

Supplementary Material for Kuerbitz et al. (cercor-bhaa323)

Supplementary Figure 1. Tshz1 gene dosage alters OB interneuron numbers but not striatal size in Sp8 misexpressing mice. A-D, Sp8 misexpressing mice show a decrease in Foxp2⁺ GL cells (C) as compared to controls (A). Tshz1 heterozygosity in Sp8 misexpressing mice significantly increases Foxp2⁺ GL cell numbers (D) but not to control levels (A). E,
Quantification of Foxp2⁺ GL cell densities in wild type and Tshz1 hets with or without Sp8 misexpression. F-I, Sp8 misexpressing mice show a decrease in Meis2⁺ GL cells (H) as compared to controls (F). Tshz1 heterozygosity in Sp8 misexpressing mice did not significantly increase Meis2⁺ GL cell numbers (I). J, Quantification of Meis2⁺ GL cell densities in wild type and Tshz1 heterosygosity in sp8 misexpressing mice did not significantly increase Meis2⁺ GL cell numbers (I). J, Quantification of Meis2⁺ GL cell densities in wild type and Tshz1 heterosygosity in sp8 misexpressing mice show a decrease in wild type and Tshz1 heterosygosity in Sp8 misexpressing mice did not significantly increase Meis2⁺ GL cell numbers (I). J, Quantification of Meis2⁺ GL cell densities in wild type and Tshz1 heterosygosity in Sp8 misexpressing mice show a decrease in wild type and Tshz1 heterosygosity in Sp8 misexpressing mice show a decrease in wild type and Tshz1 heterosygosity in Sp8 misexpressing mice did not significantly increase Meis2⁺ GL cell numbers (I). J, Quantification of Meis2⁺ GL cell densities in wild type and Tshz1 heterosygosity in Sp8 misexpressing mice show a decrease in striatal area as marked by Foxp2 and Meis2 (M) as compared to controls (K). Tshz1

heterozygosity in *Sp8* misexpressing mice did not significantly alter striatal area (**N**). **O**, Quantification of striatal area in wild type and *Tshz1* hets with or without *Sp8* misexpression. Data are represented as mean \pm SEM and stats were performed using a two-way ANOVA with a post hoc Tukey HSD, *p<0.05, **p<0.01, ***p<0.001. Ctx, cortex; GCL, granule cell layer; GL glomerular layer; Str, striatum. Scale bars: A-D, F-I, 100 µm; K-N, 500 µm.



Supplementary Figure 2. *Sp8 misexpressing mice exhibit normal numbers of Meis2*⁺ *GCL interneurons*. **A-C**, Immunofluorescence (**A,B**) and quantification (**C**) of Meis2⁺ neurons in the GCL of the postnatal olfactory bulb of *Sp8* misexpressing mice which showed no significant effect, n=4 for each group ($t_{(6)}=0.12$. p=0.91). Data are represented as mean ± SEM and stats performed using a Student's t-test. GCL, granule cell layer; GL glomerular layer. Scale bars: A,B, 100 µm.



Supplementary Figure 3. Reduced PV^+ interneurons in the EPL of Sp8 misexpressing mice. A-C, Immunofluorescence (A,B) and quantification (C) of PV^+ neurons in the EPL of the postnatal olfactory bulb of *Sp8* misexpressing mice which showed a 57% reduction of PV^+ EPL cells. DAPI was used as a counterstain. Data are represented as mean ± SEM and stats performed using a Student's t-test, **p<0.01. GCL, granule cell layer; GL glomerular layer. Scale bars: A,B, 100 µm.



Supplementary Figure 4. $Foxp2^+$ cells in the SVZ and striatum of Sp8 misexpressing mice. Immunofluorescence (A,B) and quantification of Foxp2⁺ neurons in the SVZ (C) and striatum (D) of *Sp8* misexpressing embryos which showed a 48.4% reduction of Foxp2⁺ SPNs and a trend towards an increase in the SVZ in the overexpressing animals. DAPI was used as a counterstain. Data are represented as mean ± SEM and stats performed using a Student's t-test, **p<0.01. Str, striatum. Scale bars: A,B, 200 µm.