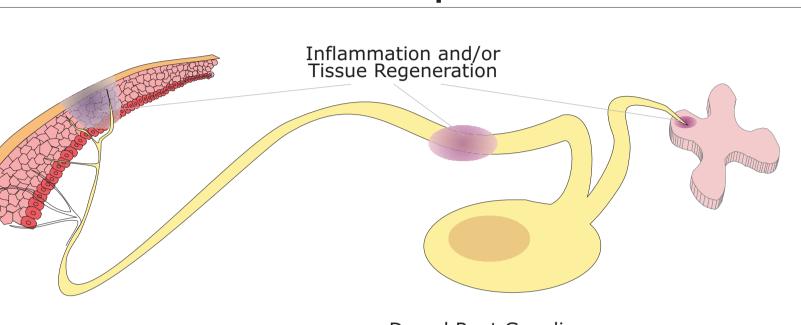
Modellierung von peripheren Schmerzschaltern (MoPS) Modelling of Pain Switches (MoPS)

R. Baron (Christian-Albrechts-Universität Kiel), M. Heiner (Brandenburgische Technische Universität Cottbus), F. Herberg (Universität Kassel), T. Hucho (Max-Planck-Institut für molekulare Genetik, Berlin), W. Marwan (Otto-von-Guericke-Universität Magdeburg), P. Reeh (Universität Erlangen-Nürnberg), H. Seitz (Max-Planck-Institut für molekulare Genetik, Berlin), C. Stein (Charité Universitätsmedizin Berlin), MicroDiscovery (GmbH, Berlin)

Characterisation of "Pain" Modules

In clinical practice pain is among the most prevalent symptoms. Treatment of 50-70% of chronic pain patients is insufficient. The current approach of therapy guided by the concurrent/symptomatic disease shows highly variable success. Thus, a shift toward a "mechanism-based" therapy is urgently needed. Progress in elucidation of molecular Target Tissue Peripheral Nerve components involved in sensitization of peripheral neurons activated by painful stimuli is enormous. Nevertheless, the fundamental question, if these components Prostaglandins Histamine Serotonin are part of a network converging onto one or Endothelin Bradykinin Epinephrine few pain, i.e. "nociceptive modules", or if they Opioids GPCR constitute discrete paralele mechanisms has hardly begun to be addressed Our project combines mathematical modelling of pre-existing data with novel data collection of a comprehensive number of signalling components to feedback into the mathematical models. Thus, we will be able to identify differential nociceptive modules. These data in turn will allow us to test the derived mathematical models electrophysiologically on signal1 primary sensory neurons and in vivo animal models.

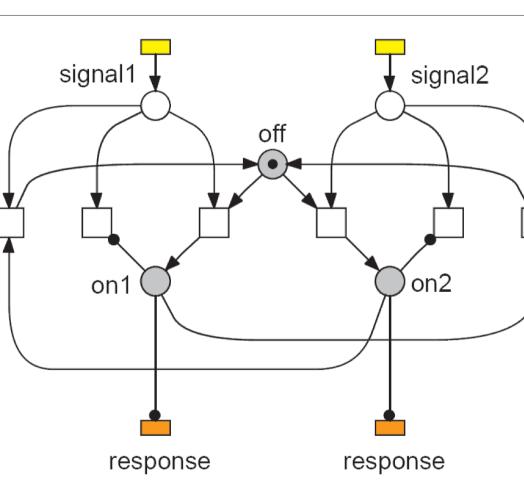
progress is the lack of an experimental cell line system. Thus we will investigate signalling events directly in primary sensory neurons. As the starting material is apply highly efficient current techniques sparse we such as BIAcore interaction meassurements, FRET/BRET technologies, cell lysate spotting and high content immunofluorescent single cell analysis. Focussing first on enzymes regulated by phosphorylation, their Dorsal Root Ganglion Spinal Cord Cell Body activation, activity kinetics as well as the relationships of various signaling molecules in response to various extracellular stimuli will be characterized. TNFa The signaling network will be nucleated by RTKs Interleukins studying well described signalling Family Other components such as cAMP, the MAP-kinases Erk1/2, PKCε, as well as PKA before enlarging the focus onto as many as possible components, known as well as novel to pain signalling. As modules around these components are in part also modelled in other cellular systems, only a minimal number of timepoints and conditions is required for the

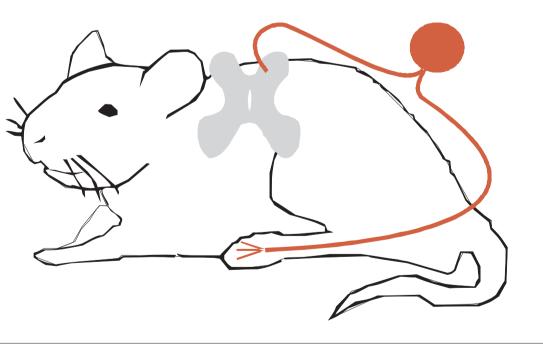


Ion Channels V3 v4 1.8 1.7 1.6 ASICs Hyperalgesia

Mathematical Modelling

The consortium includes molecular scientists, cell biologists, electrophysiologists, behavioural scientists, clinicians as well as mathematicians. Accordingly a wide variety of data qualities will have to be combined. The modelling of the data has therefore to be versatile as well as intuitively accessible for all coworkers. Thus, we use the easily visualizable approach of Petri Nets. Classicle transition nets as well as stochasticle and continous nets will be applied for qualitative and quantitative analysis of the signaling networks. Painful stimuli activated neurons are not homogenous. Instead, multiple subgroups of neurons can be differentiated. Thus, tools will be constructed for subgroup specific data analysis. As the modelling should serve as base for wet lab hypothesis building and experimental design, further tools allowing temporary partial network "knock out" and/or disturbances will be developed.



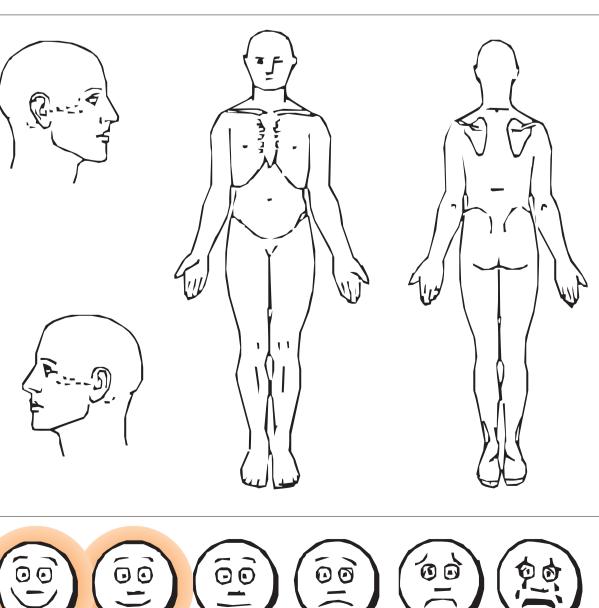


transfer to sensory neurons. This analysis will give first indications if signaling cascades converge or if they act seperately.

Validation on Animal Models and Humans

Pain modules identified through cellular as well as modelling approaches will be tested in vertebrates. Animal behaviorural studies as well as electrophysiological studies for example on nerve skin preparations will be performed to validate the cellular results. These studies will be then extended to suitable human experimental/clinical conditions. For the latter a powerful quantitative standardized sensory testing tool developed by the BMBF financed "Deutsche Forschungsverbund neuropathischer Schmerz (DFNS)" is available.

Cellular/Molecular Data Aquisition Knowledge about intracellular signalling underlying sensitization of painful- stimuliactivated neurons is still only starting to emerge. One aspect, which has hampered



Overal Aims

1) Definition of peripheral nociceptive signalling modules

2) Mechanism based differentiation between pain states

3) Guide for mechanism based therapy of current therapeutics

4) Initiation of development of novel therapeutics