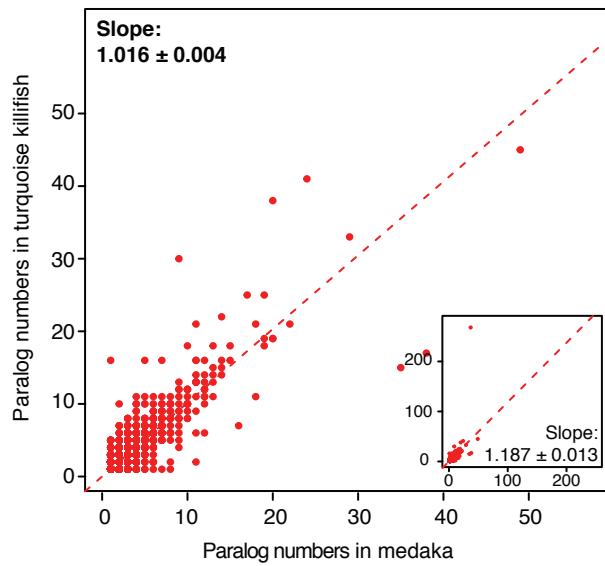
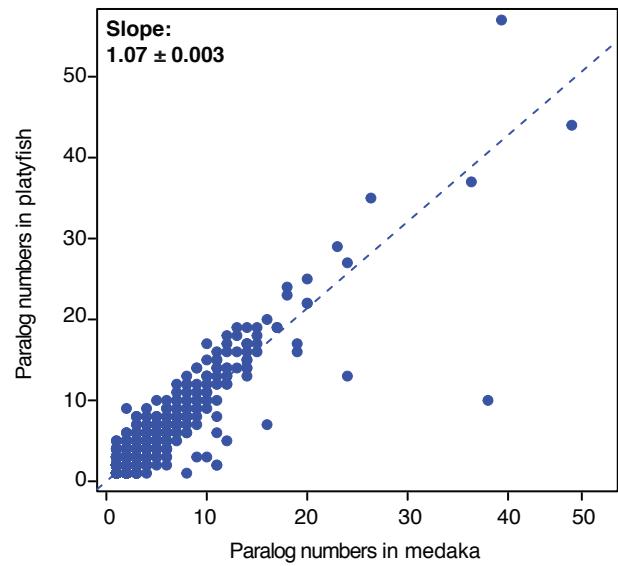


# Figure S2

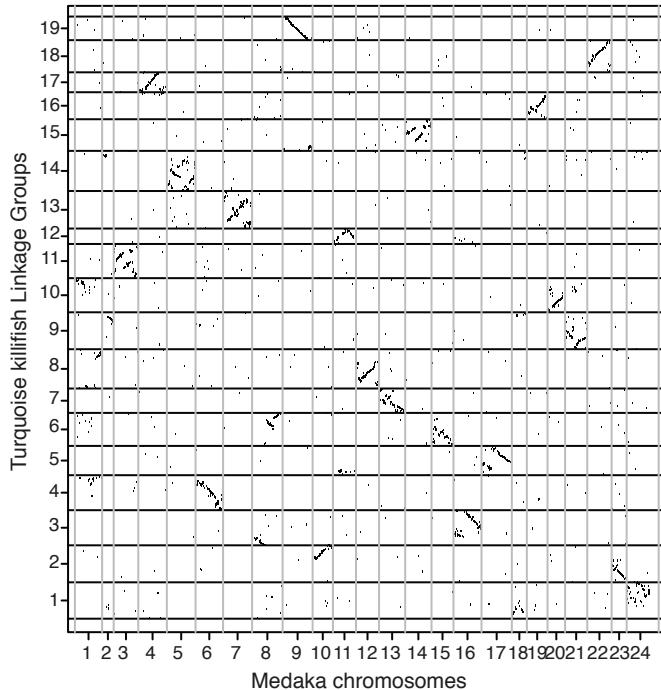
**A** Paralog numbers in turquoise killifish vs. medaka



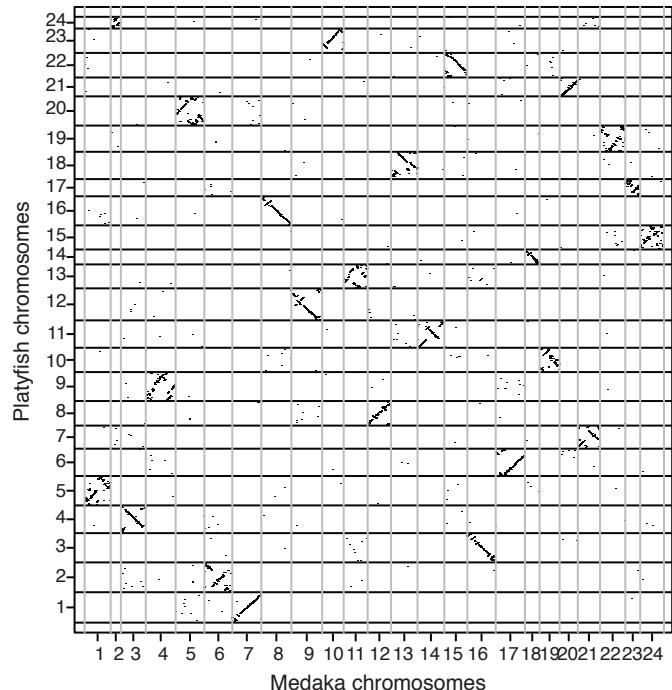
**B** Paralog numbers in platyfish vs. medaka



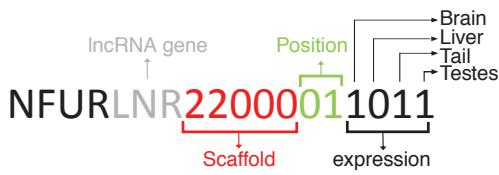
**C** Synteny between turquoise killifish and medaka



**D** Synteny between platyfish and medaka



**E** Nomenclature for lncRNA genes



**F** ncRNA gene predictions

ncRNA gene type	Number	Prediction algorithm
putative rRNA genes	200	RNAmmer*
	320	Infernal
	504	RepeatMasker
putative tRNA genes	1,841	tRNA-scan SE*
	4,344	tRNA-scan SE (pseudogenes)
	1,760	Infernal
	2,172	RepeatMasker
putative snRNA genes	230	Infernal*
	223	RepeatMasker
putative snoRNA genes	191	Infernal*
putative miRNA genes	2,029	Infernal*

\*Higher specificity predictions