Crosstalk between NMDAR Antagonists and Potassium Channels in Murine and Human Lymphocytes

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von	M.Sc. Tanima	Bose	
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Gutachter: Prof. Dr. Ursula Bommhardt Prof. Dr. Ria Baumgrass

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Table of Contents

1.	Su	mma	ary	3
2.	Zu	sam	menfassung	4
3.	Int	rodu	ction	5
3	8.1 1	The li	mmune System	5
	3.1	.1 Tł	ne Innate Immune System	5
	3.1	.2 Tł	ne Adaptive Immune System	6
3	8.2	Sigr	nalling Pathways in Lymphocytes	8
	3.2	.1	T cell Signalling	8
	3.2	.2	B cell Signalling	8
	3.2	.3	TLR Signalling	9
3	8.3	The	Function of the Membrane Potential and Ion Channels in Immune Cells	10
	3.3	.1	Reasons for lons to Cross through the Cell Membrane 1	0
	3.3	.2	Regulation of Intracellular Ca ²⁺ -level1	1
	3.3	.3	Role of Ion Channels in Maintaining the Membrane Potential 1	3
3	3.4	Con	nection between the Neuronal and Immunological Systems1	8
3	8.5	Glut	tamate Receptors1	8
	3.5	.1	Ionotropic and Metabotropic Glutamate Receptors 1	8
	3.5		Ionotroic Glutamate Receptors in the Brain1	
3	8.6	NM	DAR Inhibitors	21
3	8.7	Glut	tamatergic Regulation of Immune Cells 2	26
	3.7	.1	Presence of Glutamate Receptors in Non-neuronal Organs2	26
	3.7	.2	Glutamate Receptors in Immune Cells	27
3	8.8	Aim	of the Study:	27
4.	Ма	iteria	II and Methods 2	<u>'9</u>
4	l.1	Mate	erial 2	29
	4.1	.1	Mouse Lines	29
	4.1	.2	Cell Culture-related	29
	4.1	.3	Microscopy-related	31
	4.1	.4	DNA Constructs	34
	4.1	.5	Instruments	34
	4.1	.6	Software	35
4	.2	Neth	ods 3	6
	4.2	.1	Cell Culture-related	36
	4.2	.2	Microscopy-related	38
	4.2	.3	Statistical Analysis 4	1

5.	Results	42
5.	1 NMDAR Expression in Murine Lymphocytes is Puzzling	42
5.	2 NMDAR Antagonists Attenuate TCR/BCR- induced Ca ²⁺ -flux	45
	5.2.1 TCR-induced Ca ²⁺ -flux	45
	5.2.2 BCR-induced Ca ²⁺ -flux	45
	3 NMDAR Antagonists Modulate the Membrane Potential and Block K _v 1.3 and _{Ca} 3.1 Channels of Lymphocytes	46
	5.3.1 Mouse Primary T Cells and EL-4 Lymphoma Cells	46
	5.3.2 Murine B Cells	48
	5.3.3 Human Jurkat T- and Raji B-Lymphoma Cells	50
	5.3.4 Primary Human T Cells from Healthy Donors	55
5.	4 K _v 1.3 Channel Expression on Blood T cells from Dementia Patients	56
6.	Discussion	58
7.	References	69
8.	Abbreviations	84
9.	List of Figures and Tables	86
10.	Acknowledgements	88
11.	Erklärung	89
12.	Curriculum Vitae	90

1. <u>Summary</u>

N-methyl-D-aspartate receptors (NMDARs) are voltage- and ligand-gated ion channels of the brain. They are key glutamate receptors regulating neuronal synaptic transmission, learning and memory formation. They are also implicated in various neuronal disorders like excitotoxicity, Parkinson's disease, mood disorders or schizophrenia. NMDARs thus are attractive targets to treat neuronal diseases, and NMDAR antagonists like ifenprodil, MK801 or memantine are used to evaluate NMDAR function in vitro. Among them, only memantine is clinically used to treat Alzheimer's type of dementia as it is well tolerated. Memantine is a low-affinity, uncompetitive, voltage-dependent drug with a fast receptor kinetic. Its voltagedependency is explained by its specific blocking site near to the Mg²⁺-binding site or selectivity filter of NMDARs. As NMDARs have been described to be expressed on lymphocytes, the aim of this study was to determine how NMDAR inhibitors affect Tand B-cell function with regard to possible NMDAR activity, membrane potential, and Ca²⁺-mobilization. The uncompetitive antagonists memantine and MK801 and the non-competitive inhibitor ifenprodil reduced T-cell receptor (TCR)- and B-cell receptor (BCR)-induced Ca²⁺-flux in a concentration dependent manner indicating that these antagonists affect T- and B-cell proximal signalling and activation. Interestingly, TLR4/LPS-induced B-cell activation was also impaired by the antagonists as they depolarized the membrane potential. However, in patch clamp studies NMDAR currents on lymphocytes could not be detected. Instead, it was found that the used antagonists cross-inhibit K_v1.3 and K_{ca}3.1 K⁺-channels on primary murine and human lymphocytes and Jurkat and Raji lymphoma cell lines. K_v1.3 and K_{Ca}3.1 channels are the major K⁺-channels expressed on lymphocytes that maintain the membrane potential as important parameter for Ca²⁺-fluxes and, thereby, influence many Ca²⁺-dependent cell responses. Notably, peripheral blood T cells of Alzheimer patients treated with memantine for 1 week showed a strong reduction of K_v 1.3 channel activity, but after 12 weeks of treatment K_v 1.3 channel activity was nearly similar to the values before treatment. In conclusion, using electrophysiological and immunological techniques, this study shows substantial effects of NMDAR antagonists on lymphocytes, most likely through cross-inhibition of K_v1.3 and K_{ca}3.1 channels. The pharmacological cross-reactivity of memantine may be harmful for lymphocytes and, therefore, an immunological surveillance during memantine application is suggested.

2. Zusammenfassung

N-methyl-D-Aspartat-Rezeptoren (NMDARen) sind spannungs- und ligandengesteuerte Ionenkanäle des Gehirns. Sie spielen eine wichtige Rolle bei der synaptischen Übertragung und der Regulation von Lern- und Gedächtnisprozessen. Zu den NMDAR-assoziierten Störungen zählen Exzitotoxizität, Morbus Parkinson, affektive Störungen und Schizophrenie. Bei der Behandlung neuronaler Erkrankungen sind NMDARen daher wichtige pharmakologische Angriffspunkte. NMDAR-Antagonisten, wie Ifenprodil, MK801 und Memantin, werden bei in vitro Untersuchungen zur NMDAR-Funktionalität verwendet. Aufgrund der guten Verträglichkeit wird aber lediglich Memantin zur Behandlung von Alzheimer-Demenzen eingesetzt. Memantin ist ein niedrig-affiner, nicht kompetitiver und spannungsabhängiger Inhibitor mit schneller Rezeptorkinetik. Er blockiert die Mg²⁺-Bindungsstelle, also den Selektivitätsfilter von NMDARen und behindert derart den Ca²⁺-Einstrom in die Zelle. Die Expression von NMDARen wurde auch für Lymphozyten beschrieben. In der vorliegenden Studie wurde der Einfluss von NMDAR-Inhibitoren auf die NMDAR-Aktivität, das Membranpotential und die Ca²⁺-Mobilisierung in Lymphozyten ermittelt. Die Antagonisten Memantin, MK801 sowie Ifenprodil reduzierten den T-Zellrezeptor- und B-Zellrezeptor-induzierten Ca²⁺-Influx in konzentrationsabhängiger Weise. Dies weist darauf hin, dass die Antagonisten die proximalen Signalwege der Antigenrezeptoren und die Aktivierung von B- und T-Zellen beeinflussen. Auch die TLR4/LPS-induzierte B-Zellaktivierung wurde durch die Inhibitoren gehemmt, wobei auch hier eine Depolarisation des Membranpotentials durch die Inhibitoren nachgewiesen wurde. In den Lymphozyten wurden jedoch keine NMDAR-Ströme mittels Patch-Clamp vorgefunden. Stattdessen wurde aufgezeigt, dass die verwendeten Antagonisten K_v1.3 und K_{Ca}3.1 K⁺-Kanäle in primären Lymphozyten sowie Jurkat- und Raji-Lymphom-Zellen kreuzinhibieren. Diese Kanäle sind die wichtigsten K⁺-Kanäle der Lymphozyten zur Regulation des Membranpotentials und damit Ca²⁺-abhängiger Zellantworten. T-Zellen von Alzheimer Patienten zeigten nach einwöchiger Behandlung mit Memantin eine Reduktion der K_v1.3-Kanalaktivität, die sich nach 12-wöchiger Behandlung wieder auf das Ausgangsniveau normalisierte. Mittels elektrophysiologischer und immunologischer Techniken wurden somit substantielle Effekte von NMDAR-Inhibitoren auf Lymphozyten nachgewiesen, die wahrscheinlich auf Kreuzreaktionen mit K_v1.3 und K_{Ca}3.1 K⁺-Kanälen beruhen. Eine immunologische Überwachung bei Memantinbehandlung ist daher empfehlenswert.

3. Introduction

3.1 The Immune System

The cellular defence network of higher vertebrates against foreign antigens is a sophisticated system involving the interplay between several types of leukocytes, principally macrophages, dendritic cells (DCs), granulocytes, natural killer (NK) cells, T and B lymphocytes. The specificity, flexibility and efficiency of this security system are achieved by tightly regulated interactions between these immune cells.

The immune system is divided into two parts: the innate and adaptive arms provide the two lines of defence against invading pathogens. The cells of the innate immune system provide a first line of defence against many common microorganisms. However, they cannot always eliminate infectious agents, and some pathogens cannot be recognized due to the relatively poor diversity of specific recognition elements of innate cells. The lymphocytes of the adaptive immune system have evolved to provide a more versatile means of antigen-specific defence, which also provides increased protection against subsequent re-infection with the same pathogen. The cells of the innate immune system, however, play a crucial part in the initiation and subsequent direction of the adaptive immune response, and participate in the removal of pathogens that have been targeted by the adaptive immune response becomes effective, the innate immune response has a critical role in controlling infections during this period (Rouzaire-Dubois B et al. 2002).

3.1.1 The Innate Immune System

The innate immune system consists of cells and proteins, like the complement cascade, that are always present and ready to be mobilized to fight foreign antigens at the site of infection. The main components of the innate immune system can be classified into anatomic, physiologic, phagocytic and inflammatory mediators for occluding the entry of foreign organisms. The anatomical barriers consist of skin and mucous membrane to inhibit the entry of foreign antigens into the body. Skin is the major mechanical barrier retarding the entry of microbes. The physiological obstacles are higher temperature, low pH, and chemical mediators like lysozyme, interferons (IFs), complement, collectins, or Toll-like receptors (TLRs). TLRs can recognize microbial molecules like lipopolysaccharide (LPS) of gramnegative bacteria and send signals into the cells to secrete immune-stimulatory cytokines. Besides these two obstacles, there are cellular barriers formed by phagocytic and inflammatory branches. Blood monocytes, neutrophils, and tissue macrophages can phagocytose, digest and kill foreign microorganisms. The inflammatory process consists of a series

of events started by tissue damage and infection and is caused by the chemical mediators released by the phagocytic cells engulfing the pathogen, for instance bacteria. This whole process of tissue damage due to the phagocytosis process is characterized by four physical signs. These 'four cardinal signs of inflammation' are *rubor* (redness), *tumour* (swelling), *calor* (heat), and *dalor* (pain) as already described by the Roman physician Celsus in 1600 BC. Physiologically, this is caused by the vasodilation of the blood vessels, the increase in capillary permeability and influx of phagocytes from the capillaries into the tissues. Phagocytic cells accumulate at the site of inflammation, phagocytose bacteria, and release lytic enzymes, which can damage nearby healthy cells. Additionally, chemical mediators released due to tissue damage include serum proteins called acute-phase proteins. These, histamine, kinins, fibrin, and C-reactive protein become concentrated hugely in tissue-damaging infections and take part in the inflammation machinery (Kuby J 2003; Rouzaire-Dubois B 2002; Murphy K 2012).

3.1.2 The Adaptive Immune System

The innate immune response makes a crucial contribution to the activation of adaptive immunity. The inflammatory response increases the flow of lymph containing antigen and antigen-bearing cells like DCs into lymphoid tissue, while complement fragments on microbial surfaces and induced changes in activated phagocytes provide signals that synergize in activating lymphocytes whose antigen-receptors bind to specific microbial antigens. Macrophages that have phagocytosed bacteria and become activated can activate T lymphocytes. However, the cells that specialize in presenting antigen to T lymphocytes and are most effective in initiating adaptive immunity are the DCs.

The induction of an adaptive immune response begins when a pathogen is ingested by an immature DC in the infected tissue. These specialized phagocytic cells are resident in most tissues and are relatively long-lived, turning over at a slow rate. They derive from the same bone marrow precursor as macrophages, and migrate from the bone marrow to their peripheral locations, where their role is to survey the local environment for pathogens. After antigen-uptake, tissue-resident DCs migrate through the lymph to the regional lymph nodes where they interact with naive T lymphocytes. On activation, the immature DC matures into a highly effective antigen-presenting cell (APC) and undergoes changes that enable it to activate antigen-specific T cells in the lymph node. APCs process and present antigens bound to major histocompatibility complex (MHC) molecules which are recognized by the antigen-specific T-cell receptor (TCR) expressed on the T cells (Samelson et al. 1985). MHC class I and class II molecules are expressed on all nucleated cells and professional APCs, respectively. MHC-I molecules consist of an α -chain and β_2 -microglobulin and are

recognized by CD8 co-receptors and TCRs of cytotoxic T-cells (T_c). MHC-II molecules consist of an α - and β -chain and are recognized by CD4 co-receptors and TCRs present on T helper cells (T_H).

The adaptive immune response is mediated through humoral and cell-mediated responses. Humoral immunity is mediated by B cells, which produce antibodies after activation and differentiation into plasma cells. Antibodies are secreted into the circulation and mucosal fluids, neutralize extracellular microbes, microbial toxins and virus present in the blood and lumen of mucosal organs, such as the gastrointestinal and respiratory tracts. Defence against intracellular microbes and virus relies on cell-mediated immunity by T lymphocytes. T_H cells secrete cytokines which support phagocytosis to destroy ingested microbes and B cell differentiation. T_c cells destroy host cells harbouring intracellular virus and tumor cells. T cells recognize only processed protein antigens, whereas B cells and antibodies are able to recognize different types of antigens, including proteins, carbohydrates, nucleic acids, and lipids.

The defence system of innate immunity is effective in combatting many pathogens. However, it is constrained by relying on relatively few fixed germline-encoded receptors to recognize microorganisms which evolve more rapidly than the host cells they infect. In contrast, the antigen recognition mechanism used by lymphocytes in adaptive immune response has evolved to overcome the constraints faced by the innate immune system to specifically recognize an almost infinite diversity of antigens. The diversity and specificity of the antigen receptors of T and B cells is determined by a unique genetic mechanism, named gene recombination that operates during lymphocyte development in the bone marrow (where B cells are formed) and thymus (where T cells are generated) and generates a huge diversity of antigen-specific TCR and BCR molecules. In 1950, Macfarlane Burnet already postulated in the 'clonal selection theory' the pre-existence of many different potential antibody-producing cells in the body, each having the ability to make antibody of a different specificity (Rouzaire-Dubois 2002). Thus, although an individual lymphocyte carries antigen receptors of only one specificity, the antigen receptor specificity of each lymphocyte is different. On binding specific antigen, T and B cells are activated to divide and produce many identical progeny, known as clonal selection. Expanded B cells differentiate to produce antibodies with specificity identical to that of the BCR that triggered activation and clonal expansion (Kuby J 2003; Rouzaire-Dubois B 2002; Murphy K 2012).

A hallmark of the adaptive immune response is the generation of memory cells which get activated when the same pathogen is encountered another time and leads to a faster immune response. The immunological memory is the basis for vaccination or transplantation studies. All human memory T cells express the surface marker CD45RO⁺, whereas CD45RA⁺ and CCR7 (CD197) mark the naive and central memory cell population, respectively (Mackay 1999).

3.2 Signalling Pathways in Lymphocytes

3.2.1 T cell Signalling

TCR ligation induces a number of signalling events that ultimately determine the T cell fate through regulating cytokine production, cell survival, proliferation, and differentiation. T-cell activation is initiated by ligation of the membrane-associated TCR and 'cluster designation' molecules CD4 or CD8 by MHC-II or MHC-I molecules presenting specific peptide on the surface of an APC. TCR activation is initiated by the phosphorylation of immunoreceptor tyrosine-based motifs (ITAMs) within the TCR_c chains of the TCR/CD3 complex by lymphocyte protein tyrosine kinase (Lck). The CD45 receptor tyrosine phosphatase dephosphorylates and activates Lck and other Src family tyrosine kinases. Zeta-chain associated protein kinase (Zap-70) is recruited to the phosphorylated TCR_C chains where it becomes activated by Lck and then phosphorylates downstream adaptor or scaffold proteins like LAT (Linker for Activated T cells). Phosphorylation of SH2-domain-containing leukocyte protein of 76 kD (SLP-76) by Zap-70 promotes recruitment of Vav (a guanine nucleotide exchange factor) and inducible T cell kinase (Itk). Phosphorylation of phospholipase Cy1 (PLCy1) by Itk results in the hydrolysis of phosphatidylinositol 4,5-bisphosphate (PIP₂) to produce the second messengers diacylglycerol (DAG) and inositol-1,4,5-triphosphate (IP₃). DAG activates protein kinase $C\theta$ (PKC θ) and the mitogen-activated protein kinase (MAPK)/Erk pathway, both promoting activation of the transcription factor nuclear factor-κB (NF-κB). IP₃ triggers the release of Ca²⁺ from the endoplasmic reticulum (ER), which promotes entry of extracellular Ca²⁺ into the cell through calcium release-activated Ca²⁺ channels (CRAC) (Weiss 2009, 2010).

3.2.2 B cell Signalling

Similar to T cells, the B-cell receptor (BCR) complex consists of two modules: the antigenbinding and the signal transducing moiety. The antigen-binding moiety in essence is an immunoglobulin that is integrated into the lipid bilayer of the plasma membrane through a hydrophobic transmembrane domain. The signal-transducing moiety is a disulphide-linked heterodimer (CD79) consisting of CD79a and CD79b, also called Iga and Ig β , respectively. Binding of antigen to the BCR triggers phosphorylation of the ITAM tyrosine residues in Iga and Ig β by the tyrosine protein kinase Lyn which initiates a signalling cascade through activation of bruton tyrosine kinase (Btk) and PLC γ 2 followed by IP₃ and DAG generation and Ca²⁺-influx. Subsequent signals involve activation of phosphatidylinositol 3-kinase (PI3-K) and Akt, Erk and NF- κ B pathways, which contribute to B cell activation and proliferation (Wienands and Engels 2001).

3.2.3 TLR Signalling

Besides ligation of the BCR, B cells can be activated by TLRs. The Interleukin-1 receptor (IL-1R)/TLR superfamily of single transmembrane domain receptors comprises 24 members, including five adaptor proteins, which share a cytosolic domain named Toll/IL-1 receptor (TIR) domain. The TLR family includes 10 members, which play a key role in activating innate immune cells and B cells. TLRs recognize pathogen associated molecular patterns (PAMPs), which are highly conserved motifs of common pathogens. Each TLR has a distinct pattern-recognition specificity (Lee and Kim 2007). For example, TLR4 recognizes LPS of bacteria. However, a number of endogenous molecules (named as 'damage-associated molecular patterns' or 'danger signals') released by injured tissue can also activate the innate immune system via stimulation of certain TLRs (Bianchi 2007; Tsan and Gao 2004). IL-1R/TLR receptors are ubiquitously expressed by leucocytes (Janeway and Medzhitov 2002), epithelial cells (Yoshimoto and Nakanishi 2006) and endothelial cells (Gibson et al. 2006). Members of this family of receptors have also been shown to be expressed on CNS neurons, microglia and astrocytes (Vezzani et al. 2011). The critical role of the TLR4 pathway in treating severe sepsis and septic shock was delineated in many clinical trials (Wittebole et al. 2010).

Ligand binding of the TLRs leads to the recruitment of MyD88 and other cytosolic adaptor proteins to the TLR complex, activating IL-1R-associated kinase1/4 (IRAK1/4) and tumor necrosis factor receptor associated factor 6 (TRAF6), leading eventually to the expression of genes involved in inflammation, including the transcription factors NF-κB, activator protein-1 (AP-1) and interferon regulatory transcription factors (IRFs) (O'Neill and Bowie 2007). TLR3 and TLR4 can also signal using a MyD88-independent pathway which involves TIR-domain-containing adaptor-inducing interferon-ß (TRIF). The TRIF-dependent signalling cascade results in the activation of IRF-3, which then induces IFNα and IFNβ. Induction of PI3-K can also occur in response to TLR stimulation, presumably using a MyD88-independent pathway (Davis et al. 2006; Diem et al. 2003).

These signalling pathways are orchestrated through the interplay of different ion channels (CRAC, K_v 1.3, K_{Ca} 3.1, TRPM4 and P2X). These ion channels regulate the normal membrane potential, which when altered affects the activation of downstream signalling cascades leading to cell proliferation, differentiation or cell death.

3.3 The Function of the Membrane Potential and Ion Channels in Immune Cells

3.3.1 Reasons for lons to Cross through the Cell Membrane

Physiological processes depend on the continued flow of ions into and out of cells. Ions are more 'stable' in water than in the lipid layer inside the membrane. Thus, the hydrophobic membrane acts as a serious energy barrier for transporting ions. In a situation without biological pumps and ion channels, there can be large ion potential differences between the two sides of a biological membrane so that the predominant ions Na⁺, K⁺, Ca²⁺, and Cl⁻ can never cross it. To resolve this issue, ion pumps, ion exchangers ('active' transport) and ion channels ('passive' transport) are used in cells. An ion channel needs a single gate and ion pump works with at least two gates. A gate or a selectivity filter is considered to be a part of a protein that hinders ion movement along the translocation pathway in the prohibitive confirmation but not in the permissive confirmation. The ion channels, like voltage-gated Na⁺-, Ca²⁺- or K⁺-channels, are opened when a change of membrane potential displaces the voltage sensors connected to a cytoplasmic side of the 'activation gate'. They can be closed by reversal of those displacements ('deactivation') in response to an opposite change of membrane voltage. But even with their activation gates in the permissive position, the ion pathway through those channels can be closed by a separate gating process called 'inactivation'. Both these gates should be in a permissible position for the channel to conduct ions, and closure of either gate obstructs the ion flow (Gadsby 2009). In contrast, ion pumps are controlled by timely cohesion of two gates which are never open simultaneously (Lauger 1979). Instead, the chosen ions are allowed to enter the pathway from one side of the membrane while one of the gates is open, and then to leave at the other side of the membrane through another gate after the first one shuts down (Vidaver 1966; Gadsby 2009). Although these two transport systems work differently, ion selectivity is a prime criterion for both of them. The ion pumps generally transport ions against the electrochemical gradient with the use of energy like adenosine tri-phosphate (ATP) and a relatively slow speed. In contrast, ion channels are passive transporters of ions with a very high ion conduction rate to maintain the proper membrane potential. The membrane potential is defined as the electrical potential difference between the interior and exterior of the cell. It can be defined with Ohm's law (V = IR, where V = voltage, I = current and R = resistance). Voltage in this law is synonymous to the difference in the electrical potential, thereby, the ability to drive electric current across a resistance. The plasma membrane of a cell acts as an electrical resistor as it has low intrinsic permeability to ions. Insertion of ion-specific channels creates a membrane potential, which depends on the ion selectivity of the ion channels. Most eukaryotic cells possess selective channels for K⁺ and Cl⁻ resulting in a membrane potential between -40 to -80 mV depending on the cell type (Gouaux and Mackinnon 2005; Dubyak 2004).

The resting potential of excitable cells like neurons and muscles varies from -70 to -80 mV, whereas for non-excitable cells like immune cells it is between -40 to -50 mV. The difference in the resting potential of these cell types relates to their function (Hille B 2001). The opening and closing of transmembrane ion channels alter the resting potential. It is called depolarization, if the interior voltage becomes less negative, and hyperpolarization, if the voltage becomes more negative. In excitable cells, a sufficient depolarization of the membrane can evoke an action potential for a short time period, which transiently repolarises the membrane. In immune cells, changes in the membrane potential are less transient and depend on the activation state of the cell. Maintaining a normal membrane potential is important for any immune cell. Thus, several ion channels (described in **section 3.3.3**) are expressed in the plasma membrane to allow the modulation of the membrane potential.

In non-excitable cells including immune cells, the membrane potential plays a prominent role in setting the electrical driving force for Ca²⁺-influx. In cells where voltage-independent Ca²⁺-channels like TRPM4 and K_{2P} are present, Ca²⁺-influx only depends on the electrochemical gradient over the membrane and is stronger if the membrane potential is more negative (hyperpolarised). In contrast, Ca²⁺-influx is mediated through voltage dependent Ca²⁺-channels in excitable cells, like neurons. Functional coupling of Ca²⁺-activated K⁺- channels (K_{Ca}) with other predominant ion channels leads to a positive feedback cycle promoting a sustained Ca²⁺-influx in case of immune cells (**Figure 1**) (Gao et al. 2010) or a negative feedback to terminate the flux by hyperpolarising the membrane potential and hence closing voltage-activated Ca²⁺-channels in neurons (Fakler and Adelman 2008).

3.3.2 Regulation of Intracellular Ca²⁺-level

Na⁺ and K⁺ are the most abundant cations in biological systems. Na⁺ ions are most often present at high concentrations outside the cell, and K⁺ is present at high concentrations inside the cell. Gradients for these ions across the cell membrane provide the energy source for action potentials generated by opening Na⁺ and K⁺-channels (Murata et al. 2005; Meier et al. 2005), and for moving solutes and other ions across the cell membrane via coupled transporters. Among several ions, the gradient for Ca²⁺ ions is the largest. It helps in controlling several physiological processes like secretion, excitation, contraction, and cellular proliferation (Berridge 1995; Berridge et al. 2000). The cytosol is surrounded by two massive Ca²⁺-stores: the extracellular space, where the Ca²⁺ concentration is ~1.8 mM, and the sarco-endoplasmic reticulum (SER), where the Ca²⁺ concentration varies from 300 µM to 2 mM (Hannaert-Merah et al. 1995). In immune cells, the intracellular Ca²⁺ concentration is ~0.1 µM in the resting state, but it is increased 10-fold when the cells are activated (Feske et al. 2012; Hoth and Penner 1992). Plasma membrane Ca²⁺-channels are specifically important in cell-cycle progression and proliferation of immune cells. Accumulating evidence suggests that Ca²⁺-influx is important at different steps of the cell cycle, the progression at late G1 into S phase as well as G2/M transition (Takuwa et al. 1991; Takuwa et al. 1992; Nordstrom et al. 1992). The nature of Ca²⁺-channels involved in proliferation is well-defined and can be separated into voltageactivated (VACC), receptor-activated (RACC), store-operated (SOCC) and second messenger-operated (SMOCC) channels. RACC, SOCC and SMOCC are ubiquitous, whereas VACC is expressed in excitable cells only. VACC (e.g. L-, T-, N-, P-, Q-type Ca²⁺-channels) open when the membrane is depolarized (Tsien et al. 1995). RACC (e.g. P2X purinergic receptors) open when a ligand binds to the channel (MacKenzie et al. 1999), whereas SOCC (e.g. transient receptor potential (TRP) (Clapham et al. 2001) and archetype CRAC (Hoth and Penner 1992) are activated when the level of Ca²⁺ within the lumen of the ER drops below a threshold level (Putney 1986; Putney and McKay 1999; Putney et al. 2001). Another type, SMOCC (e.g. arachidonic acid-regulated Ca²⁺ current) is activated by intracellular second messengers like arachidonic acid (Shuttleworth 1996). The role of CRAC, TRPM4 and P2X channels are important in case of immune cells in the continuous struggle to keep Ca²⁺ at an optimal level important for the maintenance of cellular functions in parallel with ion pumps like Na^+/K^+ pumps (Mijatovic et al. 2007; Lefranc and Kiss 2008).

Among the different ion channels mentioned in **Figure 1** involved in the regulation of Ca²⁺ homeostasis, CRAC channels are the most important ones. CRAC channels have been extensively characterized (Hoth and Penner 1992; Zweifach and Lewis 1993) and are distinguished by an extremely high ion selectivity for Ca²⁺ and a low conductance (Prakriva 2009). CRAC channels are activated through the binding of the ER Ca²⁺ sensors stromal interaction molecule 1 (STIM1) and STIM2 to the CRAC channel proteins ORAI1-3 (also known as CRACM1-3) (Hogan et al. 2010). ORAI1 is a widely expressed surface glycoprotein with four predicted transmembrane domains, intracellular amino- and carboxyl-termini and no sequence homology to other ion channels except for its homologues ORAI2 and ORAI3. All three ORAI proteins form Ca²⁺-channels with broadly similar functional properties when ectopically expressed, although they differ in their inactivation characteristics, pharmacological properties and tissue expression (Lis et al. 2007; DeHaven et al. 2007). The activation of ORAI CRAC channels involves a complex series of coordinated steps, during which STIM proteins fulfil two crucial roles. Firstly, they sense the depletion of ER Ca²⁺-stores, and secondly, they communicate store depletion to the CRAC channels (Liou et al. 2005; Roos et al. 2005; Cahalan 2009). In resting cells with repleted Ca²⁺-stores. STIM proteins are diffusely distributed throughout the ER membrane. Following the depletion of Ca²⁺-stores, STIM proteins are activated, oligomerize and redistribute into discrete

punctae located in junctional ER sites that are in close proximity to the plasma membrane (Cahalan 2009; Luik et al. 2008). Lymphocytes express two closely related STIM isoforms, STIM1 and STIM2, and both mediate store-operated Ca²⁺-entry (SOCE) in B and T cells (Matsumoto et al. 2011; Oh-Hora et al. 2008). CD4⁺ and CD8⁺ T cells from ORAI1- and STIM1- deficient patients and mice show defective production of many cytokines, including IL-2, IL-17, IFN- γ and tumour necrosis factor (TNF) (Feske 2009; Feske et al. 2001). However, despite the profound defects in SOCE in B cells from ORAI1- and STIM1-deficient patients and mice, CRAC channels do not have a major role in antibody production (Feske et al. 2010).

3.3.3 Role of Ion Channels in Maintaining the Membrane Potential

The resting potential of a lymphocyte membrane is ~-50 mV (Lewis and Cahalan 1995). Membrane potential alterations mainly occur when lymphocytes get activated (Crabtree 1999). TCR engagement activates PLCy1, which catalyses the hydrolysis of PIP₂ into IP₃ and DAG. IP₃ stimulates the release of Ca²⁺ from intracellular ER stores which triggers the opening of plasma membrane CRAC channels. It is the resulting influx of extracellular Ca²⁺ that is responsible for the sustained rise in cytoplasmic Ca2+ after TCR stimulation. Ca2+ binds to the cytoplasmic Ca²⁺-dependent protein calmodulin which then activates the phosphatase calcineurin. This phosphatase dephosphorylates and activates the nuclear factor of transcription of activated T cells (NFAT), which enters the nucleus and helps to initiate interleukin-2 (IL-2) gene transcription (Serfling et al. 2007; Rao 2009). During the activation of immune cells opened CRAC-channels raise the intracellular Ca²⁺ level. To maintain the balance in membrane conductance, K_{Ca} channels get opened to hyperpolarize the membrane as this channel helps in Ca²⁺-efflux. A negative feedback loop is established until Ca²⁺ reaches high enough levels inside the cell to inhibit CRAC-channels. With the Ca²⁺dependent activation of TRPM4 channels in T cells, there is also involvement of K_v1.3 channels to repolarize the membrane (as illustrated in **Figure 1**). Along with these conventional ion channels, the two-pore K⁺-channels (K_{2P}) TASK-1 and TASK-3 are known to regulate immune cell effector functions by hyperpolarizing the membrane (Meuth et al. 2008). Although the best characterized channel for Ca²⁺-influx in T cells is CRAC, several other channels may also mediate Ca²⁺-influx in T cells, including members of the transient receptor potential (TRP) family, P2X receptors and voltage-gated Ca²⁺ (Ca_v) channels. Compared to CRAC channels, however, their contribution to TCR-induced Ca²⁺-influx in immune cells is less well-defined (Feske 2013).



Figure 1. Oscillatory changes of the membrane potential in lymphocytes. Ca^{2+} -influx in lymphocytes depends on the gradient between the extracellular Ca^{2+} concentration (~1 mM) and the intracellular Ca^{2+} concentration (~0.1 µM) and on an electrochemical gradient established by the K⁺-channels (K_v1.3, K_{Ca}3.1 and partially by K_{2P} channels) and the Na⁺-permeable channel TRPM4 (Cahalan and Chandy 2009; Chandy et al. 2004). CRAC channels are activated following the engagement of the TCR or BCR and is mediated through the activation of PLC γ , the production of IP₃ and the release of Ca²⁺ from ER Ca²⁺-stores. The ensuing activation of STIM1 and STIM2 results in the opening of ORAI1 CRAC channels and SOCE. Sustained Ca²⁺-influx through CRAC channels leads to the activation Ca²⁺-dependent enzymes and transcription factors, including calcineurin and NFAT (Lewis 2001; Hogan et al. 2010; Feske 2007). Additionally, P2X receptors (e.g. P2X4 and P2X7), which are non-selective Ca²⁺-channels, are activated by extracellular ATP mediating Ca²⁺-influx (modified from *Launay, P., 2004*) (Feske et al. 2012; Launay et al. 2004).

3.3.3.1 K⁺-channels

K⁺-channels are encoded by a superfamily of 78 genes (Harmar et al. 2009) and are involved in diverse physiological and pathological processes (Wulff et al. 2009). Structurally, K⁺-channels are classified into three major groups: six transmembrane (**Figure 2A**), two transmembrane (Figure 2B) and four transmembrane (**Figure 2C**) K⁺-channels. Voltagegated K⁺-channels (the first type) can further be subdivided into four families: K_v (shakerlike), Ether-a-go-go (EAG), KCNQ and K_{Ca} (Ca²⁺-activated K⁺-channels) (Yellen 2002; Shieh et al. 2000). K_{Ca} channels can be classified into three types: BK_{Ca} (K_{Ca}1.1), IK_{Ca} (K_{Ca}3.1) and SK_{Ca} (K_{Ca}2.1, K_{Ca}2.2, K_{Ca}2.3) (Wei et al. 2005). Among K⁺-channels, immune cells mainly express voltage-activated (K_v 1.3), calcium-activated (K_{Ca} 3.1) and two-pore K_{2P} channels (TASK-1, TASK-3). K⁺-channels protect against membrane depolarization by mediating the efflux of K^+ to depolarize the plasma membrane (Cahalan and Chandy 2009). K_v1.3 is a homotetramer of four α -subunits, each composed of six transmembrane segments (S1-S6), and is activated by membrane depolarization (Cahalan et al. 1985). Depolarization of the membrane is sensed by four arginine residues that are localized in the S4 segment and results in a conformational change that causes channel opening (Bezanilla 2008). $K_{Ca}3.1$ is a Ca²⁺-activated K⁺-channel, but it has similar membrane topology and pore architecture as K_v1.3. However, rather than containing a voltage sensor, the C- terminus of K_{Ca}3.1 is constitutively bound to calmodulin, and channel opening occurs after Ca²⁺ binds to calmodulin (Xia et al. 1998). K_{Ca}3.1 channels powerfully hyperpolarize the membrane following elevations in the intracellular Ca²⁺ concentration and thus help to sustain the driving force for Ca²⁺-entry. In addition to the requirement of Ca²⁺, K_{Ca}3.1 channel activity depends on a class II PI3-K, which increases the concentration of phosphatidylinositol 3phosphate (PtdIns(3)P) in the plasma membrane (Feske et al. 2012). TASK-1 and TASK-3 also contribute to hyperpolarizing activities (Meuth et al. 2008).

The relative contribution of $K_v 1.3$ and $K_{Ca} 3.1$ in lymphocyte Ca^{2+} -influx are determined primarily by their expression level, which depends on the lymphocyte subset and its state of activation. Under resting conditions, CCR7⁺CD45RA⁺ naïve human T cells predominantly express $K_v 1.3$ channels and depend on $K_v 1.3$ for activation (Leonard et al. 1992). Following activation, naïve human T cells up-regulate K_{Ca}3.1 expression (Ghanshani et al. 2000), and inhibition of K_{Ca}3.1 in pre-activated T cells blocks TCR-stimulated Ca²⁺-influx and proliferation (Fanger et al. 2000; Fanger et al. 2001). Furthermore, mouse T_H1 and T_H2 cells predominantly express K_{Ca}3.1 and depend on K_{Ca}3.1 for TCR-mediated Ca²⁺-influx and cytokine production, whereas $T_H 17$ cells mainly express $K_v 1.3$ and require $K_v 1.3$ for their activation and production of IL-17 (Di et al. 2010). Differential use of K⁺-channels is also observed in effector memory T cells (T_{EM}) and central memory T cells (T_{CM}) (Cahalan and Chandy 2009; Srivastava et al. 2006; Beeton et al. 2001). When activated at sites of inflammation, T_{EM} cells, which have the phenotype CCR7⁻CD62L^{low}CD45RA⁻, produce various cytokines including IFN_γ, IL-4 and IL-5, and exclusively up-regulate K_v1.3 expression. In contrast, T_{CM} cells, which are CCR7⁺CD62L^{hi}CD45RA⁻, up-regulate the expression of K_{Ca}3.1 following their activation in lymph nodes and mucosal lymphoid organs. As a result, K_v1.3 blockers are effective inhibitors of T_{EM} cells, whereas $K_{Ca}3.1$ blockers are effective at inhibiting T_{CM} cells.

Figure 2. Scheme of structural classification of K⁺-channel subunits. Structurally, K⁺-channels

are classified into three types. (A) Sixtransmembrane one-pore. These channels are composed of four subunits, each of them with six transmembrane segments (S1-S6) with a voltage-sensor domain (S4) and a conducting S5 S6. pore between and (B) Twotransmembrane one-pore. The inwardrectifying channels belong to this group with four subunits each consisting of two transmembrane segments (M1 and M2) with a conducting pore in between. (C) Fourtransmembrane two-pore. The ion channels belonging to this group consist of fourtransmembrane (M1-M4) and two-pore domains (P1 and P2) (Modified from Shieh, C.C. et al., 2000) (Shieh et al. 2000). K_v: voltagegated; IK: delayed-rectifier; K_{ir}: inward-rectifier; TWIK: Two-pore weak inward rectifier; TREK:



TWIK-related; TASK: TWIK-related acid-sensitive; TALK: TWIK-related alkaline pH activated; TRAAK: TWIK-related arachidonic acid-stimulated K⁺-channel.

The finding that $K_v1.3$ and $K_{Ca}3.1$ function to activate distinct lymphocyte subsets provides an opportunity to more selectively target lymphocyte subsets for therapeutic purposes. The relevance of these findings to humans was demonstrated by the observation of high levels of $K_v1.3$ expression by myelin-reactive T cells isolated from patients with multiple sclerosis (Wulff et al. 2003). Similar studies have shown an increase of $K_v1.3$ channels in diseaseassociated T_{EM} cells in patients with type I diabetes, rheumatoid arthritis (RA) and psoriasis, and the treatment of these diseases with $K_v1.3$ blockers lead to the amelioration of the disease (Beeton et al. 2006; Fasth et al. 2004; Friedrich et al. 2000; Gilhar et al. 2011). By contrast, inhibition of $K_{Ca}3.1$ protected mice from developing colitis in two mouse models of inflammatory bowel disorder (Di et al. 2010), suggesting that $K_{Ca}3.1$ may be a novel therapeutic target to treat patients with Crohn's disease or ulcerative colitis.

 K_v 1.3 is one of the first voltage-gated K⁺-channels reported to be modulated during apoptosis (Szabo et al. 1996) and is shown to contribute to the increased K⁺-efflux underlying the late phase of lymphocyte apoptosis. There is also proof of stimulation of these channels by death receptor CD95/Fas during apoptosis of Jurkat T lymphocytes (Storey et al. 2003). K_v 1.3 expression also shows a correlation with tumour progression as exemplified by the up-regulation of K_v 1.3 expression in diffuse human large B-cell lymphoma and glioma (Preussat et al. 2003). Mechanistically, pro-apoptotic mediator cytochrome C can activate K_v -channels while anti-apoptotic protein Bcl-2 inhibits them (Remillard and Yuan 2004). K_v 1.3 depletion decreases the expression of Caspase-3, Caspase-9 and Bad, molecules that exacerbate apoptotic cell death (Wasserman and Koeberle 2009). This mechanism indicates that inhibition of K_v 1.3 channels confers resistance to apoptosis while their over-expression favours this process.

3.3.3.2 Transient Receptor Potential (TRP) Channel

In humans, TRP channels form a large superfamily of 28 cation channels, which can be divided into 7 subfamilies (Venkatachalam and Montell 2007). T cells predominantly express channels belonging to TRPC and TRPM subfamilies, including TRPC1, TRPC3, TRPC5, TRPM2, TRPM4, and TRPM7 (Wenning et al. 2011). Most TRP channels are nonselective and permeable to several cations, including Ca^{2+} and Na^{+} (Ramsey et al. 2006; Owsianik et al. 2006). The function of TRPM4 channels is well documented in T cells and other immune cells unlike most other TRP channels. TRPM4 channels mainly conduct Na⁺ and K⁺ and, in contrast to other TRP channels, are only weakly permeable to Ca²⁺ (Vennekens and Nilius 2007). The activation of TRPM4 channels, which occurs in response to an increase in intracellular Ca²⁺ concentration, results in Na⁺-influx, membrane depolarization and a reduction in the electrical driving force for Ca²⁺-influx. TRPM4 channels thus provide a negative feedback mechanism for the regulation of SOCE and were proposed to prevent cellular Ca²⁺-overload. Given that TRPM4 and K_v channels elicit opposing effects on the membrane potential, it remains to be elucidated precisely how TRPM4 works together with K_v1.3 and K_{Ca}3.1 to regulate changes in the membrane potential and intracellular Ca²⁺ concentration (Feske et al. 2012).

3.3.3.3 Purinergic P2X Receptor Channel

P2X receptors are a family of non-selective ion channels that are activated by extracellular ATP and regulate the influx of Na⁺, Ca²⁺ and other cations (Junger 2011). At least three different P2X receptors have been implicated in Ca²⁺-influx in human T cells: P2X1, P2X4 (Woehrle et al. 2010) and P2X7 (Yip et al. 2009). Their opening, especially that of P2X7, causes Ca²⁺-influx and the activation of downstream signalling molecules such as calcineurin, resulting in the proliferation of B and T cells (Baricordi et al. 1996; Padeh et al. 1991) and IL-2 production (Adinolfi et al. 2005; Woehrle et al. 2010). Potential sources for the ATP required for P2X receptor activation include the T cells themselves, which are reported to release ATP in an autocrine manner through pannexin 1 hemichannels that co-

localize with P2X7 at the immunological synapse (Woehrle et al. 2010; Schenk et al. 2008). It has been suggested that autocrine ATP signalling in T cells via P2X receptors serves to amplify weak TCR signals, gene expression and T cell effector functions (Junger 2011).

The bi-directional connection between the neuronal and immune system is established by the presence and functional relevance of molecules and ion channels present in both systems as explained in the next chapter.

3.4 Connection between the Neuronal and Immunological Systems

Although there has been evidence for years for a crosstalk between the immune and nervous systems following injury, the 'dogma' in the field of neuro-immunology has been that the healthy central nervous system (CNS) is 'immune-privileged' because it lacks classical immune molecules (Joly et al. 1991; Murphy and Sturm 1923). However, the role of MHC-I molecules in neuronal synapse development and synaptic plasticity is quite evident in brain cells of the visual system (Huh et al. 2000), and TCR/CD3⁽ molecules contribute to the spatial learning and memory function (Xu et al. 2010). Furthermore, the role of MHC-II molecules in the degradation of neurons in Alzheimer's disease (AD) is established (McGeer et al. 1989). Several inflammatory molecules like IL-1, TLR4 and TLR3 are expressed during brain inflammation (Rogers et al. 1988) and antagonists to these inflammatory mediators were shown to protect the brain. In addition, auto-antibodies generated against transmitterreceptors or voltage-gated ion channels in the brain influence the function of neuronal ligand- and voltage-gated ion channels, leading to synaptic dysfunction, and are found in Rasmussen's encephalitis, Lambert-Eaton Myasthenic Syndrome (LEMS) or NMDARassociated encephalitis, as shown in **Table 3**. In the same way as neurons are regulated by molecules predominantly expressed in immune cells, brain molecules can regulate immune function, like regulation of immune cells by glutamate as described in section 3.7.

3.5 Glutamate Receptors

3.5.1 Ionotropic and Metabotropic Glutamate Receptors

Glutamate (Glu) can interact with multiple receptor types, which are divided into two main groups: ionotropic glutamate receptors (iGluRs), which form homo- or heteromeric ion channels from several subunits with four transmembrane domains and mediate fast excitatory glutamate responses, and metabotropic glutamate receptors (mGluRs), which have seven transmembrane domains and belong to the superfamily of G protein-coupled receptors (Nakanishi 1992). On the basis of sequence homology and agonist preference, iGluRs are classified into N-methyl-D-aspartate (NMDA), α -amino-3-hydroxy-5-methylisoxasole-4-

propionate (AMPA) and kainate (KA) receptors, which are associated with permeability to particular cations (Wisden and Seeburg 1993). The first type is highly permeable to Ca²⁺, whereas AMPA and KA receptors are mostly permeable to Na⁺ ions (Nakanishi 1992). mGluRs are classified into three subgroups (I, II, III) and there are eight family members identified so far (Pin and Duvoisin 1995). Group I contains mGlu1R and mGlu5R subtypes, which are mainly coupled to PLC, and quisqualic acid is their most potent agonist. Group II consists of mGlu2R and mGlu3R, which negatively couple to adenylate cyclase and for which L-2-(carboxycyclopropyI)-glycine is a potent agonist. Group III contains mGlu4R, mGlu6R, mGlu7R, and mGlu8R, which have the same property as type-II, but they have a different agonist, namely L-2-amino-4-phosphonobutyric acid (Pin and Duvoisin 1995; Pin and Acher 2002). This classification is schematically represented in **Figure 3**.

3.5.2 Ionotroic Glutamate Receptors in the Brain

NMDARs and AMPARs are the main ionotropic GluRs involved in glutamatergic neurotransmission in the CNS. Their functions in synaptic transmission and plasticity are well established including long term potentiation/depression and excitotoxicity. NMDARs are hetero-tetramers consisting of the obligatory GluN1 subunit and two accessory subunits named GluN2A-D, GluN3 or GluN4. Activation of NMDARs requires the binding of glutamate or aspartate, the co-agonists glycine or D-serine and membrane depolarization. The GluN2 family of NMDAR subunits contains a binding site for glutamate, the endogenous agonist, whereas the GluN1 subunit binds glycine (Johnson and Ascher 1987) as illustrated in Figure 4. There is another component, D-serine, which binds to the glycine site of classical NMDARs. Although levels of glycine are 10-fold higher than D-serine, several reports indicate that endogenous D-Serine and not glycine is the dominant exogenous co-agonist for NMDAR-mediated neurotoxicity (Shleper et al. 2005; Mothet et al. 2000; Wolosker et al. 1999). When glutamate and glycine/D-serine bind and the cell is depolarized to remove the Mg²⁺-block of NMDARs, the channel opens with consequent influx of Ca²⁺ and Na⁺ into the cell, the amount of which can be altered by higher levels of agonists and by substances binding to one of the modulatory sites on the receptor (Figure 4). The opening kinetic of NMDARs depends on the subunit composition and has profound consequences for downstream signalling pathways. Thereby, NMDARs can sense different activation patterns and trigger specific intracellular signalling pathways via the induction of intracellular Ca²⁺ changes within small domains below the neuronal membrane (Paoletti et al. 2013).



Figure 3. Classification of glutamate receptors (GluRs). GluRs are divided into two major classes, according to their differential intracellular signal transduction mechanisms as well as sequence homologies. iGluRs are sub-classified on the basis of sequence homologies and agonist preference into NMDA, AMPA and KA receptors, which are all associated with ion channels permeable to particular cations. There are three distinct subtypes of mGluRs, classified by their sensitivity to exogenous agonists and intracellular second messengers employed (modified from *Hinoi, E. et al.*) (Hinoi et al. 2004; Pacheco et al. 2007; Collingridge et al. 2009).

As NMDARs are important for several neuronal functions, they are implicated in various neuronal disorders like traumatic brain injury and ischemic stroke (Shohami and Biegon 2013), Huntington's disease (Daggett and Yang 2013), AD (Malinow 2012), schizophrenia (Paoletti et al. 2013), mood disorders (Machado-Vieira et al. 2010), and encephalitis (Finke et al. 2013). The physiological functions of NMDARs are mediated by the signalling events occurring downstream of their activation. Ca²⁺-activated kinases like Ca²⁺/calmodulin dependent protein kinase II (CaMKII) and members of the protein kinase C (PKC) family mediate Early-Long-Term-Potentiation (E-LTP). The src kinase Fyn enhances NMDAR currents by phosphorylation of GluN2 subunits. More recent studies show NMDAR-induced activation of the MAPK, Erk1/2 and PI3-K/Akt pathways. NMDAR signals culminate in the activation of a cohort of transcription factors that orchestrate specific gene expression programs guiding neuronal homeostasis, cell death or plasticity. The localization and composition of the NMDARs in the neuronal membrane is fundamental for the initiation of these intracellular signalling events (Groc et al. 2006; Cognet et al. 2006; Kahlfuss et al. 2014).

Figure 4. NMDAR model illustrating important binding sites for agonists and antagonists. The

agonists glycine (Gly)/D-serine (D-Ser) and glutamate/N-methyl-Daspartate (Glu/ NMDA) are shown bound to their respective binding sites. The binding sites for Mg²⁺ (Mori et al. 1992; Mayer et al. 1984), MK801 (Huettner and Bean 1988) and memantine (Chen and Lipton 1997; Chen et al. 1992) are within the ion channel pore region and ifenprodil binds to the GluN2B subunit (Kew et



al. 1996). The different kinds of antagonists are marked as: *competitive, +uncompetitive, -non-competitive (modified from *Lipton, S.A. et al.*) (Lipton 2006).

Excessive stimulation of NMDARs leads to excessive intracellular Ca²⁺-influx, generation of free radicals such as nitric oxide and reactive oxygen species, collapse of the mitochondrial membrane potential, loss of ATP, and eventually neuronal apoptosis or necrosis depending on the intensity of the initial insult and the extent of energy recovery. This process is termed excitotoxicity and appears to be an integral component in a final common pathway to neuronal injury in neurodegenerative disorders including HIV-associated dementia (Yeh et al. 2000).

As NMDARs are involved in many neuronal disorders, modulation of their activity is important in clinical perspective. Among the NMDAR modulators clinically used for the treatment of neuronal disorders, memantine hydrochloride was approved by the U.S. Food and Drug Administration (FDA) and European Agency for the Evaluation of Medicinal Products (EMEA) in 2003 for moderate to severe kinds of AD. Another NMDAR inhibitor ketamine, which originally was developed as an anesthetic, is able to improve depressive symptoms within hours in subjects with treatment-resistant depression (Berman et al. 2000). Experimentally, ketamine mainly acts through Erk and Akt, which activate the kinase mTOR (mammalian target of rapamycin) enabling the translation of synaptic proteins (Zunszain et al. 2013).

3.6 NMDAR Inhibitors

A competitive antagonist is defined as an inhibitor which competes with the agonist binding site present in the receptor. This kind of antagonist competes with glutamate for binding to the glutamate receptor site on the GluN2 subunit of the NMDAR. Examples for competitive

antagonists are D-APV (mostly used in neurobiological studies) and selfotel (anticonvulsant with side-effects), as illustrated in Figure 4*. A non-competitive antagonist can work via two mechanisms: it can bind to the active site of the receptor or to an allosteric site of the receptor. An allosteric site is defined as a binding site distinct from the active or agonist binding site. The end result of these two non-competitive mechanisms is guite similar. They reduce the magnitude of maximum response attained by any amount of agonist, whereas competitive antagonists reduce the quantity of agonist required to achieve a maximal response. In case of NMDAR antagonists, these two kinds of non-competitive mechanisms are exemplified by ifenprodil and ketamine. Ifenprodil binds to the GluN2B subunit and aptiganel to the Mg²⁺-binding site (active site) of NMDARs. Ketamine, which acts as an analgesic and is in clinical trial for depression and mood disorders, appears to bind to the allosteric site of the channel (Orser et al. 1997), as shown in Figure 4-. Finally, an uncompetitive antagonist is defined when receptor activation by an agonist is required before it can bind to a separate allosteric binding site of the receptor. This type of antagonist follows a kinetic rule where the 'same amount of antagonist blocks higher concentration of agonist better than the lower concentration' (Lipton 2004). The prominent examples in this group are memantine (used for the treatment of AD), MK801 (used in scientific research) and, amantidine (used for treating influenza and Parkinson's disease) as shown in Figure 4+.

Cross-reaction of an antagonist with other channels is quite common within the neuronal ion channel population. Memantine and MK801 are known to cross-react with α -7-nicotinic acetylcholine (α -7-nAchR) and serotonin receptors in heterologous expression systems and rat hippocampal neurons (Aracava et al. 2005; Amador and Dani 1991; Rammes et al. 2001; Iravani et al. 1999). Furthermore, memantine is known to block dopaminergic (D2) receptors in pituitary cells (Seeman et al. 2008), and ifenprodil blocks ionotropic serotonin receptors (5-HT3) (Barann et al. 1998), presynaptic P/Q type Ca²⁺-channels (Delaney et al. 2012) and K⁺ inward rectifier channels (Kobayashi et al. 2006).

3.6.1 Pharmacological Quantification: Hill slope, Dwell time, Type of Inactivation

In pharmacological research, there are several parameters used for showing the binding kinetic of a drug to its target. These terms are described in the following:

The Hill slope (n) provides a way to quantify whether one or more ligands bind to a receptor. It describes the co-operativity of ligand binding to its receptor in this way: If n > 1, then the receptor has affinity to bind to more than one molecule (positively co-operative binding).

If n < 1, then it does not bind to more than one molecule (negatively co-operative binding), and if n = 1, then the affinity does not depend on whether there is any chance of binding any other molecule (non-cooperative binding).

The Dwell time (τ) gives information on the kinetic processes. It defines the amount of time a channel remains in the closed position and is used to describe the amount of time an ion spends in an ion channel pore at a particular binding site. This information is important when choosing a drug to be pharmacologically important. For example, memantine is preferred among other pharmacological NMDAR inhibitors like MK801 or D-APV because of its small τ . The Dwell time describes the kinetics of binding of an inhibitor to its receptor electrophysiologically, whereas the behaviour of the ion channel can be described biochemically with rate constants K_{on} and K_{off}. The biomolecular scheme of macroscopic blocking and unblocking are hypothetically shown with this equation:

 K_{on} is dependent on the memantine concentration and K_{off} in Eqn. (1) is inhibitorindependent. The macroscopic on-rate constant (K_{on}) is related to the time for onset constant (τ_{on}) by a sum of blocking rate (K_{on}) and unblocking rate (K_{off}) constants. Here, K_{off} is the reciprocal of the measured unblocking time constant (τ_{off}). Thus, the equation is:

$$K_{on} = 1/\tau_{on} - 1/\tau_{off}$$
.....(2)
 $K_{off} = 1/\tau_{off}$(3)

 K_{on} calculated from Eqn. (2) experimentally is dependent on an increasing memantine concentration, whereas K_{off} in Eqn. (3) remains relatively constant (Nelson DL 2004).

The dissociation constant (K_i) at equilibrium for memantine action can be calculated from Eqn. (4). It is found empirically that memantine is a low-affinity (apparent affinity of ~1 μ M) open-channel blocker of the NMDAR and a major component of the affinity is determined by a K_{off} at clinically relevant concentrations in the low micromolar range (Chen and Lipton 1997).

Any voltage-gated ion channel exists in three different states: activation, deactivation and inactivation. In this context, the concept of gate should be described. The pore domain, as

explained in Figure 2 with P marked in green colour of a voltage-gated ion channel as exemplified by a K^+ -channel in **Figure 2**, acts as a permeation pathway, which is opened and closed by two distinct molecular gates: activation and inactivation gates. In most voltagegated ion channels, the activation gate is closed at normal membrane potential. Membrane depolarization causes a conformational change in voltage-sensing domains, as shown in the S4 domain of Figure 2A, that is transmitted to the pore domain, as illustrated between S5 and S6 of Figure 2A, resulting in opening of the gate. Inactivation is a non-conducting state during maintained depolarization. Conventionally, there are two kinds of inactivation: N- and C-type. The different names come from the protein parts that are involved in the inactivation process. The fast inactivating N-terminal region is involved in N-terminal inactivation, in contrast to C-type, which includes the C-terminal part of the protein. Voltageactivated K⁺-channels modulate through the slower C-type of inactivation. In case of Ninactivation, the N-terminal residues (amino acids 6-46) of the channel move into the internal vestibule, as described in the S4-S5 linker of Figure 2, to occlude the intracellular mouth of the ion-conducting pore (Isacoff et al. 1991; Hoshi et al. 1991). Once the pore is occluded, it is hard to close the pore similar to a 'foot-in-the-door' mechanism (Demo and Yellen 1991) keeping it in a deactivated state. In comparison, C-type inactivation involves a slower rearrangement of the outer mouth and specific residues in the pore region (Liu et al. 1996).

Inhibitors can change the property of an ion channel by binding to the sites which are involved in the inactivation phenomenon. For example, L-type Ca²⁺-channel blockers like nifepidine and verapamil, used for cardiovascular disorders, stabilize the inactivated closed state. This effect delays the transition to the resting phase and, thereby, inhibits the depolarization-induced Ca²⁺-influx. With this property, these drugs are clinically used for the treatment of hypertension and cardiac arrhythmias by decreasing blood pressure and cardiac contractility (Abernethy and Schwartz 1999; Striessnig et al. 1998).

3.6.2 Reasons for Selecting Memantine to Block NMDAR Activity

Memantine HCI (systematic name: 3, 5-dimethyladamantan-1-amine) was first developed by Eli Lily & Co. in 1963 as an anti-diabetic agent for lowering the blood glucose level. Merz & Co. then proved its effectiveness in the treatment of AD due to its antagonism of NMDARs (Parsons et al. 1999). This was succeeded by a series of clinical trials by French, UK and USA pharmaceutical companies. The results of these trials convinced the European Union in 2002 and the US FDA in late 2003 (Lipton 2006) to approve memantine for the treatment of AD. Among the different types of NMDAR inhibitors mentioned before, memantine is chosen as a drug of choice because of its low side-effects. To be clinically acceptable, the anti-excitotoxic therapy must block the excessive activation of NMDARs while leaving NMDAR function relatively normal. Drugs that compete with Glu or Gly at the agonist-binding sites block normal function and therefore do not meet this requirement. Consequently, these drugs have failed in clinical trials because of severe side-effects such as drowsiness, hallucination and even coma (Lipton and Rosenberg 1994; Kemp and McKernan 2002; Koroshetz and Moskowitz 1996; Hickenbottom and Grotta 1998; Lutsep and Clark 1999; Rogawski 2000; Palmer 2001).

Although memantine and MK801 are uncompetitive inhibitors, memantine was chosen for clinical trials due to its small τ . The K_{off} rate of an inhibitor is a major determinant of clinical tolerability of open-channel blockers, because an excessively slow off-rate (associated with a long τ and higher K_{off}) causes the drug to accumulate in the channels, interfere with normal neurotransmission and to produce unacceptable adverse effects as in the case of MK801. These kind of drugs make patients hallucinate (e.g. phencyclidine, also known as Angel Dust) or so drowsy that they can serve as anaesthetics (as ketamine). The relatively small K_{off} and short τ of memantine prevent the drug from accumulating in the ion channels and interfering with normal synaptic transmission. Thus, memantine can provide neuroprotection without displaying adverse side effects (Lipton 2006). The neuro-protective properties of memantine have been confirmed in a large number of in vitro studies and in vivo animal models (Lipton 2006). Neurons were protected by memantine in several areas of the brain like cerebrocortical, cerebellar and retinal regions (Vorwerk et al. 1996; Lipton 1993; Chen and Lipton 1997; Chen et al. 1992; Lipton 1992; Pellegrini and Lipton 1993; Sucher et al. 1997; Osborne 1999). A series of human clinical trials have been completed or are nearly completed testing the efficacy of memantine in AD, vascular dementia, HIVassociated dementia, diabetic neuropathic pain, depression, and glaucoma. Along with mild to moderate vascular dementia (Orgogozo et al. 2002), randomized clinical trials reported that memantine was beneficial in severely demented patients, probably representing both AD and vascular dementia (Winblad and Poritis 1999).

In terms of binding to NMDARs, memantine has two binding sites. The specific site of memantine action is presumed to be near to the Mg²⁺-binding site at the selectivity filter region of the NMDAR channel (Sakurada et al. 1993). This specific binding site manifests a slow unblocking rate, moderate voltage dependence and high affinity (Danysz and Parsons 2003). The voltage-dependency is exemplified by the increase of IC₅₀ (inhibiting NMDAR responses by 50%) with depolarization (Johnson and Kotermanski 2006; Parsons et al. 2007; Rogawski and Wenk 2003). The second unspecific binding site of memantine is reported to have a much lower affinity, minimal voltage dependence and a non-competitive mechanism of blocking (Bresink et al. 1996; Antonov and Johnson 1996).

Pharmacokinetics: Memantine (trade name: Namenda) is absorbed completely from the GI tract, and peak plasma concentrations are achieved within 6-8 h after oral intake. By repeated administration a steady-state plasma level is reached within 21 days. Under therapeutic conditions, the serum levels of memantine with daily maintenance doses of 20 mg range from 0.5 to 1.0 μ M. The plasma clearance half-life is 60-100 h. The elimination of memantine is mainly performed by the kidneys as schematically showed in **Figure 26** as unchanged substance or hydroxylated metabolite. Memantine crosses the blood-brain barrier (BBB), but cerebrospinal fluid (CSF) concentration is ~20-50% lower due to its binding to albumin in the blood serum (Kornhuber and Quack 1995).

3.7 Glutamatergic Regulation of Immune Cells

3.7.1 Presence of Glutamate Receptors in Non-neuronal Organs

Dopamine, Glu, serotonin and other neurotransmitters constitute a group of physiochemically stable molecules, which may act on target cells relatively far from where they were originally released (volume transmission). In contrast, acetylcholine belongs to the group of labile compounds which, when released, achieve effective concentrations to act near to the target cells due to their rapid degradation by cholinesterases that are abundant in tissue and plasma (Danysz and Parsons 2003). Thus, substantial amounts of the former type of neurotransmitters can be detected in extracellular fluids including plasma. The Glu concentration in plasma is relatively high (50-100 µmol/L) compared with the CNS (0.5-2 µmol/L in extracellular fluids) (Morrell et al. 2008) and is tightly regulated by peripheral Glu transporters (Hinoi et al. 2004). For example, platelets express excitatory amino acid transporters (EAATs) to clear Glu from the extracellular environment (Morrell et al. 2008). Emerging evidence suggests that Glu can play a dual role in mechanisms underlying cellular homeostasis: as an excitatory neurotransmitter in the central neurocrine system and as an extracellular autocrine and paracrine signal mediator in peripheral tissues. This leads to the assumption that Glu receptors are present on different non-neuronal cells. Accumulating evidence indicates the expression of Glu receptors in the heart, spleen, testis, kidney, pancreas, and on osteoblasts, osteoclasts, and platelets (Morrell et al. 2008). Functional relation studies reveal stimulation of insulin release from pancreatic β cells by AMPA (Bertrand et al. 1993) and regulation of platelet production from megakaryocytes by NMDA (Hitchcock et al. 2003). Several of the Glu receptors were cloned and sequenced and are identical to those found in the CNS (Hinoi et al. 2004). Over the last years evidence has emerged that immune cells including macrophages, neutrophils, T cells, and DCs release glutamate and

can be regulated by glutamate found in the blood stream, peripheral organs or CNS (Pacheco et al. 2007).

3.7.2 Glutamate Receptors in Immune Cells

Immune cells were shown to express NMDARs, AMPARs (GluA3-subunit) (Ganor et al. 2003) and metabotropic Glu receptors (group I, II and III mGluRs) (Pacheco et al. 2007), which modulate their functions. Glu receptors were found to be expressed on murine thymocyte subsets (double negative (DN) immature, double positive (DP) and mature thymocytes), peripheral T cells and human peripheral blood lymphocytes (PBLs). Glu transporters are reported to be expressed on macrophages and DCs (Pacheco et al. 2007). Although less explored, mGluRs are present on B cells (Rush et al. 2004) and DCs (Rezzani et al. 2003). Functionally, effects of NMDARs and AMPARs on the migration and apoptosis of immune cells were described. NMDARs on rodent lymphocytes seem to mediate an increase of intracellular Ca²⁺ and reactive oxygen species (Boldyrev et al. 2004), and AM-PARs may play a role in the integrin-mediated adhesion to laminin and fibronectin (Ganor et al. 2003). Furthermore, inhibitory effects of CNS Glu on myelin basic protein (MBP)- and myelin oligodendrocyte glycoprotein (MOG)-specific lymphocyte activation were described in case of multiple sclerosis (MS) patients (Sarchielli et al. 2007). In case of another autoimmune disorder, Rasmussen's Encephalitis, antibodies to AMPAR GluR3 subunits are found in paediatric patients (Levite and Hermelin 1999) and auto-antibodies to NMDARs may be involved in several neuronal disorders (Kleopa 2011). In 2011, when our studies on NMDAR function in T and B cells had already commenced, Affaticati et al. showed that NMDAR GluN1 subunits accumulate in the synaptic contact region formed between OT-II TCR transgenic (tg) thymocytes and DCs presenting cognate ovalbumin (OVA)-peptide. Inhibition of NMDARs by the antagonists MK801 or memantine altered the duration of TCRinduced Ca²⁺-flux and, thereby, influenced the apoptosis of DP thymocytes (Affaticati et al. 2011). This study further supported the idea that NMDARs are central regulators of T cell function.

However, it has to be noted that a major difference between the functional studies on neuronal and immune cells is the reported sensitivity to NMDAR-specific pharmacology. In immunological studies, NMDAR antagonists were often used in 10-fold higher concentrations than in neurons raising the question to what extent NMDARs are functional and important in lymphocytes.

3.8 Aim of the Study:

In view that NMDARs were reported to be expressed in lymphocytes and to affect human T cell function, the major aims of my thesis were

- a) to proof the expression and functionality of NMDARs in lymphocytes
- b) to determine the effects of several types of NMDAR inhibitors on the membrane potential and Ca²⁺-flux of murine and human lymphocytes.

For this analysis, electrophysiological and immunological methods were applied.

4. Material and Methods

4.1 Material

4.1.1 Mouse Lines

Experiments were conducted with wild-type (wt) C57BL/6 mice, OT-II TCR transgenic (tg) (Barnden et al. 1998) and OT-I TCR tg (Hogquist et al. 1994) mice on C57BL/6 background. Mice were at the age of 6-10 weeks. All animal work was performed in compliance with the German Guidelines for the Use of Experimental Animals. Animals were housed in the Animal Facility of the Medical Faculty, Magdeburg.

4.1.2 Cell Culture-related

4.1.2.1 Media

Amount
5% or 10%
25 ml
100 U/ml
50 mM

in 500 ml RPMI-1640

20X SC contains

Chemicals	Amount
FBS	500 ml
(heat-inactivated)	
Na-Pyruvate	12.2 mM
Non-essential amino	10 mm
acids	
Penicillin/ Streptomy-	1x10 ⁴
cin	U/ml
L-Glutamine	5%

FBS was inactivated at 56°C, aliquoted and stored at -20°C. Ingredients used for supplementation of the medium were sterile-filtered prior to use. 1X SC is used for optimal culture conditions.

4.1.2.2 Cell Isolation Kits

Kit name (cell type)	Constituents (biotinylated Abs)
mouse CD4 ⁺ /CD8 ⁺	NK 1.1 (PK136), CD8α (53-6.7), CD4 (GK1.5), I-A/I-E
T cells	(2G9), CD45R/B220 (RA3-6B2), Ter-119
human T cells	CD14, CD16, CD19, CD36, CD56, CD123, Glycophorin A
mouse B cells	CD43 (Ly48), CD4 (L3T4), Ter-119

All kits were purchased from Miltenyi Biotec (Bergisch Gladbach, Germany) and contain streptavidin-coupled microbeads along with the biotinylated Abs for isolation of the cell subsets.

4.1.2.3 Cell Stimulation

Antibodies	Company
and mito-	
gens	
CD3 (2C11)	BD Biosciences
	(Heidelberg, Germany)
CD28 (CD28.2)	BD Biosciences
SuperAvidin coated	Bangs Laboratories
microsphere beads	(IN, USA)
LPS	Sigma-Aldrich
	(Steinheim, Germany)
α-IgM (Fab') ₂ fragment	Jackson Immunoresearch
	(Hamburg, Germany)

4.1.2.4 Reagents

Cell culture reagents	Company
RPMI-1640	Biochrom AG
	(Berlin, Germany)
DMEM	Gibco AG
	(Darmstadt, Germany)
AIMV	Gibco AG
EDTA	Sigma-Aldrich
Trypsin	Gibco AG

FBS (Fetal Bovine Serum)	Pan Biotech
	(Aidenbach, Germany)
Hank's Buffer	Biochrom AG
PFA	Sigma-Aldrich
Triton X-100	Sigma-Aldrich
BSA	Sigma-Aldrich
Streptomycin	Biochrom AG
Gentamycin	Roth GmbH
	(Karlsruhe, Germany)
OVA- peptide	AnaSpec
	(Fremont, USA)
2-mercaptoethanol	Gibco AG
DMSO (Dimethylsulfoxide)	Roth GmbH
Na-pyruvate	Biochrom AG
NEA (Non-essential amino	Gibco AG
acids)	
L-Glutamine	Gibco AG
Trypan Blue	Roth GmbH
PBS	Biochrom AG
Poly-D-Lysine	Sigma-Aldrich
Poly-L-Lysine	Sigma-Aldrich
Mowiol [®] 4-88	Sigma-Aldrich

4.1.3 Microscopy-related

4.1.3.1 Immunofluorescence Abs

Antibodies	Epitope	Company
GluN1	extracellular rabbit α-mouse	Alomone labs (Jerusalem, Israel) and Synaptic systems (Göttingen, Germany)
GluN2A	extracellular rabbit α-mouse	Alomone labs
GluN2B	extracellular rabbit α-mouse	Alomone labs
GFP	monoclonal mouse-α-GFP	Roche GmbH (Penzberg, Ger- many)

lgG2b	rabbit IgG2b control	Dianova (Hamburg, Germany)
lgG2a	rabbit IgG2a control	Dianova
FITC-	secondary Ab α-mouse	Invitrogen
coupled		
Cy5-coupled	secondary Ab α-rabbit	Jackson Immuno Research

4.1.3.2 Solutions for Electrophysiological Analysis

4.1.3.2.1 Extracellular Solution

Channel	Chemical	Concentration (mM)
K _v 1.3	NaCl	160
	KCI	4.5
	HEPES	5.0
	MgCl ₂	1.0
	CaCl ₂	2.0
K _{Ca} 3.1	Na-aspartate	160
	KCI	4.5
	CaCl ₂	2.0
	MgCl ₂	1.0
	HEPES	10.0

4.1.3.2.2 Intracellular Solution

Channel	Chemical	Concentration (mM)
K _v 1.3	KF	162
	EGTA	11
	HEPES	10
	CaCl ₂	1
	MgCl ₂	2
K _{Ca} 3.1	K-aspartate	145
	CaCl ₂	8.5
	MgCl ₂	2
	EGTA	10
	HEPES	10

Extracellular and intracellular solutions were adjusted to pH7.4 and pH7.2, respectively. The osmolarity of both solutions was maintained at 300-340 mOsm.

4.1.3.3 Inhibitors

Inhibitors (blocked channel)	Company
Margatoxin (K _v 1.3)	Tocris Biosciences (Bristol, UK)
Charybdotoxin (K _{Ca} 3.1)	Tocris Biosciences
TRAM-34 (K _{Ca} 3.1)	Tocris Biosciences
Ifenprodil (GluN2B of NMDARs)	Tocris Biosciences
MK801 (open channel of NMDARs)	Tocris Biosciences and Alomone labs
Memantine (open channel of NMDARs)	Tocris Biosciences
Ketamine (open channel and allosteric site of	Tocris Biosciences
NMDARs)	
D-APV (competitive for Glu-site of NMDARs)	Tocris Biosciences

All inhibitors were reconstituted in ddH_2O and stored at -20°C for 3 months.

4.1.3.4 Ca²⁺-imaging Dyes/Chemical

Dye	Company	Activity	Use
Fura-2 AM	Molecular Probes, Invitrogen (Darmstadt,	double-excitation (at 340^* and 380^+ nm) and single-	fluorescence microscopy
	Germany)	emission (510 nm)	тистозсору
Indo-1 AM	Molecular Probes	single-excitation (350 nm) and double- emission (400 ⁺ and 475 nm*)	flow cytometry
lonomycin	Molecular Probes and Calbiochem (Merck, Darmstadt, Germany)	acts as ionophore to raise the intracellular Ca ²⁺ concentration	flow cytometry and fluores- cence micros- copy

*: Ca²⁺-free; +: Ca²⁺-bound

Both Ca²⁺-imaging dyes are ratiometric, which reduces the effects of uneven dye loading, leakage of dye, photobleaching, and problems associated with measuring Ca²⁺ in cells of unequal diameter.

4.1.3.5 Chemicals

Electrophysiological	Company
Chemicals	
NaCl	Roth GmbH
KCI	Roth GmbH
HEPES	Roth GmbH
MgCl ₂	Roth GmbH
CaCl ₂	Roth GmbH
Na-aspartate	Sigma-Aldrich
KF	Sigma-Aldrich
EGTA	Roth GmbH
K-aspartate	Sigma-Aldrich

4.1.4 DNA Constructs

Construct	Vector backbone	Insert	Source
GluN1-wt	pRcCMV	GluN1	gift from Dr.
			Paoletti
			(Bordeaux)
GluN2A-SEP (5.8 kb)	pCI (4 Kb)	GluN2A (1.1 Kb)+SEP	Addgene
		(0.7 Kb)	
GluN2B-SEP (6.3 kb)	pCI (4 Kb)	GluN2B (1.6 Kb)+SEP	Addgene
		(0.7 Kb)	
GluR2-SEP (9.6 kb)	pcDNA (not known)	GluR2 (~4 Kb)+SEP	gift from Dr.
		(0.7 Kb)	Passafaro
			(Bordeaux)

SEP = Super-ecliptic protein

4.1.5 Instruments

Instruments	Company
pH Meter	Mettler-Toledo
	(Gießen, Germany)
OsmoMeter	Osmomat 320 Genotec
	(Berlin, Germany)

Vortex	Vortex Genie2
	(New York, USA)
Light microscope	Olympus CK2
	(Tokyo, Japan)
Inverted microscope	Olympus
(for Ca ²⁺ -imaging)	
Inverted Microscope	Zeiss (Oberkochen, Germany)
(for electrophysiology)	
Epifluorescence Microscope	Zeiss
Water-bath chamber	Lauda AQUALINE AL5
	(Lauda-Königshofen, Germany)
Centrifuge	Eppendorf AG 5415R, 5810R
	(Hamburg, Germany)
EPC 10 Amplifier	Warner Instruments
	(Hamden, USA)
Electrophysiological chamber and	Warner Instruments
manipulator	
LSR II Flow Cytometer	BD Biosciences
Neubauer Chamber	Roth GmbH (Karlsruhe, Germany)

4.1.6 Software

Software	Company
PatchMaster v.2.11	HEKA Electronic (Lambrecht (Pfalz), Germany)
FitMaster v2x53	HEKA Electronic
Metamorph	Molecular Devices (CA, Germany)
IgorPro5.04B	WaveMetrics Inc. (Portland, USA)
FlowJo v3.6.1	TreeStar (Ashland, USA)
Graphpad Prism 5.0	Prism (CA, USA)
ImageJ	National Institute of Health (USA)
4.2 Methods

- 4.2.1 Cell Culture-related
- 4.2.1.1 Cell Isolation
- 4.2.1.1.1 T cell Isolation

4.2.1.1.1.1 Mouse T cell Isolation

Mice were killed with CO_2 . Spleen and lymph nodes were extracted and kept in RPMI-1640 medium supplemented with 10% FCS at RT. For preparation of single cell suspensions, spleens or lymph nodes were passed through a plastic cell strainer with a diameter of 100 μ m in Petri dishes containing the medium. An aliquot of cells was diluted in 0.05% Trypan blue in PBS and counted in a Neubauer chamber to determine the cell number. Splenic cells were treated with Gey's solution to lyse erythrocytes. This solution was pre-warmed at 37°C in the water bath before use.

CD4⁺ or CD8⁺ T cells were isolated from pooled lymph nodes of wt, OT-II or OT-I TCR tg mice by negative selection using a cocktail of biotinylated Abs: NK1.1 (PK136), CD8α (53-6.7), CD4 (GK1.5), I-A/I-E (2G9), CD45R/B220 (RA3-6B2), Ter-119 (all from BD Bioscience) and streptavidin magnetic beads (Miltenyi Biotec) according to the manufacturer's protocol. Purity of CD4⁺ or CD8⁺ T cells after MACS isolation was routinely above 90%.

CD4⁺ T cells were activated with CD3+CD28 antibodies (3+5 μ g/ml) in supplemented RPMI-1640 medium for 48 h for patch clamp recording.

4.2.1.1.1.2 T cell Isolation from Human Peripheral Blood

Fresh CD3⁺ T cells were isolated from the collected blood samples of healthy donors and dementia patients using the Pan and CD4⁺ T cell isolation kit from Miltenyi Biotec and the Ficoll-Hypaque gradient procedure according to manufacturer's protocol. The dementia patients were prescribed to have a daily 10 mg memantine dose in the first week followed by 20 mg for the next 11 weeks. The isolation kit contained biotinalyted CD14, CD16, CD19, CD36, CD556, CD123, and Glycophorin A and anti-biotin microbeads for the conjugation to anti-biotin antibodies. The cells were maintained in AIMV medium. Purity of iolated T cells was 90%. All experiments with human cells were conducted in collaboration with Prof. Dr. Ursula Bommhardt (Institute for Molecular and Clinical Immunology) and Dr. Stefan Busse (Department of Psychiatry, Magdeburg) and were approved by the local Ethics Committee.

4.2.1.1.2 B cell Isolation

Splenic or lymph node B cells were isolated with the B-cell isolation kit (Miltenyi Biotec) composed of a cocktail of biotinylated Abs: CD43 (Ly48), CD4 (L3T4) Ter119 and antibiotin microbeads. The procedure was followed according to the manufacturer's protocol and purity of B cells was 90-95%.

B cells were cultured in complete RPMI-1640 medium supplemented with 10% FCS, 50 μ M β -mercaptoethanol, 1% penicillin/streptomycin and were activated with goat anti-mouse α -IgM (Fab')₂ or LPS (10 μ g/ml each) for 48-72 h and subsequently used for the experiments.

4.2.1.1.3 DC Isolation

4.2.1.1.3.1 Generation of Bone-marrow Derived Dendritic Cells (BMDCs)

Bone marrow (BM) was collected from femur and tibia of wt mice and cells were suspended in RPMI-1640 medium reconstituted with 1% non-essential amino acids, 5% FCS, 1% Lglutamine, 0.1% gentamycine, 0.1% 2-mercaptoethanol, IL-4 (48 ng/ml) and GM-CSF (10 ng/ml) from a hybridoma supernatant (a gift of Prof. Dr. Matthias Gunzer). A total of 3x10⁶ BM cells/5 ml BMDC medium were cultured for 7 days. At day 3, 2 ml medium was replaced by fresh BMDC medium, at day 6, total medium was replaced and cells were stimulated with LPS in a concentration of 20 ng/ml for 24 h. DCs used at day 9 or 10 were restimulated with LPS 24 h before experimental onset. Maturation of BMDCs was verified at day 7 by staining cells with Abs against MHC-II (NIMR-4), CD11c (HL3), CD80 (2D10) and CD86 (all from BD Bioscience).

4.2.1.1.3.2 Activation of DCs

Mature BMDCs (MHC-II⁺ CD11c⁺ CD80⁺ CD86⁺) were pulsed with OVA-peptide (aa 323-339, 10 µg/ml or aa 257-264, 5 µg/ml (SIINFEKL), AnaSpec) for 2 h and cultured with OT-II CD4⁺ or OT-I CD8⁺ T cells, respectively. CD4⁺ T cells ($0.5x10^{5}$ - $1x10^{5}$) were stimulated with CD3 Abs (3 or 10 µg/ml) or CD3 pus CD28 Abs (3 and 5 µg/ml, respectively).

For single-cell Ca²⁺-flux measurement, a total of 1x10⁶ OT-II CD4⁺ or OT-I CD8⁺ T cells were left untreated or were activated with pOVA-loaded BMDCs in a DC-T cell ratio of 1:10. NMDAR-inhibitors were added during recording. Matured BMDCs were pulsed with the OVA-peptide for overnight.

4.2.1.2 Maintenance of Cell Lines

EL-4 (mouse T cell), JE6.1 (human T cell) and Raji (human B cell) lymphoma cells were grown in RPMI-1640 medium containing 10% FCS supplemented with nutrients at 37°C and 5% CO₂. Cells were diluted 1:10 after 3 days to maintain an optimal cell density $(1\times10^5-1\times10^6$ cells/ ml). For freezing, the cells were diluted in DMSO at a ratio of 1:10.

Adherent HEK-293T cells were grown in Dulbecco's MEM (DMEM) containing 10% FCS, 100 units/ml penicillin and 100 μ g/ml streptomycin at 37°C and 5% CO₂. Cells were diluted 1:10 in 3 day cycles to avoid confluence of the cell layer. For splitting, adherent cells were washed once with PBS and subsequently treated with 5 ml Trypsin/EDTA solution (0.25% trypsin/0.02% EDTA in PBS without Ca²⁺, Mg²⁺) for 1 min at RT for cell detachment. Cells were suspended and diluted 1:10 in fresh medium for culture.

4.2.1.3 Determination of Cell Number

Determination of the cell number was done with the Neubauer chamber, which is subdivided into four quadrants (each having a surface area of 1 mm²). Each quadrant is subdivided into 16 smaller quadrants. When cells are added to the chamber, each big quadrant receives a volume of 0.1 μ l (0.1 mm³). The cell suspension was diluted 1:50 in Trypan blue solution and counted with a light microscope of 20x magnification. Bright live cells in the quadrants were counted excluding the blue coloured ones representing dead cells. Calculation of cell count = counts of 4 bigger quadrants)/4 x dilution factor x 10,000 = cell count/ml.

4.2.1.4 Transfection Protocol

4.2.1.4.1 Fugene Method

600 μl of PBS was mixed with 2 μg of DNA and 50 μl Fugene (Roche, Berlin, Germany). This mixture was kept for 15 min at RT and then added to HEK-293T cells according to the manufacturer's protocol. Transfected cells were immuno-stained after 24-48 h. This method was used for transfecting the cell line to test NMDAR Ab specificity.

4.2.2 Microscopy-related

4.2.2.1 Immunofluoroscence Method

For live extracellular staining, HEK-293T cells were incubated with primary Abs (GluN1-wt, GluN2A-SEP and GluN2B-SEP) in DMEM medium for 30 min at 37°C. Then, the cells were fixed with 4% PFA and blocked with a solution containing 10% FCS in PBS, 1% Triton X-100. Then, the cells were washed three times with PBS. Afterwards, secondary Abs in

PBS/1% BSA were added to the cells and kept for 1 h in the dark. Finally, the cells were washed three times and mounted in mowiol.

For staining of fixed cells, the cells were initially fixed with 4% PFA and then blocked with 10% FCS/PBS/1% Triton X-100. Then, the cells were incubated with primary Abs in PBS for 1 h. This was followed by washing the cells three times with PBS/1% BSA. Finally, the cells were incubated with secondary Abs for 1 h in the dark followed by three times washing in PBS. At the end, the cells were mounted in mowiol. BSA (bovine serum albumin) inhibits the non-specific binding of Abs. Thus, its use in immunofluorescence stainings is preferred.

4.2.2.2 Electrophysiological Methods (Patch clamp)

All experiments were carried out in the whole-cell configuration of the patch-clamp technique using an EPC10 amplifier and PatchMaster v.2.11 (HEKA Electronic) at RT (20-24°C).

4.2.2.2.1 Voltage Clamp

Patch pipettes from borosilicate glass used for recordings had a resistance between 3-5 M Ω . K_v1.3 and K_{Ca}3.1 currents were recorded with external and internal solutions mentioned in **section 4.1.3.2**. Osmolarity was set to 300-340 mOsM by the Osmomat instrument. K_v1.3 currents were measured with depolarizing voltage steps up to +60 mV from a holding potential of -80 mV every 30 s in steps of 20 mV. K_{Ca}3.1 currents were elicited by a 200 ms voltage ramp from -120 to +40 mV from a holding potential of -80 mV every 15 s. Sampling rate was 50 kHz in case of K_v1.3 and 20 kHz in case of K_{Ca}3.1. The antagonists ifenprodil, MK801, memantine, ketamine or D-APV (Tocris) were added during the recording with a constant inhibitor concentration. As a positive control for the measured currents, margatoxin and TRAM-34 (Tocris) were used.

4.2.2.2.2 Current Clamp

For membrane potential experiments, activated mouse T or B cells and JE6.1 or Raji cells were recorded in the current clamp mode with 0 pA holding current immediately after establishment of the whole-cell configuration. Ifenprodil, memantine or D-APV were added during the recording with a constant inhibitor concentration to determine the membrane depolarization in the presence of the inhibitors.

4.2.2.3 Ca²⁺-flux Measurement

4.2.2.3.1 Whole-cell Ca²⁺-flux

For T cells, lymph node cells from wt mice were stained with 4 μ M Indo-1 AM (Invitrogen, Molecular Probes) for 45 min at 37°C. After being washed with PBS, the cells were stained for CD8 and B220 or CD4 and B220 surface expression for 15 min, washed and resuspended in Hank's buffer (Biochrom AG) supplemented with 1 mM CaCl₂. CD3-biotin Abs (145.2C11, 10 μ g/ml) plus streptavidin (25 μ g/ml, Dianova) were added to induce Ca²⁺-flux. The NMDAR antagonist ifenprodil (10 or 30 μ M) was added for 5 min before CD3 Ab and streptavidin treatment. Additionally, external NMDA (100 μ M) was added to induce Ca²⁺-flux.

For B cells, splenocytes were stained with 4 μ M Indo-1 AM for 45 min at 37°C. Cells were washed, stained for CD8 and CD4 surface expression and suspended in Hank's buffer supplemented with 1 mM CaCl₂. NMDAR antagonist ifenprodil (10 or 30 μ M) or memantine (30 or 50 μ M) was added for 5 min before B cells were activated with α -IgM (10 μ g/ml) to induce Ca²⁺-flux.

Towards the end of each measurement, ionomycin (2 μ M, Calbiochem) was added as a positive control for cell reactivity. Ca²⁺-flux was measured on a LSRII flow cytometer (BD Biosciences). Data files were transferred to FlowJo V3.6.1, mean Ca²⁺-flux was determined for unlabelled CD4⁺ or CD8⁺ T cells and unlabelled B cells and data were further processed with IgorPro5.04B software. For each graph, Δ Ca²⁺-flux was defined as the difference between the maximum and minimum value of Ca²⁺-intensity.

4.2.2.3.2 Single-cell Ca²⁺-flux

Freshly-isolated CD4⁺ or CD8⁺ T cells were loaded with 2-4 μ M Fura-2 AM for 45 min at 37°C. These Fura-loaded cells were measured with the MetaMorph Program (Molecular Devices) under a fluorescence microscope (Olympus) in 40x magnification along with the prepared DCs. Mean Ca²⁺-flux was determined by plotting the 340 and 380 nm values in Excel and each trace was assessed in IgorPro 5.04B software. The T cells were stimulated with either OVA-presenting DCs or CD3/CD28 coated microbeads.

4.2.2.3.2.1 Bead Stimulation Protocol

Beads (mean diameter: 10.14 μ m, 33 μ l for 1x10⁶ cells) were incubated with CD3 and CD28 Abs (both 10 μ g/ml) for 30 min at 37°C to activate the cells. Then, they are washed with PBS and added to the used cell culture medium.

4.2.3 Statistical Analysis

Data are given as mean values \pm standard deviation (SD), if not stated otherwise. Student's *t* test and other statistical measurements were performed in GraphPad Prism 5.04B. Statistical significance was set as *p<0.05, **p<0.01 and ***p<0.001.

5. <u>Results</u>

5.1 NMDAR Expression in Murine Lymphocytes is Puzzling

In previous work GluR expression was detected in murine thymocytes and human peripheral lymphocytes by intracellular staining and flow cytometry, and localization of GluN1 subunits in the thymocyte-DC contact zone was shown by confocal microscopy (Affaticati et al. 2011; Ganor et al. 2003; Lombardi et al. 2001). Inspired by this work, experiments within our collaborative research project apparently confirmed the results for NMDAR subunit expression, namely expression of GluN1, GluN2A and GluN2B subunits in thymocytes and peripheral T cells (Kahlfuss et al., 2014). Upon co-culture of OT-II CD4⁺ T cells with pOVApresenting DCs, GluN1 and GluN2B subunits were detected in the immunological synapse, as shown by confocal microscopy. However, in Western blot analyses, GluN1 protein in thymocytes and CD4⁺ T cells appeared at a lower molecular weight than GluN1 protein in brain lysate. To proof whether the detected protein is GluN1, we performed analyses on thymocytes obtained from newborn GluN1 knock out (ko) mice. PCR and RT-PCR analyses showed the deletion of GluN1 at the DNA and mRNA level in thymocytes, and GluN1 protein was absent in brain samples, but not in thymocytes. These observations strongly suggest that GluN1 protein is not detectable in thymocytes. Intracellular staining with two different GluN1 Abs and flow cytometry as well as immunohistochemistry also showed identical staining for wt and GluN1 ko thymocytes, although the used Abs for GluN1, GluN2A and GluN2B showed specificity for the subunits in transfected HEK-293T cells (Figure 5). Hence, on the protein level, there is no evidence for expression of the obligatory GluN1 subunit of NMDARs in murine thymocytes and T cells (Kahlfuss et al. 2014).



Figure 5. NMDAR subunit transfection for testing the specificity of Abs used to detect NMDAR subunit expression. Fluorescent images of HEK-293T cells transfected with the NMDAR subunits GluN1-wt, GluN2A-SEP and GluN2B-SEP were analysed by an epifluroscent microscope taken in

63x magnification and processed with ImageJ. Scale bar = 4 μ m. Transfected cells were analysed for GFP(SEP)-expression to detect the individual subunits and were stained with (C) mouse-anti-GluN1 (Synaptic System), (D) rabbit-anti-GluN2A (Alomone labs), (E) rabbit-anti-GluN2B (Alomone labs), (A) mouse-IgG2b, (B) rabbit-IgG2a, and secondary PE-labelled Abs. (A) and (B) were used as controls for mouse-anti-GluN1 and rabbit-anti-GluN2A/GluN2B, respectively. The immunohistochemical stainings show strong overlap with GFP-expression indicating that the Abs are specific for the individual NMDAR subunits. Differential interference contrast (DIC) images show the position and shape of an individual cell. Arrows point towards a representative transfected HEK-293T cell.

Although expression of NMDARs at the protein levels was elusive, we detected Ca²⁺-flux in lymph node (LN) CD4⁺ and CD8⁺ T cells and LN and splenic B cells upon addition of the NMDAR agonist NMDA (100 μ M). NMDA-induced Ca²⁺-flux in T cells was reduced by 10 μ M ifenprodil to ~80% and it was nearly abrogated with 30 μ M ifenprodil (**Figure 6A**). In case of B cells, the reduction in Ca²⁺-flux was even more prominent, as Ca²⁺-flux was reduced to ~40% in LN and to ~20% in splenic B cells by 10 μ M ifenprodil (**Figure 6B**).



Figure 6. NMDAR antagonists attenuate NMDA-induced Ca²⁺-flux. Indo-1 AM-loaded (A) LN CD4⁺ and CD8⁺ T cells and (B) LN and splenic B cells were activated with NMDAR agonist NMDA (100 μ M) in the absence or presence of ifenprodil (Ifen.) and Ca²⁺-flux was determined with flow cy-tometry. Ionomycin (IO) was added towards the end of each measurement to control cell reactivity. Left histograms in (A, B) show a representative experiment for LN CD4⁺ T cells and B cells; right graphs show the relative Δ Ca²⁺-flux for CD4⁺ and CD8⁺ T cells and B cells calculated from 3-4 ex-

periments. ΔCa^{2+} -flux from cells activated without ifenprodil (none) was set as 1. Data represent mean values <u>+</u> standard error of mean (SEM) from 3-4 independent experiments.

Although NMDA-induced Ca²⁺-flux was detected in T and B cells, the presence of NMDARs should be further examined by performing Ca²⁺-flux experiments in thymocytes from GluN1 ko mice. It should also be mentioned that external Glu (100 μ M) did not induce any Ca²⁺-flux. Thus, we hypothesize that NMDA either unspecifically binds to some other unknown ion channel(s) on lymphocytes or that NMDARs are expressed only transiently and in very small numbers and therefore are not detectable under the used experimental conditions.

Along with whole-cell Ca²⁺-flux in mature T cells, single-cell Ca²⁺-influx was performed, as *Affaticati et. al., 2011* showed Ca²⁺-influx at the immune cell synapse formed between OT-II TCR tg CD4⁺/CD8⁺ thymocytes and OVA-presenting DCs. We used the DC system and CD3/CD28 Ab-coated microbeads to activate T cells. OT-I TCR tg CD4⁺ and OT-II TCR tg CD8⁺ T cells were loaded with 4 μ M Fura-2 AM to monitor TCR-induced Ca²⁺-flux changes using the MetaMorph Program and an inverted microscope. Matured DCs were loaded with cognate OVA-peptide for 24 h before use. A transient Ca²⁺-flux was detected in OT-II CD4⁺ and OT-I CD8⁺ T cells after CD3/CD28 bead-stimulation as shown in **Figure 7A and C**. For DC/antigen-stimulated OT-II CD4⁺ T cells, transient as well as sustained Ca²⁺-fluxes were observed (**Figure 7B**). However, there was no reliable detection of a reduction of single-cell Ca²⁺-flux in the T-DC or T-bead contact zone after addition of ifenprodil (data not shown). This was unexpected and puzzling and could be due to the more homogenous cell population using TCR tg T cells instead of heterogeneous whole cell populations used for the measurement of Ca²⁺-flux at all or due to technical limitations.



Figure 7. Single-cell Ca²⁺-flux of TCR tg T cells in the presence of antigen-presenting DCs or CD3/CD28-coated beads. Single cell Ca²⁺-flux was recorded for 500 s in OT-II CD4⁺ T cells

stimulated with (A) CD3/28-coated beads or (B) OVA-presenting DCs and in (C) OT-I CD8⁺ cells after stimulation with CD3/28-coated beads. The data represent averaged Ca²⁺-fluxes from 14-22 sin-

gle cells of n = 3-4 experiments out of 8-9 preparations from OT-I/II TCR tg mice. Insets show the representative flux for different stimulation conditions mentioned.

5.2 NMDAR Antagonists Attenuate TCR/BCR- induced Ca²⁺-flux

5.2.1 TCR-induced Ca²⁺-flux

In order to understand how NMDAR antagonists influence T-cell activation, their effect on proximal T-cell signalling was analyzed. I concentrated on the analysis of Ca²⁺-flux, as this is pivotal for the activation of many Ca²⁺-regulated proteins which guide T-cell activation. LN CD4⁺ and CD8⁺ T cells, loaded with Indo-1 AM to monitor intracellular Ca²⁺-changes by flow cytometry, responded to TCR ligation by cross-linked CD3 Abs with a rapid mobilization of Ca²⁺. The Ca²⁺-flux was significantly reduced by 10 μ M and almost entirely blocked by 30 μ M ifenprodil as shown in **Figure 8**.



Figure 8. NMDAR antagonists attenuate TCR-induced Ca²⁺-flux. (A) Indo-1 AM-loaded CD4⁺ and CD8⁺ T cells were activated with CD3 Abs (10 µg/ml) in absence or in presence of 10 and 30 µM ifenprodil (Ifen.) and Ca²⁺-flux was determined by flow cytometry. IO was added towards the end of each measurement to control cell reactivity. The histogram in (A) shows a representative Ca²⁺-flux measurement for CD4⁺ T cells and the graphs in (B) shows the statistics for CD4⁺ and CD8⁺ T cells. Data in the the graphs provide the relative Δ Ca²⁺-flux as mean <u>+</u> SEM calculated from 3 experiments each. The obtained Δ Ca²⁺-flux of cells activated in the absence of Ifen. (none) was set as 1.

5.2.2 BCR-induced Ca²⁺-flux

As TCR-induced Ca²⁺-flux was modulated by ifenprodil, it was intriguing to study B cells as several autoimmune disorders, as described in **Table 3**, involve B cells producing autoantibodies, for instance against NMDARs. Ligation of the BCR with IgM Abs (α -IgM) and LPS were used for polyclonal stimulation of B cells. Indo-1 AM-labelled splenic B cells showed a concentration-dependent inhibition of BCR-induced Ca²⁺-flux when they were treated with ifenprodil or memantine. **Figure 9A** shows a reduction of Ca²⁺-flux to ~80% of untreated cells in the presence of 10 μ M and to ~30% by 30 μ M ifenprodil. **Figure 9B** depicts the reduction of Ca²⁺-flux to ~80% in the presence of 30 μ M and to ~40% by 50 μ M memantine. Thus, NMDAR antagonists also impair BCR-induced B-cell activation.



Figure 9. Reduced Ca²⁺-flux in BCR-activated B cells in the presence of NMDAR antagonists. Indo-1 AM-labelled B cells were stimulated with α -IgM (10 µg/ml) in the presence or absence of (A) ifenprodil (Ifen.) and (B) memantine (Mem.) and Ca²⁺-flux was determined with flow cytometry. (A) shows the reduction of BCR-induced Ca²⁺-flux by 10 and 30 µM Ifen. and (B) by 30 and 50 µM Mem. The Δ Ca²⁺-flux from cells activated without Ifen. or Mem. (none) was set as 1. Corresponding graphs represent the Δ Ca²⁺-flux from three experiments each.

5.3 NMDAR Antagonists Modulate the Membrane Potential and Block $K_\nu 1.3$ and $K_{Ca} 3.1$ Channels of Lymphocytes

5.3.1 Mouse Primary T Cells and EL-4 Lymphoma Cells

In view of the strong effects of NMDAR pharmacology on Ca²⁺-flux in lymphocytes, but the elusive expression of NMDARs at protein level, we hypothesized that the inhibitor's targets could involve K_v1.3 and K_{Ca}3.1 potassium channels (Partiseti et al. 1992; Partiseti et al. 1993; Lewis and Cahalan 1995). These channels regulate T-cell activation by controlling the membrane potential and, hence, the Ca²⁺-flux into T cells (Desai et al. 2000; Lam and Wulff 2011; Conforti 2012). Indeed, current-clamp recordings of CD4⁺ T cells activated with CD3+CD28 Abs showed that ifenprodil (30 μ M) and memantine (50 μ M) depolarized the membrane potential from ~-50 mV to ~-15 mV and ~-20 mV, respectively, as shown in **Figure 10**.

Figure 10. NMDAR antagonists change the membrane potential. The membrane potential of murine $CD4^+$ T cells, activated with CD3+CD28 Abs (3+5 µg/ml) for 24-48 h, was determined with current clamp recording. Addition of ifenprodil or memantine in concentrations as indicated lead to a membrane depolarization. KCI treatment served as a positive control for cell integrity and membrane



depolarization. All data were calculated from 5-6 cells of three experiments in each case and are represented as mean <u>+</u> SEM.

Since the membrane potential was strongly affected by NMDAR antagonists, I next addressed whether this involves K_v 1.3 and K_{Ca} 3.1channels, the major and most abundant K⁺channels in lymphocytes. K_v 1.3 and K_{Ca} 3.1 channels currents were recorded by voltage clamp method and dose-response curves were constructed from the transient maximal current amplitudes. The obtained Hill slopes and 50% inhibitory concentrations (IC₅₀) for K_v 1.3 channels on T cells in the presence of ifenprodil and memantine were ~1.5 and ~1.9 and ~35 μ M and ~45 μ M, respectively, as shown in **Figure 11 A, B**. For K_{Ca} 3.1 channels, Hill slope and IC₅₀ values were ~1.2 and ~15 μ M for ifenprodil and ~1.6 and ~30 μ M for memantine, respectively (**Figure 11 C, D**).



Figure 11. NMDAR antagonists inhibit $K_v 1.3$ and $K_{Ca}3.1$ channel activity in activated T cells. $K_v 1.3$ and $K_{Ca}3.1$ channel-mediated currents were recorded by voltage-clamp method from T cells activated with CD3+CD28 Abs (3+5 µg/ml) for 24-48 h. Voltage clamp protocol for $K_v 1.3$ recording and current traces are shown in (A). (B) shows the curves formed by plotting the transient amplitude of the recorded currents versus the indicated inhibitor concentrations. (C) Ramp protocol for measuring $K_{Ca}3.1$ recorded current and example traces are shown and the dose-response inhibition curves formed by the recorded currents is shown in (D). Each data point in the dose-response inhibition curves represents mean calculated from 5-7 cells and vertical lines show mean ± SEM.

A competitive antagonist can determine the functional expression of a channel as it is competing with the agonist site of the channel. In case of NMDARs, the competitive antagonist frequently used in neuronal systems is D-APV, which is competing with the Glu-binding site. Therefore, the effect of D-APV on K_v1.3 channels in activated CD4⁺ T cells was determined. As shown in **Figure 12A**, D-APV blocked K_v1.3 channels, but only at ~20-fold higher concentrations (1 mM) than needed by the other tested inhibitors. In addition to activated primary murine CD4⁺ T cells, the inhibitory effects of ifenprodil, memantine, MK801, ketamine, and D-APV on K_v1.3 channels were analysed in murine EL-4 lymphoma T cells. All NMDAR inhibitors strongly blocked K_v1.3 channel currents, but in case of D-APV, ~15-fold higher concentrations were needed to reach an inhibition similar to that obtained with ifenprodil (**Figure 12B**). Thus, the employed concentrations of NMDAR antagonists, which were similar to those used in previous publication by others (Affaticati et al. 2011), non-specifically inhibit two K⁺-channels, which reportedly modulate many Ca²⁺-mediated processes in T cells (Lam and Wulff 2011).



Figure 12. NMDAR antagonists decrease K_v 1.3 channel activity in activated CD4⁺ T cells and EL-4 lymphoma cells. K_v 1.3 channel currents were determined in (A) activated murine CD4⁺ T cells and (B) EL-4 lymphoma cells in the absence and presence of the indicated NMDAR antagonists. The currents were recorded by voltage-clamp with the protocol given in Figure 11A. The data in the bar graphs represent the relative inhibition of recorded K_v 1.3 transient currents from 5-6 cells and all data represent mean ± SEM determined by unpaired Student's *t* test.

5.3.2 Murine B Cells

K⁺-channels are also expressed on B cells and their inhibition was found to differentially influence B-cell activation and proliferation (Partiseti et al. 1992; Partiseti et al. 1993; Lewis and Cahalan 1995). Since NMDAR inhibitors downregulate BCR-induced Ca²⁺-flux in B cells, their effects on the membrane potential and K_v1.3 and K_{Ca}3.1 channels were alalyzed. Ifenprodil (20 μ M) and memantine (30 μ M) reduced the membrane potential of α-IgM- or LPS/TLR4-activated B cells from ~-40 mV to ~-20 mV and ~-10 mV, respectively as shown in **Figure 13**. Next, K_v1.3 channel-mediated currents from activated B cells were recorded and the dose-response curves in the presence of the inhibitors were calculated from maximal transient current amplitudes. Ifenprodil and memantine markedly reduced the K_v1.3 currents of B cells, irrespective of whether they were stimulated with α -IgM or LPS (**Figure 14 A,B**). IC₅₀ and Hill slope values for α -IgM-activated B cells were ~20 μ M and ~1.3 for ifenprodil and ~40 μ M and ~1.8 for memantine. For LPS-activated B cells, IC₅₀ and Hill slope values were ~18 μ M and ~1.4 for ifenprodil and ~45 μ M and ~1.2 for memantine. K_{Ca}3.1 channel-mediated currents were also recorded for B cells activated by BCR ligation (**Figure 14B**), whereas K_{Ca}3.1 currents were not detected in LPS-stimulated B cells. IC₅₀ values for ifenprodil and memantine were ~30 μ M and ~50 μ M and Hill slopes were ~1.4 and ~1.6. Notably, the competitive NMDAR antagonist D-APV, which blocks neuronal NMDARs at the 1 μ M range, had no effect on K_v1.3 and K_{Ca}3.1 channels, even at 300 μ M (**Figure 14C**). Thus, K_v1.3 and K_{Ca}3.1 channels, whose specific blockade abolishes B-cell activation (Amigorena et al. 1990; Wulff et al. 2004), are at least partially inhibited by the non-competitive NMDAR antagonists ifenprodil and memantine.



Figure 13. NMDAR antagonists reduce the B cell membrane potential. B cells were activated with α -IgM or LPS (10 µg/ml each) for 24-48 h and analyzed by current-clamp recordings for changes membrane potential upon addition of ifenprodil and memantine in concentrations as indicated. KCl treatment served as a positive control for checking the cell integrity. Each data point contains data from 5-6 cells of four experiments.



Figure 14. NMDAR antagonists lower K⁺-channel activity of B cells. Splenic B cells, activated with α -IgM or LPS, were recorded by voltage-clamp to measure K_v1.3 and K_{Ca}3.1 currents. Dose-inhibition curves of K_v1.3 in (A) and K_{Ca}3.1 in (B) were plotted in the presence of Ifen. or Mem. from the recorded maximal transient currents. Insets show one particular trace of control and inhibited current along with the protocols used for measuring K_v1.3- and K_{Ca}3.1-mediated currents. Data in the graphs in (C) represent the relative inhibition of K_v1.3- and K_{Ca}3.1-mediated currents in the presence of the competitive NMDAR antagonist D-APV in a concentration ~10 times higher than used for the other inhibitors. All data were calculated from 5-6 cells of four experiments.

5.3.3 Human Jurkat T- and Raji B-Lymphoma Cells

As ~2-fold increased plasma Glu levels act as a prognostic marker in certain cancers (e.g. breast cancer and colorectal carcinoma) (Ollenschlager et al. 1989) and the expression of GluN1 subunits seems to correlate with the prognosis of other types of cancers like oral squamous cell carcinoma and retinoblastoma (Choi et al. 2004; Stepulak et al. 2009), it was also interesting to analyze Jurkat and Raji human lymphoma cells. The question was whether lymphoma cells would express NMDARs induced by plasma Glu or whether NMDAR inhibitors would modulate K⁺-channels as found for primary murine T and B cells.

5.3.3.1 Jurkat T-Lymphoma Cells

At first, the membrane potential was determined by current clamp analysis. The membrane potential of Jurkat cells was reduced from \sim -50 mV to \sim -25 mV and \sim -20 mV in the presence of ifenprodil (30 μ M) and memantine (50 μ M), respectively, as shown in **Figure 15**.

Figure 15. NMDAR antagonists change the membrane potential of Jurkat cells. Human Jurkat T-lymphoma cells were analysed by current clamp for determining the membrane potential. In the presence of Ifen. or Mem., in concentrations as indicated, the membrane potential was reduced. KCl treatment served as a positive control for cell integrity. All data were calculated from 5-6 cells of three experiments and are represented as mean \pm SEM.



Next, the recorded currents (I) were plotted versus the used voltage (V) potentials. **Figure 16A** displays the I-V relationship measured at potentials ranging from -80 mV to 60 mV (in 20 mV steps) in control and in presence of ifenprodil (30 μ M). The activation of K_v1.3 channel differs distinctly in presence of ifenprodil. **Figure 16B** shows the mean current densities (in pA/pF) calculated as maximal current (I_{max}) divided by the cell capacitance (C_{slow}). With this calculation, standardization of the current to the cell size is achieved. Under control condition, K_v1.3 value was ~150 pA/pF and it was decreased to ~70 pA/pF in the presence of ifenprodil (30 μ M).





Finally, the major K⁺-channels (K_v1.3 and K_{Ca}3.1) taking part in the depolarizing events of immune cells were shown to inhibit in Jurkat cells in the presence of the tested NMDAR inhibitors ifenprodil and memantine (**Figure 17**). The K_v1.3 and K_{Ca}3.1 measuring voltageclamp protocols were mentioned in **Figure 17A** and **Figure 17C** respectively. Furthermore, the current traces measured from this population were depicted here on the lower part of the **Figure 17A** and 17C. As a result, the Hill slope and IC₅₀ values for K_v1.3 in presence of ifenprodil and memantine were ~2, ~1.5 and ~30 μ M, ~45 μ M. For K_{Ca}3.1, channel Hill slope and IC₅₀ values were ~1.2 and ~20 μ M for ifenprodil and ~1.4 and ~30 μ M for memantine.

In addition to these inhibitors mentioned in **Figure 17**, MK801, ketamine and D-APV were also tested. The constructed curves gave Hill slopes and IC₅₀ values for K_v1.3 current are respectively ~1.5, ~1.3, ~1.7 and ~70 μ M, ~200 μ M, ~550 μ M for MK801, ketamine and D-APV as mentioned in **Figure 18**. D-APV, a competitive antagonist in case of Jurkat cells needed ~10-fold more concentration to be inhibited in comparison to the other non-competitive inhibitors like ifenprodil and memantine as like murine cells shown before.



Figure 17.

Electrophysiological analysis in Jurkat lymphoma T cells in the presence of lfen. and Mem. The Voltage clamp protocol for K_v1.3 channel recording and current traces are shown in (A) and the graphs formed by plotting the transient amplitude of the recorded currents versus the indicated inhibitor concentrations are shown in (B). The protocol for K_{Ca}3.1ramp recorded currents and example traces are shown in (C) and the dose-response inhibition curves in (D). Each data point in the dose-response curves represents the mean ±

SEM calculated from 5-7 cells.

Figure 18. K_v 1.3 current recorded from Jurkat cell lines in presence of mentioned inhibitors. K_v 1.3 currents from Jurkat cells in the presence of MK801, ketamine and D-APV were recorded according to the voltage-clamp protocol shown in Figure 17A and plotted with the measured maximal amplitude values versus the inhibitor concentrations. Data represent mean \pm SEM and each vertical line represents 5-7 cells from 4 experiments.



A very interesting aspect in terms of channel physiology was observed with the two inhibitors ifenprodil and memantine: K_v 1.3-mediated tonic currents were also modulated, along with the transient currents (**Figure 19**). The inactivation constant (τ 1) for the control population was ~20 ms measured with the 1-exponential model (HEKA FitMaster Program). This was increased to ~60 ms and to ~80 ms in the presence of ifenprodil and memantine, respectively. Thus, these inhibitors increase the inactivation kinetics similar to the other traditional K⁺-channel blockers like Tetra Ethyl Ammonium (TEA) or 4-Amino Pyridine (4-AP) (Leung 2012). Thus, it will be interesting to determine whether these inhibitors act at the same amino acid modulating sites as the other K⁺-channel blockers. The K_v1.3 channel behavior of Jurkat cells in the presence of different NMDAR inhibitors is summarized in **Table 1**.



Figure 19. Inactivation current differences in K_v1.3-mediated currents in control and inhibitor population. Steady-state current were differed in case of Ifen. and Mem. as shown in Figure (A) that the τ 1 was increased in presence of Ifen. (10 µM) and Mem. (50 µM). For the inactivation protocol, the currents were recorded with a slow ramp of 1 s (as shown in the insets in B). The recorded current traces were shown in (B). The current traces were recorded in HEKA PatchMaster and traces

were developed in IgorPro 5.04B program. Data represent 7-8 cells in each case and mean mean ± SEM.

features	ifenprodil	MK801	memantine	ketamine	D-APV
Hill slope	~2	~1.5	~1.4	~1.3	~1.7
IC ₅₀ (μm)	~30	~70	~45	~200	~550
effect on transient current	yes	yes	yes	yes	yes, only at higher conc.
effect on tonic current	only in low con- centration	no	at every concentration (from low-to-high)	no	no
mode of action	non-competitive, GluN2B blocker	uncompetitive, open-channel blocker	uncompetitive, open-channel blocker	uncompetitive and allosteric binding site blocker	competitively inhibits Glu -binding site of NMDARs

Table 1: $K_{\nu} 1.3$ Channel Behaviour in Jurkat Cells in Presence of the Tested NMDAR Inhibitors

5.3.3.2 Raji B-Lymphoma Cells

In parallel to Jurkat T cells, the effects of NMDAR antagonists were analyzed on Raji Blymphoma cells. For Daudi B-lymphoma cells, it was shown that the B cell lymphoma drug Rituximab can work through K_v 1.3 channels (Wang et al. 2012). Thus, modulation of K_v 1.3mediated currents in Raji cells through NMDAR inhibitors would suggest that these drugs can potentially act as an adjuvans in treating B-cell lymphoma.

The membrane potential of Raji cells was found to be reduced from ~-45 mV to ~-20 mV in the presence of ifenprodil (20 μ M) and memantine (30 μ M) (**Figure 20A**). K_v1.3 currents were recorded from Raji cells in the presence of ifenprodil and memantine (**Figure 20B**) and the competitive blocker D-APV (**Figure 20C**). The IC₅₀ and Hill slopes values for ifenprodil, memantine and D-APV were ~25 μ M, ~40 μ M and ~700 μ M and ~1.4, ~1.2 and ~1.7, respectively. Thus, the competitive antagonist D-APV only inhibits at a much higher concentration than the open channel and uncompetitive blockers tested. This strongly suggests that NMDAR inhibitors do not act via NMDARs potentially expressed on (T and B) lymphoma cells but most likely through cross-inhibition of K_v1.3-channels.

Figure 20. Effects of NMDAR antagonists on the membrane potential and K⁺-channel activity

of Raji B-lymphoma cells. (A) NMDAR antagonists Ifen. and Mem. induce a membrane depolarization as recorded by currentclamp. K_v1.3 currents were measured by voltage-clamp in the presence of non-competitive Ifen., uncompetitive Mem. (B) and competitive NMDAR antagonist D-APV (C). Dose-response curves for $K_v 1.3$ currents were plotted from the recorded maximal transient currents. All data were calculated from 5-6 cells of four experiments and are represented as mean ± SEM.



5.3.4 Primary Human T Cells from Healthy Donors

Finally, I analysed primary human T cells isolated from peripheral blood of healthy donors. Isolated CD4⁺ T cells were left untreated or were activated with CD3 Ab (MEM-92) and patched for recording K_v1.3 current. It was observed that stimulated T cells needed a higher concentration of antagonist memantine to be inhibited to the same degree as unstimulated T cells. As a result, IC₅₀ and Hill slope values are ~20 μ M, ~40 μ M and ~1.2 and ~1.6 for unstimulated and stimulated populations, respectively as shown in **Figure 21A**.

In addition, $CD4^+$ cells isolated from young (20-30 years) and old donors (55-80 years) were patched to verify the differential expression of K_v1.3 channels on T cell populations of younger and aged persons as already indicated in the literature before by others (Den Braber I, et al. 2012). Interestingly, it is shown that the expression of K_v1.3 was ~20% less in T cells of older population from the younger ones as shown in **Figure 21B**.



Figure 21. Electrophysiological analysis of primary human T cells. $CD4^+$ T cells from blood of normal healthy human donors, unstimulated or stimulated with MEM-92 Ab, were patched in the presence or absence of Mem. to record K_v1.3-mediated currents according to the protocol described in Figure 17A. The amplitude of transient currents was plotted against the inhibitor concentrations to obtain dose-response inhibition curves as shown in (A). IC₅₀ values are indicated. The data in (B) represent K_v1.3 currents expression recorded from CD4⁺ T cells isolated from young and old donors. Data represent the mean <u>+</u> SEM from 4-5 cells recorded from 5-6 cell isolations.

5.4 K_v1.3 Channel Expression on Blood T cells from Dementia Patients

Since NMDAR antagonist memantine affected K_v 1.3 channel activity in primary human T cells, it was highly interesting to investigate how the drug influences T cells of patients diagnosed with Alzheimer's disease, who were treated with memantine. Z1, Z2 and Z3 patients were designated as untreated, treated for 1 week or for 12 weeks with memantine, respectively. The patients were on medication of 10 mg of memantine twice a day.

5.4.1 Altered K_v1.3 Channel Expression in Alzheimer's Patients

To analyse whether NMDAR-low-affine inhibitor memantine applied *in vivo* has effects on $K_v1.3$ -channels, CD4⁺ T cells were isolated from the blood of Z1, Z2 and Z3 Alzheimer patients and $K_v1.3$ currents were recorded. Interestingly, in Z2 patients, who had been treated with memantine for only 1 week, the $K_v1.3$ current of T cells was strongly reduced by ~50%. In Z3 patients, the average amplitude values (70%) had returned to the values of untreated patients (77%). Thus, *in vivo* application of memantine, at least transiently, will have a strong effect on $K_v1.3$ -channel activity of T cells and, thereby, may alter the activation/differentiation processes of T cells.



Figure 22. Memantine *in vivo* treatment affects K_v 1.3-channel activity of T cells. CD4⁺ T cells isolated from blood of Alzheimer patients before (Z1) and after the treatment with Mem. for 1 week (Z2) and 12 weeks (Z3) were recorded for measuring the K_v 1.3 channel activity. The recorded transient current traces shown in (B) were normalized to the highest recorded amplitude values (set as 100%) and are shown in (A) as mean <u>+</u> SEM. The recorded single cells (n) from the number of patients (N) are indicated in the inset.

6. Discussion

6.1 The Expression of Functional NMDARs in Immune Cells is Not Feasible

Several reports have described the presence of NMDARs in immune cells as briefly described in **section 3.7**. As the concentration of Glu in blood is higher than in the CSF, scientists since years have presumed the presence of Glu receptors in immune cells and a role for Glu in regulating immune function. Excited by these observations, we analyzed the function of T and B lymphocytes in the presence of different NMDAR inhibitors used to prove the functional impact of NMDARs in neurons and immune cells (Xiao et al. 2004; Andine et al. 1999). In our collaborative project, we showed that several NMDAR inhibitors reduced antigen-specific T-cell proliferation, cytotoxicity of CD8⁺ T cells, the migration of lymphocytes towards chemokines and TCR-induced signalling (Kahlfuss et al. 2014). However, the presumptive protein expression of NMDARs in T cells was inconclusive due to the lack of electrophysiological recordings of these receptors and detection of protein bands of the obligatory GluN1 subunit at the exact size (~120 kD) as found in brain protein lysates. The first observation was in line with what was mentioned in the publication by Affaticati (Affaticati et al. 2011) and there is no further literature showing the presence of this receptor in lymphocytes at the electrophysiological level.

However, we detected NMDAR expression at the mRNA level (Kahlfuss et al. 2014) and through Ca²⁺-influx after application of external NMDA (**Figure 6**). There can be at least two explanations for the described NMDA-induced Ca²⁺-influx. Firstly, NMDAR expression could be very transient, for instance in stress conditions, as the NMDA-induced Ca²⁺-influx was only observed in HBSS buffer, but not in the cell culture medium RPMI-1640. In connection with this hypothesis, it was observed that GluN1 mRNA expression is increased in the medulla and cortical region of the adrenal gland 24 h after stress exposure (Pirnik et al. 2001). Secondly, NMDA as reagent could cross-react with the pore forming subunit of K⁺-channels and, thereby, induce Ca²⁺-influx in T and B cells. Quantitative RT-PCR analysis should be applied to more clearly determine the differences in NMDAR expression under different cellular conditions.

6.2 NMDAR Inhibitors can be Enlisted in the Therapeutic Arena of K^+ -channels of Immune cells

As there was no GluN1 protein expression detectable in wild-type thymocytes when compared with GluN1 ko thymocytes (Kahlfuss et al. 2014), we thought of candidates that could be modulated by the used NMDAR inhibitors. There are a number of ion channels having a role in maintaining the membrane potential of an immune cell as described in **section 3.3.3**. We have chosen $K_v1.3$ - and $K_{Ca}3.1$ -channels because of their similar structure with NMDAR subunits and their well-described conductive and gating properties. In addition, there is a similarity between the pore-forming subunit of $K_v1.3$ - and $K_{Ca}3.1$ -channels between S5 and S6 (as illustrated in **Figure 2A**) and the binding sites between the obligatory GluN1 and GluN2A/B/C subunits of NMDARs, as illustrated in **Figure 4**. Thus, there is a chance of overlap-binding of NMDAR antagonists to these K⁺-channels. Inspired by this fact, we measured the membrane potential by current-clamp and recorded K_v1.3- and K_{Ca}3.1-currents by voltage-clamp method in murine and human lymphocytes. We detected strong inhibitory effects of both non-competitive and uncompetitive NMDAR inhibitors on the overall membrane potential and on the two described K⁺-channels of immune cells. My experiments spanned from mouse and human primary T cells to mouse (EL-4) and human (Jurkat) lymphoma cells as shown in **Figure 10-12, 15, 17, 18,** and **21** and primary B-cells and Raji lymphoma B-cells depicted in **Figure 14-15, 20**.

As shown in the following **Table 2**, inhibition of $K_v 1.3$ - and $K_{Ca}3.1$ -channels leads to the amelioration several immune disorders, ranging from transplantation rejection, leukaemia to autoimmune disorders like colitis, Rheumatoid Arthritis and asthma. Thus, based on our results, NMDAR inhibitors can be added in the list K⁺-channel inhibitors. As NMDAR antagonists are already in use to treat Alzheimer's disease and are promising candidates for therapy of various other neuronal diseases like Parkinson's disease, depression or stroke (Olivares et al. 2012), the use of these pharmaceuticals also necessitates thorough evaluation of their possible effects on lymphocytes. As NMDAR drugs are administered orally in case of memantine, they will at minimum affect blood lymphocytes while on their way to the brain.

We speculate that NMDAR inhibitors also have side effects on other ion channels. In this respect, specfically the two-pore gated K⁺-channels should be studied as they have two-pore forming subunits (P1 and P2 pores as shown in **Figure 2C**), which can simulate the NMDAR tetrameric structure. They are 'background' K⁺-channels playing a crucial role in setting the resting membrane potential and regulating cell excitability. They are activated by physical and chemical factors (e.g. lipid, pH, heat, and volatile anaesthetics) and are involved in many physiological processes (Kim 2005). Furthermore, they always remain open, are affected even at low membrane potential and regulate lymphocyte osmotic volume (Bobak et al. 2011; Andronic et al. 2013), apoptosis and tumour genesis (Williams et al. 2013; Patel and Lazdunski 2004).

of ion	disease	references	
channel			
	chronic lymphoid leukaemia	Leanza et al. 2013	
	treatment		
	allo-transplantation studies	Hautz et al. 2013	
	inhibition of T _c cells	Hu et al. 2013	
K _v 1.3	Rheumatoid arthritis	Toldi et al. 2013	
	amelioration		
	experimental autoimmune	Li et al. 2012	
	encephalitis treatment		
	immunosuppression strategy in	Grgic et al. 2009b	
	kidney allograft		
	ischemia treatment improvement	Chen et al. 2011	
	helping in lowering blood pressure	Damkjaer et al. 2012	
	T-cell mediated colitis	Di et al. 2010	
	renal fibrosis	Grgic et al. 2009a	
	immunosuppression strategy in	Grgic et al. 2009b	
	kidney allograft		
	prevention of obliterative airway	Hua et al. 2013	
K _{Ca} 3.1	diseases		
	blockage of pancreatic cell growth	Jager et al. 2004	
	target for hypertension disease	Kohler et al. 2010	
	target for Alzhimer's Disease	Maezawa et al. 2012	
	C .	Tharp et al. 2008	
	limiting stenosis	Toyama et al. 2008	
	blocking atherogenesis		
	human osteoclast shock recovery	Weskamp et al. 2000	
	cardiovascular targets	Wulff and Kohler 2013	
	RA amelioration	Toldi et al. 2013	
	idiopathic pulmonary fibrosis	Roach et al. 2013	
	prevention of allograft vasculopathy	Chen et al. 2013	

increases tumor cell killing	Koshy et al. 2013
chronic asthma	Girodet et al. 2013

6.3 NMDAR Surface Expression under Tight Control

As discussed, there could be a transient expression of NMDARs in immune cells. It should be considered that even in neuronal cells the expression of NMDARs is hard to detect. In the neuronal synapse, only few NMDARs are expressed, often less than 10 per synapse (Racca et al. 2000). This suggests that few NMDARs expressed on the cell surface are functionally important. In addition, a single NMDAR (30-50 pS) conducts more Ca²⁺ than a single Stim-ORAI1 complex (0.024-0.4 pS) (Hogan et al. 2010; Feske 2007). Thus, the surface expression of NMDARs on immune cells should be under very tight control. In our experiments, we could not reliably detect NMDAR surface expression on live resting or activated T cells or on thymocytes with routine FACS staining and flow cytometry using two different GluN1 Abs (from Synaptic Systems and Alomone labs), which bind to extracellular epitopes of GluN1.

Figure 23. Differential conductivity levels of ion channels present in immune cells. The different ion channels present in immune cells have different conductive properties as indicated. K_v 1.3- and K_{Ca} 3.1-channels have more or less the same conductive properties, whereas ORAI1 complex has very less conductivity in comparison with other ion channels. Along with the conductive properties, they have differences in the gating and opening kinetics. NMDARs are hypothetically shown in the figure to illustrate the differences explained in the text.



In addition, experiments for over-expressing these receptors in immune cells using several transfection procedures were not successful in terms of transfection efficiency. Among the different transfection protocols used for transfecting immune cells, the Amaxa Nucleofector procedure was best in immune cells showing in ~5% of cells GluN1-GluN2B double transfection (GluN1-wt from *Paoletti, P.* and GluN2B-SEP from Addgene). Since these two constructs showed the best results among the different tested NMDAR constructs, but still had a very low transfection efficiency, the two constructs were subcloned into another vector, pXJ41, known to carry bigger inserts. Although the subcloning procedure worked with this vector, the transfection efficiency for NMDAR subunits in immune cells was still not improved. This indicates that lentiviral vectors might be needed, which are efficient for gene transfer into human T cells (Verhoeyen et al. 2009).

6.4 NMDAR Inhibitors Modulate the Steady-state Current of K_v1.3-channels

 K_v 1.3- and K_{Ca} 3.1-channels are similar with regard to their conductance properties as both channels are highly selective and have similar single-channel conductance in the order of 10-14 pS as shown in **Figure 23**. However, they are remarkably different in their gating and blocker sensitivity. K_v 1.3-channels are activated by depolarization of the plasma membrane. The activation threshold of the channel is close to the resting potential of the immune cells of ~-50 mV. This current is quickly activated and then inactivated by a slow C-type inactivation. On the contrary, K_{Ca} 3.1-channels are solely activated by the increase of cytosolic Ca²⁺-concentration and they are not deactivated after prolonged exposure to increased cytosolic Ca²⁺-concentration (Cahalan and Chandy 2009; Panyi et al. 2006).

Ifenprodil blocks pre-synaptic L-type Ca²⁺ channels at a concentration of >10 μ M (Delaney et al. 2012). We saw ifenprodil at lower concentration and memantine from low to high concentration has substantial effects on both the amplitude and steady-state parameters of K_v1.3-channel kinetics as described in **Table 1** for Jurkat cells. It was shown in **Figure 19** that ifenprodil and memantine have quantitative effects on the tonic current properties of Jurkat cells. This can hypothetically be simulated as shown **Figure 24**, where MK801 block in immune cells can easily be released as it is only affecting the amplitude of K_v1.3 channel transient current. In contrast, a block with ifenprodil is difficult to release immediately as the steady-state or tail current parameters are also affected along with the amplitude. Although, 20 min wash-out with the extracellular solution can reverse back the normal K_v1.3 current in an experimental situation.



Figure 24. Hypothetical model for the regulation of K_v 1.3-channel by ifenprodil and memantine. As explained in section 3.6.1, K_v 1.3 generally follows C-type inactivation, which is characterized by a slow type of inactivation. Ifenprodil blocks the pore region so that the channel can not be inactivated anymore. A channel generally follows three steps of opening and closing: open, inactivated, deactivated and closed. It is easy to change an inactivated state of a channel to a closed and open stage, but it is hard to close a channel completely when it reaches a deactivated stage. In case of ifenprodil, K_v 1.3 hypothetically reaches to a deactivated stage. Thus, this inhibitor is hard to re-

lease from the pore region of the bound channel. In contrast, MK801 block can easily be released as the state reached after its release is an inactivated stage (modified from *Barros, F., 2012*) (Barros et al. 2012).

The inactivation parameters of $K_v 1.3$ -channels are interesting to study in terms of $K_v 1.3$ channel inhibitors. Increasing amount of evidence suggests that there are certain compounds which inhibit K_{v} -currents not directly by the K⁺-conduction pathway (S4), but by accelerating the selectivity filter destabilization (between S5 and S6) once the channel opens (as illustrated in **Figure 25**). This mode represents an alternative mechanism of K_{v} -channel inhibition. Mutation of these amino acids to arginine, lysine, alanine or glutamate accelerates C-type inactivation, while mutation to valine or tyrosine residues retards it. In addition, the interaction via hydrogen bonds between Shaker 438 and 447 is important in controlling the C-type inactivation gate. However, how pivotal these residues determine the C-type inactivation rate is still unknown. Notably, the role of T449, a residue at the external mouth has been studied in details for this kind of inactivation (Cordero-Morales et al. 2011; Lopez-Barneo et al. 1993). The traditional K⁺-channel blocker Tetra-ethyl ammonium (TEA) and 4-Aminopyridine (4-AP) have been shown to hasten the slow C-type inactivation. It was already shown that TEA possibly binds to T449 preventing the selectivity filter constriction in a 'foot-in-the-door' fashion (Leung 2012). Thus, it would be interesting to study whether the NMDAR inhibitors acting on the tonic current also modulate the T449 site like traditional K⁺channel blockers or any other site of the K_v 1.3-channel unknown until now. This inactivity phenomenon of shaker K⁺-channels by NMDAR antagonists could be very complex as they could also interplay with accessory β subunits as shown for K_v1.5- channels (Decher et al. 2005).



channels have a 6-transmembrane topology along with a one poreforming subunit (forming the gate) and a charge-transfer subunit (S4). It is shown how the Cterminal is occluding the intracellular mouth of the ion-conducting pore in case of K_v 1.3 inactivation



(modified from *Buckley, C,2005* and *Barros, F., 2012*) (Barros et al. 2012; Buckley and Vincent 2005).

6.5 NMDAR Inhibitors Affect B-Cell Activation

Since there exist Abs against different ion channels present in neuronal cells as described in **Table 3**, it is clear that B cells take part in neuronal disorders like MS (Meinl et al. 2006; Ritchie et al. 2004). Therefore, we also analysed B-cell activation in the presence of different NMDAR inhibitors. Notably, they inhibited BCR-induced Ca²⁺-flux as shown in **Figure 9**, lowered the total membrane potential and inhibited K⁺-channels in activated primary murine B cells and Raji lymphoma cells (**Figure 13, 14, 20**).

In general, B cells contribute to neuronal disorders via four mechanisms: by production of Abs causing tissue damage either via complement activation or Ab-dependent-cellmediated cytotoxicity; by acting as APCs for the expansion of cytotoxic T cells and Th cell cytokine production; by production of pro-inflammatory cytokines like IL-6 and TNF, which activate macrophages and T cells enhancing tissue damage; and by formation of ectopic germinal centres in the intermeningeal place in a process called neolymphogenesis (Dalakas 2008). In specific, Abs against NMDARs were detected in women with ovarian tumour, young children and men without tumour (Dalmau et al. 2007; Novillo-Lopez et al. 2008). The major symptoms in women were memory loss, seizures and decreased consciousness, whereas children and men showed a diffuse lymphocytic meningoencephalitis and acute juvenile non-herpetic encephalitis (Dale et al. 2009; lizuka et al. 2008). The main epitope targeted by the Abs lies in the extracellular N-terminal domain of the GluN1 subunit (Dalmau et al. 2008). Thus, application of non-competitive NMDAR antagonists during chronic treatments of neurological disorders like Morbus Alzheimer may not only involve neuronal NMDARs, but may have additive side-effects by targeting B cells, in addition to T cells, which are assumed to contribute to these disorders (Yanaba et al. 2008; Danysz and Parsons 2012; Cunningham 2013; Frodl and Amico 2014). The inhibitory side-effects of NMDAR inhibitors on B cells might also be beneficial in treating sepsis (Roger et al. 2009) as we have seen a reduction in LPS/TLR4-induced B-cell signalling and proliferation (Figure 9 and data not shown but in revision for publication) in the presence of memantine and other NMDAR inhibitors.

B cells express $K_v 1.3$ and $K_{Ca}3.1$ channels and their mode of Ca^{2+} -influx is through Stim-ORAI complexes as in T cells. Blockade of these K⁺-channels by specific K⁺-channel inhibitors results in immunosuppression (Wulff et al. 2004; Matsumoto et al. 2011). The K_v1.3and K_{Ca}3.1-specific blockers Shk and TRAM-34 are used in treating vasculopathy or kidney allograft rejection (Chen et al. 2013; Grgic et al. 2009a). K_v1.3 inhibitor Shk-186 had been started to use in human trial stage I for treating diseases by Kineta. In addition to BCR- and LPS-induced activation of primary B cells, we found that memantine reduced K_v1.3-channel currents recorded from Raji lymphoma cells (**Figure 20**). In this respect, memantine may therapeutically compete with Rituximab, which has been shown to act through the inhibition of K_v 1.3 channels in human B lymphoma cells (Wang et al. 2012). Further studies are required to determine the drug's suitability in diverse *in vivo* settings.

channel	disease	references	
voltage-gated Ca ²⁺	Lambert Eaton Myasthenic	Pellkofer et al. 2008	
channel	Syndrome (LEMS)		
Voltage-gated K ⁺	Acquired Neuromyotonia,	Hart et al. 1997; Liguori et al.	
channel	Morvan's Syndrome (MoS),	2001; Bataller et al. 2007;	
	Limbic Encephalitis (LE),	Dedek et al. 2001	
	Epilepsy		
α-3 ganglionic	antibodies to α -3 AchR	Vernino et al. 2000	
acetylcholine recep-	containing α 3 and β 4		
tor			
Glu and GABA	Encephalitis associated with	Novillo-Lopez et al. 2008; lizuka	
receptors	NMDAR, AMPAR and GABAR	et al. 2008; Dale et al. 2009; Lai	
		et al. 2009; Rogers et al. 1994;	
		Lancaster et al. 2010	
Aquaporin-4	Neuromyelitic Optica (NMO) or	Lennon et al. 2005	
	Devic's Syndrome		

 Table 3. Autoimmune Disorders or Abs Related with Neuronal Channels

6.6 Surveillance of Immune Status is Required during Memantine Treatment

A new therapeutic concept generally passes through a long journey of clinical trials to accomplish the final verification for using it in humans. NMDAR antagonists like memantine may have a wide range of potential therapeutic applications, spanning from acute neurodegeneration in stroke and trauma, chronic neurodegeneration in Morbus Parkinson and Alzheimer, amyotrophic lateral sclerosis, and Huntington's disease to symptomatic treatments in epilepsy, anxiety, or chronic pain (Parsons et al. 1995; Meldrum 1992a, b; Muller et al. 1996). The high affinity competitive inhibitor D-APV cannot be used *in vivo* as it impairs normal synaptic transmission creating serious side-effects. Therefore, the major challenge was to develop an antagonist that prevents the pathological activation of NMDARs but maintains their physiological activity. Memantine (1-amino-3, 5-dimethyl-adamantane) as an uncompetitive low-affinity inhibitor has fulfilled the criteria. It was registered in Germany for a variety of CNS indications in 1978, but its most likely therapeutic mechanism was only discovered 10 years later (Bormann 1989; Kornhuber et al. 1989; Kornhuber et al. 1991). It was first synthesized by researchers at Eli Lilly company as an agent to lower blood sugar levels (Gerzon et al. 1963). But, since it is devoid of such activity, Merz & Co. applied for a patent in 1972 for the treatment of several cerebral disorders. Since then, several *in vivo* studies were performed to determine its effects and Merz filed an international application in 1989 claiming memantine as ideal for the treatment of cerebral ischemia and Alzheimer's dementia. Since then, clinical research has focused on the treatment of dementia (Parsons et al. 1999).



Figure 26. Scheme of memantine trafficking in the body. A schematic pathway of the traversing route of memantine when administered in patients. Memantine is mainly administered orally because of stability issues. It is taken up by the gut and passes into the blood stream where it can act on blood lymphocytes. Memantine then crosses the BBB, gets absorbed into the liver or is eliminated by the kidney as unchanged substance or hydroxylated metabolites. Bioavailability is thought to reach ~100% in case of this drug.

Memantine is prescribed orally as a tablet or solution. In our experiments, we have analysed T cells from patients who are taking a 10 mg tablet twice a day. We found that K_v1.3-mediated currents were lowered by 50% on T cells isolated from patients treated for 1 week with memantine, compared to T cells from those patients before drug treatment. K_v1.3-mediated currents were nearly normal in T cells from patients treated for 12 weeks with the drug. Orally uptaken memantine crosses the BBB to act on neurons and is metabolized in the liver through first-pass metabolism or is excreted through the kidney as an unchanged substance or hydroxylated metabolites. Thus, it is obvious that memantine could affect blood lymphocytes. No major side-effects of memantine on lymphocytes were reported until the Food and Drug Administration (FDA)'s October 2013 survey summary. This summary describes memantine's adverse reactions on blood cells and the lymphatic system as post-marketing drug exposure experience.

In this context, our results with the attenuation of K_v 1.3-channel activity in Z2 patients is very striking. Although K_v 1.3-channel activity was almost "normal" in Z3 patients, T cells from Z3 patients are non-reactive to stimulation (personal communication of unpublished results by T. Lowinus/U. Bommhardt).



Figure 27. Hypothetical model of K_v**1.3 counter-adaptive action upon memantine exposure.** A hypothetical model to determine the action of K_v**1.3** channels during first and second drug exposure. During first drug exposure the immune cells become suppressed, whereas during continued or second exposure lymphocytes regain their function through the increase of K_v**1.3** channel density (*modified from Ghezzi, A.*) (Ghezzi et al. 2010).

The observed "re-expression" of K_v1.3 currents in Z3 patients thus may represent some counter-adaptive role of K_v-channels to resist to the drug's long-term exposure. Such counter-adaptive drug tolerance was described as triggering adaptive homeostatic changes that oppose the drug's effects to produce drug tolerance (Koob and Bloom 1988). Upon drug clearance, the same changes become counter-adaptive and produce symptoms of dependence. Our idea of memantine-induced K_v-channel tolerance can hypothetically be described by a model as illustrated in **Figure 27**. In this case, the immune system or lymphocytes become suppressed with the first exposure to memantine after 7 days (**Figure 27B**). Then, there is thought to be the induction of higher K_v1.3 transcription in the drug removal phase (**Figure 27C**). Thus, the immune system reaches to a tolerance level in the 2nd exposure of the drug after 12-weeks (**Figure 27D**). In addition, there should be a time point recorded in between these two extreme time points (1 and 12 weeks) to validate the hypothesis of increased channel expression described in **Figure 27C**. Furthermore, single channel current should be recorded to support our idea of K_v channel counter-adaptive effect.

6.7 Good News for Neurons

The physiological role of K⁺-channels is evident in microglia, resident macrophages in CNS (Villalonga et al. 2010). Expression of K_v1.3 and K_{Ca}3.1 is similar in macrophages to peripheral T and B cells along with the Ca²⁺-influx through CRAC channels. Microglia has dichotomous functions when activated, either causing neuronal damage by releasing cytotoxic substances and pro-inflammatory cytokines or being neuro-protective by releasing neurotrophic factors and clearing amyloid aggregates and debris from degenerated neurons

(McGeer and McGeer 2010; El Khoury and Luster 2008; Cameron and Landreth 2010; Hickman et al. 2008). K_{Ca}3.1-mediated Ca²⁺-entry has been shown to be neurodegenerative in case of microglia when induced by amyloid aggregates causing oxidative burst, nitric oxide production, and neuronal death (Kaushal et al. 2007; Schilling et al. 2004; Khanna et al. 2001). Soluble A β oligomers (A β O) are small and early stage amyloid aggregates (Maezawa et al. 2011). In experimental conditions, ABO was found to initiate an increase of intracellular Ca²⁺ of microglia either directly by forming a membrane pore or indirectly through interaction with an unknown membrane receptor. This leads to a hyperpolarisation of the membrane potential by effluxing K⁺ through K_{Ca}3.1 channel. This hyperpolarisation provides the driving force of Ca²⁺-entry through store-operated inward-rectifier CRAC channels, thus sustaining the Ca²⁺-signals necessary for selective Ca²⁺-activated pathways (Maezawa et al. 2012). In this case, K_{Ca}3.1 channel inhibitors should be neuro-protective by deactivating the microglial functions involved in degenerating neurons. Indeed, the K_{Ca}3.1 inhibitor TRAM-34 inhibits ABO-induced microglial activation and microglia-mediated neuronal toxicity (Maezawa et al. 2011). In conclusion, if memantine would inhibit microglial K_{Ca}3.1 currents as well as pathologic NMDAR functions, the drug would have a double benefit for dementia patients. However, this interesting hypothesis needs further experimentation.

7. <u>References</u>

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8. Abbreviations

- aa: amino acid
- Abs: antibodies
- AchR: acetyl choline receptor
- AD: Alzheimer's disease
- APC: antigen-presenting cell
- AMPA: α-amino-3-hydroxy-5-methylisoxasole-4-propionate
- BBB: blood-brain barrier
- BCR: B-cell receptor
- BSA: bovine serum albumin
- Cav: voltage-activated calcium channel
- CRAC: calcium release-activated Ca2+-channels
- CD: cluster of differentiation
- CLL: chronic lymphoid leukemia
- CNS: central nervous system
- CSF: cerebro-spinal fluid
- DC: dendritic cell
- DMSO: *Dimethylsulfoxide*
- DP: double positive
- DN: double negative
- ER: endoplasmic reticulum
- EAE: experimental autoimmune encephalomyelitis
- FACS: fluorescence activated cell sorter
- FBS: fetal bovine serum
- FITC: Fluorescein IsoThio Cyanate
- Glu: glutamate
- HBSS: Hanks' Balanced Salt Solution
- 5-HT3: serotonin receptor
- Ifen.: *ifenprodil*
- iGluR: ionotropic glutamate receptor
- KO: knockout
- K_v: voltage-activated potassium channel
- K_{Ca}: calcium-activated potassium channel
- LN: lymph node

LPS: lipopolysaccharide Mem.: memantine MEM: minimum essential medium eagle MAPK: mitogen-activated protein kinase MOG: myelin oligodendrocyte glycoprotein mGluR: metabotropic glutamate receptor mTOR: mammalian target of rapamycin NMDAR: N-methyl D-aspartate receptor NEA: non-essential amino acids NF-κB: nuclear factor-κB NFAT: nuclear factor of transcription of activated T-cells OVA: ovalbumin PBS: phosphate buffered saline PBL: peripheral blood lymphocytes PFA: paraformaldehyde PBMCs: peripheral blood mononuclear cells rpm: revolutions per minute RACC: receptor-activated calcium channel RT: room temperature **RPMI:** Roswell Park Memorial Institute Medium SOCC: store-operated calcium channel SOCE: store-operated Ca²⁺-entry SMOCC: second messenger-operated calcium channel STIM: stromal interaction molecule tg: transgenic T_H: T helper cell T_C: cytotoxic T cell TCR: T-cell receptor TLR: Toll-like receptor TRP: transient receptor potential T_{CM}: central memory T cell T_{EM}: effector memory T cells

9. List of Figures and Tables

9.1 Figures

- 9.1.1 Oscillatory changes of the membrane potential in lymphocytes (p.14)
- 9.1.2 Scheme of structural classification of K⁺-channel subunits (p. 16)
- 9.1.3 Classification of glutamate receptors (GluRs) (p. 20)
- 9.1.4 NMDAR model illustrating important binding sites for agonists and antagonists (p. 21)
- 9.1.5 NMDAR subunit transfection for testing the specificity of Abs used to detect NMDAR subunit expression (p. 42-43)
- 9.1.6 NMDAR antagonists attenuate NMDA-induced Ca²⁺-flux (p. 43-44)
- 9.1.7 Single-cell Ca²⁺-flux of TCR tg T cells in the presence of antigen-presenting DCs or CD3/CD28-coated beads (p. 44-45)
- 9.1.8 NMDAR antagonists attenuate TCR-induced Ca²⁺-flux (p. 45)
- 9.1.9 Reduced Ca²⁺-flux in BCR-activated B cells in the presence of NMDAR antagonists (p. 46)
- 9.1.10 NMDAR antagonists change the membrane potential (p. 46-47)
- 9.1.11 NMDAR antagonists inhibit K_v 1.3 and K_{Ca} 3.1 channel activity in activated T cells (p. 47)
- 9.1.12 NMDAR antagonists decrease K_v 1.3 channel activity in activated CD4⁺ T cells and EL-4 lymphoma cells (p. 48)
- 9.1.13 NMDAR antagonists reduce the B cell membrane potential (p. 49)
- 9.1.14 NMDAR antagonists lower K⁺-channel activity of B cells (p. 50)
- 9.1.15 NMDAR antagonists change the membrane potential of Jurkat cells (p. 51)
- 9.1.16 Comparison of recorded K_v1.3 current properties between control and Ifen.-treated
 (30 μM) Jurkat cells (p. 51)
- 9.1.17 Electrophysiological analysis of Jurkat T cells in the presence of Ifen. and Mem. (p. 52)
- 9.1.18 K_v1.3 current recorded from Jurkat cell lines in presence of mentioned inhibitors (p. 53)
- 9.1.19 Inactivation current differences in K_v1.3-mediated currents in control and inhibitor population (p. 53-54)
- 9.1.20 Effects of NMDAR antagonists on the membrane potential and K⁺-channel activity of Raji B-lymphoma cells (p. 55)
- 9.1.21 Electrophysiological analysis of primary human T cells (p. 56)
- 9.1.22 Memantine in vivo treatment affects Kv1.3-channel activity of T cells (p. 57)
- 9.1.23 Differential conductivity levels of ion channels present in immune cells (p. 61)

- 9.1.24 Hypothetical model for the regulation of K_v1.3-channel by ifenprodil and memantine (p. 62)
- 9.1.25 The K_v 1.3-channel inactivation pathway (p. 63)
- 9.1.26 Scheme of memantine trafficking in the body (p. 66)
- 9.1.27 Hypothetical model of $K_\nu 1.3$ counter-adaptive action upon memantine exposure (p. 67)

9.2 Tables

- 9.2.19 K_v1.3 Channel Behaviour in Jurkat Cells in Presence of the Tested NMDAR Inhibitors (p. 54)
- 9.2.20 The disease pathologies related with $K_v 1.3$ and $K_{Ca} 3.1$ channels (p. 60-61)
- 9.2.21 Autoimmune disorders or Abs related with neuronal channels (p. 65)

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11. <u>Erklärung</u>

M.Sc. / Bose, Tanima Phone Nr. : +49-(0)-17662668893 (Mob.), +49-(0)-3912427262 Adresse: Leibniz Institute for Neurobiology, Brennecke Strasse 6, D-39118, Magdeburg.

E-Mail-Adresse: tbose@lin-magdeburg.de, tanimabose@gmail.com

Erklärung

Hiermit erkläre ich, dass ich die von mir eingereichte Dissertation zum dem Thema

'Crosstalk between NMDAR Antagonists and Potassium Channels in Murine and Human Lymphocytes'

selbständig verfasst, nicht schon als Dissertation verwendet habe und die benutzen Hilfsmittel und Quellen vollständig angegeben wurden.

Weiterhin erkläre ich, dass ich werder diese noch eine andere Arbeit zur Erklangung des akademischen Grades Dr. rer. nat. an anderen Einrichtungen eingereicht habe.

Unterschrift

12. <u>Curriculum Vitae</u>

Name: Date of Birth:	Tanima Bose 10.01.1985
E-mail:	tbose@lin-magdeburg.de
	tanimabose@gmail.com
Address:	Rudolf-wolf Strasse 11, D-39112, Magdeburg
Gender:	Female
Nationality:	Indian
Languages known:	English (TOEFLiBT), German (A1 level from Goethe Institute),
	Hindi, Bengali

PhD Project

\blacktriangleright	04/2010- continuing	'Crosstalk between NMDAR Antagonists and Potassium Channels in Lymphocytes',	
		Supervisors: Dr. Martin Heine and Prof. Dr. Ursula Bommhardt, Leibniz Institute for Neurobiology	

Training experience

	10/2009-03/2010	Molecular Neurobiology in Univ. of Greifswald, Germany
	05/2009- 09/2009	Calcium Signaling and Cellular Nanodomains, Homburg, Germany
4	09/2008- 04/2009	'Role of WISP3 in Cartilage', Supervisor: Dr. Malini Sen, Indian Institute of Chemical Biology, Kolkata, India (Master Thesis)

Academic qualification

Examinations	10 th Standard	10+2	B.Sc.	M.Sc.
Year of Passing	2001	2003	2006	2008
Institution Name		Bethune Collegiate School, India	Chemistry, Zool- ogy, Mathematics from Visva-Bharati University, India	

Marks secured	85.6%	71.2%	74.6%	Aggregate:
				5.24 CGPA (out of 6)

Publications

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'Insight into the molecular basis of the unique T cell signaling induced by the CD28 superagonistic antibody TGN1412.' Rudolph J.M., Wiechec-Los E., Bartholomäus P., Merten C., Ramonat A., Steiner M., Bose T., Heine M., Fischer K.-D., Kliche S., Kalinke U., Schraven B. - selected for oral presentation+
 +both presented in German Society for Immunology (DGfI, Mainz, 2013).

91

Other Appearances

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