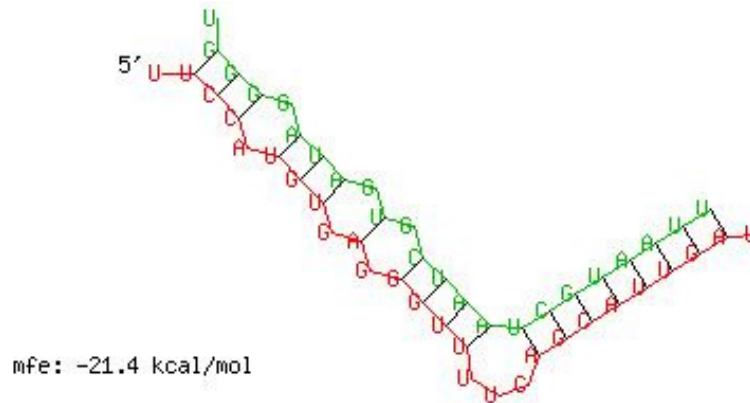


Site 1



Site 2

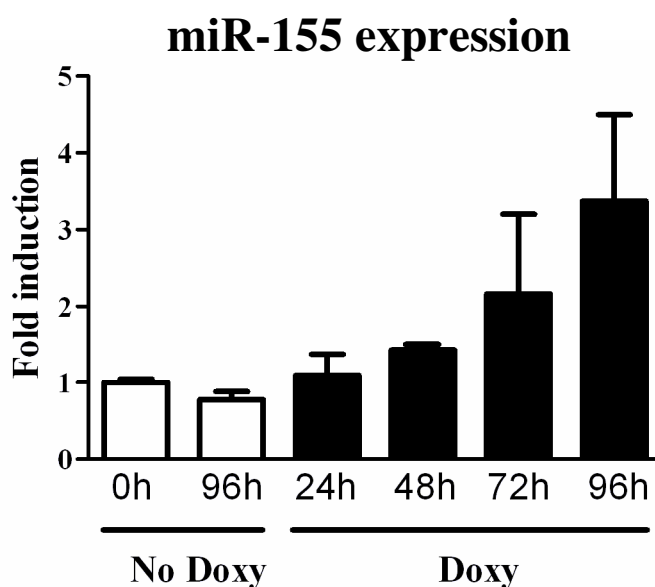
Supplement Figure S1. Predicted structure for miR-155 pairing to each site in the 3'UTR of IL13R α 1. Also shown is the correspondent predicted minimum free energy hybridisation. Site 1 shows miR-155 hybridization to positions 1049-1071 in the 3'UTR of IL13R α 1; Site 2 shows miR-155 hybridization to positions 1399 and 1424 in the 3'UTR of IL13R α 1.

<i>H. sapiens</i>	TTT----CTTCACCTCTGCTACTCAAG T--AGCATT TACTGTGTCTTTGGTTT
<i>P. troglodytes</i>	TTT----CTTCACCTCTGCTACTCAAG T--AGCATT TACTGTGTCTTTGGTTT
<i>P. pygmaeus</i>	TTT----CTTCACCTCTGCTACCCAAG T--GGCATT TACTGTGTCTTTGGTTT
<i>M. mulatta</i>	TTT----CTTCACCTCTGCTACTCAAG T--AGCATT TACTGTGTCTTTGGCTT
<i>C. jacchus</i>	TGT----CTTCACCTCTGCTACTCAAT T--AGTATT TACTGTGTCTTTGGCTT
<i>M. musculus</i>	GCCACCTTTTCATTTCTGCTATTCAA----- GTTTT GACTATGTATCTAGCTT
<i>R. norvegicus</i>	GCTTACCTTTTCATTTCTGCTACTCAAG TCAAGTATAG ACTATGTATCTAGCTT
<i>B. taurus</i>	CTT----CTTCACCACTGCTATTTCAGA TCAAGTATT TAGCACACGTCGGATT
<i>C. lupus familiaris</i>	CTT----CTTAATCTCTGCCGATCA----- GATATT CACCAGGTGT---GCCT
<i>E. caballus</i>	TGT----CTTCACCTCTGCTACTCAGA TCAAGTACT CACTATGTGTCTAGATT

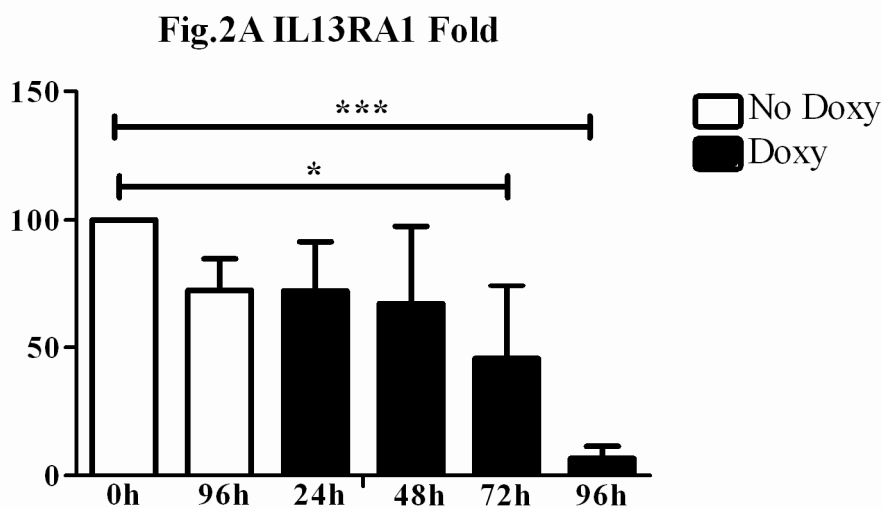
Supplement Figure S2. Conservation of miR-155 targeting “Site1” in the 3'UTR's of different species.

<i>H. sapiens</i>	TTCCATGTGAG-----GGTTTTTC <u>AGCA-TTGA</u> -TATT----T-----GTGC
<i>P. troglodytes</i>	TTCCATGTGAG-----GGTTTTTC <u>ATCA-TTGA</u> -TATT----T-----GTGC
<i>P. pygmaeus</i>	TTCCATGTGAG-----GGTTTTTC <u>AGCA-TTGA</u> -TATT----T-----GTGC
<i>M. mulatta</i>	TTCCATGTGAG-----GGTTTTTC <u>AGCA-TTGA</u> -TATT----T-----GTGC
<i>C. jacchus</i>	TTCTGTGTGAG-----GGTTTTTC <u>AGCA-TTGA</u> -TATT----T--ATCTATGC
<i>M. musculus</i>	TTACCTGTGAGCCTGCAATTATTTGAGGTTTTTC <u>AGCAATTGG</u> -TGTT----TCT----ATGT
<i>R. norvegicus</i>	TTCCCTGTGAGCCTGCAA---TTTGAGATTTTCA <u>AGCAATTGG</u> -TGTT----TCT----GTGT
<i>B. taurus</i>	TTCTATGTGAGTCTGCATTTATTTGAAGGTTTTTC <u>AGCA-TTTA</u> CTGTT----T-----ATGC
<i>C. lupus f.</i>	TTCTGCGTGAGTCTGCATTTCTTTGAAGGTTTTTC <u>AGCA-TTTA</u> CTAAT----T-----ATGC
<i>E. caballus</i>	TGTTGTGTGAGTCTGCATTTCTTTGAAGGTTTTTC <u>AGCA-TTTA</u> CTTTTTCTCT-----ACGC

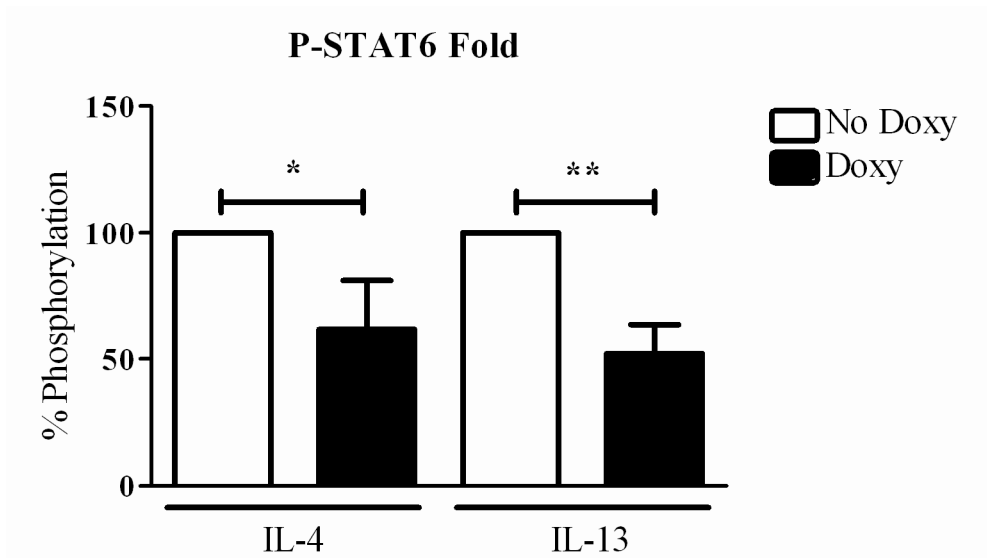
Supplement Figure S3. Conservation of miR-155 targeting “Site2” in the 3’UTR’s of different species.



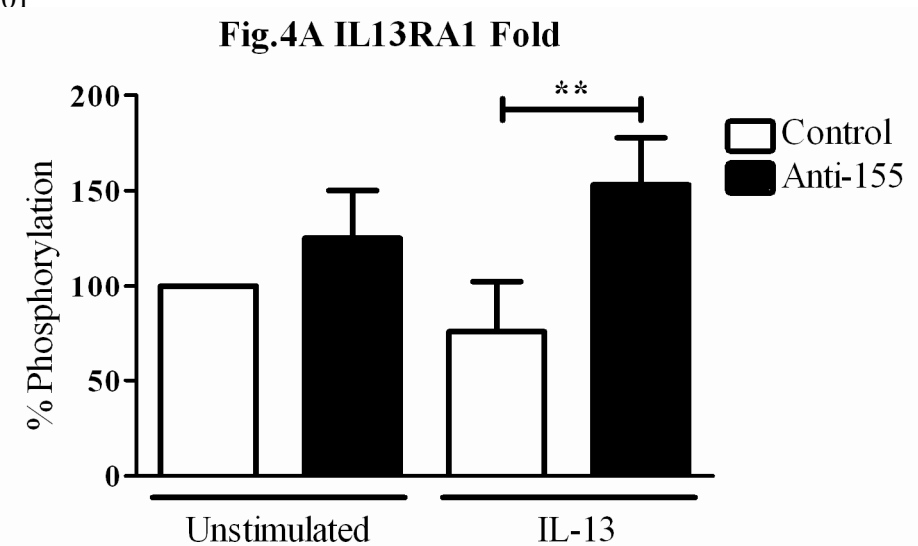
Supplement Figure S4. MiR-155 fold increase upon addition of Doxycycline to THP1-155 cells.
Control experiment for Fig 2 and Fig 3.



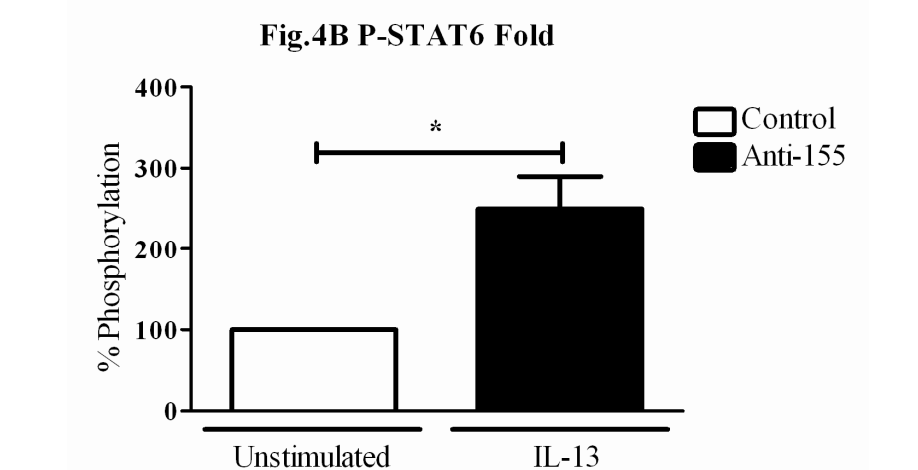
Supplement Figure S5. IL13RA1 protein levels in THP1-155 cells after doxycycline treatment. Statistical analysis of three independent western blot experiments that match results shown in Fig 2A. * $p \leq 0.05$; *** $p \leq 0.001$



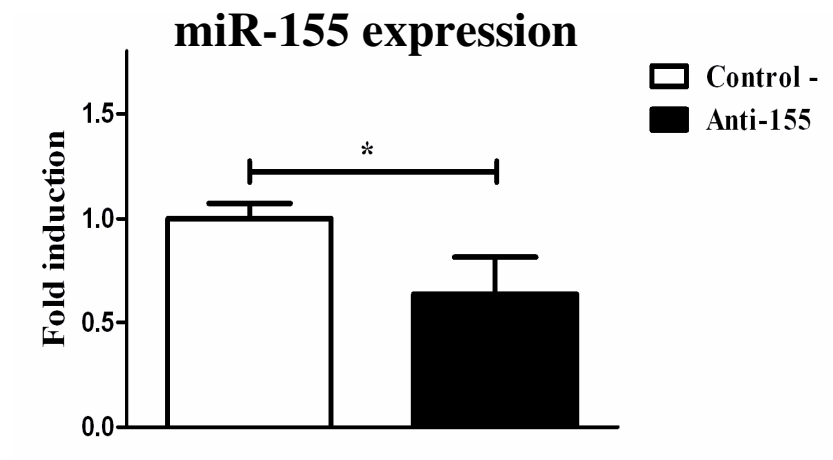
Supplement Figure S6. P-STAT6 levels in THP1-155 cells after doxycycline treatment and IL4 or IL13 stimulation. Statistical analysis of three independent western blot experiments that match results shown in Fig 3A. * $p \leq 0.05$; ** $p \leq 0.01$



Supplement Figure S7. IL13RA1 protein levels in macrophages after anti-miR-155 transfection. Statistical analysis of three independent western blot experiments that match results shown in Fig 4A. ** $p \leq 0.01$



Supplement Figure S8. P-STAT6 levels in macrophages after anti-miR-155 transfection. Statistical analysis of three independent western blot experiments that match results shown in Fig 4B. * $p \leq 0.05$



Supplement Figure S9. MiR-155 knock down efficiency in human macrophages at day 3 before IL-13 treatment. Control experiment for Figs. 4 and 5.