## Schwann cell precursors represent a neural crest-like hub state with biased multipotency Louis Faure\*1, Maria Eleni Kastriti\*1, Dorothea von Ansen\*1, Francois Lallemend<sup>2</sup>, Saida Hadiab<sup>2</sup>, Idor Adamevko<sup>1</sup> <sup>1</sup>Medical University of Vienna - Center for Brain Research, Dpt. of Neuroimmunology, Wien, Austria, \*: equal contribution Der Wissenschaftsfonds. <sup>2</sup>Karolinska Institute, Department of Neuroscience, Stockholm, Sweden Bifurcation analysis reveals early biasing-Scientific quests **Enteric** · What are the molecular portrait of NC & SCPs lineages? neurons ( . What is the structure of cell fate decision? dlia eneurons Endoneurial Are SCPs multipotent neural crest-like cell state? fibroblasts -Data sampling & analysis-Neuro-muscular iunction SC Early genes Early genes Chromaffin myelinating Cells are sampled using lineage tracing over various embryonic Ckb Fabp7 stages and locations, and were sequenced using SmartSeg2 cells Sox8+ TIx2 Mist . From the count matrices were generated the differentiation tree and CvtoTRACE measurements. From spliced/unspliced matrices were generated RNA velocity mapping non myelin. Hub cells contains several bifurcations toward terminal fates such as ChC, enteric glia/neurons, and Serpine2+ Hub state -Investigation of SC heterogeneityiSCs F12.5 DE analysis is performed to identify branch specific genes, which are then further classified according to their pseudotime activation: before fork=early, after fork=late Cranial NCC . In all bifurcations, the following phenomenon occurs: F9.5 (1) co-activation of branch-specific gene modules mSCs prior to the fork/bifurcation (2) repulsion of these modules while reaching the fork Trunk neural crest cells (NCC) pseudotime (3) commitment after the bifurcations രം Coordinated metaregulons maintain the NC-like state in hub cells dataset Poust1 Mp: Pmp22 Mai IMAP 2 **Mesenchyme** Pmp2 . The "hub"/SCPs might be defined as a cell 02 state with re-activation of metaregulon 5, Pmp22\*\*\* characterized by Ets1 LIMAP 1 · Metaregulons 1 and 3 are linked to neural Spatio-temporal aspects crest properties and cell reprogramming. Metaregulon 1 Metaregulon 2 Metaregulon 3 Metaregulon 4 Metaregulon 5 Metaregulon 2 is linked to the terminal · Focused trajectory analysis shows transcriptional and regulatory paths differentiation of Schwann cells towards myelinating, non-myelinating, neuro-muscular junction terminal Ets1 Phox2b/ I eff . The "hub" state is distinct in terms of Msx3 Rest SCs and endoneurial fibroblasts Tfap2a mad3 Gbx2 Nfia/b Phy1 Acel1 regulation from the majority of the neural Sox2/8 Pax3 Tap2b Sox10 Hdac2 . Pou3f1, known to play a crucial role in myelination, emerged as a crest cells where metaregulons 1 and 3 are Nfatc1/2 candidate biasing factor towards terminal neuro-muscular junction SCs. dominating.