



1 Supplementary materials



Table S1. Primers for Real-Time PCR

Gene	Forward (5'-3')	Reverse (5'-3')	Genbank acc. no.
il1b	GACATGCTCATGGCGAACG	GCAAATCGTGCATTGCAAGACG	NM_212844
il6	GTGAAGACACTCAGAGACG	GTTAGACATCTTTCCGTGCTG	NM_001261449
il8	TTGAAGGAATGAGCTTGAGAGG	TCATGGAGCAGAGGGGTC	XM_009306855
mmp9	CATTAAAGATGCCCTGATGTATCCC	AGTGGTGGTCCGTGGTTGAG	NM_213123
<i>mmp</i> 13	ATGGTGCAAGGCTATCCCAAGAGT	GCCTGTTGTTGGAGCCAAACTCAA	NM_001290479
β -actin	TGGGTATGGAATCTTGCGGT	GTGGGGCAATGATCTTGATCT	NM_181601





- Figure S2. (A) Image of the tail of a 3 dpf zebrafish larvae, showing the position of the caudal tail amputation
- (red line) and the area designated for counting neutrophils number. (B) Representative fluorescence microscopy
- images of wound-induced migration of neutrophils, in the four experimental conditions.





Figure S3. Morphological parameters analyzed on 5 dpf zebrafish larvae after co-culture with *Phormidium* at initial concentration of 0.3 (OD value) for either 5 days or 2 days. A: eye area; B: body length; C: swim bladder area. To avoid redundancy only results obtained with the higher *Phormidium* concentration used are shown. The data represent the mean±SD of three independent experiments each conducted with 12-15 larvae. Statistical analysis was performed using GraphPad Prism 7 (One-way ANOVA followed by Tukey's multiple comparison test).





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Figure S4. Copper treatment results in NF-κB pathway activation. Induction was analysed with the zebrafish
transgenic line NFκB:GFP,Luc at three time points and was calculated by dividing luciferase values from treated
larvae by values of untreated siblings. The data represent the mean±SEM of a single experiment conducted with

30 20 larvae. Statistical analysis was performed using GraphPad Prism 5 (Unpaired t test with Welch's correction).

31 Statistical significance: * $p \le 0.05$, ** $p \le 0.01$.



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33 Figure S5. Caudal fin amputation results in NF-κB pathway activation. Induction of NF-κB activity was analyzed

34 with the zebrafish transgenic line NFkB:GFP,Luc at three time points and was calculated by dividing luciferase

35 values from treated larvae by values of untreated siblings. The data represent the mean±SEM of a single

36 experiment conducted with 20 larvae. Statistical analysis was performed using GraphPad Prism 5 (Unpaired t

37 test with Welch's correction). Statistical significance: ** $p \le 0.01$.