

A

		WT (cells/mm ²)	<i>Maf</i> cDKO (cells/mm ²)
Cortex	<i>Pnoc</i> ⁺	80.17 ± 10.35	173.62 ± 25.37**
	<i>Pnoc-tdTomato</i> ⁺	23.65 ± 3.13	80.88 ± 9.98****
dmCortex	<i>Pnoc</i> ⁺	44.03 ± 8.52	341.84 ± 38.08****
	<i>Pnoc-tdTomato</i> ⁺	19.17 ± 3.59	186.70 ± 20.58****
Hippocampus	<i>Pnoc</i> ⁺	765.51 ± 67.20	1241.42 ± 73.20**
	<i>Pnoc-tdTomato</i> ⁺	295.41 ± 25.44	724.58 ± 47.78****
CA1	<i>Pnoc</i> ⁺	646.06 ± 67.43	939.43 ± 91.05*
	<i>Pnoc-tdTomato</i> ⁺	200.92 ± 42.28	518.50 ± 45.29***
	<i>Pnoc-Htr3a</i> ⁺	44.74 ± 8.74	50.62 ± 5.55
CA3	<i>Pnoc</i> ⁺	671.60 ± 48.86	1327.51 ± 46.62****
	<i>Pnoc-tdTomato</i> ⁺	298.53 ± 19.63	848.62 ± 49.45****
	<i>Pnoc-Htr3a</i> ⁺	55.15 ± 3.19	50.44 ± 7.47
DG	<i>Pnoc</i> ⁺	989.45 ± 85.01	1570.45 ± 111.25**
	<i>Pnoc-tdTomato</i> ⁺	401.58 ± 21.69	881.10 ± 76.07***
	<i>Pnoc-Htr3a</i> ⁺	28.65 ± 7.55	49.90 ± 4.13

B

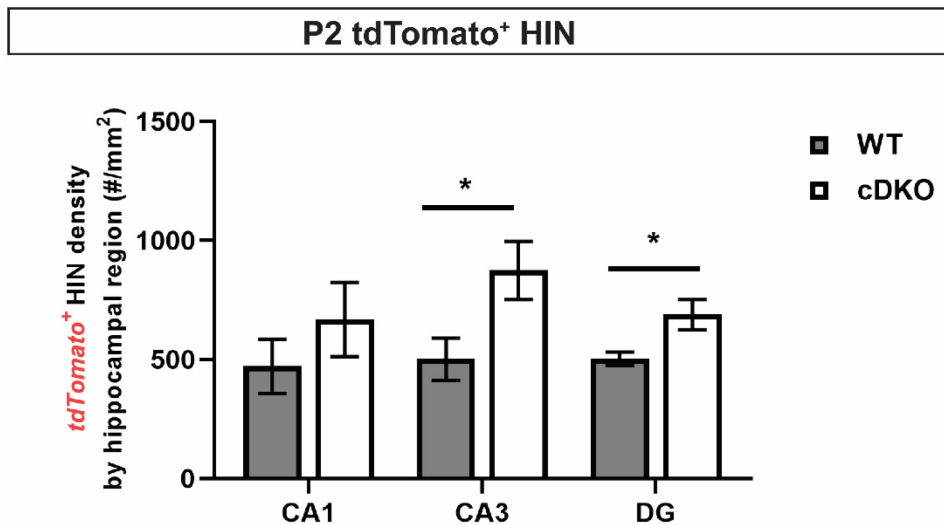


Figure 5 – figure supplement 1. Quantification of *Pnoc*⁺ INs by region and by IN subgroups/ Quantification of tdTomato⁺ HINs by hippocampal region

(A) Quantification of *Pnoc*⁺ INs (cell density ± standard error) in the P2 neocortex and hippocampus by genotypes, and by IN subtypes. MGE origin: *tdTomato*-*Pnoc*⁺ INs; CGE origin: *tdTomato*-*Htr3a*⁺ INs. (B) Quantification of tdTomato⁺ HIN density by hippocampal regions. Unpaired t-test; *p<0.05, **p<0.01, ***p<0.0001, ****p<0.00001