

Appendix figures for Marinaro et al., MicroRNA-independent functions of DGCR8 are essential for neocortical development and TBR1 expression

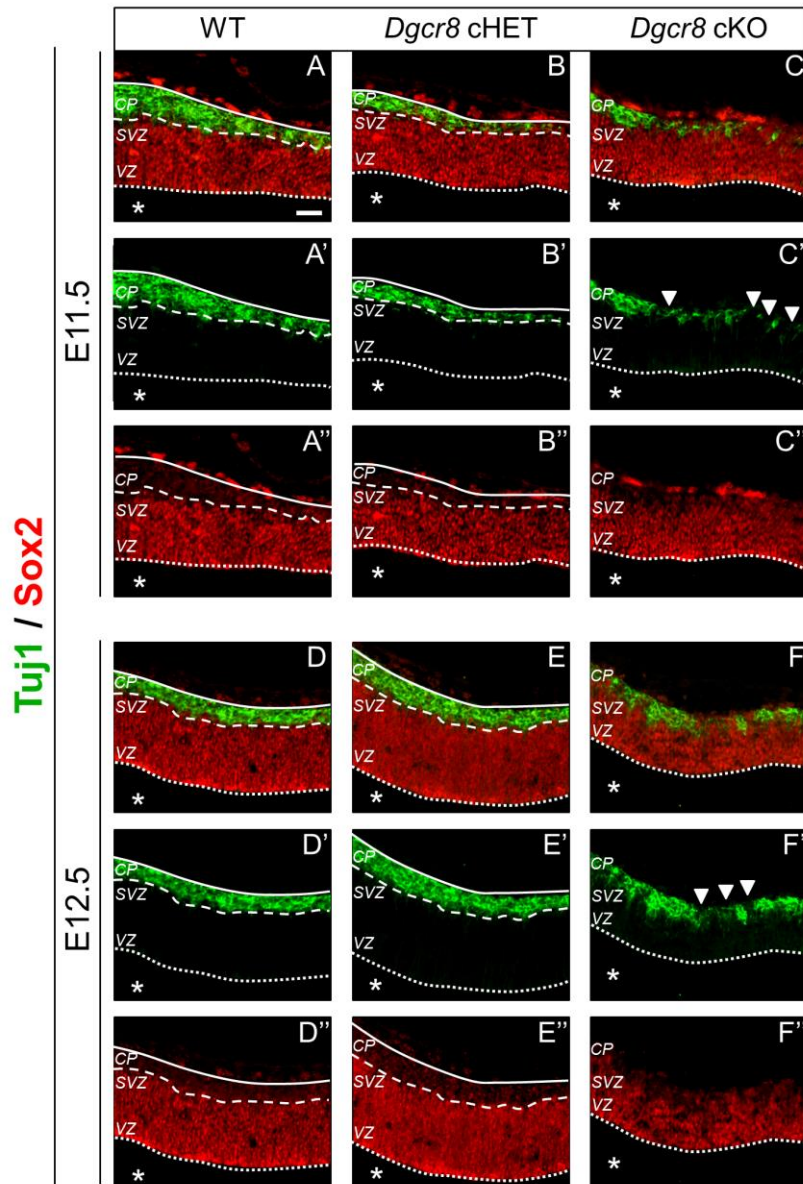
Table of contents

Page 2 **Appendix Figure S1**

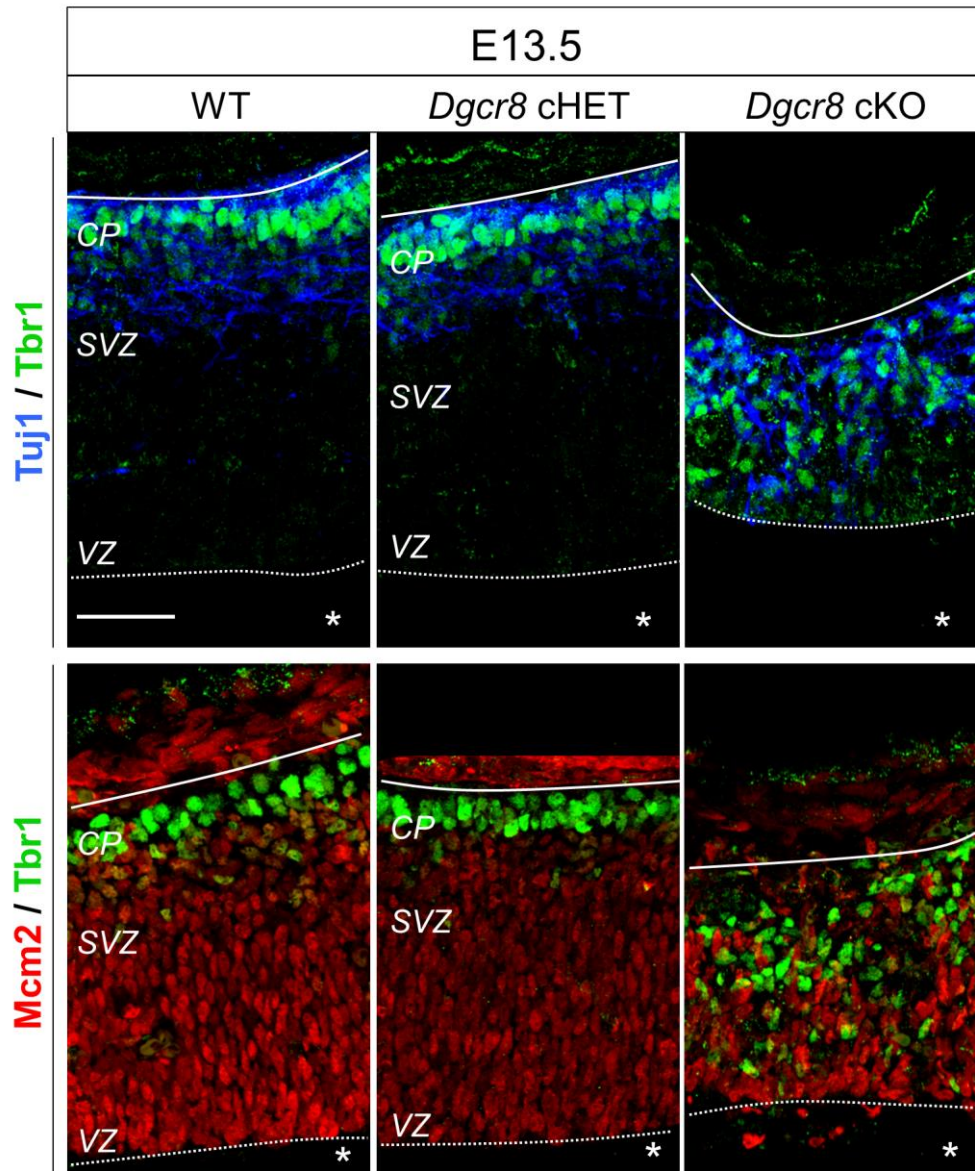
Page 3 **Appendix Figure S2**

Page 4 **Appendix Figure S3**

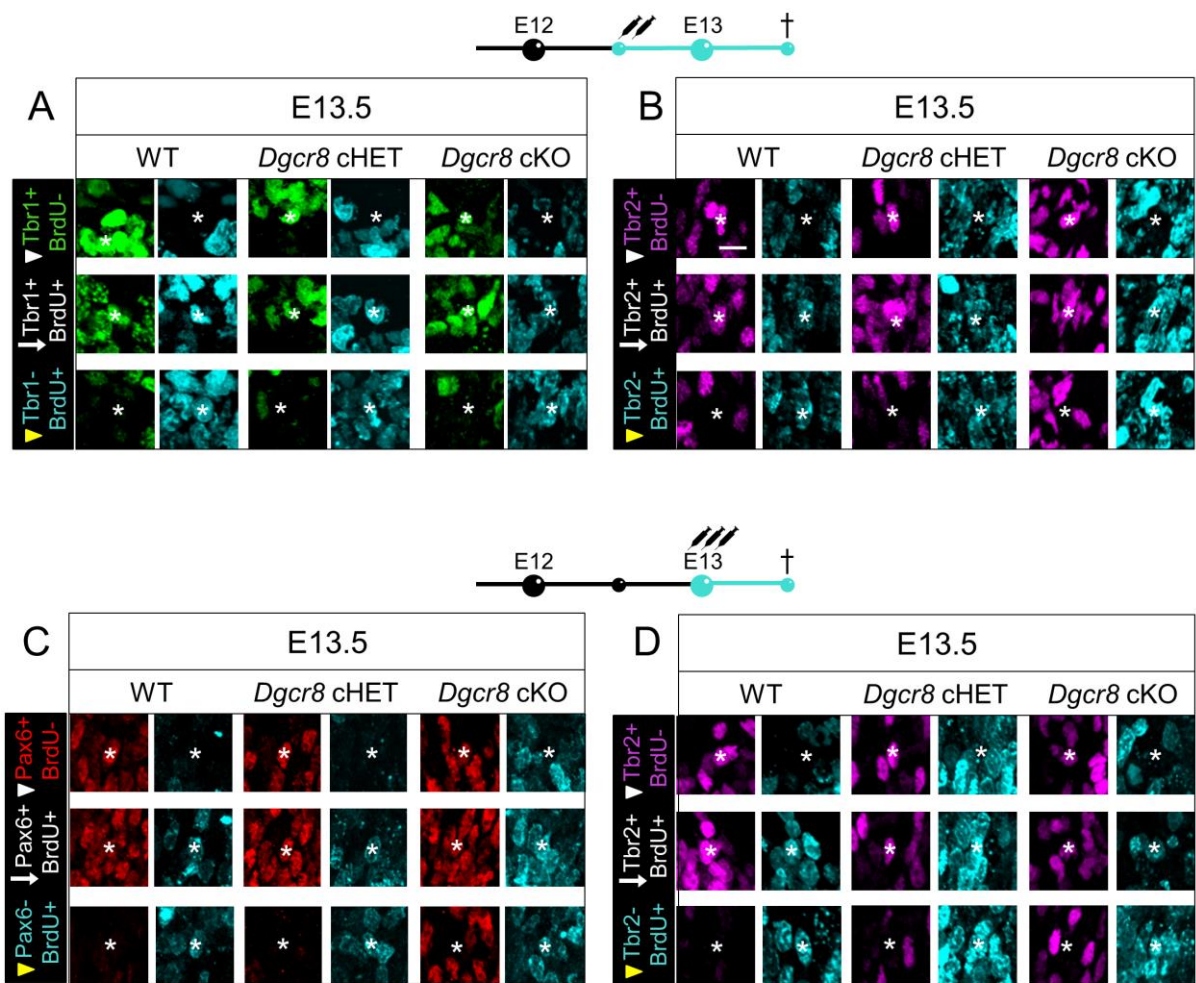
Page 5 **Appendix Figure S4**



Appendix Figure S1. Altered cortical lamination in the dorsal telencephalon of *Dgcr8* cKO embryos from E11.5. Tuji1 and Sox2 immunofluorescence microscopy images of coronal cryosections through the dorsal telencephalon of E11.5 (A-C''), E12.5 (D-F'') of WT (A-A'', D-D''), *Dgcr8* cHET (B-B'', E-E'') and *Dgcr8* cKO (C-C'', F-F'') embryos. Note at E11.5 the discontinuous staining of Tuji1 (arrowheads) in *Dgcr8* cKO, compared to WT and *Dgcr8* cHET cortices. At E12.5 note the intermixing of Tuji1 and Sox2 staining in *Dgcr8* cKO compared to WT and *Dgcr8* cHET cortices. Scale bar: 50 μ m. Asterisks indicate the ventricular lumen. Solid and dashed lines indicate cortex boundaries. Cortical plate (CP); subventricular zone (SVZ); ventricular zone (VZ).

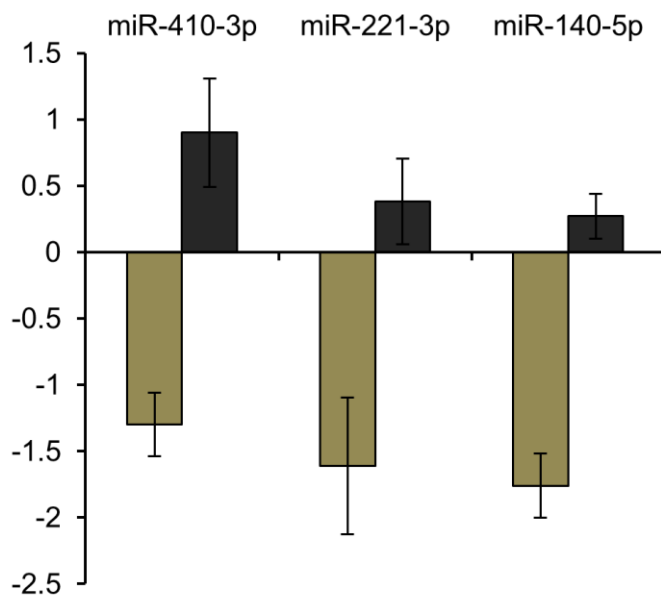
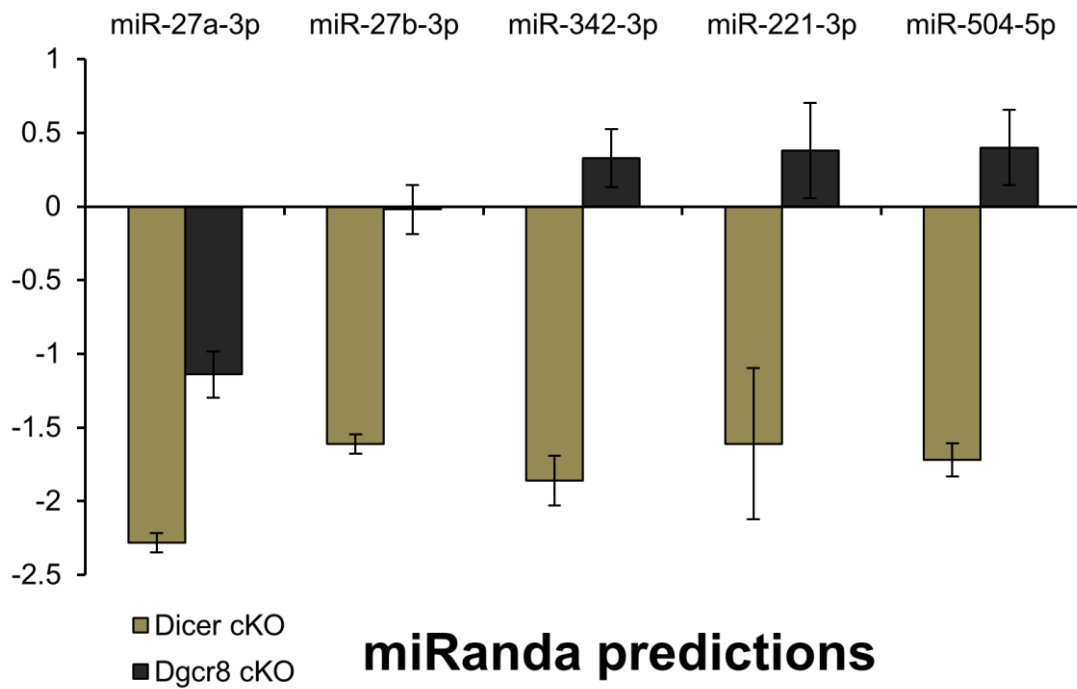


Appendix Figure S2. Analysis of Tbr1⁺ postmitotic neurons in E13.5 neocortices. Tuj1/Tbr1 (upper panel) and Mcm2/Tbr1 immunofluorescence microscopy images of coronal cryosections through the dorsal telencephalon of E13.5 WT, *Dgcr8* cHET and *Dgcr8* cKO embryos.



Appendix Figure S3. Categories of cells counted in Figures 3 and 4. Littermate mouse embryos were subjected to BrdU injection and sacrificed as in Figure 3 (A, B) or as Figure 4 (C, D), respectively. **A)** White arrowheads indicate Tbr1+/BrdU- nuclei; arrows indicate Tbr1+/BrdU+ nuclei; yellow arrowheads indicate Tbr1-/BrdU+ nuclei. **B, D)** White arrowheads indicate Tbr2+/BrdU- nuclei; arrows indicate Tbr2+/BrdU+ nuclei; yellow arrowheads indicate Tbr2-/BrdU+ nuclei. **C)** White arrowheads indicate Pax6+/BrdU- nuclei; arrows indicate Pax6+/BrdU+ nuclei; yellow arrowheads indicate Pax6-/BrdU+ nuclei.

TargetScan predictions



Appendix Figure S4. Read counts of miRNAs predicted to target *Tbr1* in E13.5 Dicer cKO and Dgcr8 cKO neocortices.