

## Hyperoxygenation during mid-neurogenesis accelerates cortical development in the fetal mouse brain

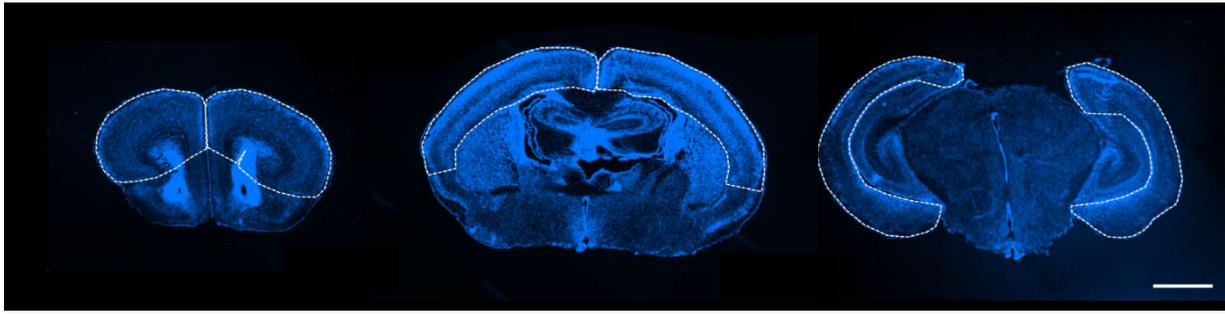
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### Supplementary Figures:

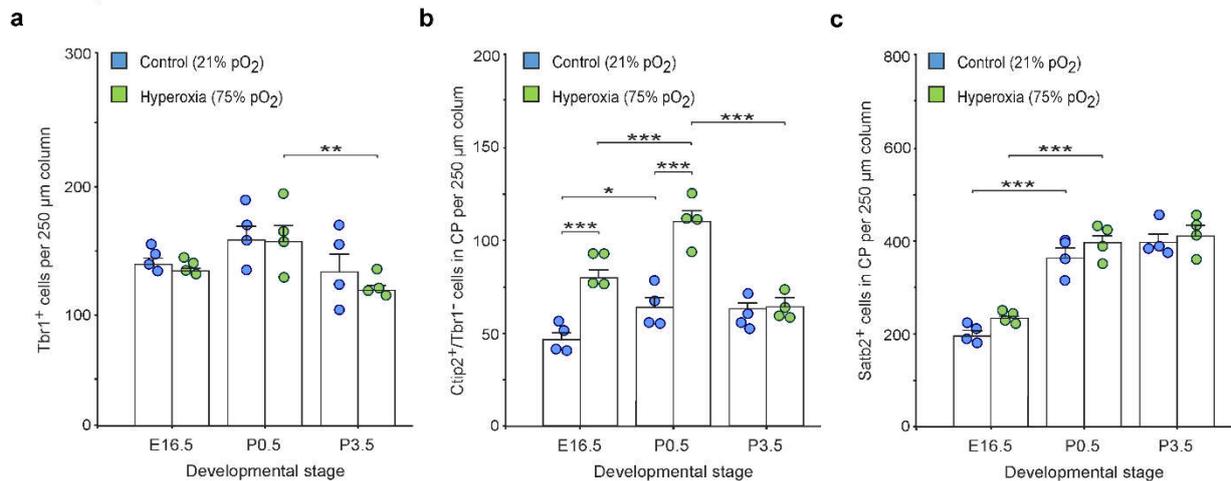
- **Supplementary Figure S1.** Example for the determination of the volume of the cortical plate.
- **Supplementary Figure S2.** Effects of maternal hyperoxygenation on the absolute number of layer specific neurons.
- **Supplementary Figure S3.** Effects of maternal hyperoxygenation on the distribution of microglia in a P16.5 and P3.5 mouse cortex.
- **Supplementary Figure S4.** Effects of hyperoxygenation on the total number of microglia within the developing cortex.
- **Supplementary Figure S5.** Iba1<sup>+</sup> cells are able to target Satb2<sup>+</sup> cells.

### Supplementary Tables:

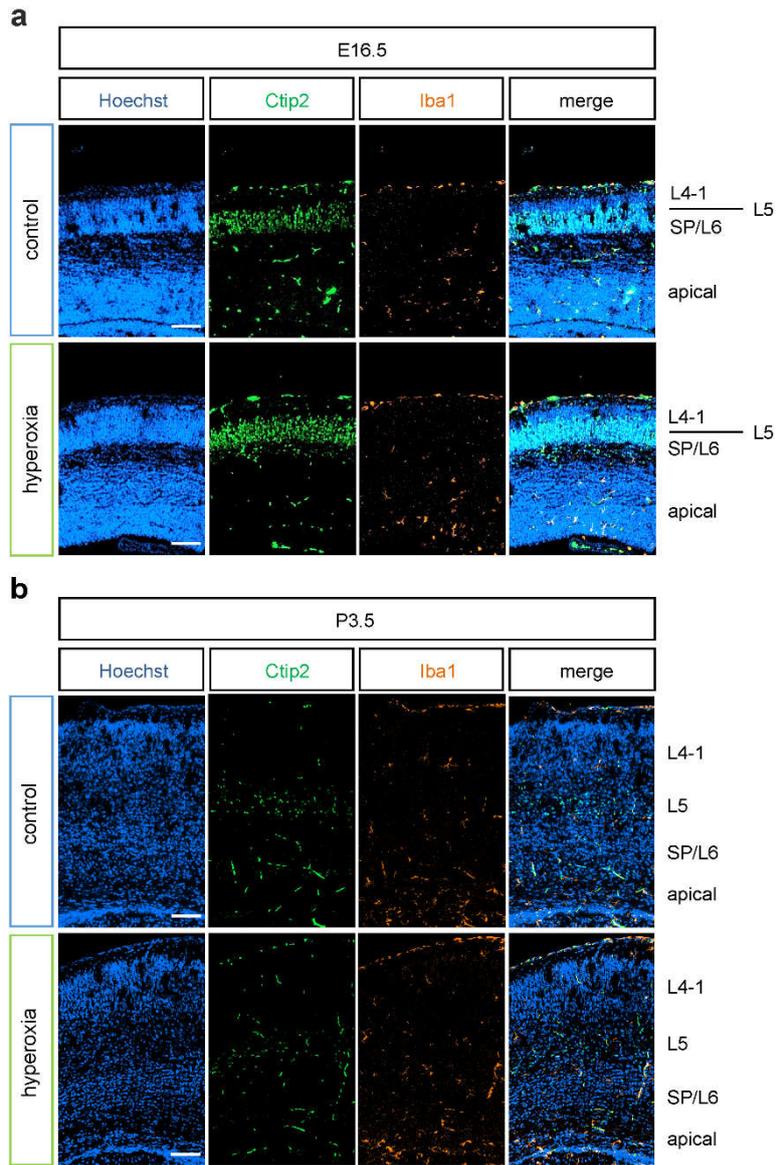
- **Supplementary Table S1.** Statistics determined for NeuN<sup>+</sup> cortical neurons.
- **Supplementary Table S2.** Statistics determined for Tbr1<sup>+</sup> cortical neurons.
- **Supplementary Table S3.** Statistics determined for Ctip<sup>+</sup>/Tbr1<sup>-</sup> neurons.
- **Supplementary Table S4.** Statistics determined for Satb2<sup>+</sup> cortical neurons.
- **Supplementary Table S5.** Statistics determined for apical Iba1<sup>+</sup> cells.
- **Supplementary Table S6.** Statistics determined for subplate/layer 6 (SP/L6) Iba1<sup>+</sup> cells.
- **Supplementary Table S7.** Statistics determined for layer 5 (L5) Iba1<sup>+</sup> cells
- **Supplementary Table S8.** Statistics determined for layer 4-1 (L4-1) Iba1<sup>+</sup> cells.
- **Supplementary Table S9.** Statistics determined for absolute CC3<sup>+</sup> cell counts.
- **Supplementary Table S10.** Statistics determined for vGluT2<sup>+</sup> synapses in L5.
- **Supplementary Table S11.** Statistics determined for absolute Tbr1<sup>+</sup> neuron counts.
- **Supplementary Table S12.** Statistics determined for absolute Ctip<sup>+</sup>/Tbr1<sup>-</sup> neuron counts.
- **Supplementary Table S13.** Statistics determined for absolute Satb2<sup>+</sup> neuron counts.
- **Supplementary Table S14.** Statistics determined for total Iba1<sup>+</sup> cell counts.



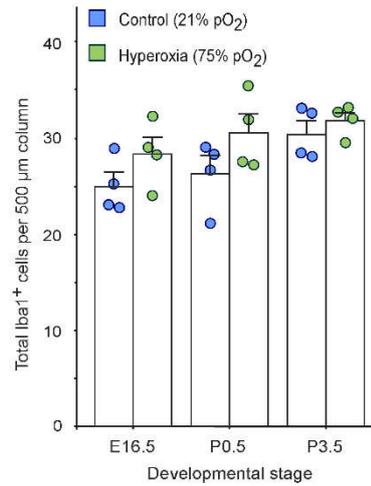
**Supplementary Figure S1.** Example for the determination of the volume of the cortical plate (CP). Every 6<sup>th</sup> Hoechst stained slice of a mouse brain was outlined as shown in the figure (left to right: rostral, middle and caudal section) and used to calculate the volume. Scale bar, 1000  $\mu\text{m}$ .



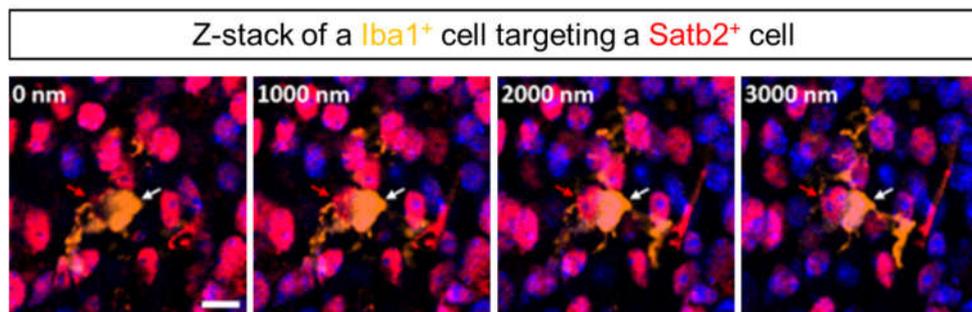
**Supplementary Figure S2.** Effects of maternal hyperoxygenation on the absolute number of layer specific neurons. Quantification of absolute Tbr1<sup>+</sup>, Ctip2<sup>+</sup>/Tbr1<sup>-</sup> and Satb2<sup>+</sup> cells within 250  $\mu\text{m}$  wide cortical columns of E16.5, P0.5 and P3.5 mice. Data are means  $\pm$  s.e.m. (n = 4). \*  $p < 0.05$ , \*\*  $p < 0.01$ , \*\*\*  $p < 0.001$  from two-way ANOVA with *post-hoc* two-sided t-test with Bonferroni correction. For full statistics, see **Supplementary Tables S11-S13**.



**Supplementary Figure S3:** Effects of fetal brain hyperoxygenation on the distribution of microglia in E16.5 and P3.5 mouse cortex. Representative fluorescent images of Iba1<sup>+</sup> cells (orange) from (a) E16.5 and (b) P3.5 in the middle cortical sections along the rostro-caudal axis—of hyperoxia treated and control mice showed no differences. Ctip2<sup>+</sup> (green) was used for layer determination and Hoechst (blue) was used to stain cell nuclei. Scale bars represent 100  $\mu$ m.



**Supplementary Figure S4.** Effects of hyperoxygenation on the total number of microglia within the developing cortex. Quantification of the total number of Iba1<sup>+</sup> microglia showed no differences with respect to hyperoxia treatment. Data are means±s.e.m. (n = 4). \*  $p < 0.05$ , \*\*  $p < 0.01$ , \*\*\*  $p < 0.001$  from two-way ANOVA with *post-hoc* two-sided t-test with Bonferroni correction. For full statistics, see **Supplementary Table S14**.



**Supplementary Figure S5.** Iba1<sup>+</sup> cells are able to target Satb2<sup>+</sup> cells. Representative z-stack images of a microglia cell (white arrow) targeting Satb2<sup>+</sup> cells (red arrow) in a P0.5 mouse cortex. Scale bars represent 10 μm.

## Supplementary Tables

**Supplementary Table S1.** Statistics determined for NeuN<sup>+</sup> cortical neurons in various development stages (E16.5, P0.5, P3.5) after different oxygen exposures during mid-neurogenesis (**Figure 1e**). Two-way ANOVA with *post-hoc* t-test and Bonferroni adjustment with atmospheric oxygen concentrations and development stage as fixed factors revealed that atmospheric oxygen concentration and developmental stage have a significant interaction effect on NeuN<sup>+</sup> neuron counts ( $p=0.028$ , F-value=4.2) and significant differences among atmospheric oxygen concentrations ( $p=0.006$ , F-value=8.8) and developmental stages ( $p<0.001$ , F-value=9.3). Displayed are Bonferroni-adjusted *P*-values (E16.5: n=4 [control], n=3 [hyperoxia]; P0.5: n=8 [control], n=6 [hyperoxia]; P3.5: n=4 [control], n=6 [hyperoxia]). **(A)** Significances among the different atmospheric oxygen concentrations. **(B)** Significances among the different developmental stages. Bold values indicate significant differences.

**A**

	<b>E16.5</b>	<b>P0.5</b>	<b>P3.5</b>
<b>Normoxia (21% O<sub>2</sub>) vs. Hyperoxia (75% O<sub>2</sub>)</b>	<b>0.005</b>	<b>0.013</b>	0.581

**B**

	<b>Normoxia (21% O<sub>2</sub>)</b>	<b>Hyperoxia (75% O<sub>2</sub>)</b>
<b>E16.5 vs. P0.5</b>	<b>0.0459</b>	1
<b>E16.5 vs. P3.5</b>	<b>&lt; 0.001</b>	0,769
<b>P0.5 vs. P3.5</b>	<b>0.011</b>	1

**Supplementary Table S2.** Statistics determined for Tbr1<sup>+</sup> cortical neurons in various development stages (E16.5, P0.5, P3.5) after different oxygen exposures during mid-neurogenesis (**Figure 2b**). Two-way ANOVA with *post-hoc* t-test and Bonferroni adjustment with atmospheric oxygen concentrations and development stage as fixed factors revealed that atmospheric oxygen concentration and developmental stage have no significant interaction effect on Tbr1<sup>+</sup> neuron counts ( $p=0.210$ , F-value=1.7), but significant differences among atmospheric oxygen concentrations ( $p=0.006$ , F-value=9.8) and developmental stages ( $p<0.001$ , F-value=84.3). Displayed are Bonferroni-adjusted *P*-values ( $n = 4$ ). **(A)** Significances among the different atmospheric oxygen concentrations. **(B)** Significances among the different developmental stages. Bold values indicate significant differences.

**A**

	<b>E16.5</b>	<b>P0.5</b>	<b>P3.5</b>
<b>Normoxia (21% O<sub>2</sub>) vs. Hyperoxia (75% O<sub>2</sub>)</b>	<b>0.004</b>	0.333	0.278

**B**

	<b>Normoxia (21% O<sub>2</sub>)</b>	<b>Hyperoxia (75% O<sub>2</sub>)</b>
<b>E16.5 vs. P0.5</b>	<b>&lt; 0.001</b>	<b>0.002</b>
<b>E16.5 vs. P3.5</b>	<b>&lt; 0.001</b>	<b>&lt; 0.001</b>
<b>P0.5 vs. P3.5</b>	<b>0.005</b>	0.004

**Supplementary Table S3.** Statistics determined for Ctip<sup>+</sup>/Tbr1<sup>-</sup> cortical neurons in various development stages (E16.5, P0.5, P3.5) after different oxygen exposures during mid-neurogenesis (**Figure 2c**). Two-way ANOVA with *post-hoc* t-test and Bonferroni adjustment with atmospheric oxygen concentrations and development stage as fixed factors revealed that atmospheric oxygen concentration and developmental stage have a significant interaction effect on Ctip<sup>+</sup>/Tbr1<sup>-</sup> cortical neuron counts ( $p=0.002$ , F-value=9.5) and significant differences among atmospheric oxygen concentrations ( $p<0.001$ , F-value=49.2) and developmental stages ( $p=0.002$ , F-value=44.9). Displayed are Bonferroni-adjusted *P*-values ( $n = 4$ ). **(A)** Significances among the different atmospheric oxygen concentrations. **(B)** Significances among the different developmental stages. Bold values indicate significant differences.

**A**

	<b>E16.5</b>	<b>P0.5</b>	<b>P3.5</b>
<b>Normoxia (21% O<sub>2</sub>) vs. Hyperoxia (75% O<sub>2</sub>)</b>	<b>&lt; 0.001</b>	<b>&lt; 0.001</b>	0.629

**B**

	<b>Normoxia (21% O<sub>2</sub>)</b>	<b>Hyperoxia (75% O<sub>2</sub>)</b>
<b>E16.5 vs. P0.5</b>	0.135	0.092
<b>E16.5 vs. P3.5</b>	<b>0.003</b>	<b>&lt; 0.001</b>
<b>P0.5 vs. P3.5</b>	0.309	<b>&lt; 0.001</b>

**Supplementary Table S4.** Statistics determined for Satb2<sup>+</sup> cortical neurons in various development stages (E16.5, P0.5, P3.5) after different oxygen exposures during mid-neurogenesis (**Figure 2d**). Two-way ANOVA with *post-hoc* t-test and Bonferroni adjustment (n = 4) with atmospheric oxygen concentrations and development stage as fixed factors revealed that atmospheric oxygen concentration and developmental stage have no significant interaction effect on Satb2<sup>+</sup> neuron counts ( $p=0.922$ , F-value=0.1), no significant differences among atmospheric oxygen concentrations ( $p=0.922$ , F-value=3.6), but significant differences among developmental stages ( $p=0.048$ , F-value=3.6). Displayed are Bonferroni-adjusted *P*-values (n = 4). **(A)** Significances among the different atmospheric oxygen concentrations. **(B)** Significances among the different developmental stages. Bold values indicate significant differences.

**A**

	<b>E16.5</b>	<b>P0.5</b>	<b>P3.5</b>
<b>Normoxia (21% O<sub>2</sub>) vs. Hyperoxia (75% O<sub>2</sub>)</b>	0.411	0.748	0.437

**B**

	<b>Normoxia (21% O<sub>2</sub>)</b>	<b>Hyperoxia (75% O<sub>2</sub>)</b>
<b>E16.5 vs. P0.5</b>	0.141	0.369
<b>E16.5 vs. P3.5</b>	0.720	0.774
<b>P0.5 vs. P3.5</b>	1	1

**Supplementary Table S5.** Statistics determined for apical Iba1<sup>+</sup> cells in various development stages (E16.5, P0.5, P3.5) after different oxygen exposures during mid-neurogenesis (**Figure 4b**). Two-way ANOVA with *post-hoc* t-test and Bonferroni adjustment with atmospheric oxygen concentrations and development stage as fixed factors revealed that atmospheric oxygen concentration and developmental stage have no significant interaction effect on apical Iba1<sup>+</sup> cell counts ( $p=0.465$ , F-value=0.8) and no significant differences among atmospheric oxygen concentrations ( $p=0.363$ , F-value=0.9), but significant differences among developmental stages ( $p<0.001$ , F-value=20.9). Displayed are Bonferroni-adjusted *P*-values ( $n = 4$ ). **(A)** Significances among the different atmospheric oxygen concentrations. **(B)** Significances among the different developmental stages. Bold values indicate significant differences.

**A**

	<b>E16.5</b>	<b>P0.5</b>	<b>P3.5</b>
<b>Normoxia (21% O<sub>2</sub>) vs. Hyperoxia (75% O<sub>2</sub>)</b>	0.146	0.819	0.744

**B**

	<b>Normoxia (21% O<sub>2</sub>)</b>	<b>Hyperoxia (75% O<sub>2</sub>)</b>
<b>E16.5 vs. P0.5</b>	0.605	<b>0.020</b>
<b>E16.5 vs. P3.5</b>	<b>0.003</b>	<b>&lt; 0.001</b>
<b>P0.5 vs. P3.5</b>	<b>0.049</b>	0.064

**Supplementary Table S6.** Statistics determined for subplate/layer 6 (SP/L6) Iba1<sup>+</sup> cells in various development stages (E16.5, P0.5, P3.5) after different oxygen exposures during mid-neurogenesis (**Figure 4c**). Two-way ANOVA with *post-hoc* t-test and Bonferroni adjustment with atmospheric oxygen concentrations and development stage as fixed factors revealed that atmospheric oxygen concentration and developmental stage have no significant interaction effect on SP/L6 Iba1<sup>+</sup> cells counts ( $p=0.295$ , F-value=1.3) and no significant differences among atmospheric oxygen concentrations ( $p=0.203$ , F-value=1.7), but significant differences among developmental stages ( $p<0.001$ , F-value=38.6). Displayed are Bonferroni-adjusted *P*-values ( $n = 4$ ). **(A)** Significances among the different atmospheric oxygen concentrations. **(B)** Significances among the different developmental stages. Bold values indicate significant differences.

**A**

	<b>E16.5</b>	<b>P0.5</b>	<b>P3.5</b>
<b>Normoxia (21% O<sub>2</sub>) vs. Hyperoxia (75% O<sub>2</sub>)</b>	0.942	0.054	0.772

**B**

	<b>Normoxia (21% O<sub>2</sub>)</b>	<b>Hyperoxia (75% O<sub>2</sub>)</b>
<b>E16.5 vs. P0.5</b>	0.096	<b>&lt; 0.001</b>
<b>E16.5 vs. P3.5</b>	<b>&lt; 0.001</b>	<b>&lt; 0.001</b>
<b>P0.5 vs. P3.5</b>	<b>0.005</b>	0.211

**Supplementary Table S7.** Statistics determined for layer 5 (L5) Iba1<sup>+</sup> cells in various development stages (P0.5, P3.5) after different oxygen exposures during mid-neurogenesis (**Figure 4d**). Two-way ANOVA with *post-hoc* t-test and Bonferroni adjustment with atmospheric oxygen concentrations and development stage as fixed factors revealed that atmospheric oxygen concentration and developmental stage have a significant interaction effect on L5 Iba1<sup>+</sup> cells counts ( $p=0.014$ , F-value=8.1) and significant differences among atmospheric oxygen concentrations ( $p=0.003$ , F-value=13.2) and developmental stages ( $p<0.001$ , F-value=37.5). Displayed are Bonferroni-adjusted *P*-values (n = 4). **(A)** Significances among the different atmospheric oxygen concentrations. **(B)** Significances among the different developmental stages. Bold values indicate significant differences.

**A**

	<b>P0.5</b>	<b>P3.5</b>
<b>Normoxia (21% O<sub>2</sub>) vs. Hyperoxia (75% O<sub>2</sub>)</b>	<b>&lt; 0.001</b>	0.588

**B**

	<b>Normoxia (21% O<sub>2</sub>)</b>	<b>Hyperoxia (75% O<sub>2</sub>)</b>
<b>P0.5 vs. P3.5</b>	<b>&lt; 0.001</b>	<b>0.039</b>

**Supplementary Table S8.** Statistics determined for layer 4-1 (L4-1) Iba1<sup>+</sup> cells in various development stages (P0.5, P3.5) after different oxygen exposures during mid-neurogenesis (**Figure 4e**). Two-way ANOVA with *post-hoc* t-test and Bonferroni adjustment with atmospheric oxygen concentrations and development stage as fixed factors revealed that atmospheric oxygen concentration and developmental stage have no significant interaction effect on L4-1 Iba1<sup>+</sup> cells counts ( $p=0.945$ , F-value=0.0) and no significant differences among atmospheric oxygen concentrations ( $p=0.945$ , F-value=0.0), but significant differences among developmental stages ( $p<0.001$ , F-value=85.8). Displayed are Bonferroni-adjusted *P*-values (n = 4). **(A)** Significances among the different atmospheric oxygen concentrations. **(B)** Significances among the different developmental stages. Bold values indicate significant differences.

**A**

	<b>P0.5</b>	<b>P3.5</b>
<b>Normoxia (21% O<sub>2</sub>) vs. Hyperoxia (75% O<sub>2</sub>)</b>	0.920	1.000

**B**

	<b>Normoxia (21% O<sub>2</sub>)</b>	<b>Hyperoxia (75% O<sub>2</sub>)</b>
<b>P0.5 vs. P3.5</b>	<b>&lt; 0.001</b>	<b>&lt; 0.001</b>

**Supplementary Table S9.** Statistics determined for CC3<sup>+</sup> cell counts in various development stages (E16.5, P0.5, P3.5) after different oxygen exposures during mid-neurogenesis (**Figure 7b**). Robust ANOVA using raov function from Rfit package with *post-hoc* unpaired Wilcoxon-test and Bonferroni adjustment (n = 3) with atmospheric oxygen concentrations and development stage as fixed factors revealed that atmospheric oxygen concentration and developmental stage have a significant interaction effect on CC3<sup>+</sup> cell counts ( $p=0.001$ , F-value=9.1) and significant differences among atmospheric oxygen concentrations ( $p=0.004$ , F-value=10.4) and developmental stages ( $p<0.001$ , F-value=30.1). **(A)** Significances among the different atmospheric oxygen concentrations. **(B)** Significances among the different developmental stages. Bold values indicate significant differences.

**A**

	<b>E16.5</b>	<b>P0.5</b>	<b>P3.5</b>
<b>Normoxia (21% O<sub>2</sub>) vs. Hyperoxia (75% O<sub>2</sub>)</b>	0.564	<b>0.008</b>	1.000

**B**

	<b>Normoxia (21% O<sub>2</sub>)</b>	<b>Hyperoxia (75% O<sub>2</sub>)</b>
<b>E16.5 vs. P0.5</b>	<b>0.014</b>	<b>0.024</b>
<b>E16.5 vs. P3.5</b>	0.075	0.107
<b>P0.5 vs. P3.5</b>	0.276	0.786

**Supplementary Table S10:** Statistics determined for vGluT2<sup>+</sup> synapses in L5 (P0.5, P3.5) after different oxygen exposures during mid-neurogenesis (**Figure 8**). Two-way ANOVA with *post-hoc* t-test with atmospheric oxygen concentrations and developmental stage as fixed factors revealed that atmospheric oxygen concentration and developmental stage have a significant interaction effect on VGlut2<sup>+</sup> synapses ( $p=0.046$ , F-value=4.7) and significant differences among atmospheric oxygen concentrations ( $p=0.030$ , F-value=5.6), but no significant differences among developmental stages ( $p=0.371$ , F-value=0.8). Displayed are *P*-values ( $n = 5$ ). **(A)** Significances among the different atmospheric oxygen concentrations. **(B)** Significances among the different developmental stages. Bold values indicate significant differences.

**A**

	<b>P0.5</b>	<b>P3.5</b>
<b>Normoxia (21% O<sub>2</sub>) vs. Hyperoxia (75% O<sub>2</sub>)</b>	<b>0.006</b>	0.881

**B**

	<b>Normoxia (21% O<sub>2</sub>)</b>	<b>Hyperoxia (75% O<sub>2</sub>)</b>
<b>P0.5 vs. P3.5</b>	0.394	<b>0.045</b>

**Supplementary Table S11.** Statistics determined for absolute Tbr1<sup>+</sup> cortical neuron counts in various development stages (E16.5, P0.5, P3.5) after different oxygen exposures during mid-neurogenesis (**Supplementary Figure S2a**). Two-way ANOVA with *post-hoc* t-test and Bonferroni adjustment with atmospheric oxygen concentrations and development stage as fixed factors revealed that atmospheric oxygen concentration and developmental stage have no significant interaction effect on Tbr1<sup>+</sup> neuron counts ( $p=0.780$ , F-value=0.3) and no significant differences among atmospheric oxygen concentrations ( $p=0.338$ , F-value=1.0), but significant differences among developmental stages ( $p=0.014$ , F-value=5.5). Displayed are Bonferroni-adjusted *P*-values (n = 4). **(A)** Significances among the different atmospheric oxygen concentrations. **(B)** Significances among the different developmental stages. Bold values indicate significant differences.

**A**

	<b>E16.5</b>	<b>P0.5</b>	<b>P3.5</b>
<b>Normoxia (21% O<sub>2</sub>) vs. Hyperoxia (75% O<sub>2</sub>)</b>	0.654	0.895	0.278

**B**

	<b>Normoxia (21% O<sub>2</sub>)</b>	<b>Hyperoxia (75% O<sub>2</sub>)</b>
<b>E16.5 vs. P0.5</b>	0.567	0.327
<b>E16.5 vs. P3.5</b>	1.000	0.852
<b>P0.5 vs. P3.5</b>	0.261	<b>0.036</b>

**Supplementary Table S12.** Statistics determined for absolute Ctip2<sup>+</sup>/Tbr1<sup>-</sup> neuron counts in various development stages (E16.5, P0.5, P3.5) after different oxygen exposures during mid-neurogenesis (**Supplementary Figure S2b**). Two-way ANOVA with *post-hoc* t-test and Bonferroni adjustment with atmospheric oxygen concentrations and development stage as fixed factors revealed that atmospheric oxygen concentration and developmental stage have a significant interaction effect on Ctip2<sup>+</sup>/Tbr1<sup>-</sup> neuron counts ( $p < 0.001$ , F-value=10.8) and significant differences among atmospheric oxygen concentrations ( $p < 0.001$ , F-value=53.2) and developmental stages ( $p < 0.001$ , F-value=19.1). Displayed are Bonferroni-adjusted *P*-values ( $n = 4$ ). **(A)** Significances among the different atmospheric oxygen concentrations. **(B)** Significances among the different developmental stages. Bold values indicate significant differences.

**A**

	<b>E16.5</b>	<b>P0.5</b>	<b>P3.5</b>
<b>Normoxia (21% O<sub>2</sub>) vs. Hyperoxia (75% O<sub>2</sub>)</b>	<b>&lt; 0.001</b>	<b>&lt; 0.001</b>	0.574

**B**

	<b>Normoxia (21% O<sub>2</sub>)</b>	<b>Hyperoxia (75% O<sub>2</sub>)</b>
<b>E16.5 vs. P0.5</b>	<b>0.047</b>	<b>&lt; 0.001</b>
<b>E16.5 vs. P3.5</b>	0.208	0.057
<b>P0.5 vs. P3.5</b>	1.000	<b>&lt; 0.001</b>

**Supplementary Table S13.** Statistics determined for absolute Satb2<sup>+</sup> cortical neuron counts in various development stages (E16.5, P0.5, P3.5) after different oxygen exposures during mid-neurogenesis (**Supplementary Figure S2c**). Two-way ANOVA with *post-hoc* t-test and Bonferroni adjustment (n = 4) with atmospheric oxygen concentrations and development stage as fixed factors revealed that atmospheric oxygen concentration and developmental stage have no significant interaction effect on Satb2<sup>+</sup> neuron counts ( $p=0.828$ , F-value=0.2), no significant differences among atmospheric oxygen concentrations ( $p=0.066$ , F-value=3.8), but significant differences among developmental stages ( $p<0.001$ , F-value=76.9). **(A)** Significances among the different atmospheric oxygen concentrations. **(B)** Significances among the different developmental stages. Bold values indicate significant differences.

**A**

	<b>E16.5</b>	<b>P0.5</b>	<b>P3.5</b>
<b>Normoxia (21% O<sub>2</sub>) vs. Hyperoxia (75% O<sub>2</sub>)</b>	0.153	0.225	0.527

**B**

	<b>Normoxia (21% O<sub>2</sub>)</b>	<b>Hyperoxia (75% O<sub>2</sub>)</b>
<b>E16.5 vs. P0.5</b>	<b>&lt; 0.001</b>	<b>&lt; 0.001</b>
<b>E16.5 vs. P3.5</b>	<b>&lt; 0.001</b>	<b>&lt; 0.001</b>
<b>P0.5 vs. P3.5</b>	0.573	1.000

**Supplementary Table S14.** Statistics determined for total Iba1<sup>+</sup> cell counts (E16.5, P0.5, P3.5) after different oxygen exposures during mid-neurogenesis (**Supplementary Figure S4**). Two-way ANOVA with *post-hoc* t-test and Bonferroni adjustment with atmospheric oxygen concentrations and development stage as fixed factors revealed that atmospheric oxygen concentration and developmental stage have no significant interaction effect on total Iba1<sup>+</sup> cell counts ( $p=0.643$ , F-value=0.5), but significant differences among atmospheric oxygen concentrations ( $p=0.030$ , F-value=5.5) and significant differences among developmental stages ( $p=0.034$ , F-value=4.1). Displayed are Bonferroni-adjusted *P*-values (n = 4). Significances among the different developmental stages. Bold values indicate significant differences.

**A**

	<b>E16.5</b>	<b>P0.5</b>	<b>P3.5</b>
<b>Normoxia (21% O<sub>2</sub>) vs. Hyperoxia (75% O<sub>2</sub>)</b>	0.144	0.069	0.541

**B**

	<b>Normoxia (21% O<sub>2</sub>)</b>	<b>Hyperoxia (75% O<sub>2</sub>)</b>
<b>E16.5 vs. P0.5</b>	1.000	0.989
<b>E16.5 vs. P3.5</b>	0.072	0.411
<b>P0.5 vs. P3.5</b>	0.234	1.000