

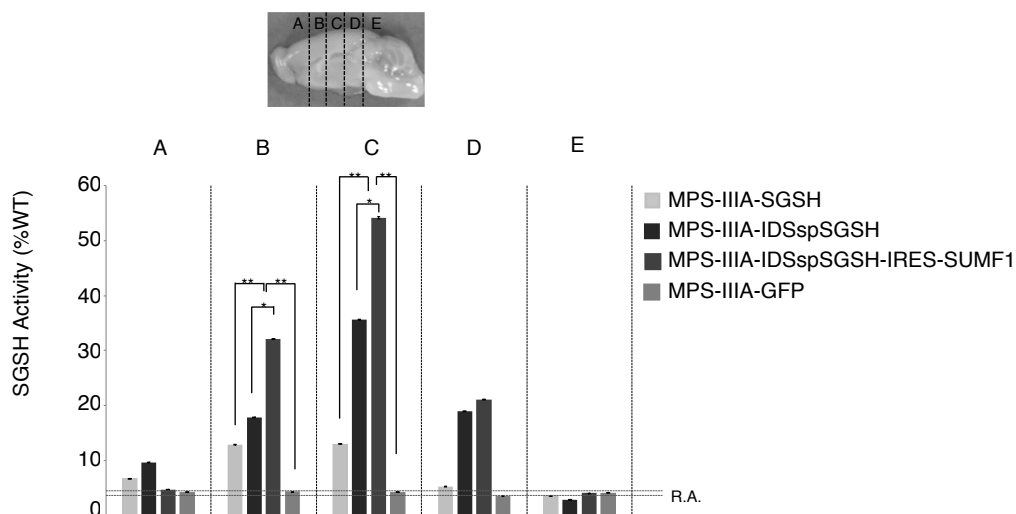
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## **Supplemental Information**

### **Enhancing the Therapeutic Potential of Sulfamidase for the Treatment of Mucopolysaccharidosis IIIA**

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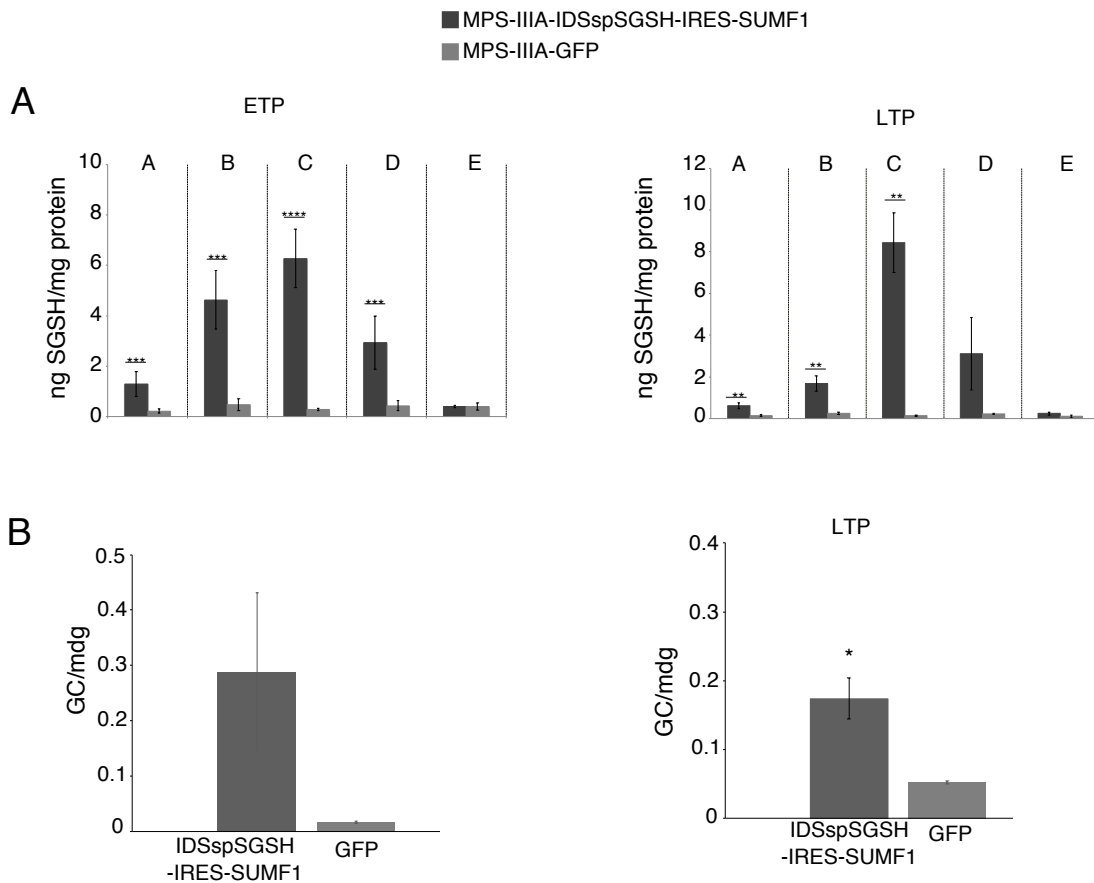
# Supplementary Figure 1



## Supplementary Figure 1. Sulfamidase activity in the brain of MPS-III A mice injected with AAV9 vectors carrying different SGSH expression cassettes

P60 MPS-III A mice were intra-CSF injected (via lateral ventricle administration: ICV) with  $5.4 \times 10^{12}$  GC/Kg of AAV9 encoding under the CMV promoter the following expression cassettes: *GFP*, *SGSH* WT, *IDSspSGSH*, or *IDSspSGSH-IRES-SUMF1*. The brain and the first region of the spinal cord of treated mice was divided in five slices (A-E) covering the main representative area of the CNS (A: olfactory bulb and prefrontal cortex, B: frontal cortex, lateral septum and basal ganglia regions, C: parietal cortex, hippocampus, striatum, thalamus, D: occipital cortex, pons, hippocampus; E: cerebellum, medulla oblongata, cervical region of spinal cord). One month after injection sulfamidase activity was measured in these areas and expressed as the percentage of the activity found in control GFP-treated WT mice. N = 3-4 animals per group. Data represent mean  $\pm$  SEM. \* $P < 0.05$ , \*\* $P < 0.01$  MPS-III A-IDSspSGSH-IRES-SUMF1 VS MPS-III A-GFP, MPS-III A-IDSspSGSH-IRES-SUMF1 VS MPS-III A-IDSspSGSH, MPS-III A-IDSspSGSH-IRES-SUMF1 VS MPS-III A-SGSH. One-way ANOVA followed by Tukey's post hoc test.

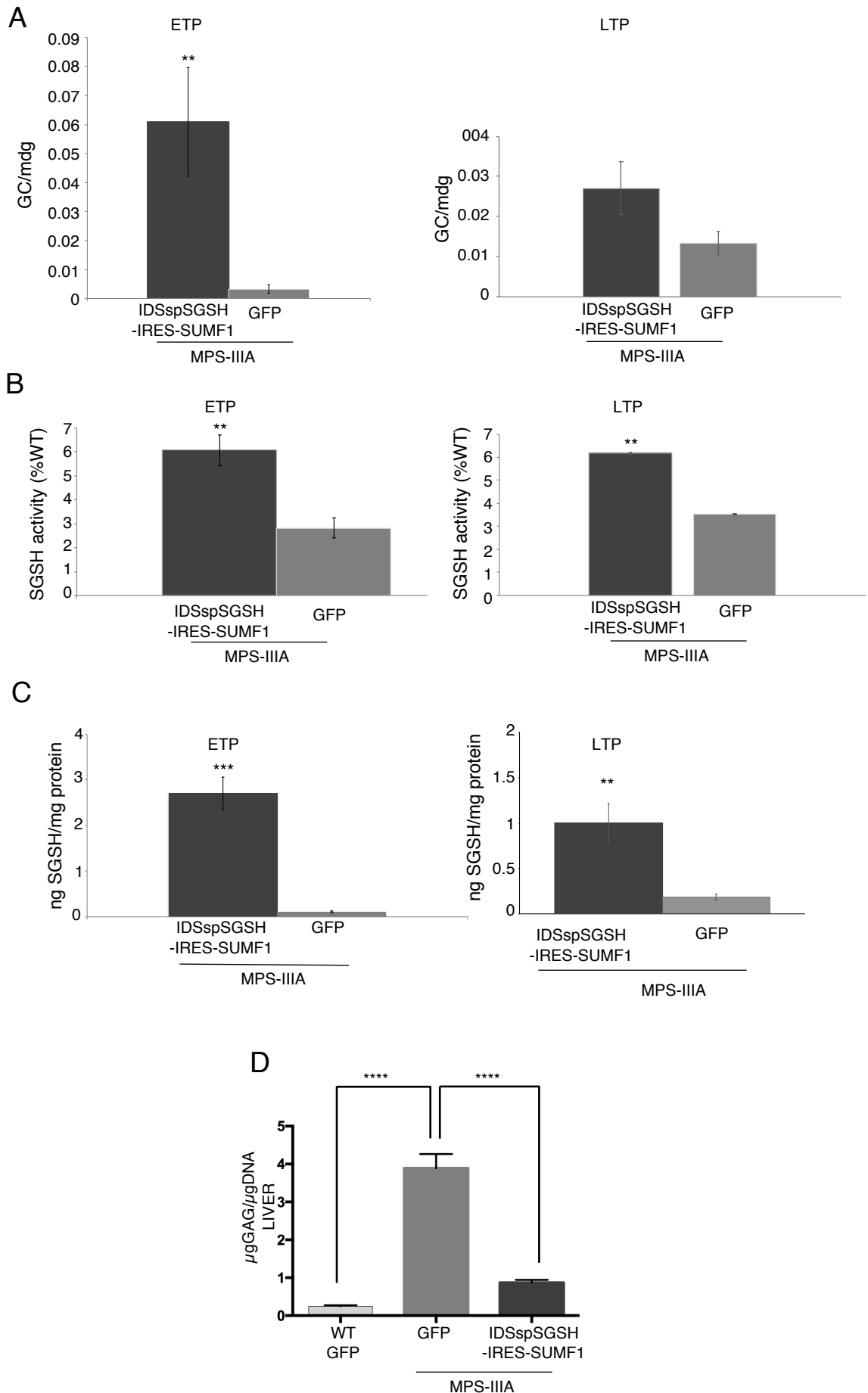
## Supplementary Figure 2



### Supplementary Figure 2. Sulfamidase protein and vector copy numbers quantitation in the brain of MPS-III A mice injected with AAV9 bearing the IDSpSGSH-IRES-SUMF1 transgene

(A) Sulfamidase protein was immuno-quantified by ELISA and expressed as ng of SGSH/mg protein in the five CNS slices (A-E; as described in the supplementary figure 1) of the indicated experimental groups of mice at ETP and LTP. Age-matched WT and MPS-III A mice ICV injected with AAV9 encoding for GFP were used as control. Data represent mean  $\pm$  SEM. N= 5-7 animals for each group. \*\* $P < 0.01$ , \*\*\* $P < 0.001$ , \*\*\*\* $P < 0.0001$  VS MPS-III A-GFP. One-way ANOVA followed by Tukey's post hoc test. (B) Vector genome copy number (expressed as GC/mouse diploid genome; mdg) were measured in the whole brain samples from MPS-III A mice ICV injected with AAV9 encoding IDSpSGSH-IRES-SUMF1 and age-matched MPS-III A mice ICV injected with AAV9 encoding GFP at ETP and LTP. Data represent mean  $\pm$  SEM. N= 5-7 animals for each group. \* $P < 0.05$ , Student T-test.

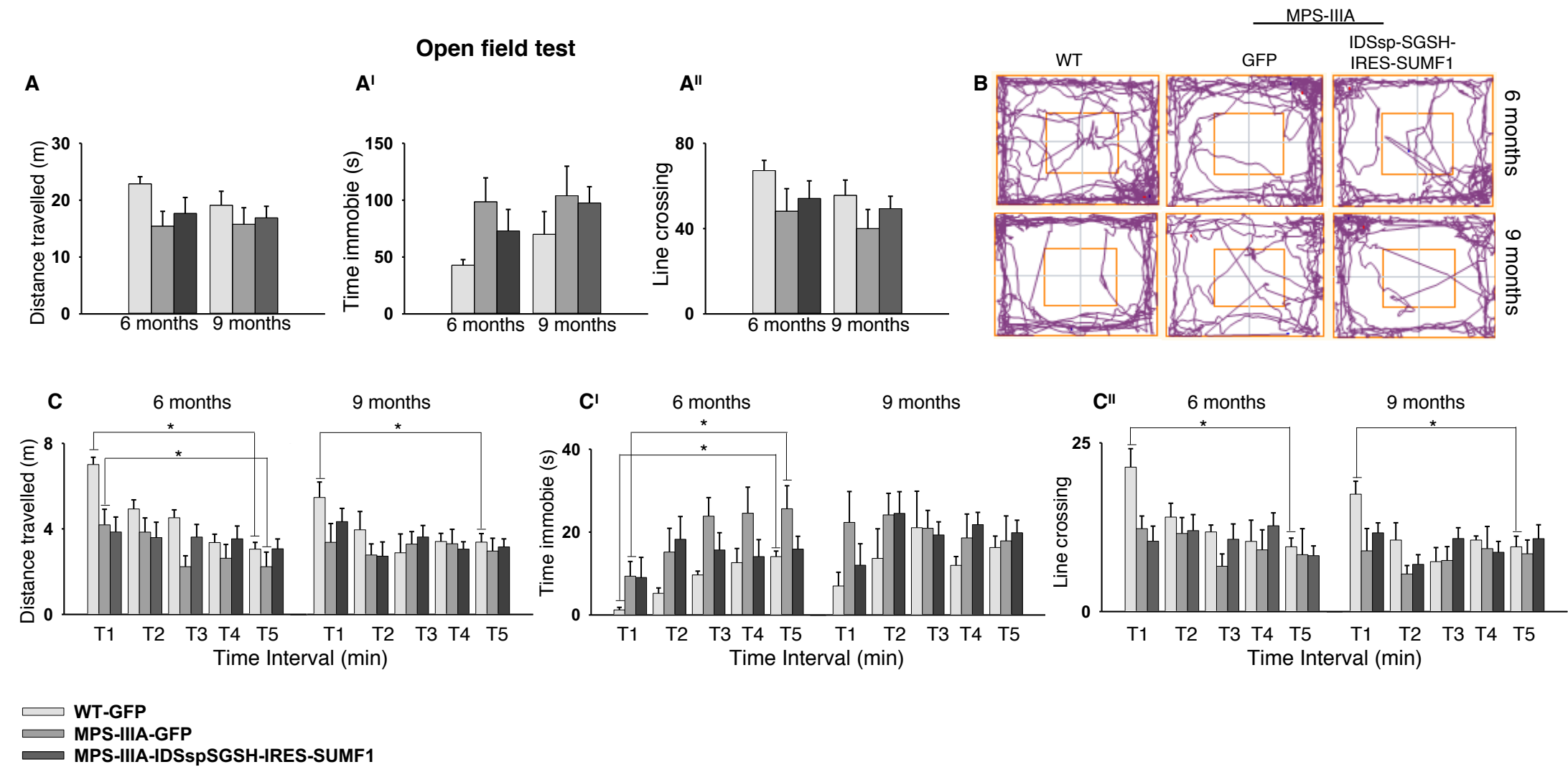
# Supplementary Figure 3



### Supplementary Figure 3. Liver transduction in MPS-IIIa mice injected with AAV9 bearing *IDSspSGSH-IRES-SUMF1* transgene

(A-C) Vector genome copy number (expressed as GC/mouse diploid genome; mdg) (A), sulfamidase activity (expressed as the percentage of WT sulfamidase activity) (B) and ELISA immunoquantification of the sulfamidase protein (expressed as ng of SGSH/mg protein) (C) were measured in liver samples from MPS-IIIa mice ICV injected with AAV9 encoding *IDSspSGSH-IRES-SUMF1* and age-matched WT and MPS-IIIa mice ICV injected with AAV9 encoding GFP at ETP and LTP. Data represent mean  $\pm$  SEM. N= 6-8 animals for each group. \*\*P<0.01, \*\*\*P<0.001. *vs* MPS-IIIa GFP-treated. Student T-test. (D) Quantitative analysis of GAG content ( $\mu$ gGAG/ $\mu$ gDNA) in liver samples collected at LTP in MPS-IIIa mice ICV injected with AAV9 encoding *IDSspSGSH-IRES-SUMF1*. Age-matched WT and MPS-IIIa mice ICV injected with AAV9 encoding GFP were used as controls. N = 6 animals per group. Data represent mean  $\pm$  SEM; \*\*\*\*P<0.0001. One-way ANOVA followed by Tukey's post hoc test.

# Supplementary Figure 4



**Supplementary Figure 4. Assessment of exploratory activity in MPS-IIIa mice injected with AAV9 bearing IDSspSGSH-IRES-SUMF1 transgene.**

MPS-IIIa mice and relative controls (WT) were tested at 6 and 9 months of age in the open field test. (A- AII) There were no significant differences between groups at any of the testing age in the total distance travelled (m) [Group (F2/16=1.16; p=0.22); Distance (F1/16=0.65; p=0.43); Group x Distance x Age (F2/16=0.45; p=0.64)] (A), total number of line crossings [Group (F2/16=1.94; p=0.17); Line crossing (F1/16=1.77; p=0.20); Group x Line crossing x Age (F2/16=0.24; p=0.78)] (AI) and total immobility time (sec) [Group (F2/16=1.46; p=0.25); Immobility time (F1/16=2.31; p=0.14); Group x Immobility time x Age (F2/16=0.12; p=0.88)] (AII). (C-CII) A deeper analysis of the results considering 1 min time intervals (T1-T5) evidenced that at 6 months of age MPS-IIIa mice, as compared to WT littermates, showed reduced distance travelled [Time interval (F4/64=17.9; p<0.0001); Time intervals x Group (F4/64=3.16; p=0.004), Age x Time intervals x Group (F4/64=2.6; p=0.01)] (C), increased immobility time [Time intervals (F4/64=2.4; p=0.05); Group x Age x Time interval (F8/64=2.04; p=0.05)] (CI) and reduced line crossing frequency [Time interval (F4/64=6.8; p=0.0001); Group x Time intervals (F4/64=2.66; p=0.01)] (CII) mainly present in the very first minute of the task; these behavioral defects, however, were not anymore detectable at 9 months of ageing probably due to a test-retest habituation effect observed in WT animals [distance, age x time interval (F8/64=3.73; p=0.0085); time immobile, age x time intervals (F4/64=1.61; p=0.05), line crossing, age x time intervals (F4/64=2.4; p=0.05)]. Representative track-plots of the trajectory in the open field (B). \* p<0.05, Duncan post hoc analysis.

# Supplementary Table 1

| Pvalue of SGSH activity in CNS areas of wild type pigs |                                    |  |                               |                              |
|--|------------------------------------|--|-------------------------------|------------------------------|
| CNS regions  | IDSsp-SGSH-IRES-SUMF1 vs IDSspSGSH | IDSsp-SGSH-IRES-SUMF1 vs SGSH-IRES-SUMF1 | IDSsp-SGSH-IRES-SUMF1 vs SGSH | IDSsp-SGSH-IRES-SUMF1 vs PBS |
| Frontal cortex   | ns                                 | ns                                       | *                             | **                           |
| Accumbens  | **                                 | **                                       | ****                          | ****                         |
| Parietal ccortex                                       | ns                                 | **                                       | ***                           | ****                         |
| Amigdala   | ns                                 | *  | **                            | ***                          |
| Hypotalamus  | ns                                 | **                                       | *                             | **                           |
| N.caudatus   | ns                                 | *  | *                             | ***                          |
| Putamen  | ns                                 | **                                       | *                             | ***                          |
| Sub.Nigra  | ns                                 | *  | *                             | **                           |
| Stria  | ns                                 | *  | *                             | ***                          |
| Hippocmpus   | ns                                 | **                                       | **                            | ***                          |
| Thalamus   | *                                  | ***                                      | **                            | ****                         |
| C.callosum   | *                                  | ****                                     | ****                          | ****                         |
| Subcallosus  | *                                  | **                                       | ***                           | ****                         |
| N.Pontis   | ns                                 | ns                                       | ns                            | **                           |
| Culliculli   | *                                  | **                                       | *                             | ***                          |
| Occipital cortex                                       | ns                                 | **                                       | **                            | ***                          |
| Cerebellum   | ns                                 | *  | *                             | ***                          |
| Pons   | **                                 | *  | **                            | ****                         |
| Med. Ablong.   | ns                                 | *  | ns                            | ***                          |
| Spinal cord  | ns                                 | ns                                       | *                             | ***                          |

P ≤0,05; \*\* P ≤0,01; \*\*\* P ≤0,001; \*\*\*\* P ≤0,001; ns: not significant.  
 No significant differences of SGSH activity in CNS regions among SGSH-IRES-SUMF1 VS SGSH experimental groups.