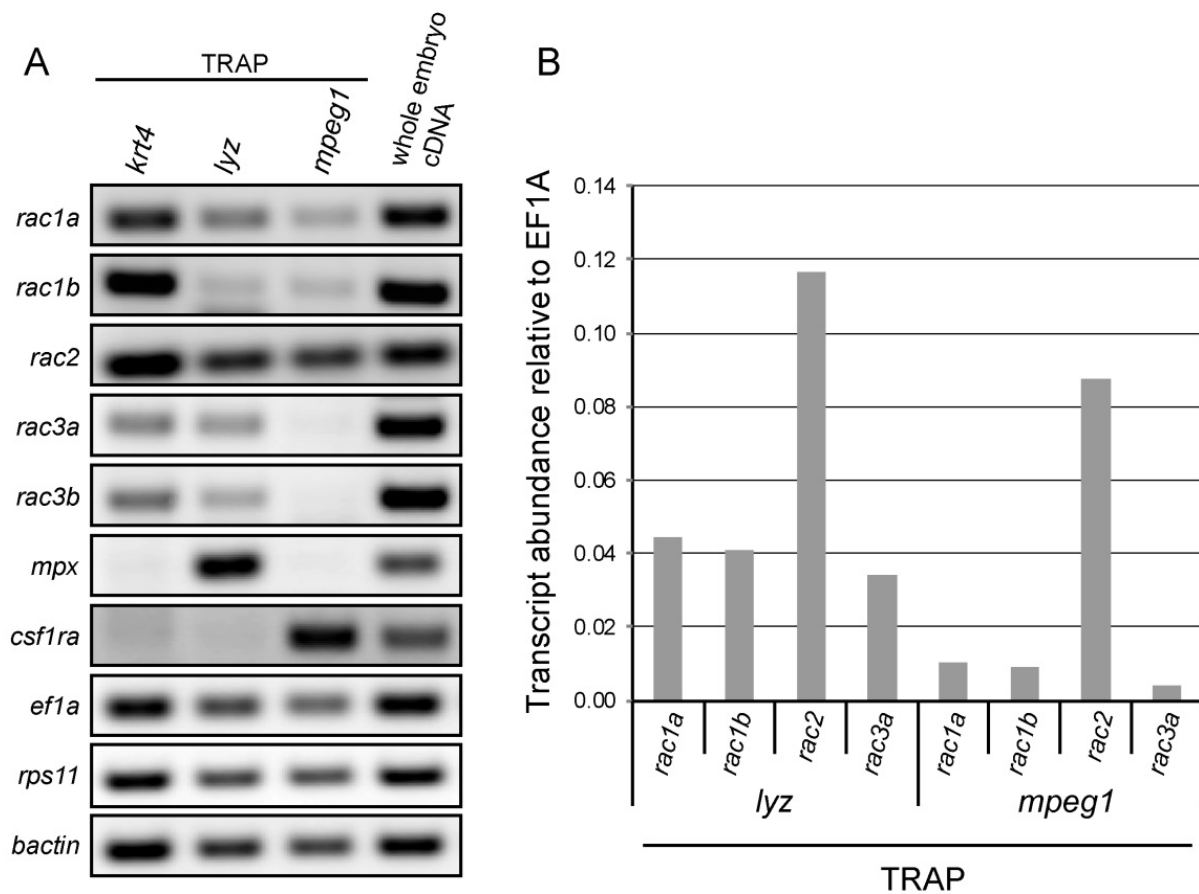
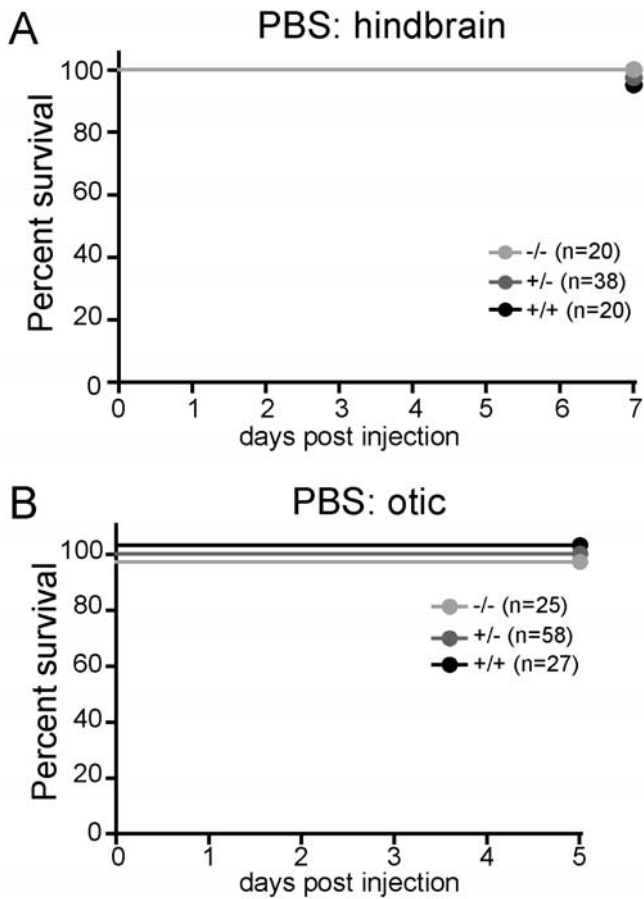


**Supplemental Figure 1. Rho GTPase expression in *rac2*<sup>-/-</sup> larvae.** RT-qPCR was performed on RNA from individual larvae from a *rac2*<sup>+/-</sup> in-cross. Data are from three pooled independent experiments; bars indicate calculated LS-adjusted means and SE. These experiments were done on the exon 5-B mutant line. Transcripts from *cdc42l2* and *rhoga* could not be amplified from samples.



**Supplemental Figure 2. *rac* expression in zebrafish larval neutrophils and macrophages.** TRAP was performed on pools of ~50 3 dpf larvae expressing EGFP-tagged L10a under the epithelial-specific *krt4* promoter, neutrophil-specific *lyz* promoter, or macrophage-specific *mpeg1* promoter. Total RNA was also isolated from whole embryos. Expression of genes was measured by RT-PCR (A) or RT-qPCR (B). This experiment was done three times, representative data are shown for both (A) and (B).



**Supplemental Figure 3. *rac2*<sup>-/-</sup> larvae do not succumb to control PBS injections.** **A.** Larvae 2 dpf from a *rac2*<sup>+/-</sup> in-cross were injected with PBS into the hindbrain. Survival was monitored for seven days. **B.** Larvae 3 dpf from a *rac2*<sup>+/-</sup> in-cross were injected with PBS into the otic vesicle and survival was monitored for five days. Data shown for each experiment are from three pooled independent replicates.

**Supplemental Table I. Primer sequences.**

Primer	Sequence (5' to 3')	Ref. (if previously published)
rac2_ex3_F	AACCAGTCAACCTGGGACTCT	
rac2_ex3_R	TGGTGTTTACCGTCTGCGG	
rac2_ex5_F	CACTGCCCTTCCACTCCAAT	
rac2_ex5_R	TCGGTGCCAGTTTCTTCTCC	
rac2_ex5_intronR	GACAAATCCAGCTCTGCCAC	
irf8_F	ACATAAGGCGTAGAGATTGGACG	(Shiau, Kaufman et al. 2015)
irf8_R	GAAACATAGTGCGGTCCCTCATCC	
qrac1a_F	TCCGGCCTCATTTGAAAACG	
qrac1a_R	GTGTCTTGTTCATCTCGCAA	
qrac1b_F	CTCTCTCGTACCCTCAGACG	
qrac1b_R	TGGGAGTGGTTTGACAGTGA	
qrac2_F	AATACCTGGAGTGTTCGGCC	
qrac2_R	GCCCTTCTTCTTGACCTTGG	
qrac3a_F	CCCAGGAGAGTATATACCCACAG	
qrac3a_R	GGTATGAAAGTGGTCTAAGGCG	
qrac3b_F	CCCCGGCGAGTACATCCCTACAG	
qrac3b_R	GGTATGAGAGTGGACGGAGGCG	
qcdc42_F	TCTCAGTTGTTTACCTTCTTCA	
qcdc42_R	GATCAATCTGCGTCCCCAC	
qcdc42l_F	AGTTATCCGCAGACAGACGT	
qcdc42l_R	AATGGAGTGCCTGGACAATG	
qrhoaa_F	TCCTGAGGTTTACGTTCCCA	
qrhoaa_R	TGGCCAGCTGTATCCCATAG	
qrhoab_F	ACAGGCTTCGTCCTCTTTCA	
qrhoab_R	AATGTTTGACCTCAGGCGTC	
qrhogb_F	CTACTAACTTCCACAGCCGC	
qrhogb_R	CCACCACCACACACTTGATG	
qrhogc_F	CGCTGCGATGATGGAAACTT	
qrhogc_R	CTGCACCATCACCAACAACA	
qmpx_F	AGGTGTTGCTGAGCCTTTTG	
qmpx_R	TGGTGAACACTCCATTGCTC	
qcsf1a_F	ATGTCCAGACCTGACTTCGC	
qcsf1a_R	TCGGATGTTTCTCCAGCAT	
qef1a_F	TGCCTTCGTCCCAATTTTCAG	(Oehlers, Flores et al. 2010)
qef1a_R	TACCCTCCTTGCGCTCAATC	
qrps11_F	TAAGAAATGCCCTTCACTG	(de Oliveira, Reyes-Aldasoro et al. 2013)
qrps11_R	GTCTCTTCTCAAAACGGTTG	
qbactin2_F	GTGCCATCTACGAGGGTTA	(de Oliveira, Reyes-Aldasoro et al. 2013)
qbactin2_R	TCTCAGCTGTGGTGGTGAAG	

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