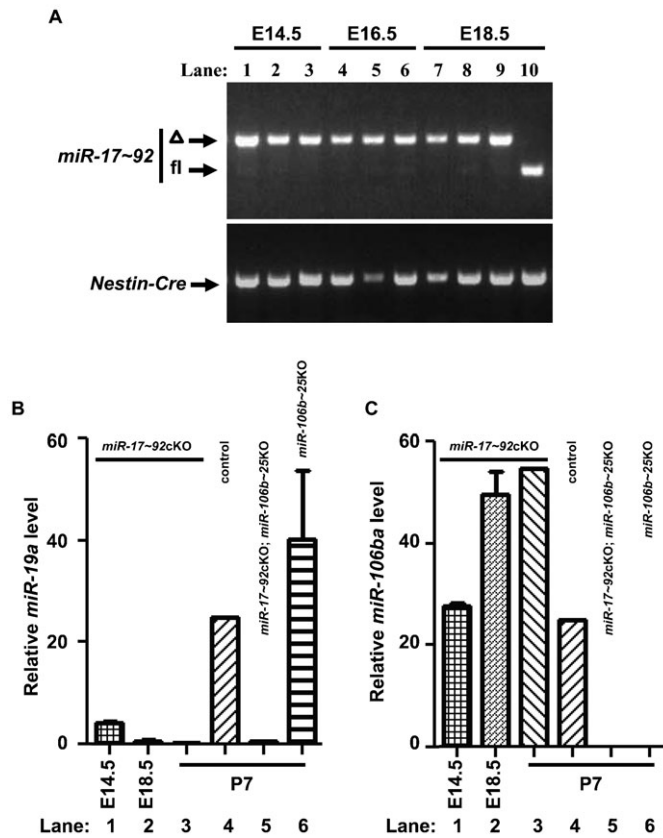


## Supplementary Material

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**Fig. S1. Relative expression of *miR-19a* and *miR-106b* in *miR-17~92cKO; miR-106b~25KO* mice.** (A) PCR for *miR-17~92* and *Nestin-Cre* was performed on DNA extracted from total cerebellum (lanes 1–9) or tail (lane 10) from *miR-17~92cKO* mice (lanes 1–10) at E14.5 (lanes 1–3), E16.5 (lanes 4–6) and E18.5 (lanes 7–10). (fl) floxed and (Δ) deleted *miR-17~92* alleles. Relative levels of mature microRNA *miR-19a* (B) and *miR-106b* (C) were determined by Q-RT-PCR on total RNAs extracted from total cerebella (lanes 1, 2, 5, 6) or purified granule cell precursors (lanes 3, 4). Lanes 1, 2 and 3 correspond to *miR-17~92cKO* at E14.5, E18.5 and P7, respectively, lane 4 to *miR-17~92cKO; miR-106b~25KO* at P7, lane 5 to *miR-17~92cKO*; *miR-106b~25KO* at P7 and lane 6 to *miR-106b~25KO* at P7.