

1 16 Supplementary FIGURE LEGENDS

2 **Supplementary Figure 1:** Diminished PNN organization in the retrosplenial-, secondary visual- and
3 auditory cortex of quadruple knockout mice. **(A-F)** Immunohistochemical staining of PNNs in murine
4 coronal brain slices with WFA (green) and anti-aggrecan (red). Images of WFA-positive and aggrecan-
5 positive PNN-enwrapped neurons were taken and counted in the retrosplenial-, secondary visual- and
6 auditory cortex. **(G)** A significantly reduced number of WFA-positive cells in the retrosplenial-,
7 secondary visual- and auditory cortex of quadruple knockout mice could be noticed ($p < 0.001$, $N = 7$).
8 **(H)** Also, the number of aggrecan-positive cells was significantly reduced in retrosplenial-, secondary
9 visual- and auditory cortex of quadruple Knockout mice ($p < 0.001$, $N = 7$). **(I, J)** The number of WFA-
10 positive and aggrecan-positive processes per PNN were counted in retrosplenial-, secondary visual-,
11 primary visual- and auditory cortex. Both, WFA-positive and aggrecan-positive processes were
12 significantly reduced in the examined cortical areas ($p < 0.001$, $N = 7$); 4xKO = quadruple knockout,
13 Aud = auditory cortex, V1 = primary visual cortex, V2 = secondary visual cortex, RSC = retrosplenial
14 cortex, WFA = Wisteria floribunda agglutinin, WT = wildtype, *** = $p < 0.001$ data are shown as
15 mean \pm SEM and SD, scale bar = 20 μ m.

16 **Supplementary Figure 2:** Inhibitory synaptic elements in the V1 of quadruple knockout mice. **(A)**
17 Western blot analysis of gephyrin protein levels in the V1. **(B)** No significant differences in the
18 gephyrin protein band intensity were detectable in visual cortex tissue of wildtype and quadruple
19 knockout mice ($p = 0.16$, $N = 8$). **(C)** RT-qPCR analyses revealed a comparable *Gephn* mRNA
20 expression in the visual cortex of wildtype and quadruple knockout mice ($p = 0.07$, $N = 6$). **(D)** Western
21 blot analysis of gephyrin protein levels in the V1. **(E)** Comparable VGAT protein band intensity in
22 visual cortex tissue of wildtype and quadruple knockout ($p = 0.14$, $N = 8$). **(F)** RT-qPCR analyses
23 revealed a significant lower *Slc32a1(VGAT)* mRNA expression in the visual cortex of quadruple
24 knockout mice ($p < 0.001$, $N = 6$); 4xKO = quadruple knockout, *Gephn* = *Gephyrin*, V1 = primary
25 visual cortex, VGAT = vesicular GABA transporter, WT = wildtype, * = $p < 0.05$ data are shown as
26 mean \pm SEM and SD.

27 **Supplementary Figure 3:** Excitatory synaptic elements in the V1 of quadruple knockout mice. **(A)**
28 Western blot analysis of PSD95 protein levels in the V1. **(B)** No significant differences in the PSD95
29 protein band intensity were detectable in visual cortex tissue of wildtype and quadruple knockout mice
30 ($p = 0.85$, $N = 8$). **(C)** RT-qPCR analyses revealed a comparable *Dlg4* mRNA expression in the visual
31 cortex of wildtype and quadruple knockout mice ($p = 0.10$, $N = 6$). **(D)** Western blot analysis of
32 VGLUT1 protein levels in the V1. **(E)** Comparable VGLUT1 protein band intensity in visual cortex
33 tissue of wildtype and quadruple knockout ($p = 0.46$, $N = 8$). **(F)** RT-qPCR analyses revealed
34 comparable *Slc17a7 (VGLUT1)* mRNA expression in the visual cortex of wildtype and quadruple
35 knockout mice ($p = 0.40$, $N = 6$); 4xKO = quadruple knockout, *Dlg4* = *postsynaptic density protein*
36 *95*, PSD95 = postsynaptic density protein 95, *Slc17a7* = vesicular glutamate transporter 1, V1 =
37 primary visual cortex, VGLUT1 = vesicular glutamate transporter 1, WT = wildtype, * = $p < 0.05$ data
38 are shown as mean \pm SEM and SD.

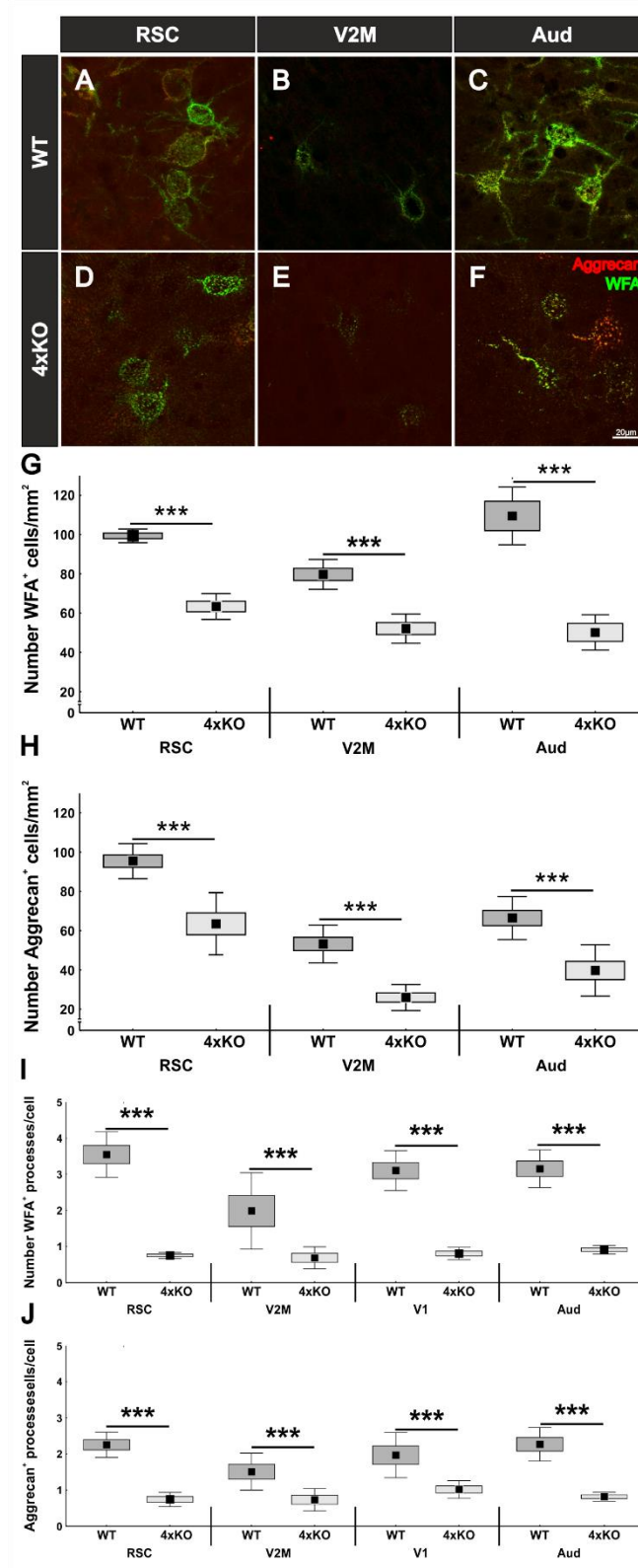
39 **Supplementary Figure 4:** Analyses of parvalbumin-positive interneuron populations in the
40 retrosplenial-, secondary visual- and auditory cortex of wildtype and quadruple knockout mice. **(A-F)**
41 Representative coronal cortical brain slices of wildtype and quadruple KO double-labeled using a
42 specific antibody against parvalbumin and WFA. **(G)** The number of parvalbumin-positive cells was
43 comparable in the retrosplenial cortex in wildtype and quadruple knockout mice ($p = 0.67$, $N = 8$).
44 Furthermore, the number of parvalbumin-positive cells was comparable in the secondary visual cortex
45 ($p = 0.11$, $N = 8$) and the auditory cortex of wildtype and quadruple knockout mice ($p = 0.61$, $N = 8$).

46 4xKO = quadruple knockout, *Pvalb* = parvalbumin, WT = wildtype, WFA = *Wisteria floribunda*
47 agglutinin, * = $p < 0.05$, data are shown as mean \pm SEM and SD, scale bar = 200 μm .

48 **Supplementary Figure 5:** Analyses of parvalbumin and calretinin-positive interneuron populations in
49 the retrosplenial-, secondary visual- and auditory cortex of wildtype and quadruple knockout mice. (A-
50 F) Representative coronal cortical brain slices of wildtype and quadruple KO double-labeled using a
51 specific antibody against calretinin and WFA retrosplenial-, secondary visual- and auditory cortex of
52 wildtype and quadruple knockout mice. (G) The number of calretinin-positive cells was comparable in
53 the retrosplenial cortex between wildtype and quadruple knockout mice ($p = 0.63$, $N = 8$). Furthermore,
54 the number of calretinin-positive cells was comparable in the secondary visual cortex ($p = 0.53$, $N =$
55 8) and the auditory cortex of wildtype and quadruple knockout mice ($p = 0.19$, $N = 8$). 4xKO =
56 quadruple knockout, *Pvalb* = parvalbumin, WT = wildtype, WFA = *Wisteria floribunda* agglutinin, *
57 = $p < 0.05$, data are shown as mean \pm SEM and SD, scale bar = 200 μm .

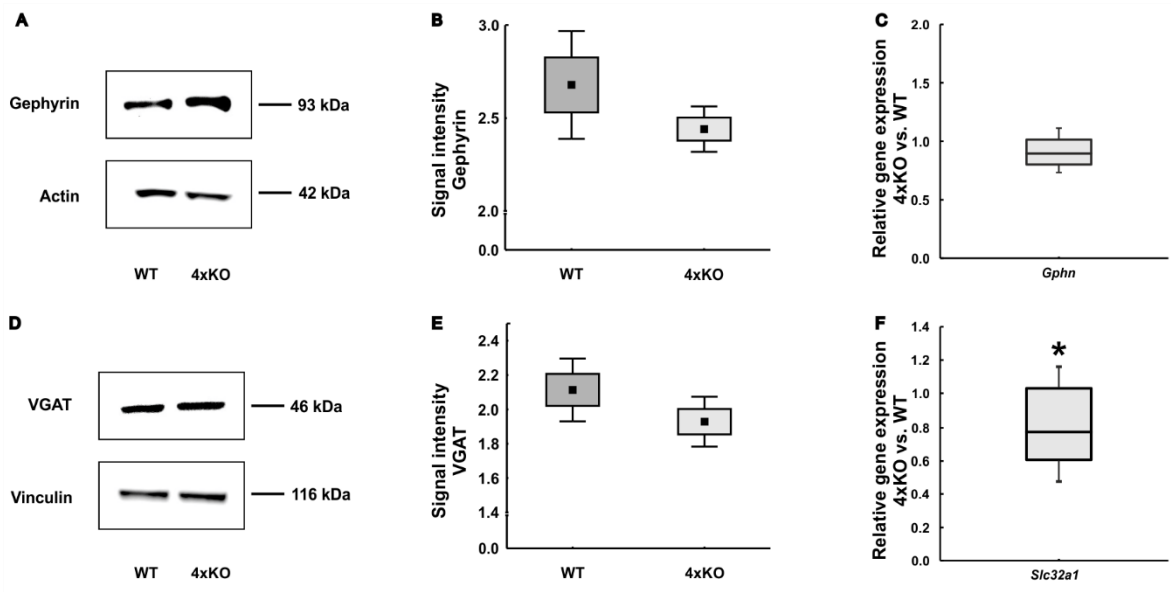
58 17 SUPPLEMENTARY FIGURES

59 Supplementary Figure 1:



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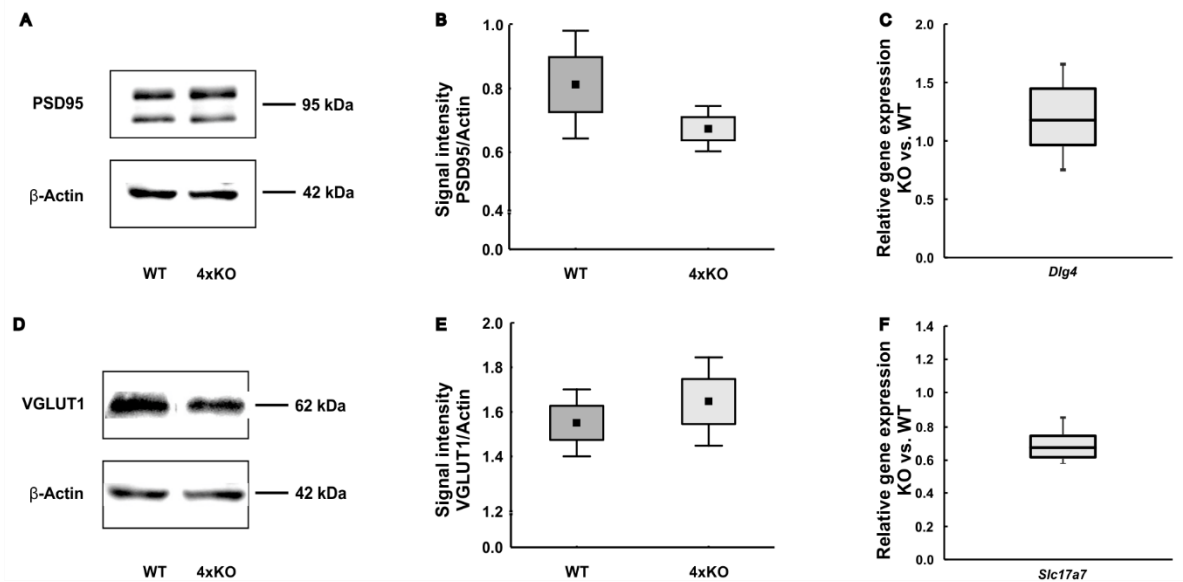
61 **Supplementary Figure 2:**



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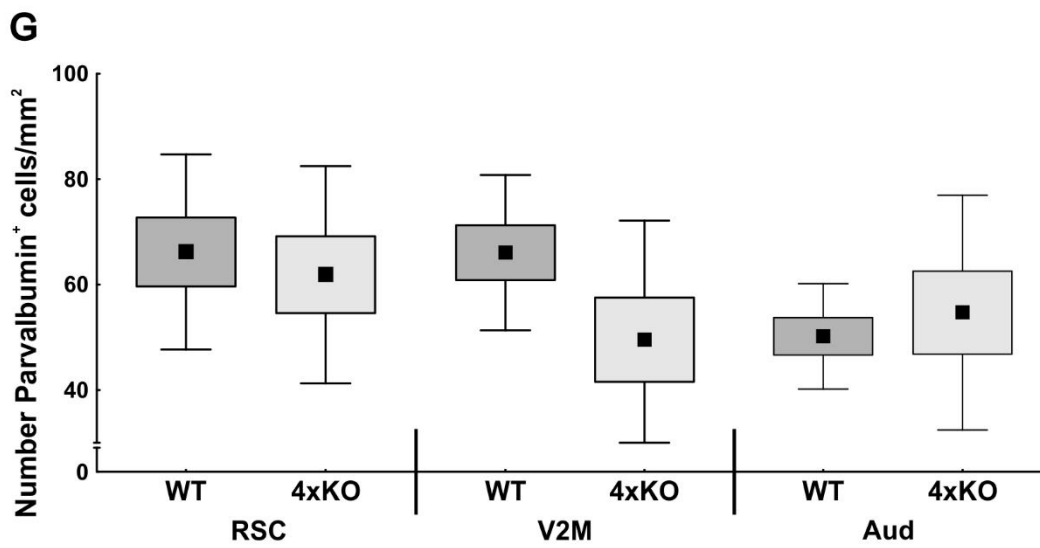
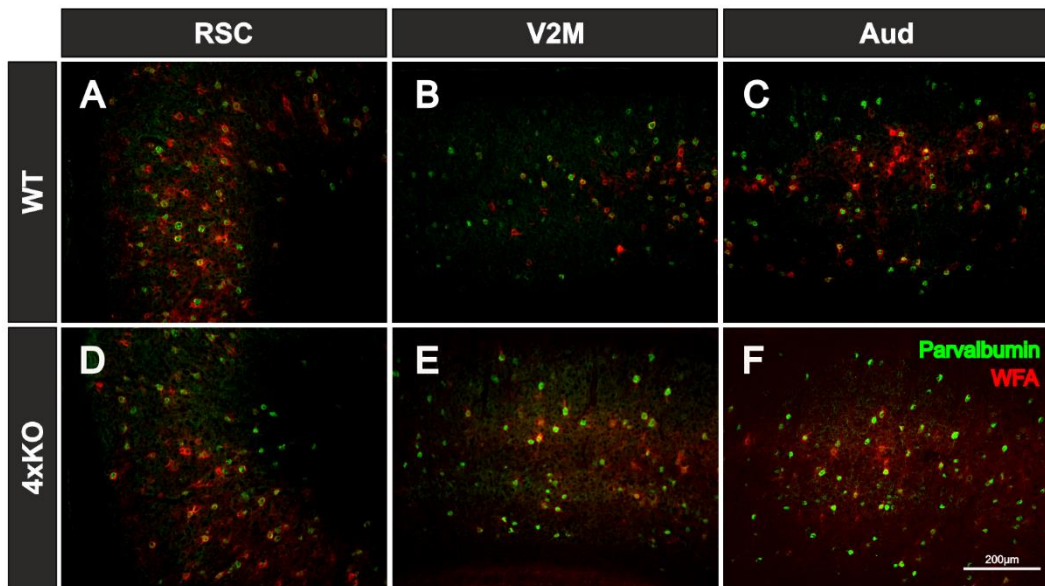
65 **Supplementary Figure 3:**



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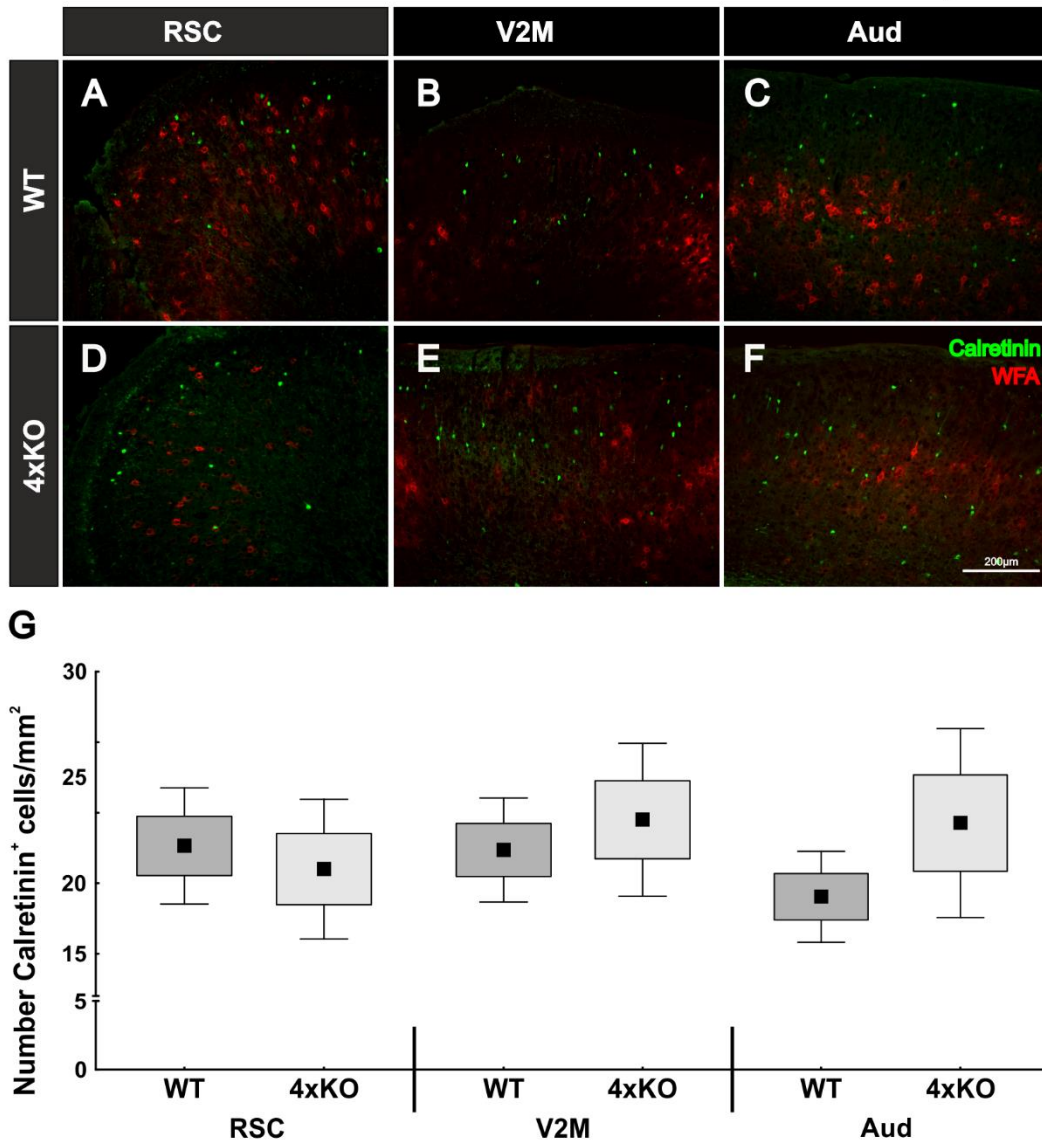
68 **Supplementary Figure 4:**



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71 **Supplementary Figure 5:**



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