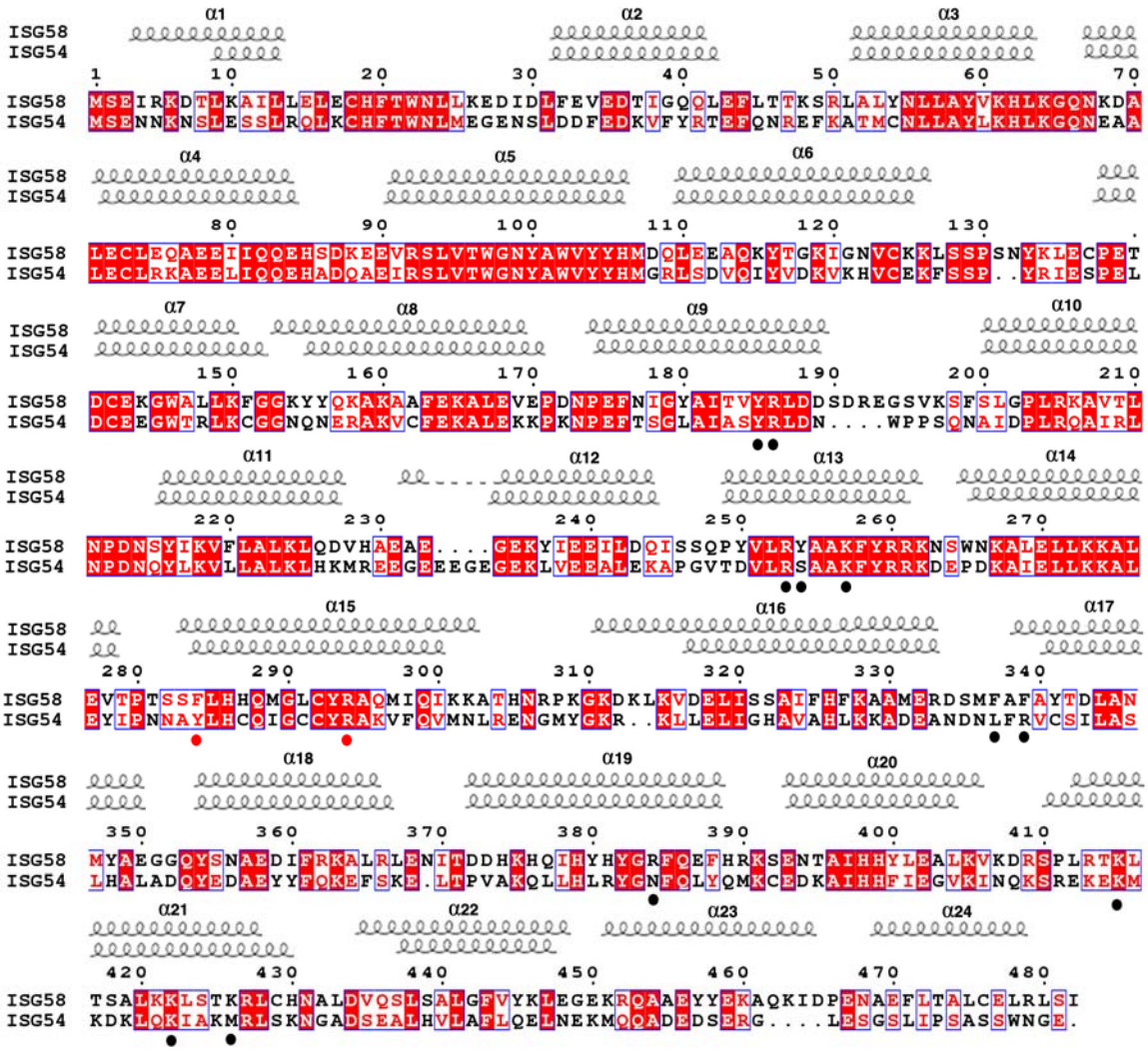

Supplementary information, Figure S2 Sequence alignment of human ISG58 and ISG54 using ClustalW2 [2]. Secondary structures of both ISG58 and ISG54, and residues important for RNA and DNA binding, as identified from the mutagenesis studies are labelled.The figure was prepared by ESPript [3].