

Supporting Figure S3. Neighbor joining tree to show similarities of human pre-miR-155 to 18 other species.

'Homo sapiens' forms the root of the tree. The tree was constructed using the NCBI Blast tree viewer (<https://blast.ncbi.nlm.nih.gov/Blast.cgi>) based on BLASTN analysis of the annotated miR-155 precursor sequence for each species.

