

### **Expanded View Figures**

### Figure EV1. MZgrk2 shows severe Hh defects.

Numbers of Prox1a<sup>+ve</sup> cells in wild-type (WT), zygotic *grk2* mutants, MZ*grk2*, MZ*grk2* injected with Grk2-GFP and Grk2 ATG morphant embryos at 30hpf (n = 8). The error bars indicate SD. Asterisks indicate statistically significant differences based on unpaired Student's *t*-test. \*\*\*\*P < 0.0001.



- A Western blot analysis of wild-type and mutant forms of Smo expressed from bicistronic myc-tagged mSmo and GFP constructs injected into wild-type zebrafish embryos. The histogram shows the quantification of Myc-tagged mSmo vs. GFP (n = 3). The error bars indicate SD.
- B Normalised expression of *ptch2* in 18hpf wild-type (AB), Cyc(A)-treated embryos and embryos injected with mRNA encoding Shh or various Smo mutants. Individual data points (blue crosses) are shown; mean is indicated by thick solid line, and SD by thin bars. Unpaired Student's *t*-test was used to determine the statistical significance between control (AB) and experimental embryos. \*\*\*P < 0.001: \*\*P < 0.01: and n.s. (not significant).
- experimental embryos. \*\*\**P* < 0.001; \*\**P* < 0.01; and n.s. (not significant). C Prox1a and Eng expression in *smo<sup>h1640</sup>* mutant embryos injected with mRNA encoding mSmoSD (*n* = 7) and mSmoKRA (*n* = 7). These mutant embryos show full recovery of SSFs and MPs. Embryos shown are at 30hpf. Scale bar, 50  $\mu$ m.







## Figure EV3. Smo translocates to the primary cilium independently of $\ensuremath{\mathsf{Grk2}}$ .

Accumulation of endogenous Smo in the PC of wild-type and  $Grk2^{-/-}$  FIp-In-3T3 cells after 4 h of Shh-N and SAG treatment. The medians with interquartile range of the fluorescence intensities of endogenous Smo are shown, quantified from multiple cilia (n = 150). Data were analysed using Kruskal–Wallis non-parametric ANOVA. \*\*\*\*P < 0.001 and n.s. (not significant).



# $\frac{mGrk2}{44} \xrightarrow{mGrk3}{44}$ $\frac{mGrk3}{44} \xrightarrow{\text{C}}{1000}$ 1000- 500-

# Figure EV4. MZgrk2 is fully rescued by mRNA encoding Grk2<sup>K220R</sup> and Grk2 domain swapping assay.

- A *In situ* hybridisation of *ptch2* and *nkx2.2a* in 24hpf wild-type (left column) and MZ*grk2* embryos expressing Grk2<sup>K220R</sup>-GFP (right column) (n = 30 for each sample). Scale bar, 200 µm.
- B Prox1a (green) and Eng (red) expression in the myotome of 30hpf MZgrk2 embryos injected with mRNA encoding Grk2<sup>ΔRGS</sup>-GFP, Grk2<sup>Δkinase</sup>-GFP, Grk2<sup>ΔPH</sup>-GFP, Grk2<sup>ΔCmargin</sup>-GFP and Grk2<sup>ΔNmargin</sup>-GFP. Merge images show nuclei labelled with DAPI (blue) (n = 20). Scale bar, 50 µm.

Figure EV5. Grk3 is not expressed in NIH3T3 cells. RT–PCR assay of Grk2 and Grk3 expression in MEF and NIH 3T3 cells.