

Supplementary material 1

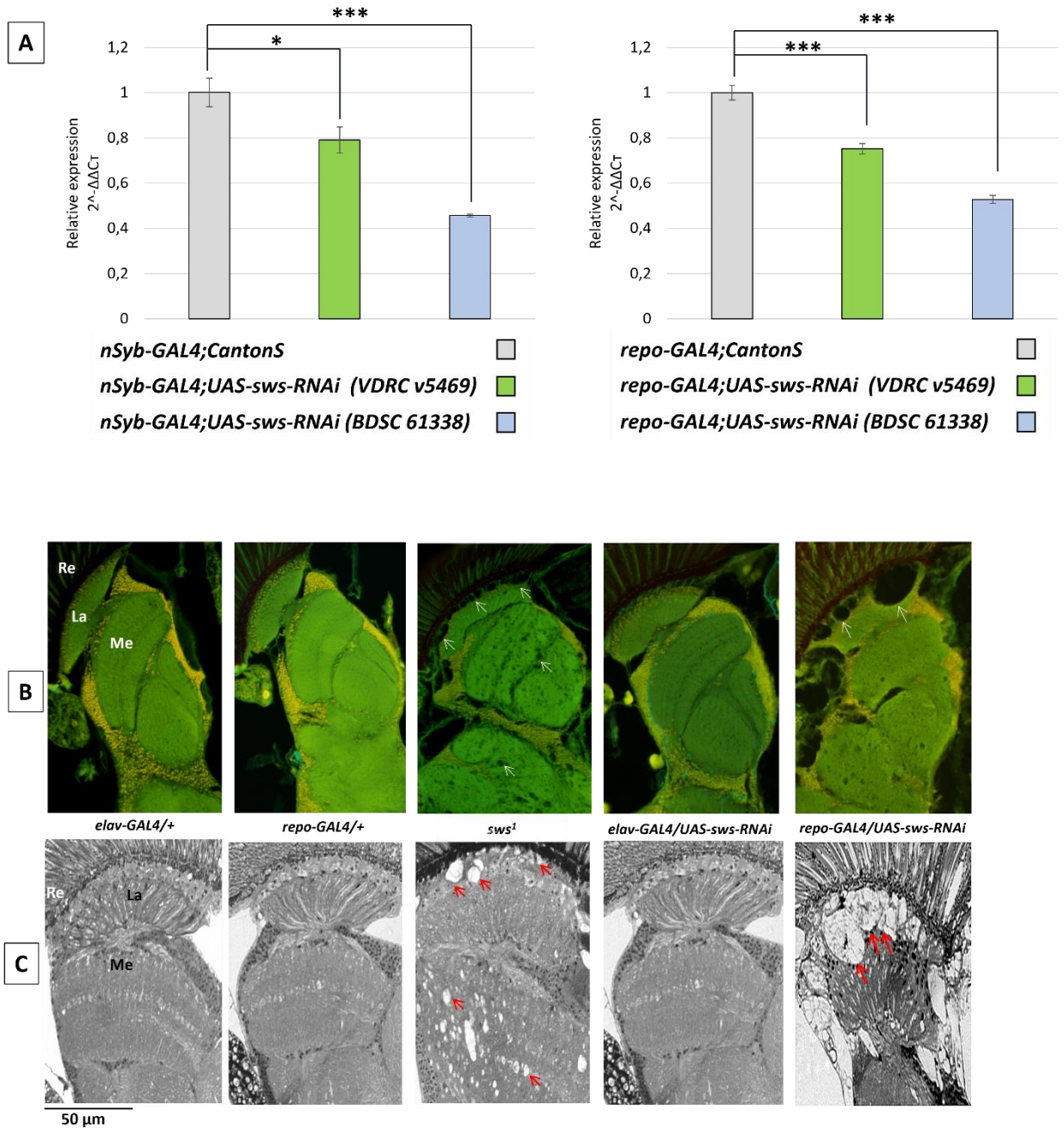


Figure S1. Analysis of the *UAS-*sws*-RNAi^{v5469}* (VDRC № v5469) transgene effects. **(A)** Relative *sws* mRNA level in heads of flies with *sws* knockdown in neurons (*nSyb-GAL4;UAS-*sws*-RNAi*) or in glia (*repo-GAL4;UAS-*sws*-RNAi*) and control flies (F1 males obtained from crossing a *CantonS* female and a respective *GAL4* male). The level of *sws* mRNA in the control is taken as 1,00 in each sample. The mean value of $2^{-\Delta\Delta CT}$ and 95% CI, Student t-test, * $p < 0.05$, *** $p < 0.001$, $N = 3$. **(B)** Horizontal 7 μm paraffin and **(C)** horizontal 1 μm plastic semithin brain sections of 20-day-old *sws¹* mutants, flies with the neuronal (*elav-GAL4;UAS-*sws*-RNAi^{v5469}*) and glial (*repo-GAL4;UAS-*sws*-RNAi^{v5469}*) *sws* knockdown or without it (*elav-GAL4/+* and *repo-GAL4/+*). Re – retina, La – lamina, Me – medulla. Scale bar: 50 μm. Brain tissue neurodegenerative vacuoles are marked with arrows. Flies of the *elav-GAL4;UAS-*sws*-RNAi^{v5469}* showed no neurodegenerative phenotype in contrast to *repo-GAL4;UAS-*sws*-RNAi^{v5469}*.

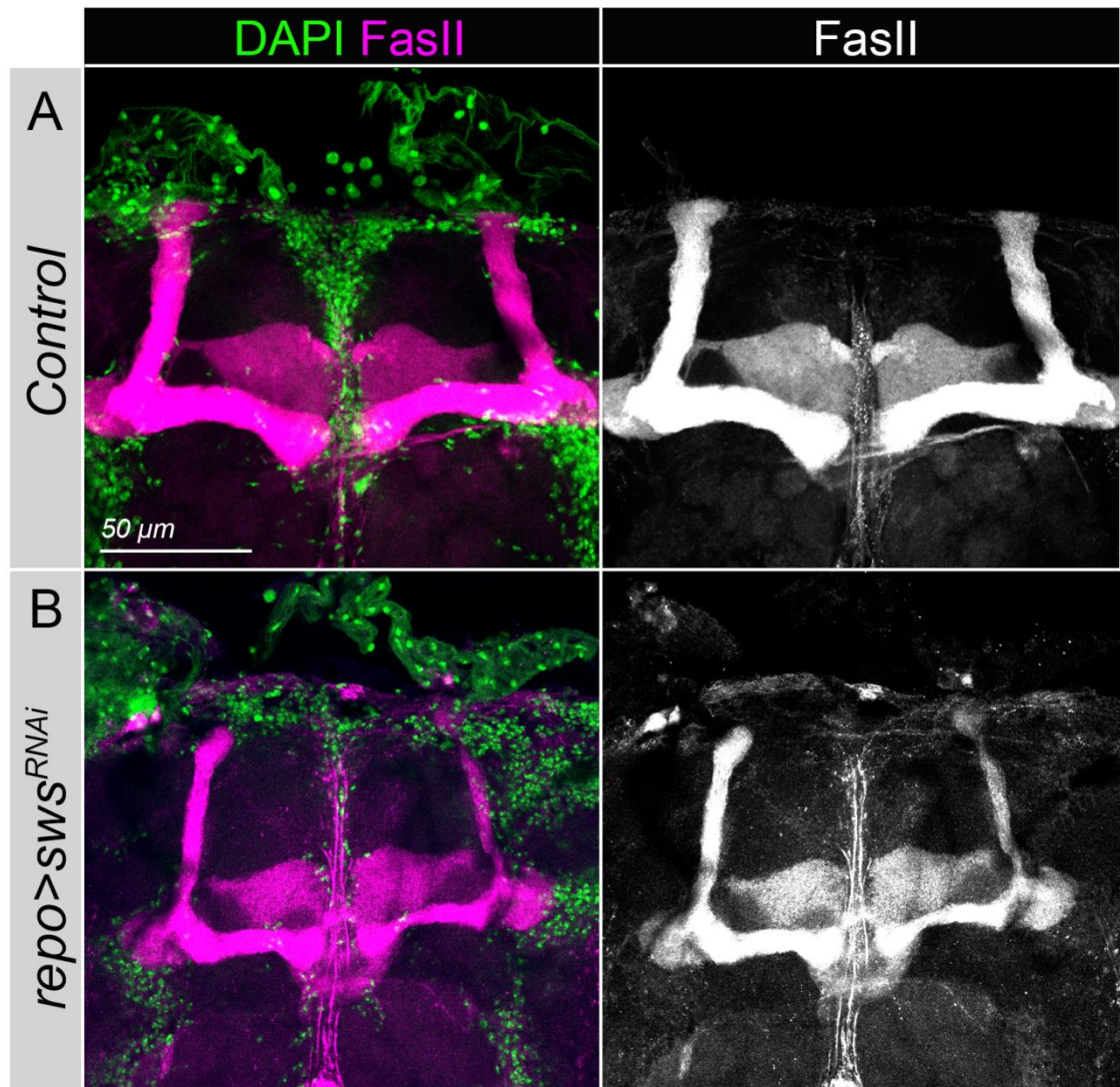


Figure S2. Downregulation of *sws* in glia (*repo-GAL4;UAS-sws-RNAi*) results in the mushroom body defects. (A) Confocal images of adult MB lobes in control flies (*repo-GAL4/+*) marked by FasII antibodies (magenta). (B) Upon *sws* downregulation in glia (*repo-GAL4;UAS-sws-RNAi*), MB lobes appear to be reduced.

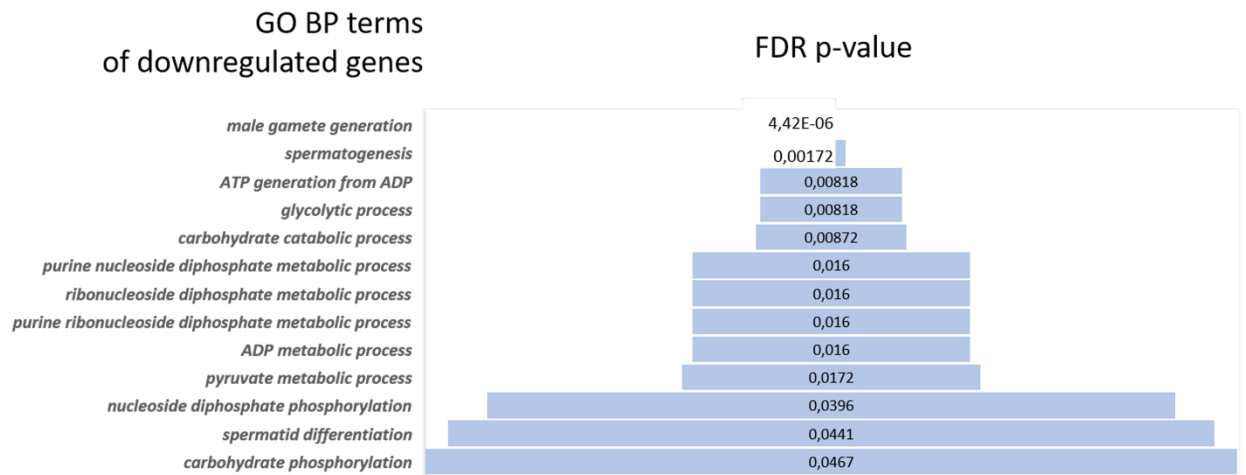


Figure S3. GO BP processes under the control of downregulated genes in 25-day-old neuronal *sus* knockdown males (*elav-GAL4;UAS-sus-RNAi*) compared to *CantonS* control and respective FDR-adjusted p-values of functional enrichment analysis in g:Profiler software.

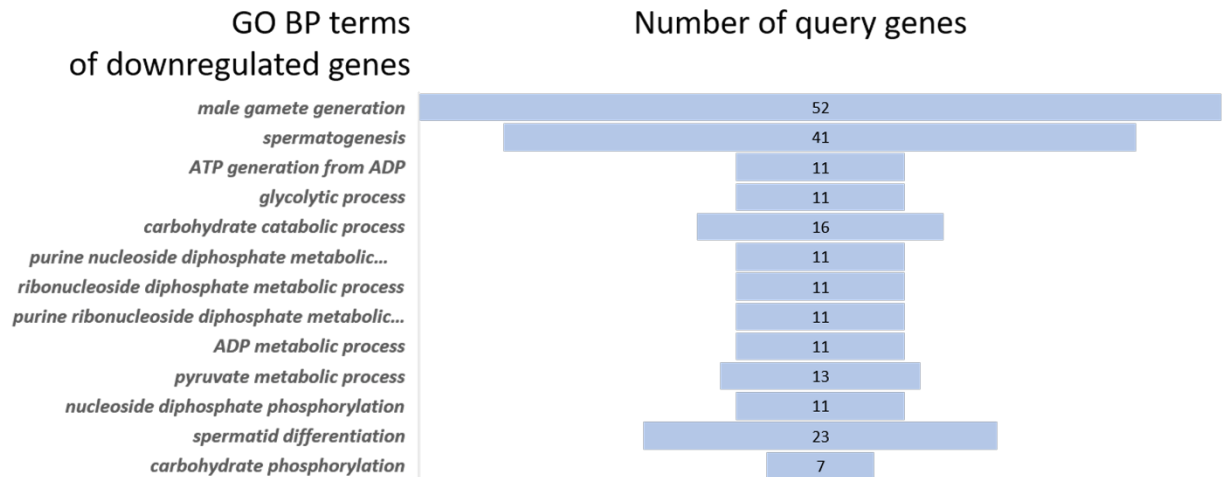


Figure S4. GO BP processes under the control of downregulated genes in 25-day-old neuronal *sus* knockdown males (*elav-GAL4;UAS-sus-RNAi*) compared to *CantonS* control and respective query gene number from functional enrichment analysis in g:Profiler software.

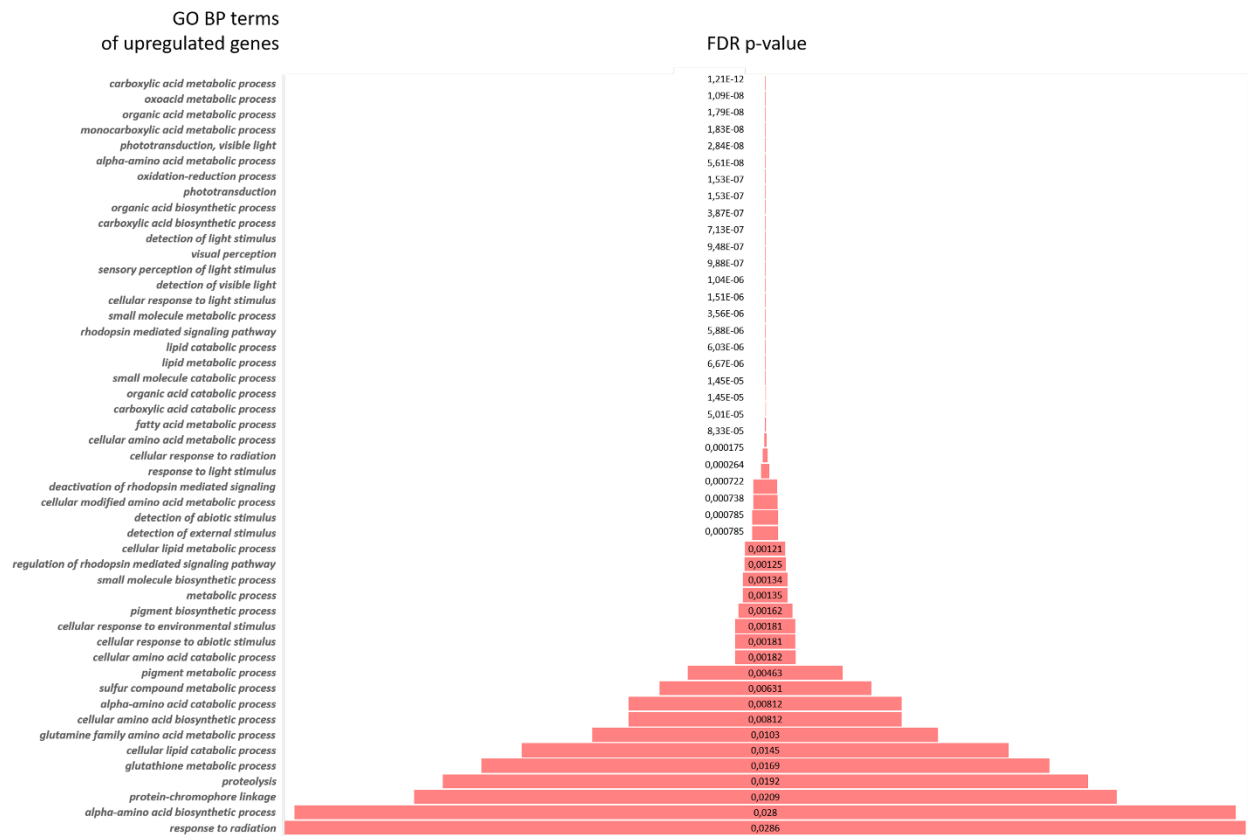


Figure S5. GO BP processes under the control of upregulated genes in 25-day-old neuronal *sws* knockdown males (*elav-GAL4;UAS-sws-RNAi*) compared to *CantonS* control and respective FDR-adjusted p-values of functional enrichment analysis in g:Profiler software.

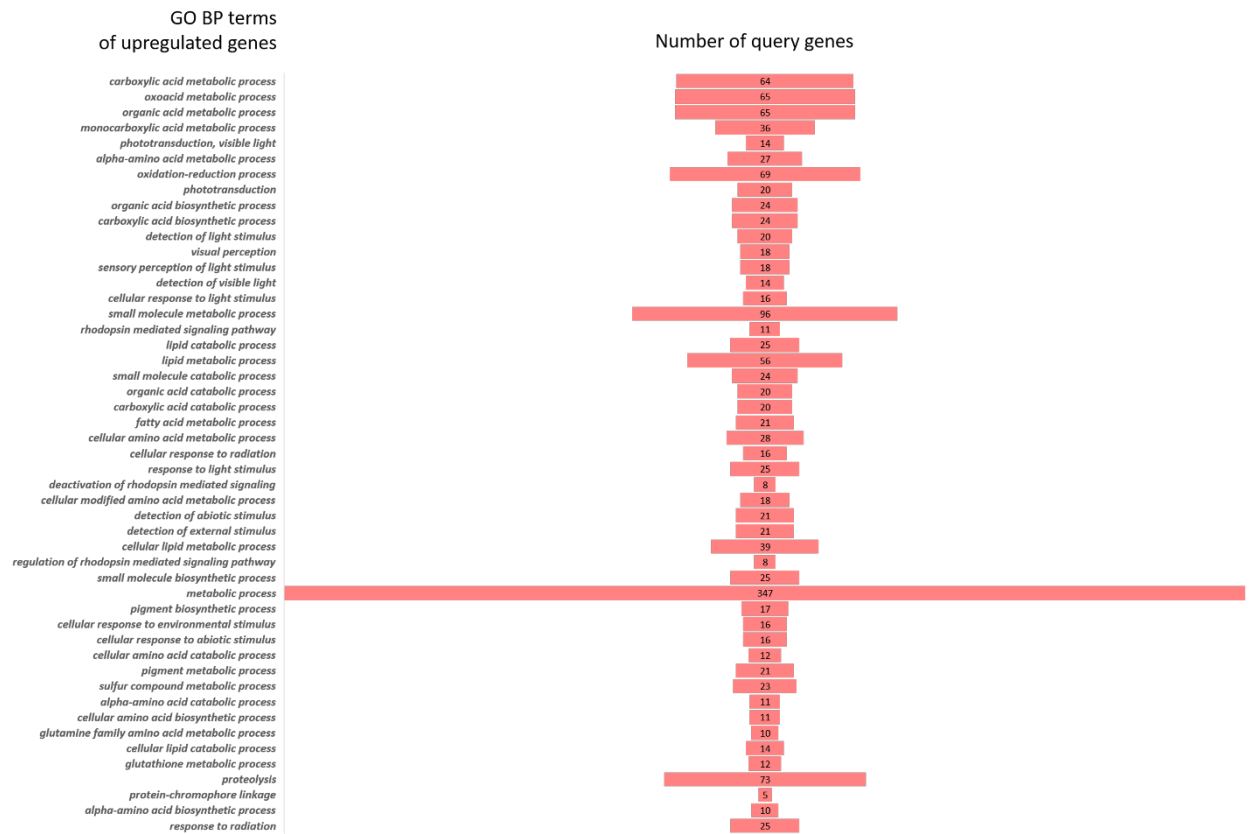
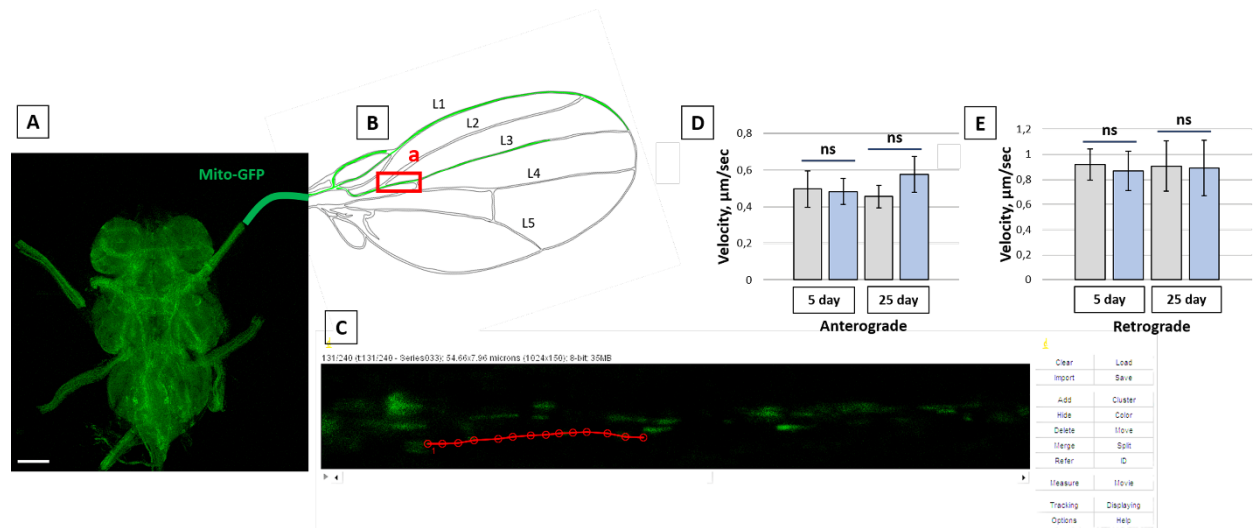


Figure S6. GO BP processes under the control of upregulated genes in 25-day-old neuronal *sws* knockdown males (*elav-GAL4;UAS-sws-RNAi*) compared to *CantonS* control and respective query gene number from functional enrichment analysis in g:Profiler software.



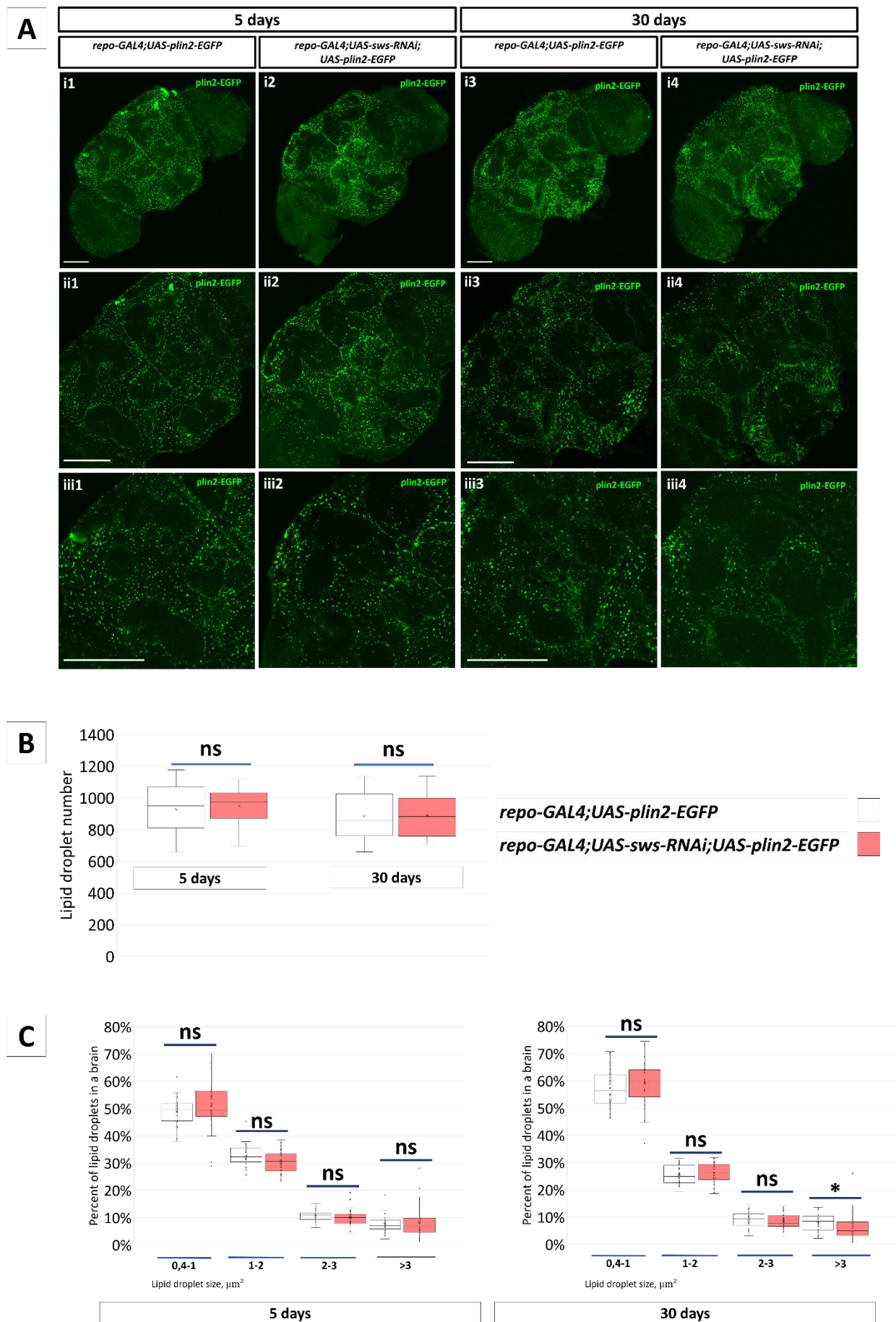


Figure S8. Analysis of lipid droplets in glia of the fly brain. (A) Single confocal images of the central brain of the control (i-iii1, i-iii3, *repo-GAL4;UAS-plin2-EGFP*) and knockdown (i-iii2, i-iii4, *repo-GAL4;UAS-sws-*

RNAi;UAS-plin2-EGFP) flies of 5 (**i-iii1, i-iii2**) and 30-day-old (**i-iii3, i-iii4**) age. Green – GFP embedded in lipid droplets (marked with arrows) via the Plin2 protein in glia. Scale bar: 100 μm . **(B)** The total lipid droplet number in the brain glia of the control (*repo-GAL4;UAS-plin2-EGFP*, white boxes) and knockdown (*repo-GAL4;UAS-sws-RNAi;UAS-plin2-EGFP*, red boxes) flies of 5 and 30-day-old age. Mann-Whitney test, ns – no significant difference ($p > 0.05$), $N = 30$. **(C)** Distribution of LD size (μm^2) in the brain glia of the control (*repo-GAL4;UAS-plin2-EGFP*, white boxes) and knockdown (*repo-GAL4;UAS-sws-RNAi;UAS-plin2-EGFP*, red boxes) flies of 5 and 30-day-old age. Mann-Whitney test, *** $p < 0.001$, * $p < 0.05$, ns – no significant difference ($p > 0.05$), $N = 30$.