensitive)	418.4	50mM (23.9mg in 1.14 ml EtD)	1mM (dilute SS1 1:49 in EtD)	Tocris 600
PMA Phorbol 12-myristate 13-acetate	616.8	1.5 mM (0.925 mg/ml in EtD)	50 μ M (dilute SS1 1:29 in EtD)	Tocris 1201
Pyr3	456.6	20mM (10mg in 1094 μ l EtD)	1mM (dilute SS1 1:19 in EtD)	Tocris 3751
SKF 96365	402.9	25 mM (10 mg/ml in EtD)	5 mM (dilute SS1 1:4 in EtD)	Tocris 1147
SNX-482	4495.0	100 μ M (10 μ g in 22.25 μ l H ₂ O)		Biotrend BP0376
U73 122	469.15	4mM (1.88mg/ml in EtD)	1mM (dilute SS1 1:3 in EtD)	Tocris 1268

8.2.2. The role of dendritic tree size in excitotoxicity:

- Pharmacological treatments inhibiting dendritic growth of Purkinje cells were started after 3 days in vitro (DIV3) and washed out at DIV10.
- The cultures were incubated for 24 hours without treatment.
- At DIV12 the cultures were challenged with AMPA for 4 hours and kept in culture for two more days without treatment.

8.2.3. Pathway analysis of mGluR1- and PKC-induced dendritic growth inhibition

- All pharmacological treatments were started at DIV2-3 and maintained for 7 days, until the end of the culture period.
- To ensure a thorough blockade of Ca²⁺ channels, the substances DHPG and PMA were added 4 hours after the channel inhibitors. This would make sure that the channel inhibitors reach their binding sites and establish a channel inhibition before DHPG or PMA could trigger a cellular reaction limiting dendritic growth.
- The P/Q- and N-type Ca²⁺ channel inhibitor ω -conotoxin MVIIC was an exception to this rule as it was added 4 hours after the other channel inhibitors at the same time with DHPG or PMA, because ω -conotoxin MVIIC only supports P/Q-type inhibition achieved by ω -agatoxin IVA if ω -agatoxin IVA reaches the binding site first (McDonough *et al.*, 2002).

8.3. Immunocytochemistry

- At the end of the culture period cultures were fixed in 4% paraformaldehyde in 100 mM phosphate buffer (PB) overnight at 4°C.
- The slices were washed with PB 3 times for 10 min.
- The plastic feet at the culture inserts were cut off to limit the required amount of antibody solution to 800 μ l.
- The 1. antibody solution was prepared in PB as follows:
 - 0.5 % Triton X-100, in order to permeabilize the tissue and prevent non-specific antigen binding
 - 3 % normal goat serum, in order to block non-specific antigen binding
 - rabbit anti-Calbindin D-28K (Swant) 1:1000, to visualize Purkinje cells
 - monoclonal anti-NeuN (Chemicon, Millipore) 1:500, to visualize CGNs
- The slices were incubated in 1. antibody solution at 4°C overnight
- The slices were washed with PB 3 times for 10 min.
- The 2. Antibody solution was prepared in PB as follows:
 - 0.1 % Triton X-100, in order to prevent non-specific antigen binding

- goat anti-rabbit Alexa 546 or 568 (Molecular Probes, Invitrogen) 1:500
- goat anti-mouse Alexa 488 (Molecular Probes, Invitrogen) 1:500
- The slices were incubated in 2. antibody solution at room temperature for 1-4 hours.
- The slices were washed with PB 3 times for 10 min.
- Stained slices were mounted on glass slides (Thermo Scientific Menzel-Gläser Superfrost Plus, Art. No. J1800AMNZ) and coverslipped with Mowiol.
- Cultures were viewed on an Olympus AX-70 microscope equipped with a Spot digital camera.
- Recorded images were adjusted for brightness and contrast with Photoshop image processing software

8.4. Genotyping of TRPC3^{-/-} mice

TRPC3^{-/-} mice were a kind gift from the Arthur Konnerth group. To confirm that those mice indeed contained the dysfunctional *Trpc3* gene which lacks exon 7 (Hartmann *et al.*, 2008), we performed a PCR.

8.4.1. Protocol for isolation of DNA

- Cut 1 mm mouse tail tip after decapitation of the mouse pup for cerebellar slice culture, store at -20°C immediately in an Eppendorf tube (1.5 mm).
- Add to each tail tip:
 - 40 µl Proteinase K (Merck, Stock solution of 10 mg/ml)
 - 700 µl 'Tail Buffer' (10 mM Tris. Cl, pH8.0, 50 mM EDTA, 100 mM NaCl, 0.5 % SDS)
- Shake at 55°C over night
- Add 175 µl NaCl (6M), shake at 55°C for 15 min
- Centrifuge at 13'000 for 20 min (Heraeus Biofuge pico)
- Transfer 660 µl supernatant into fresh Eppendorf tube (1.5 mm)
- Add 660 µl Isopropanol, shake shortly by hand
- Centrifuge at 4°C at 20 000 g, 22 min (Heraeus Biofuge stratus)
- Discard supernatant, keep pellet

- Add to pellet 660 µl ethanol (75 %, stored at 4°C), shake shortly by hand
- Centrifuge at 4°C at 20 000 g, 22 min (Heraeus Biofuge stratus)
- Discard supernatant, put open Eppendorf tube at 55°C for 15 min to let the pellet dry
- Dissolve pellet in 200 µl sterile water

8.4.2. PCR

The primer sequences for the murine *Trpc3* gene (chosen to flank missing exon 7) were chosen according to (Hartmann *et al.*, 2008) and produced by Microsynth.

- 5'-AGA ATC CAC CTG CTT ACA ACC ATG TG-3'
- 5'-GGT GGA GGT AAC ACA CAG CTA AGC C-3'

The primers were resolved in sterile water to a concentration of 50 µM.

PCR Master Mix (for 1 sample), prepare on ice

- 19.5 µl water
- 1.5 µl PCR Buffer 10x (invitrogen)
- 1.5 µl MgCl₂ 50 mM (invitrogen)
- 1.5 µl Primer mix (containing both primers at 0.5 µM in water)
- 1.5 µl ddNTP (invitrogen)
- 0.5 µl Taq Polymerase (5 units/µl, invitrogen)

For 1 sample pipet 28 µl Master Mix + 2 µl DNA solution.

PCR (36 cycles):

- 94°C 2 min 30 s
- 94°C 40 s
- 58°C 50 s (annealing)
- 72°C 50 s (elongation)

8.4.3. Gel electrophoresis

- Mix samples with sample loading buffer (Laemmli 2x concentrate, Sigma) and load 15 µl per lane, one lane contains 1kb DNA Ladder (BioLabs)

- Run 1 % agarose gel (0.3 % ethidium bromide) at 100 V for 80 min,
- photograph under UV light (see Fig. 33)

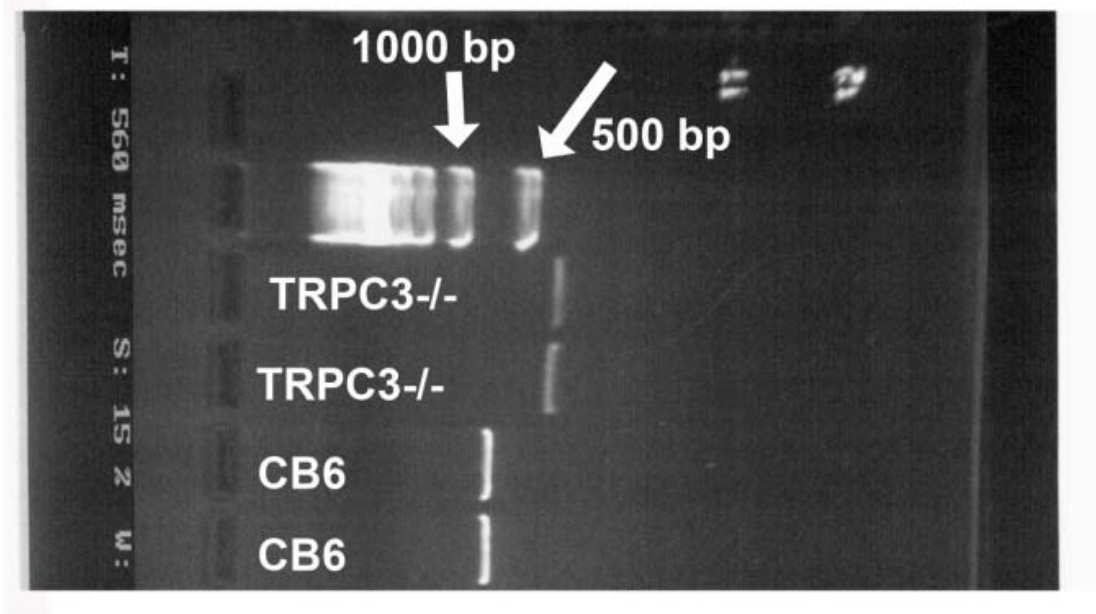


Figure 33: photograph of an agarose gel after electrophoresis, showing analysis of PCR results. Top lane shows DNA ladder with the 2 fastest bands at 1000 and 500 bp. The lanes marked TRPC3^{-/-} contain DNA from TRPC3^{-/-} mice, with bands below 500 bp. The lanes marked CB6 contain DNA from CB6 mice, with bands between 1000 and 500 bp. According to Hartmann *et al.*, the band for TRPC3^{-/-} mice is supposed to be at 300 bp, and the band for wild type mice at 700 bp (Hartmann *et al.*, 2008). This is in line with our results.

8.5. Quantitative analysis of cultured Purkinje cells

- Purkinje cells which had a dendritic tree which could be isolated from its surroundings were selected for analysis.
- Cells were viewed with a 25x lens of a POLYVAR microscope and photographed with a digital camera (Coolsnap-Pro, Photometrics, Canada).
- Dendritic branch points were counted manually.
- The image analysis program Image Pro Plus was used to trace the outline of the Purkinje cell dendritic trees (see Fig. 34) by selecting the area using the magic wand tool. This yielded the area covered by the dendritic tree.
- Cells were acquired from three independent experiments with an average number of 20 cells per experiment.
- The data were analyzed using GraphPad Prism software.

- The statistical significance of differences in parameters was assessed by non-parametric analysis of variance (Kruskal-Wallis test) followed by Dunn's post test. For comparisons of single data columns, Mann-Whitney's non-parametric test was used. Confidence intervals were 95 %, statistical significance when $p < 0.05$.
- The mean value of the dendritic tree area and number of branch points of untreated control cells were set to 100 % and the results were expressed as percentage of controls.

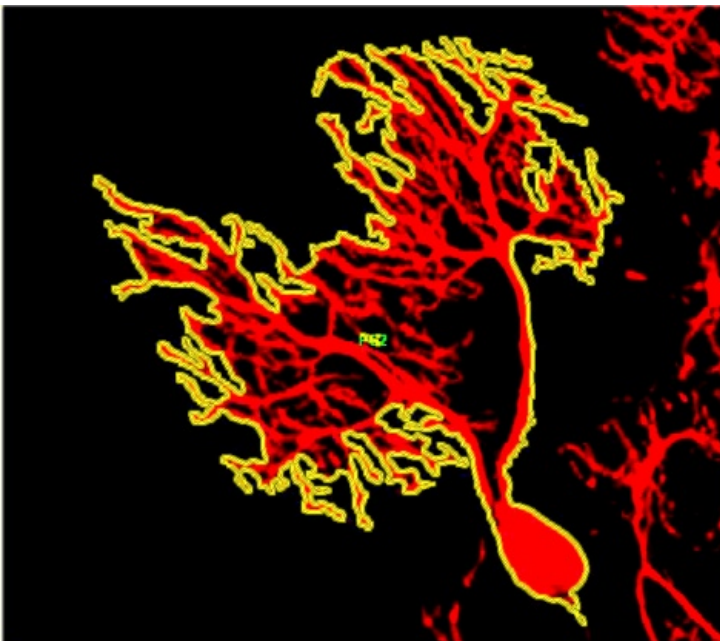


Figure 34: Example for measurement of the area covered by a dendritic tree. Image Pro Plus was used to trace the outline of a dendritic tree.

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This work is dedicated to Dahli F. von Reding - Meichtry (13.12.1955 - 28.7.2011).
Vaya con Dios.

Curriculum Vitae

Personal Details

Name Olivia Sarah Gugger
Address Niederaamtstrasse 3, CH-4656 Starrkirch-Will
Telephone +41 78 745 40 61
E-mail olivia.gugger@unibas.ch
Date of Birth 6th October, 1981
Nationality Swiss

Education & Training

Since Nov. 2007 **Ph.D. Studies in Developmental Neurobiology, Department of Biomedicine, University of Basel**

Thesis subject:

Mechanisms and Consequences of Reduced dendritic growth of cerebellar Purkinje cells

Supervisor: Prof. Josef Kapfhammer

2001- 2006 **Studies of Pharmaceutical Sciences at the Swiss Federal Institute of Technology (ETH) in Zurich**

- Master of Science ETH in Pharmaceutical Sciences
- Swiss federal Pharmacists' license

2005-2006

Practical training as a pharmacist, "Passage Apotheke" in Frauenfeld (TG)

- Customer services, Execution and checking of prescriptions
- Establishment of standard operational procedures (SOPs) for the incoming components inspection according to up-to-date guidelines
- Adaptation of the existing hygiene directives to up-to-date guidelines; including employee training in hygiene-awareness
- Development of an SOP for triage in cases of cystitis (handling and evaluation of a urine test, decision diagrams for pharmacists, actions in case of emergency and general advice for customers)

Supervisor: Susanne Hagebeucker

March – September 2005

Diploma Thesis:

“Phenotype screen with mutants of RNA-binding proteins in yeast”;

- 21 knock out mutant yeast strains were grown under several different stress conditions in order to find a specific phenotype.
- One of the found effects was chosen for further studies using DNA microarrays and immunopurification to analyze changes in gene expression of RNAs usually binding to the protein which is absent in the mutant.

Performed at the Institute of Pharmaceutical Sciences, Pharmacogenomics, ETH

Supervisor:

Dr. André Gerber

2001

Matura, Kantonsschule Rychenberg, Winterthur (ZH)

Further activities and training:

- **WIN09 (Women into Industry) mentee:** mentoring program between University of Basel and Novartis (2009-2010). Mentor: Kirsten Watson (Pharma Development HR)
- **Lab Animal Experiments Licence (CH):** LTK Module 1 (University of Zürich, 2008)

Work Experience:

March – Sept. 2007 Pharmacist at “Coop Vitality Apotheke” in Winterthur and Volketswil (ZH); Customer services, Execution and checking of prescriptions, Incoming components inspection

March-July 2001 Employee at IP Multimedia (Schweiz) AG, in Küsnacht (ZH); Receipt and forwarding of telephone calls, Informing customers in German, English and French, general administrative office assistance

Languages

German: Mother tongue
English: Fluent in speaking and writing; Cambridge Certificate of Proficiency (March 2011)
French: Basic skills

Publications

Gugger, O.S., Kapfhammer, J.P. (2010) Reduced size of the dendritic tree does not protect Purkinje cells from excitotoxic death. *J Neurosci Res*, **88**, 774-783.

Publication in Progress:

Gugger, O.S., Hartmann, J., Birnbaumer, L., Kapfhammer, J.P. P/Q-type and T-type calcium channels, but not TRPC3 channels, are involved in inhibition of dendritic growth after chronic mGluR1 and PKC activation in cerebellar Purkinje cells. *Article accepted for Publication in the Eur J Neurosci. October 2011.*

Conference presentations

June 2011	“The Plastic Brain” meeting, joint Neurex and ENSN workshop, Basel, Switzerland, poster and short plenary presentation
March 2011	SSN Annual Meeting (Swiss Society for Neuroscience), Basel, Switzerland, poster presentation A5
February 2011	USGEB Zürich, Switzerland. Poster No. 58 and short presentation in symposium “Brain Imaging”.
July 2010	FENS forum Amsterdam, Netherlands, poster presentation A17 (supported by a travel grant from SSAHE – Swiss Society of Anatomy, Histology and Embryology)
February 2010	USGEB Lugano, Switzerland, poster presentation T02-20
July 2008	FENS forum Geneva, Switzerland, poster presentation 142.10