

Table S1 Eichenlaub and Ettwiller 2011

Candidate RR Region	" ttc29^{RR} "	" dock9^{RR} "	" ccdc46^{RR} "	" fam44b^{RR} "
Human Gene ENSEMBL ID	ENSG00000137473 (TTC29)	ENSG00000088387 (DOCK9)	ENSG00000154240 (CCDC46)	ENSG00000145919 (FAM44B)
Human exon location (GRCh37)	chr4:147830178-147830401:-1	chr13:99738477-99738660:-1	chr17:63898270-63898452:-1	chr5:173036241-173036437:-1
Human exon length	224 bp	184 bp	183 bp	197 bp
Stickleback BLASTZ hit (BROADS1)	groupIX:6710625-6710804:1	groupXVI:2860453-2860626:-1	groupIX:12578736-12578860:1	groupIV:13966044-13966125:-1
Stickleback BLASTZ hit length	180 bp	174 bp	125 bp	82 bp
% ID to hs exon (BLASTZ hit)	70%	66%	72%	73%
Medaka RR (MEDAKA1)	chr1:22443500-22443723:-1	chr21:9260151-9260338:1	chr1:4650130-4650315:-1	chr10:16717951-16718149:-1
Medaka RR length	224 bp	188 bp	186 bp	199 bp
% ID to hs exon (Medaka RR)	68%	69%	68%	60%