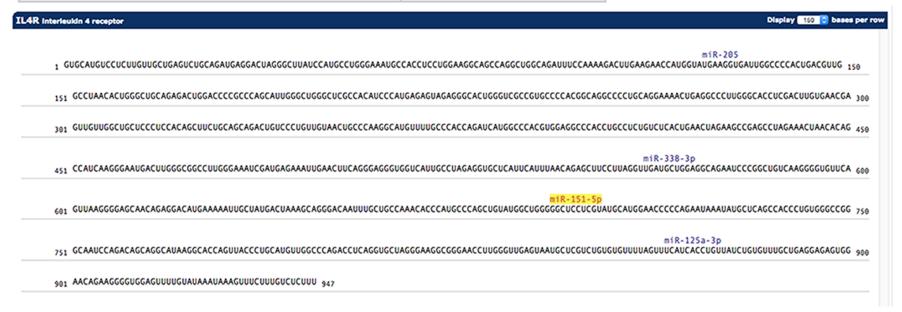
Α

Score	Expect	Identities	Gaps	Strand
42.1 bits(21)	6e-11	21/21(100%)	0/21(0%)	Plus/Plus
has miP 151 5p years		macm 01		
has-miR-151-5p ccacc mmu-miR-151-5p ccacc		TAGT ZI		

В

hsa-miR-151-5p/IL4R Alignment				
3' ugAU-CUGACACU : : 681:5' ugUAUGGCUGGGG		mirSVR score: -0.0840 PhastCons score: 0.4808		



Supplementary Figure 2. miR-151-5p characterization. (**A**) Human and mouse miR-151-5p have identical RNA sequences. (**B**) miR-151-5p was predicted to bind a site in the IL4Rα RNA sequence.