

Supplementary Materials for

Aberrant Rac1-cofilin signaling mediates defects in dendritic spines, synaptic function, and sensory perception in fragile X syndrome

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Somatosensory Cortex Synaptosomes (1wk)

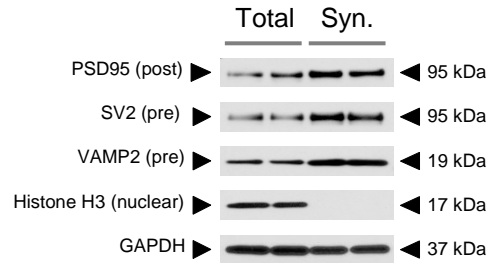


Figure S1. Validation of somatosensory cortex synapse enrichment. Representative western blot from a 1-week-old wild-type (WT) mouse showing enrichment of presynaptic (VAMP2 and SV2) and postsynaptic (PSD95) markers as well the absence of a nuclear marker (Histone H3) in somatosensory cortex synaptosome fractions versus total lysates.

Synaptosomes (1 - 2 months)

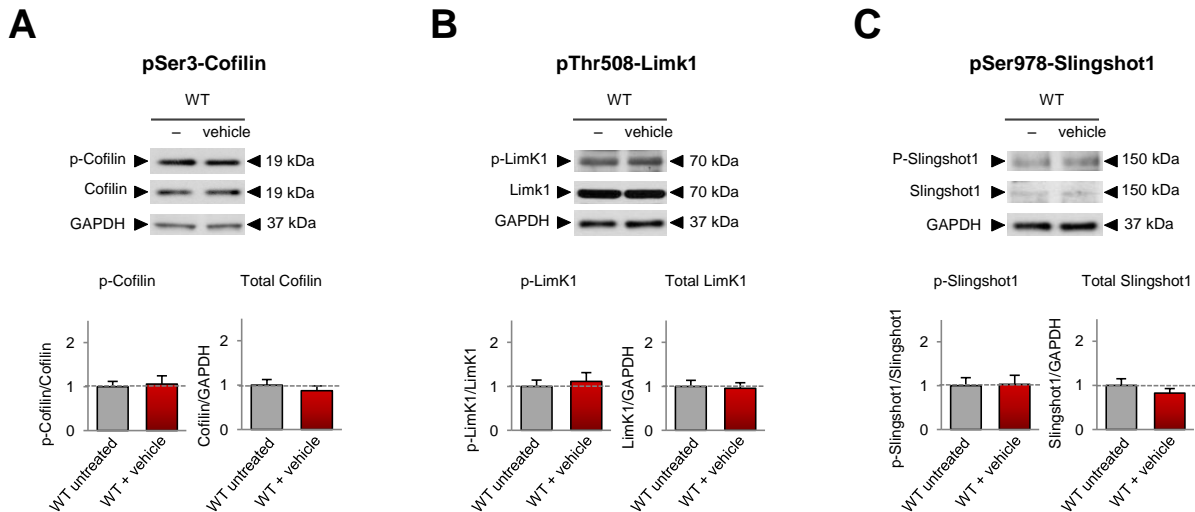


Figure S2. Subcutaneous administration of hydroxypropyl- β -cyclodextrin (vehicle) has little to no effect on cofilin signaling. (A to C) Representative Western blots (top) and summary data (bottom) from somatosensory synaptosomes of 1- to 2-month-old wild-type mice, either untreated (left) or treated with hydroxypropyl- β -cyclodextrin (vehicle; right). Synaptosomes from vehicle-treated animals exhibited no change in the abundance of phosphorylated (p-) cofilin (A), p-LIMK1 (B), and p-Slingshot1 (C) relative to those from untreated animals ($n = 7$ animals per group).

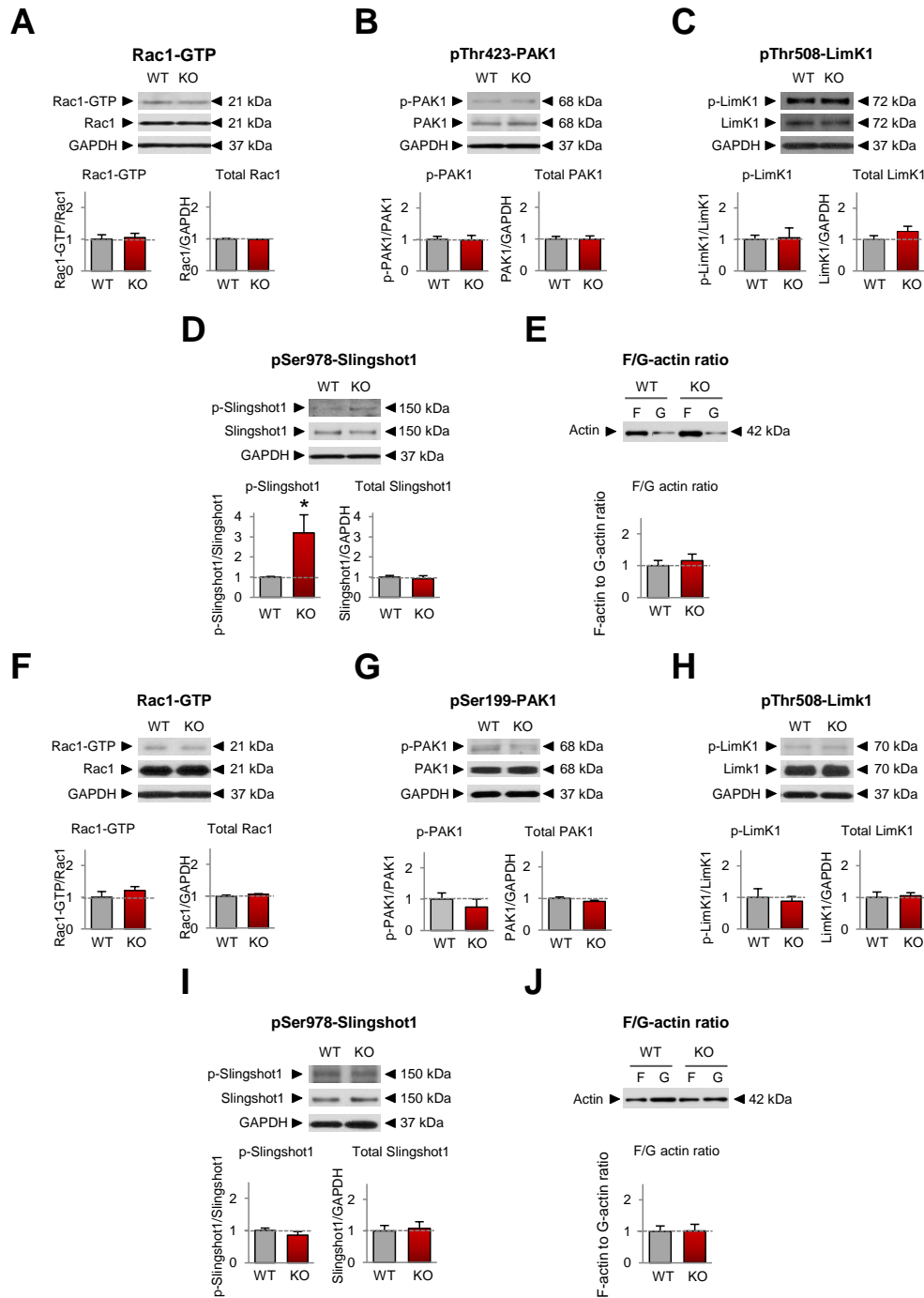


Figure S3. Mature *Fmr1* KO mice exhibit normal Rac1 signaling at somatosensory cortex synapses. (A to E) Representative Western blots (top) and summary data (bottom) from somatosensory lysates (A) or synaptosomes (B to E) of 4-week-old WT and *Fmr1* KO mice showing relative abundance of active, phosphorylated or total proteins in the Rac1 signaling pathway: (A) Rac1-GTP, (B) p-PAK1, (C) p-LIMK1, (D) p-Slingshot1, and (E) F-actin/G-actin ratio ($n = 5 - 12$ animals per group). (F to J) Representative western blots (top) and summary data (bottom) from somatosensory lysates (F) or synaptosomes (G to J) of 2- to 5-month-old WT and *Fmr1* KO mice showing relative abundance of active, phosphorylated or total proteins in the Rac1 signaling pathway, as in (A to E) ($n = 4 - 10$ animals per group). Data are mean \pm SEM. * $p < 0.05$.

Synaptosomes (2 - 5 months)

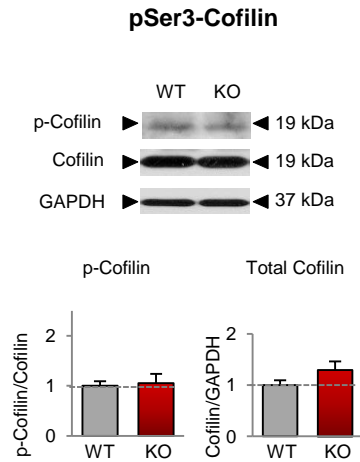


Figure S4. Mature *Fmr1* KO mice exhibit normal cofilin phosphorylation and activity at somatosensory cortex synapses. (A) Representative Western blots and summary data of cofilin phosphorylation in somatosensory synaptosomes from 2- to 5-month-old WT and *Fmr1* KO mice (9 – 14 mice per group). Membranes were probed with antibodies to p-cofilin and total cofilin.

Table S1. Summary of statistics for Fig. 1.

Figure	N	Mean \pm SEM	p value	Statistical Test
1A	WT (p-Cofilin) = 12 animals	1.000 \pm 0.2277	–	Two-tailed unpaired <i>t</i> -test, WT vs KO
	KO (p-Cofilin) = 12 animals	1.764 \pm 0.2814	*, 0.046	
	WT (Cofilin) = 12 animals	1.000 \pm 0.05896	–	
	KO (Cofilin) = 12 animals	1.080 \pm 0.05266	0.327	
1B	WT (p-Cofilin) = 7 animals	1.000 \pm 0.1544	–	
	KO (p-Cofilin) = 7 animals	1.115 \pm 0.2765	0.729	
	WT (Cofilin) = 7 animals	1.000 \pm 0.1190	–	
	KO (Cofilin) = 7 animals	0.9104 \pm 0.1131	0.5952	
1C	WT (p-Cofilin) = 12 animals	1.000 \pm 0.1458	–	
	KO (p-Cofilin) = 10 animals	1.958 \pm 0.4158	*, 0.030	
	WT (Cofilin) = 12 animals	1.000 \pm 0.0918	–	
	KO (Cofilin) = 10 animals	0.7626 \pm 0.0823	0.073	
1D	WT (p-Cofilin) = 10 animals	1.000 \pm 0.1355	–	
	KO (p-Cofilin) = 9 animals	1.035 \pm 0.2629	0.9038	
	WT (Cofilin) = 10 animals	1.000 \pm 0.1106	–	
	KO (Cofilin) = 9 animals	1.087 \pm 0.1015	0.5710	
1E	WT (F/G actin) = 5 animals	1.000 \pm 0.4336	–	
	KO (F/G actin) = 6 animals	3.495 \pm 0.7284	*, 0.021	
1G (% Filopodia)	WT = 3 animals / 12 dendrites	100.0 \pm 4.957	–	
	KO = 3 animals / 13 dendrites	126.2 \pm 5.838	***, 0.0009	
1G (% Density)	WT = 3 animals / 12 dendrites, 154 segments, 660 spines KO = 3 animals / 13 dendrites, 182 segments, 933 spines	100.0 \pm 7.941 131.1 \pm 11.82	– *, 0.0429	

Table S2. Summary of statistics for Fig. 2.

Figure	N	Mean \pm SEM	p value	Statistical Test
2A	WT (Rac1-GTP) = 4 animals	1.000 \pm 0.0396	–	Two-tailed unpaired <i>t</i> -test, WT vs KO
	KO (Rac1-GTP) = 4 animals	1.673 \pm 0.0396	****, <0.0001	
	WT (Rac1) = 4 animals	1.000 \pm 0.0327	–	
	KO (Rac1) = 4 animals	1.065 \pm 0.0320	0.207	
2B	WT (p-PAK1) = 12 animals	1.000 \pm 0.0904	–	
	KO (p-PAK1) = 15 animals	1.791 \pm 0.3295	*, 0.047	
	WT (PAK1) = 12 animals	1.000 \pm 0.0865	–	
	KO (PAK1) = 15 animals	1.109 \pm 0.1615	0.327	
2C	WT (p-PAK4) = 8 animals	1.000 \pm 0.1862	–	
	KO (p-PAK4) = 7 animals	1.133 \pm 0.3728	0.745	
	WT (PAK4) = 8 animals	1.000 \pm 0.1111	–	
	KO (PAK4) = 7 animals	1.126 \pm 0.1913	0.567	
2D	WT (p-LimK1) = 10 animals	1.000 \pm 0.0929	–	
	KO (p-LimK1) = 9 animals	2.047 \pm 0.4434	*, 0.020	
	WT (LimK1) = 10 animals	1.000 \pm 0.0730	–	
	KO (LimK1) = 9 animals	0.9765 \pm 0.1525	0.887	
2E	WT (p-SSH1) = 8 animals	1.000 \pm 0.1109	–	
	KO (p-SSH1) = 5 animals	1.764 \pm 0.3759	*, 0.037	
	WT (SSH1) = 8 animals	1.000 \pm 0.1424	–	
	KO (SSH1) = 5 animals	1.148 \pm 0.3567	0.673	

Table S3. Summary of statistics for Fig. 3.

Figure	N	Mean ± SEM	p value (p < 0.05)	Statistical Test
3A	WT vehicle (p-LimK1) = 13 animals	1.000 ± 0.2463	–	Two-way ANOVA with Tukey's post-test (WT vehicle vs KO vehicle, WT vehicle vs WT FRAX486 and KO vehicle vs KO FRAX486)
	KO vehicle (p-LimK1) = 12 animals	2.1853 ± 0.3478	*, 0.012	
	WT FRAX486 (p-LimK1) = 13 animals	0.9886 ± 0.2433	0.999	
	KO FRAX486 (p-LimK1) = 11 animals	0.9751 ± 0.1843	*, 0.016	
	WT vehicle (LimK1) = 13 animals	1.000 ± 0.1247	–	
	KO vehicle (LimK1) = 12 animals	0.9795 ± 0.1466	0.999	
	WT FRAX486 (LimK1) = 13 animals	1.0094 ± 0.1750	0.999	
	KO FRAX486 (LimK1) = 11 animals	1.1066 ± 0.2149	0.953	
3B	WT vehicle (p-SSH1) = 12 animals	1.000 ± 0.3312	–	
	KO vehicle (p-SSH1) = 10 animals	2.3299 ± 0.2033	**, 0.003	
	WT FRAX486 (p-SSH1) = 8 animals	1.3092 ± 0.2307	0.847	
	KO FRAX486 (p-SSH1) = 10 animals	1.1526 ± 0.2792	*, 0.013	
	WT vehicle (SSH1) = 12 animals	1.000 ± 0.3930	–	
	KO vehicle (SSH1) = 10 animals	0.8570 ± 0.3334	0.986	
	WT FRAX486 (SSH1) = 8 animals	0.7212 ± 0.1630	0.910	
	KO FRAX486 (SSH1) = 10 animals	0.7692 ± 0.1949	0.999	
3C	WT vehicle (p-Cofilin) = 14 animals	1.0000 ± 0.2270	–	
	KO vehicle (p-Cofilin) = 14 animals	2.0817 ± 0.2463	**, 0.006	
	WT FRAX486 (p-Cofilin) = 15 animals	0.9440 ± 0.2045	0.998	
	KO FRAX486 (p-Cofilin) = 9 animals	0.7251 ± 0.2293	**, 0.002	
	WT vehicle (Cofilin) = 14 animals	1.0000 ± 0.2013	–	
	KO vehicle (Cofilin) = 14 animals	1.0703 ± 0.2069	0.995	
	WT FRAX486 (Cofilin) = 15 animals	0.8696 ± 0.1850	0.966	
	KO FRAX486 (Cofilin) = 9 animals	1.0339 ± 0.2354	0.999	
3D	WT vehicle (F/G actin) = 10 animals	1.0000 ± 0.1430	–	
	KO vehicle (F/G actin) = 9 animals	1.9193 ± 0.3043	*, 0.012	
	WT FRAX486 (F/G actin) = 10 animals	1.0028 ± 0.1529	0.999	
	KO FRAX486 (F/G actin) = 7 animals	1.0919 ± 0.1543	*, 0.048	

Table S4. Summary of statistics for Fig. 4.

Figure	N	Mean ± SEM	p value (p < 0.05)	Statistical Test
4B (cultured neurons)	<i>3 independent experiments</i> No virus (F/G actin) = 14 wells GFP (F/G actin) = 13 wells WT-Cofilin (F/G actin) = 15 wells Cofilin S3A (F/G actin) = 13 wells Cofilin S3D (F/G actin) = 15 wells	1.0000 ± 0.1266 1.1017 ± 0.1602 0.9950 ± 0.1625 0.4280 ± 0.0750 1.0938 ± 0.1421	– 0.986 0.998 *, 0.042 0.988	One-way ANOVA with Tukey's post-test (No virus vs Cofilin S3A, No virus vs GFP, No virus vs Cofilin S3A, No virus vs Cofilin S3D)
4D (Spine length)	WT (GFP) = 10 animals, 21 neurons, 1286 spines, 196 segments of 10 µm KO (GFP) = 8 animals, 20 neurons, 1381 spines, 170 segments of 10 µm WT (WT. Cof.) = 7 animals, 15 neurons, 765 spines, 119 segments of 10 µm KO (WT. Cof.) = 9 animals, 15 neurons, 911 spines, 123 segments of 10 µm WT (S3A) = 6 animals, 16 neurons, 887 spines, 132 segments of 10 µm KO (S3A) = 7 animals, 17 neurons, 1119 spines, 151 segments of 10 µm	1.1442 ± 0.0532 1.3326 ± 0.0494 1.0087 ± 0.0403 1.2496 ± 0.0519 0.9462 ± 0.0364 0.9298 ± 0.0520	– *, 0.043 0.370 *, 0.019 *, 0.047 ****, <0.0001	Two-way ANOVA with Tukey's post-test (WT (GFP) vs KO (GFP), WT (GFP) vs WT (S3A), WT (GFP) vs WT (WT. Cof.), KO (GFP) vs KO (S3A), and WT (WT. Cof.) vs KO (WT. Cof.))
4E (Spine head width)	WT (GFP) – same as above KO (GFP) WT (WT. Cof.) KO (WT. Cof.) WT (S3A) KO (S3A)	0.7354 ± 0.0129 0.6509 ± 0.0221 0.6911 ± 0.0139 0.6046 ± 0.0146 0.7188 ± 0.0249 0.6356 ± 0.0191	– **, 0.009 0.541 *, 0.033 0.987 *, 0.031	
4F (Length-width ratio)	WT (GFP) – same as above KO (GFP) WT (WT. Cof.) KO (WT. Cof.) WT (S3A) KO (S3A)	1.6570 ± 0.0709 2.2756 ± 0.0868 1.6022 ± 0.0709 2.2876 ± 0.1141 1.4411 ± 0.0607 1.5550 ± 0.0773	– ****, <0.0001 0.899 ****, <0.0001 0.398 ****, <0.0001	
4G (%Stubby/Mushroom)	WT (GFP) – same as above KO (GFP) WT (WT. Cof.) KO (WT. Cof.) WT (S3A) KO (S3A)	0.7823 ± 0.0204 0.6568 ± 0.0177 0.7892 ± 0.0193 0.6106 ± 0.0218 0.8405 ± 0.0136 0.8109 ± 0.0222	– ****, <0.0001 0.999 ****, <0.0001 0.269 ****, <0.0001	
4H (%Thin/Filopodia)	WT (GFP) – same as above KO (GFP) WT (WT. Cof.) KO (WT. Cof.) WT (S3A) KO (S3A)	0.2127 ± 0.0201 0.3417 ± 0.0175 0.2108 ± 0.0193 0.3881 ± 0.0220 0.1595 ± 0.0136 0.1808 ± 0.0214	– ****, <0.0001 0.999 ****, <0.0001 0.355 ****, <0.0001	
4I (Spine density)	WT (GFP) – same as above KO (GFP) WT (WT. Cof.) KO (WT. Cof.) WT (S3A) KO (S3A)	6.8827 ± 0.3208 8.1235 ± 0.3485 6.1092 ± 0.3035 8.3008 ± 0.3801 6.7197 ± 0.3395 6.5762 ± 0.2974	– *, 0.046 0.583 ***, 0.0005 0.999 **, 0.009	

Table S5. Summary of statistics for Fig. 5.

Figure	N	Mean ± SEM	p value (p < 0.05)	Statistical Test
5B (NMDA/AM PA ratio)	WT vehicle = 13 animals	0.669 ± 0.054	–	Two-way ANOVA with Tukey's post-test (WT vehicle vs. KO vehicle, KO FRAX486)
	KO vehicle = 11 animals	1.531 ± 0.288	** , 0.0014	
	WT FRAX486 = 11 animals	0.727 ± 0.042	–	
	KO FRAX486 = 6 animals	0.503 ± 0.710	0.8297	
5D (mEPSC amplitude)	WT vehicle = 10 animals	25.65 ± 2.273	–	
	KO vehicle = 10 animals	21.60 ± 1.317	0.4386	
	WT FRAX486 = 11 animals	26.28 ± 1.369	–	
	KO FRAX486 = 9 animals	29.36 ± 2.521	* , 0.0365	
5D (mEPSC frequency)	WT vehicle = 10 animals	1.014 ± 0.131	–	Two-way ANOVA with Tukey's post-test (WT vehicle vs KO vehicle, WT vehicle vs WT FRAX486, KO vehicle vs KO FRAX486)
	KO vehicle = 10 animals	0.650 ± 0.095	0.1577	
	WT FRAX486 = 11 animals	0.471 ± 0.111	* , 0.0119	
	KO FRAX486 = 9 animals	0.655 ± 0.141	0.9999	
5E (NMDA/AM PA ratio)	WT vehicle = 15 animals	0.606 ± 0.068	–	
	KO vehicle = 7 animals	0.431 ± 0.065	0.3704	
	WT FRAX486 = 11 animals	0.571 ± 0.048	0.9821	
	KO FRAX486 = 10 animals	0.599 ± 0.089	0.4678	
5E (mEPSC amplitude)	WT vehicle = 18 animals	29.21 ± 1.273	–	
	KO vehicle = 7 animals	27.79 ± 1.251	0.9301	
	WT FRAX486 = 16 animals	22.67 ± 1.379	** , 0.004	
	KO FRAX486 = 12 animals	22.14 ± 1.635	0.1254	
5E (mEPSC frequency)	WT vehicle = 18 animals	2.472 ± 0.426	–	
	KO vehicle = 7 animals	2.454 ± 0.212	0.9999	
	WT FRAX486 = 16 animals	2.873 ± 0.761	0.9452	
	KO FRAX486 = 12 animals	1.920 ± 0.391	0.9505	

Table S6. Summary of statistics for Fig. 6.

Figure	N	Mean ± SEM	p value (p < 0.05)	Statistical Test
6B	WT (familiar) = 13 animals	42.05 ± 3.276	–	Two-tailed unpaired <i>t</i> test for WT (familiar) vs WT (novel) and KO (familiar) vs KO (novel)
	WT (novel) = 13 animals	58.96 ± 2.663	*** , 0.0005	
	KO (familiar) = 7 animals	48.54 ± 4.097	–	
	KO (novel) = 7 animals	51.46 ± 4.097	0.6259	
6C	WT (familiar) = 10 animals	47.50 ± 2.435	–	
	WT (novel) = 10 animals	52.60 ± 2.449	0.1569	
6D	WT (familiar) = 7 animals	52.54 ± 4.100	–	
	WT (novel) = 7 animals	47.46 ± 4.100	0.3989	
6E	WT (familiar) = 7 animals	38.13 ± 2.872	–	
	WT (novel) = 7 animals	61.87 ± 2.872	**** , <0.0001	
	KO (familiar) = 8 animals	41.08 ± 3.631	–	
	KO (novel) = 8 animals	58.92 ± 3.631	** , 0.003	
6F	WT-vehicle (familiar) = 15 animals	43.57 ± 2.705	–	
	WT-vehicle (novel) = 15 animals	56.43 ± 2.705	** , 0.0022	
	KO-vehicle (familiar) = 10 animals	51.79 ± 3.446	–	
	KO-vehicle (novel) = 10 animals	48.71 ± 2.997	0.5089	
	WT-FRAX486 (familiar) = 9 animals	41.44 ± 4.219	–	
	WT- FRAX486 (novel) = 9 animals	58.83 ± 4.344	* , 0.0111	
	KO- FRAX486 (familiar) = 10 animals	40.19 ± 3.188	–	
	KO- FRAX486 (novel) = 10 animals	59.81 ± 3.188	*** , 0.0003	

Table S7. Summary of statistics for fig. S2.

Figure	N	Mean ± SEM	p value (p < 0.05)	Statistical Test
Supp. 2A	WT untreated (p-Cofilin) = 7 animals	1.000 ± 0.1259	–	Two-tailed unpaired <i>t</i> test for WT untreated vs WT vehicle treated animals
	WT vehicle (p-Cofilin) = 7 animals	1.060 ± 0.2010	0.8031	
	WT untreated (Cofilin) = 7 animals	1.000 ± 0.1360	–	
	WT vehicle (Cofilin) = 7 animals	0.8786 ± 0.1109	0.5023	
Supp. 2B	WT untreated (p-LimK1) = 7 animals	1.000 ± 0.1525	–	
	WT vehicle (p-LimK1) = 7 animals	1.116 ± 0.2125	0.6659	
	WT untreated (LimK1) = 7 animals	1.000 ± 0.1458	–	
	WT vehicle (LimK1) = 7 animals	0.9594 ± 0.1327	0.8402	
Supp. 2C	WT untreated (p-SSH1) = 7 animals	1.000 ± 0.1953	–	
	WT vehicle (p-SSH1) = 7 animals	1.028 ± 0.2255	0.9270	
	WT untreated (SSH1) = 7 animals	1.000 ± 0.1571	–	
	WT vehicle (SSH1) = 7 animals	0.8207 ± 0.1127	0.3718	

Table S8. Summary of statistics for figs. S3 and S4.

Figure	N	Mean ± SEM	p value	Statistical Test
3A	WT (Rac1-GTP) = 8 animals	1.000 ± 0.1437	–	Two-tailed unpaired <i>t</i> test for WT vs KO
	KO (Rac-GTP) = 7 animals	1.052 ± 0.1341	0.7984	
	WT (Rac1) = 8 animals	1.000 ± 0.01174	–	
	KO (Rac1) = 7 animals	0.9846 ± 0.0082	0.3136	
3B	WT (p-PAK1) = 8 animals	1.000 ± 0.0937	–	
	KO (p-PAK1) = 8 animals	0.9798 ± 0.1460	0.9089	
	WT (PAK1) = 8 animals	1.000 ± 0.0786	–	
	KO (PAK1) = 8 animals	0.9904 ± 0.1071	0.9435	
3C	WT (p-LIMK1) = 10 animals	1.000 ± 0.1349	–	
	KO (p-LIMK1) = 9 animals	1.050 ± 0.3280	0.8849	
	WT (LIMK1) = 10 animals	1.000 ± 0.1205	–	
	KO (LIMK1) = 9 animals	1.253 ± 0.1664	0.2291	
3D	WT (p-SSH1) = 8 animals	1.000 ± 0.0716	–	
	KO (p-SSH1) = 7 animals	3.127 ± 0.9302	*, 0.0293	
	WT (SSH1) = 8 animals	1.000 ± 0.0795	–	
	KO (SSH1) = 7 animals	0.9207 ± 0.1603	0.6537	
3E	WT (F/G actin) = 5 animals	1.000 ± 0.0543	–	
	KO (F/G actin) = 5 animals	1.156 ± 0.1955	0.4592	
3F	WT (Rac1-GTP) = 7 animals	1.000 ± 0.1767	–	
	KO (Rac-GTP) = 5 animals	1.213 ± 0.1235	0.3944	
	WT (Rac1) = 7 animals	1.000 ± 0.3615	–	
	KO (Rac1) = 5 animals	1.054 ± 0.0246	0.2852	
3G	WT (p-PAK1) = 4 animals	1.000 ± 0.2757	–	
	KO (p-PAK1) = 6 animals	0.7542 ± 0.2850	0.5723	
	WT (PAK1) = 4 animals	1.000 ± 0.0708	–	
	KO (PAK1) = 6 animals	0.9089 ± 0.0372	0.2461	
3H	WT (p-LIMK1) = 8 animals	1.000 ± 0.2764	–	
	KO (p-LIMK1) = 5 animals	0.8735 ± 0.1535	0.7419	
	WT (LIMK1) = 8 animals	1.000 ± 0.1722	–	
	KO (LIMK1) = 5 animals	1.055 ± 0.0918	0.8034	
3I	WT (p-SSH1) = 9 animals	1.000 ± 0.0783	–	
	KO (p-SSH1) = 7 animals	0.8557 ± 0.1082	0.2863	
	WT (SSH1) = 9 animals	1.000 ± 0.1692	–	
	KO (SSH1) = 7 animals	1.079 ± 0.2167	0.7750	

3J	WT (F/G actin) = 9 animals	1.000 ± 0.1697	–	
	KO (F/G actin) = 10 animals	1.054 ± 0.2098	0.9599	
4A	WT (p-Cofilin) = 13 animals	1.000 ± 0.0925	–	
	KO (p-Cofilin) = 9 animals	1.052 ± 0.1864	0.7860	
	WT (Cofilin) = 13 animals	1.000 ± 0.0933	–	
	KO (Cofilin)= 9 animals	1.295 ± 0.1668	0.1097	