

MOL231: Characterization of Δ Unc13 in the multicellular stage of the choanoflagellate, *Salpingoeca rosetta*

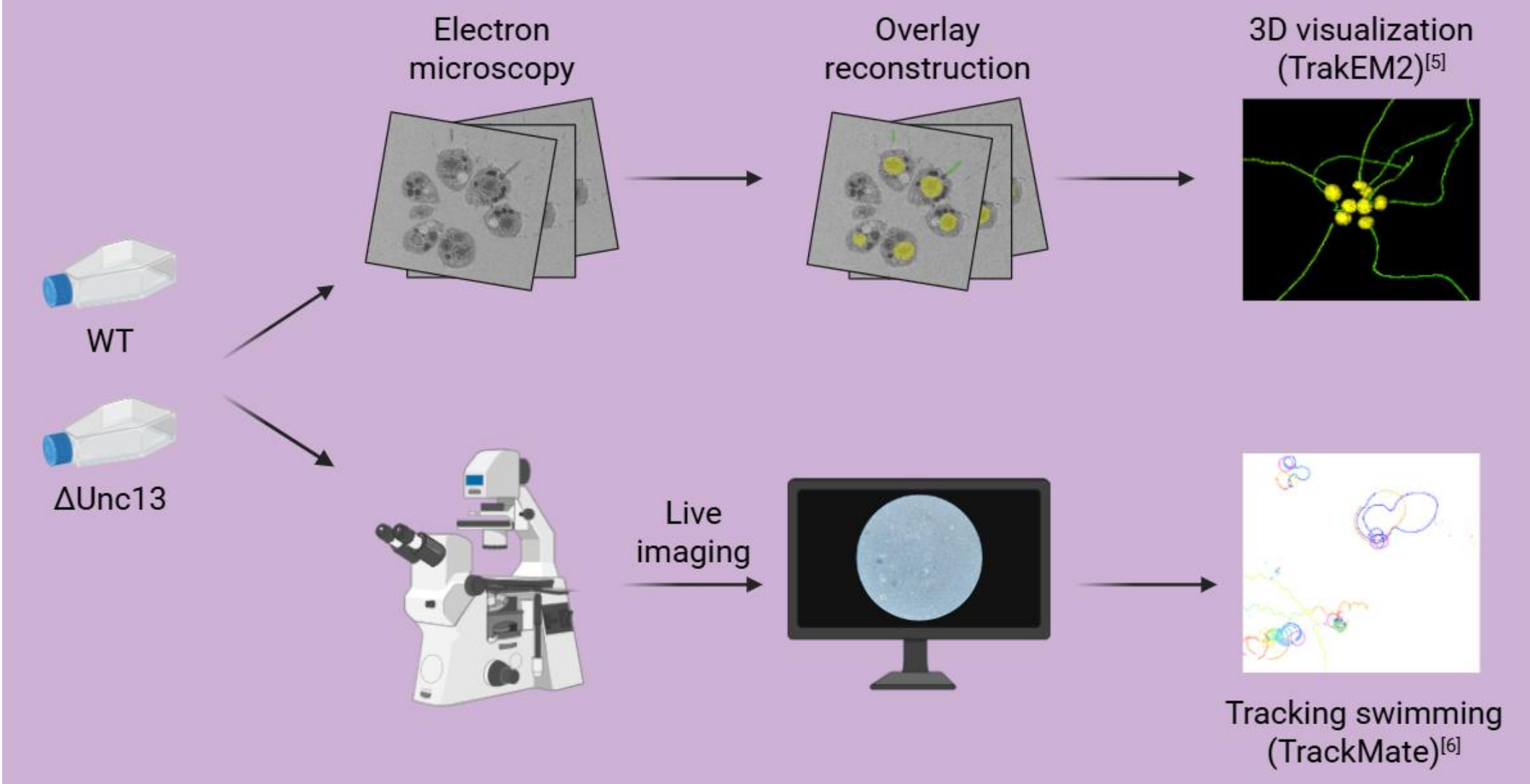
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Background

Choanoflagellates are the sister group to animals and possess many genes once thought to be animal specific^[1]. One such gene encodes the Unc13 protein, which is a key synaptic protein in animals^[2]. The choanoflagellate *Salpingoeca rosetta* exists as single cells which contain an apical feeding structure, composed of a flagellum, surrounded by a microvillar collar^[3]. Certain environmental cues can trigger the formation of clonal rosette colonies^[4], which are believed to increase feeding efficiency. Knockout of Unc13 in *S. rosetta* results in severe defects in the apical structure^[3]. The goal of this study was to elucidate the role of Unc13 in *S. rosetta* colonies.

Workflow



Δ Unc13 results in absence of flagella in some cells and change in the mobility of the colonies

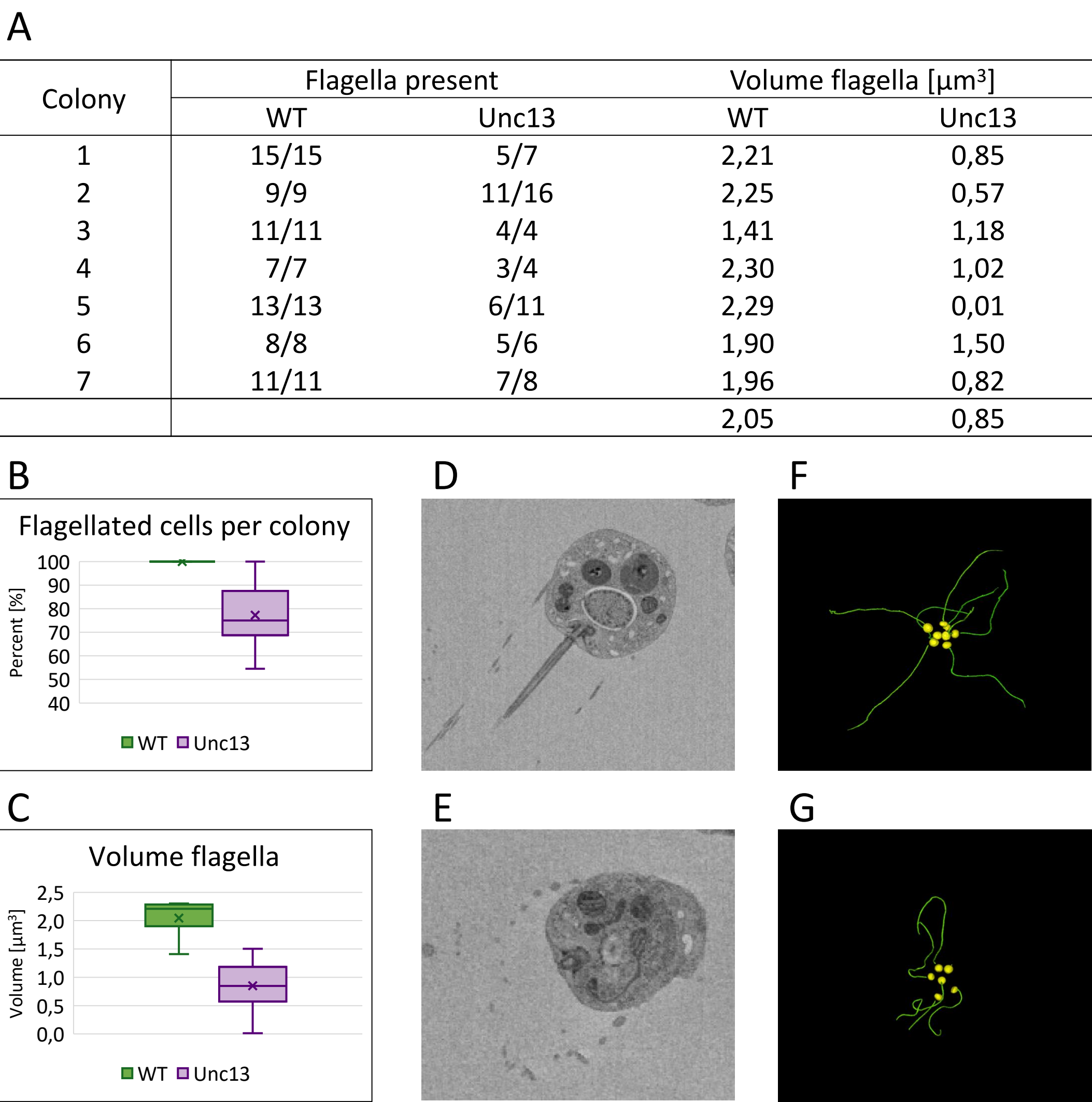


Figure 1. Δ Unc13 colonies show fewer and smaller flagella. (A) Overview of the number and average volume of full reconstructed flagella in *S. rosetta* colonies. (B) Percentage of flagellated cells per colony. (C) Flagellar volume measurements of fully reconstructed flagella in each colony. (D–E) Electron micrographs of a wild-type (WT) colonial cell (D) and a Δ Unc13 colonial cell (E). (F–G) 3D reconstructions of a WT colony (F) and a Δ Unc13 colony (G). Green: flagella, yellow: nuclei

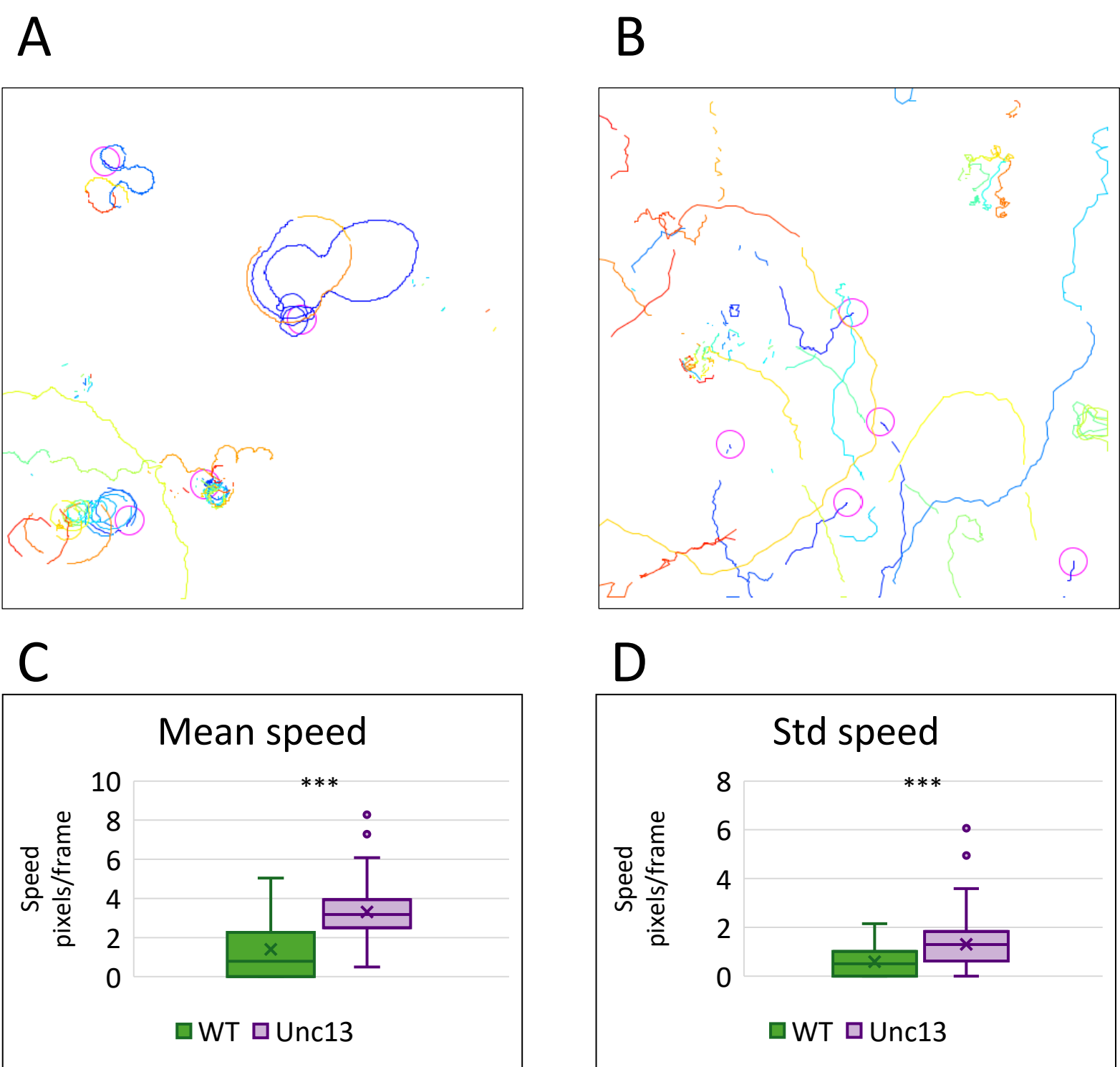
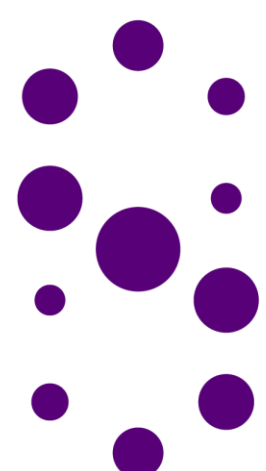


Figure 2. Δ Unc13 have a stochastic and faster movement. (A–B) Trajectories of WT colonies (A) and Δ Unc13 colonies (B). (C) Mean speed per colony over the course of imaging shows a significant increase in mutant colonies, $P < 0.001$. (D) Standard deviation speed per colony over the course of imaging shows a significant increase in mutant colonies, $P < 0.001$.

Conclusion

The electron microscopy shows that in colonies, some Δ Unc13 cells lack a flagellum and those that are present have a reduced volume, indicating shorter flagella. Furthermore, the live imaging shows a mobility change in the Δ Unc13 colonies. These findings are consistent with Unc13 playing an important role in formation or maintenance of the flagella in *S. rosetta*, providing valuable insights into the evolution of synaptic proteins^[3].



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References:
1. Fairclough et al (2013). Premetazoan genome evolution and the regulation of cell differentiation in the choanoflagellate *Salpingoeca rosetta*. *Genome biology*, 14(2), R15.; 2. Dittman, J. S. (2019). Unc13: a multifunctional synaptic marvel. *Curr Opin Neurobiol*, 57, 17-25; 3. Ravi, A. (2024). Polarized Recruitment of Secretory Vesicles in the Choanoflagellate *Salpingoeca rosetta*: Insights into the Origin of Neurosecretion; 4. Dayel et al (2011). Cell differentiation and morphogenesis in the colony-forming choanoflagellate *Salpingoeca rosetta*. *Developmental biology*, 357(1), 73–82. 5. Cardona et al (2012). TrakEM2 Software for Neural Circuit Reconstruction. *PLoS ONE*, 7(6); 6. Tinevez et al (2017). TrackMate: An open and extensible platform for single-particle tracking. *Methods*, 115, 80–90

