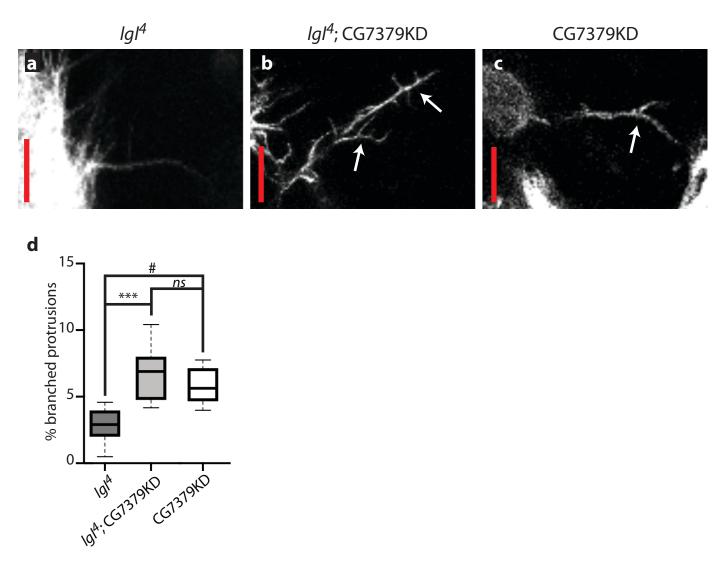
## Figure S1



## Figure S1: Abnormal protrusion morphology following CG7379KD

(**a-c**) Example of typical basal protrusion morphology from *IgI*<sup>4</sup> cells (a), and examples of abnormal branched protrusions from *IgI*<sup>4</sup>; CG7379KD (b) and CG7379KD (c) cells. Images represent *z* stack projections of relevant basal confocal slices. Arrows highlight branched protrusions. (**d**) Quantification of branched protrusion occurrence. n $\geq$ 100 protrusions/animal from a minimum of 7 animals/genotype. One-way ANOVA was performed to determine statistical significance. Scale bar = 10µm. \* p<0.05; \*\*p<0.01; \*\*\*p<0.001; #p<0.0001.