Supplemental material

Title: Olig2 regulates Purkinje cell generation in the early developing mouse cerebellum

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Supplemental Figure 1:



Figure S1. The NTZ Olig2⁺ cells represent a distinct subpopulation of DCN neurons in the embryonic cerebellum. Mid-sagittal sections of the E14.5 (A-C) and coronal sections of the E18.5 (D) wild-type cerebella are immunostained. The NTZ Olig2 and Pax6 (to label GNPs) show a non-overlapping pattern (A). Between the $Olig2^+$ and $Pax6^+$ domains, there are a group of cells that are negative for both markers, but positive for Tbr1 (to label a subset of DCN neurons) (B). Higher magnification views of the boxed regions are shown as inserts in (A and **B**). Although the $Olig2^+$ and $Ibr1^+$ domains are closer in position with double positive cells being occasionally seen (**B**, insert, arrowhead), they are largely non-overlapping. In addition, the NTZ Olig2⁺ cells are nearly all postmitotic as indicated by their inability to take up BrdU in a BrdU-pulse labeling experiment (C). Notably, a number of $BrdU^+$ cells can still be found in the VZ, and vet Olig2 expression has already disappeared in this region at this developmental stage. At E18.5, while Tbr1⁺ neurons concentrate to the medial (Med) part of the DCN, Olig2 expression is more towards the lateral (Lat) region (**D**). Two morphologically distinct Olig2⁺ cells are found with smaller cells being located in the periphery of the Olig2⁺ domain and bigger cells being in the core (**D**, inserts). Non-specific staining signals from the choroids plexus and a representative blood vessel are indicated by asterisks in (C) and (D). VZ, ventricular zone; RL, rhombic lip; NTZ, nuclear transitory zone. Scale bar: 80 um in A-C; 160 μm in **D**.