

Expanded View Figures

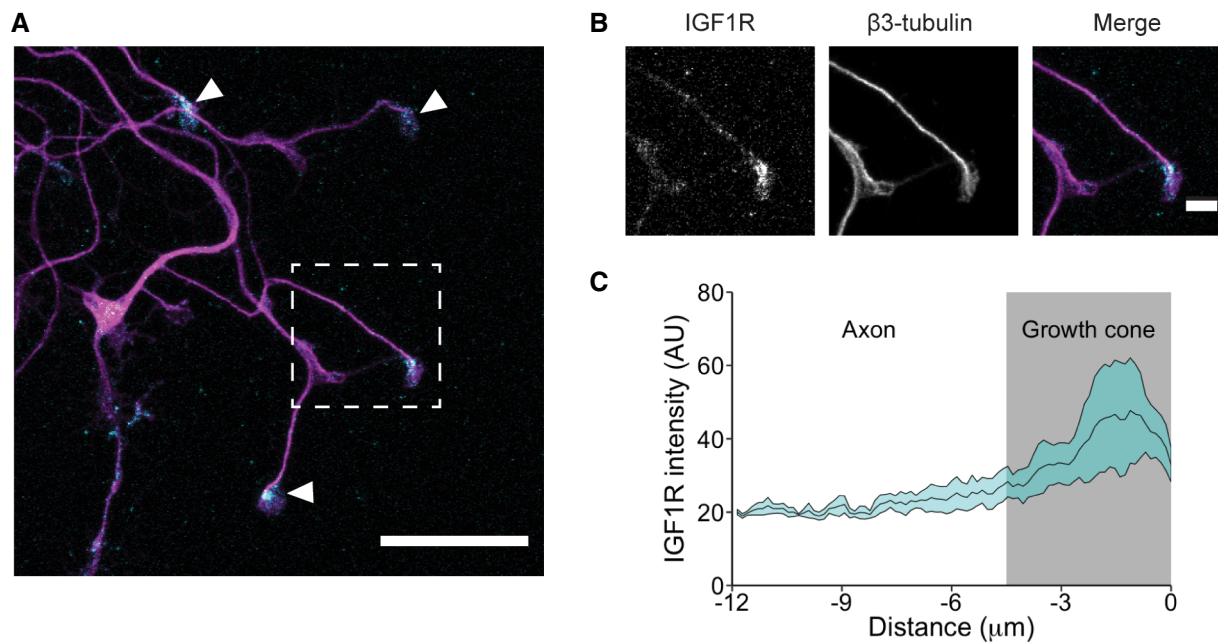


Figure EV1. IGF1R localizes to the axonal growth cone.

- A Example image of DIV6 PMN labelled with $\beta 3$ -tubulin and IGF1R. White arrowheads point to axonal growth cones. Box shows zoomed region in B). The scale bar length is 20 μm .
- B IGF1R staining at the growth cones of axons. The scale bar length is 4 μm .
- C Quantification of IGF1R intensity along the axon. IGF1R accumulates at the growth cone of axonal processes (72 cells, $N=3$ independent experiment). All data shown are mean \pm SEM.

Figure EV2. PPP downregulates Akt signalling.

- A pAkt staining in PMN treated with PPP or IGF1. The scale bar is 10 μm .
- B Image quantification. PPP treatment reduces the mean pAkt intensity in PMN ($*P = 0.046$, Student's t -test, 32 cells for each condition, $N = 3$ independent experiment). IGF1 caused no significant change in pAkt levels in PMN ($P = 0.25$, Student's t -test, 32 cells for each condition, $N = 3$ independent experiments). All data shown are mean \pm SEM.
- C Representative Western blot showing the level of TSC2 phosphorylation in PMN and N2A cells after 1 μM Capiasertib and Ipatasertib treatment for 60 min. GAPDH was used as a loading control.
- D Quantification of Western blot data. PMN and N2A cells were treated with either 1 nM, 10 nM, 100 nM or 1 μM of Capiasertib and Ipatasertib for 60 min. PMNs show decreased level of TSC2 phosphorylation after treatment. Capiasertib and Ipatasertib (1 μM) are $N = 2$ independent experiments. All other conditions are $N = 3$ independent experiments; data shown are mean \pm SEM.
- E Quantification of Western blot data. N2A shows decreased level of TSC2 phosphorylation after treatment with Capiasertib and Ipatasertib. Capiasertib 10 nM and 100 nM are $N = 3$ independent experiments; all other conditions are $N = 4$ independent experiments; data shown are mean \pm SEM.

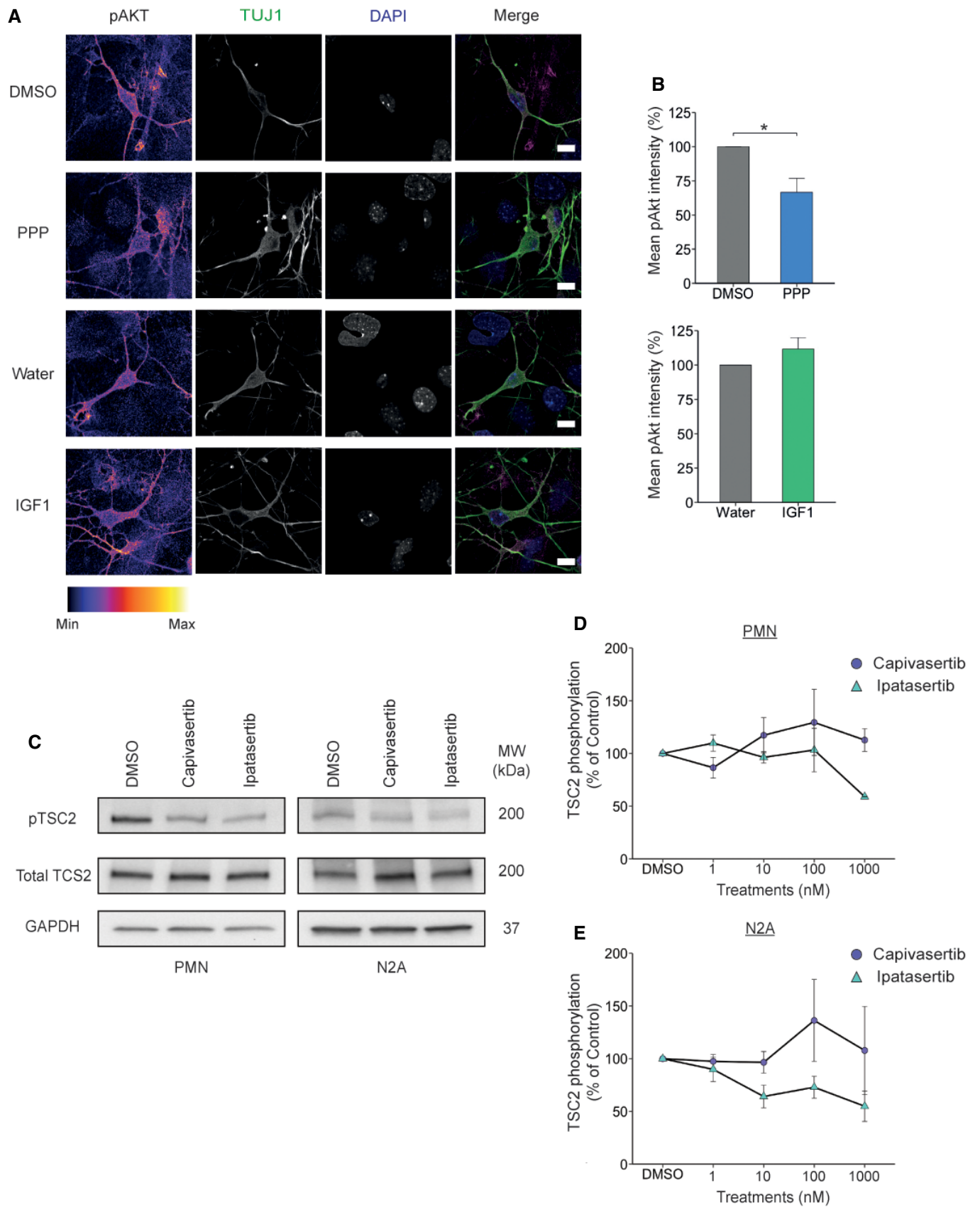


Figure EV2.

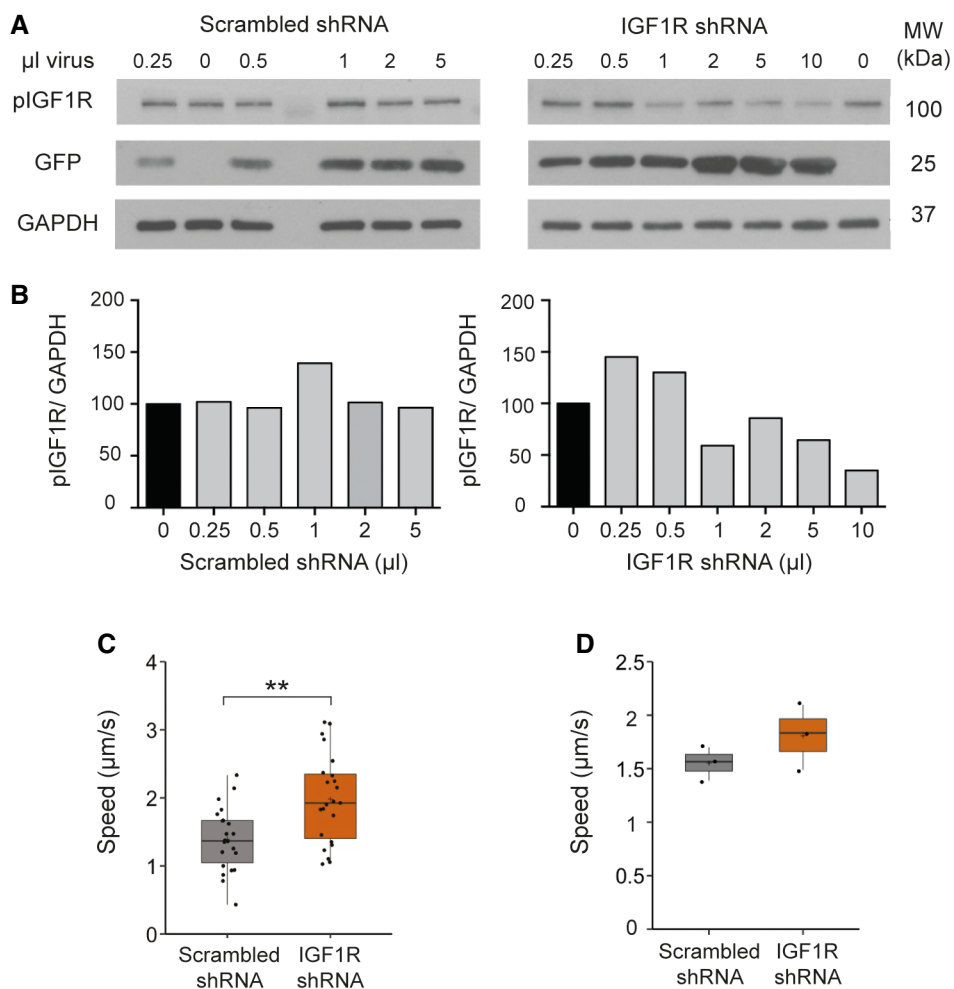


Figure EV3. shRNA knockdown of IGF1R in PMN.

- A Western blot of pIGF1R after lentiviral delivery of a shRNA targeting IGF1R or scrambled control in PMN. Cells were lysed 6 days after treatment.
- B The quantification of the Western blots in A). The shRNA viral construct effectively decreases levels of pIGF1R compared to scrambled control ($N = 1$ independent experiments).
- C Graph shows the average velocity of H₂T-containing organelles per axon after IGF1R knockdown (** $P = 0.001$, Student's t -test, $N = 3$ independent experiments; boxplot shows median, first and third quartiles. Upper/lower whiskers extend to 1.5 * the interquartile range).
- D Graph shows the average velocity of H₂T-containing organelles per experiment after IGF1R knockdown ($P = 0.26$, Student's t -test, $N = 3$ independent experiments; boxplot shows median, first and third quartiles. Upper/lower whiskers extend to 1.5 * the interquartile range).

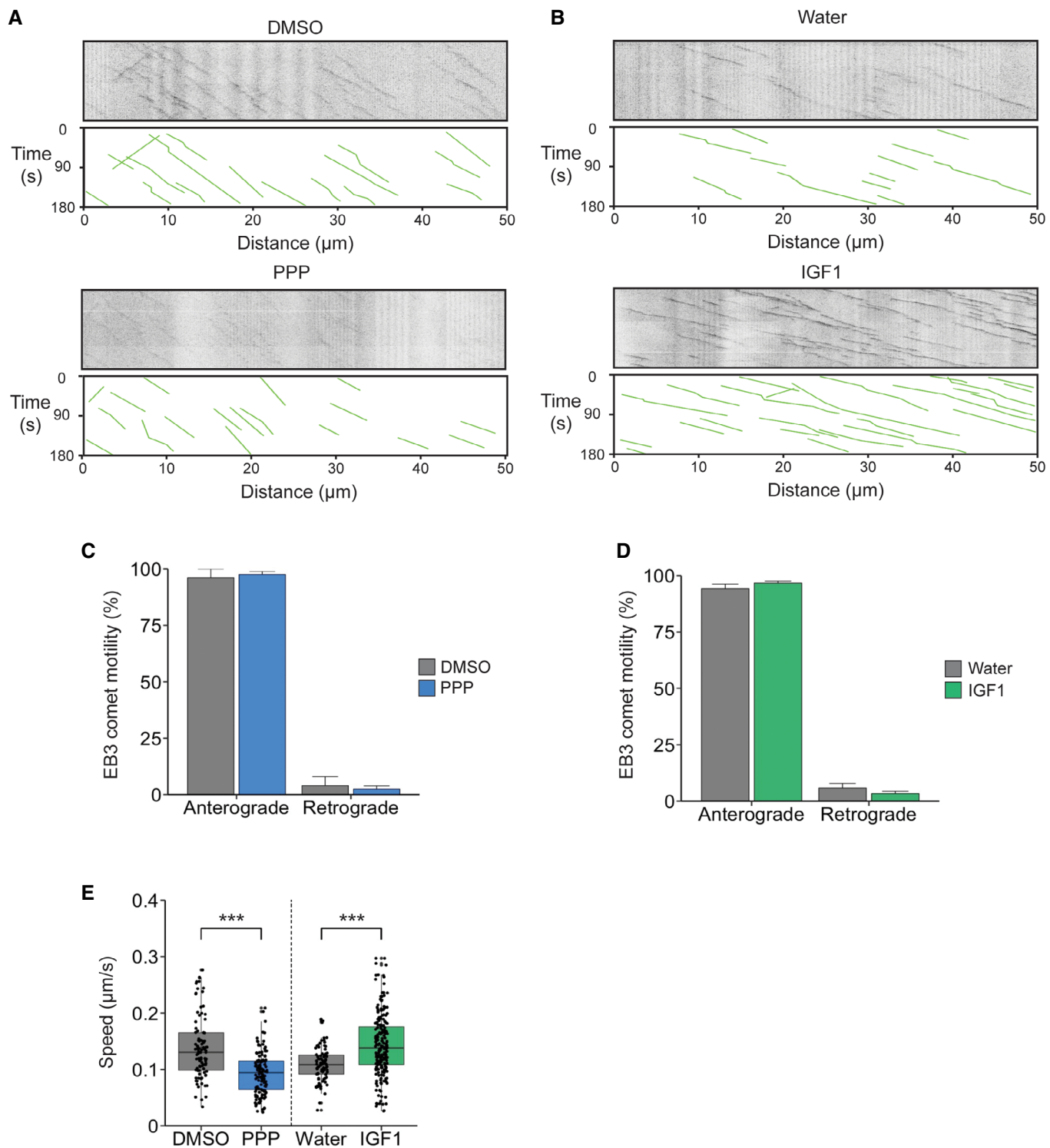


Figure EV4.

Figure EV4. IGF1R influences microtubule dynamics but does not alter polarity.

- A Kymographs of GFP-EB3 comets after treatment with 1 μ M PPP or DMSO for 45 min.
- B Kymographs of GFP-EB3 comets after treatment with 50 ng/ml IGF1 or water for 45 min.
- C Quantification of EB3-GFP directionality after PPP treatment. There was no change in microtubule polarity (anterograde: DMSO- 96.1 \pm 3.9%; PPP- 97.6 \pm 1.3%, $P = 0.77$; retrograde: DMSO-3.9 \pm 3.9% versus 2.4 \pm 1.3%, $P = 0.77$, Student's t -test, $N = 4$ independent experiments (PPP), $N = 2$ independent experiments (DMSO); data shown are mean \pm SEM).
- D Quantification of EB3-GFP directionality after IGF1 treatment. IGF1 also caused no change in polarity of new microtubules (anterograde: water- 96.6 \pm 1.0%; IGF1- 94.2 \pm 2.0%, $P = 0.23$; retrograde: water- 3.4 \pm 1.0% versus 5.8 \pm 2.0%, $P = 0.23$, Student's t -test, $N = 4$ independent experiments (IGF1), 2 independent experiments (water); data shown are mean \pm SEM).
- E Analysis of GFP-EB3 comet velocity after treatment with PPP or IGF1. PPP significantly reduced the average velocity of GFP-EB3 comets compared to controls (DMSO, 91 EB3 comets, 10 axons; PPP, 138 comets, 23 axons; *** $P = 2.6 \times 10^{-10}$, $N = 4$ independent experiments, Student's t -test). IGF1 increased the growth rate of GFP-EB3 comets compared to controls (water, 85 comets, 14 axons; IGF1, 213 comets, 25 axons; *** $P = 1.7 \times 10^{-10}$, $N = 4$ independent experiments, Student's t -test; boxplot shows median, first and third quartiles. Upper/lower whiskers extend to 1.5 * the interquartile range).

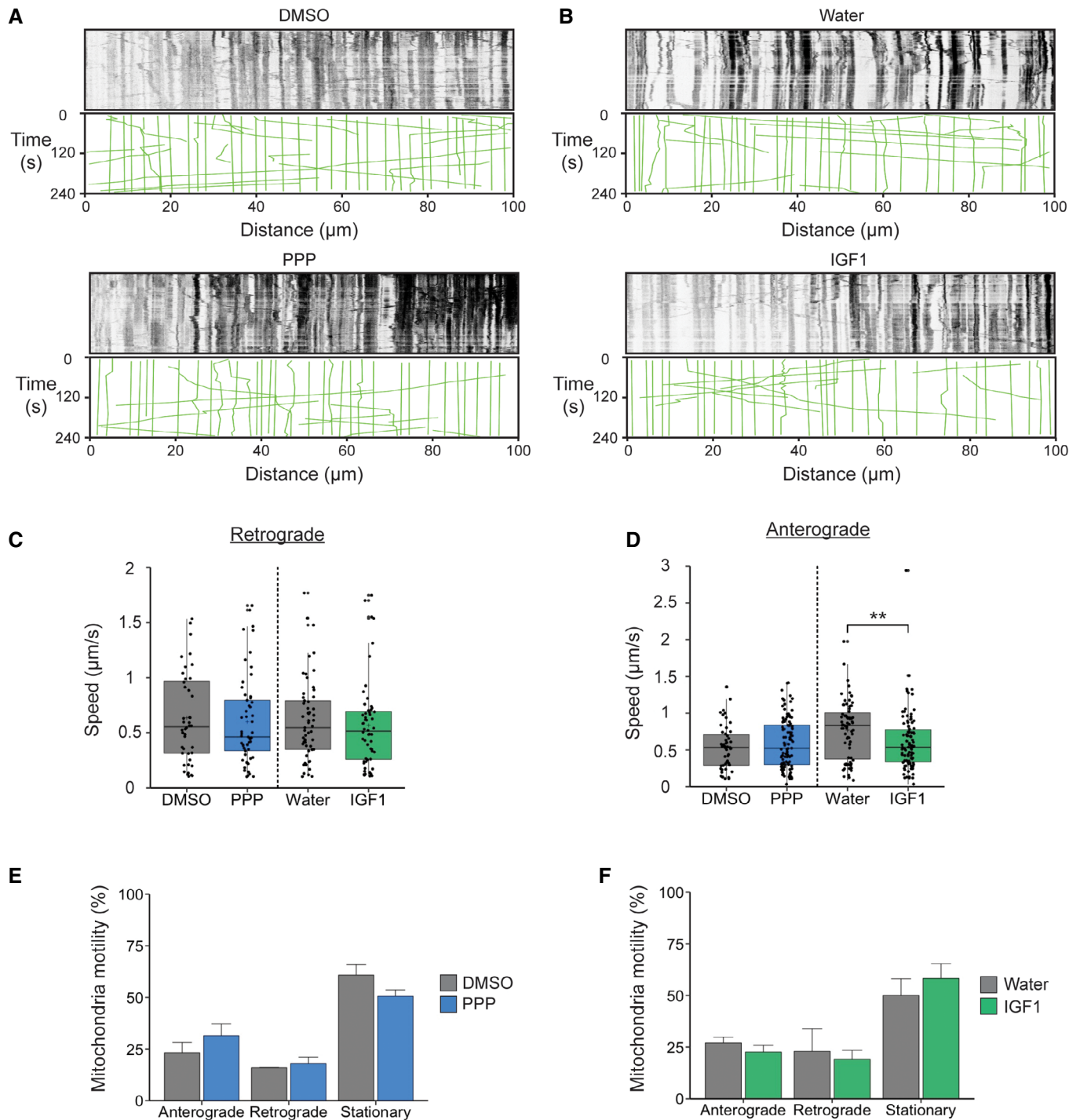


Figure EV5.

Figure EV5. IGF1R activity does not impact on the axonal trafficking of mitochondria.

- A Kymographs of TMRM-labelled mitochondria in DIV 6 PMN after treatment with 1 μ M PPP or DMSO for 45 min.
- B Kymographs of TMRM-labelled mitochondria in DIV 6 primary motor neurons after treatment with 50 ng/ml IGF1 or water for 45 min.
- C IGF1R modulation does not impact on the retrograde transport of mitochondria (DMSO: 0.64 ± 0.06 μ m/s, 45 organelles, $N = 2$ independent experiments; PPP: 0.60 ± 0.05 μ m/s, 57 organelles, $N = 3$ independent experiments; $P = 0.66$, Student's t -test; water: 0.61 ± 0.05 μ m/s, 57 organelles, $N = 3$ independent experiments; IGF1: 0.57 ± 0.05 μ m/s, 56 organelles, $N = 3$ independent experiments, $P = 0.56$, Student's t -test; boxplot shows median, first and third quartiles. Upper/lower whiskers extend to 1.5 * the interquartile range).
- D Effect of IGF1R treatment on anterograde mitochondrial velocity. PPP caused no significant change in transport rates (DMSO: 0.54 ± 0.04 μ m/s, 50 organelles, $N = 2$ independent experiments; PPP: 0.59 ± 0.03 μ m/s, 105 organelles, $N = 3$ independent experiments; $P = 0.32$, Student's t -test). In contrast, IGF1 treatment significantly slowed down anterograde mitochondria moving anterogradely (water: 0.76 ± 0.05 μ m/s, 75 organelles, $N = 3$ independent experiments; IGF1: 0.60 ± 0.04 μ m/s, 98 organelles; ** $P = 0.009$, Student's t -test, $N = 3$ independent experiments; boxplot shows median, first and third quartiles. Upper/lower whiskers extend to 1.5 * the interquartile range).
- E Quantification shows that the directionality of mitochondria transport after PPP treatment is not affected (anterograde: DMSO, $23.17 \pm 5.0\%$, 93 organelles, $N = 2$ independent experiments; PPP, $31.46 \pm 5.6\%$, 177 organelles, $N = 3$ independent experiments, $P = 0.35$; retrograde: DMSO, $16.02 \pm 0.13\%$, 64 organelles, $N = 2$ independent experiments, PPP, $17.98 \pm 3.0\%$, 105 organelles, $N = 3$ independent experiments, $P = 0.58$; stationary: DMSO, $60.8 \pm 5.1\%$, 248 organelles, $N = 2$ independent experiments; PPP, $50.55 \pm 3.1\%$, 291 organelles, $N = 3$ independent experiments, $P = 0.24$, Student's t -test; data shown are mean \pm SEM).
- F Quantification shows that IGF1 treatment does not alter the directionality of mitochondria (anterograde: water, $26.99 \pm 2.9\%$, 119 organelles, $N = 3$ independent experiments; IGF1, $22.58 \pm 3.4\%$, 136 organelles, $N = 3$ independent experiments, $P = 0.39$; retrograde: water, $22.94 \pm 10.9\%$, 95 organelles, $N = 3$ independent experiments; IGF1, $19.2 \pm 4.3\%$, 115 organelles, $P = 0.79$; stationary: water, $50.1 \pm 8.0\%$, 222 organelles, $N = 3$ independent experiments; IGF1, $58.25 \pm 7.1\%$, 340 organelles, $N = 3$ independent experiments, $P = 0.51$, Student's t -test; data shown are mean \pm SEM).