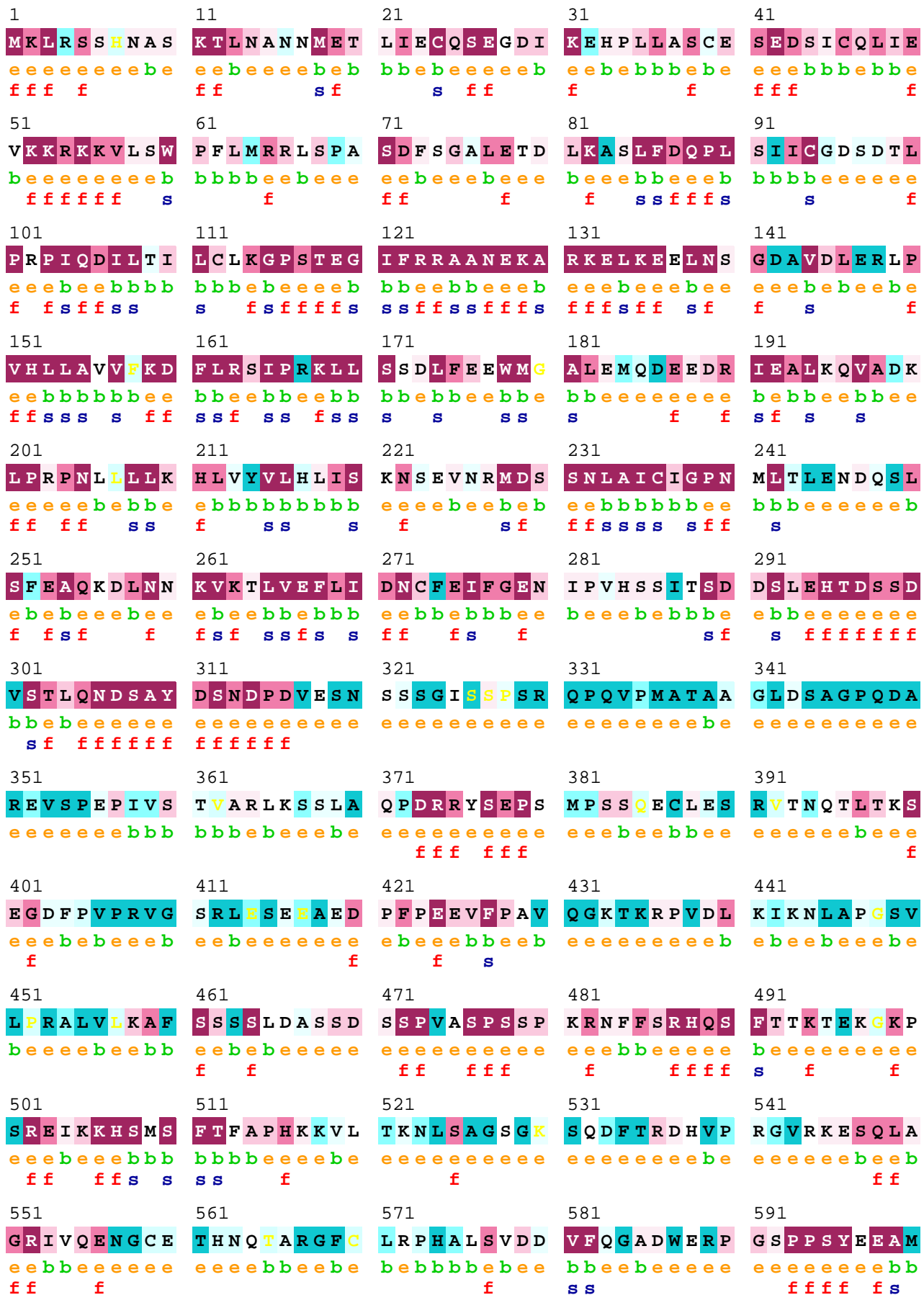
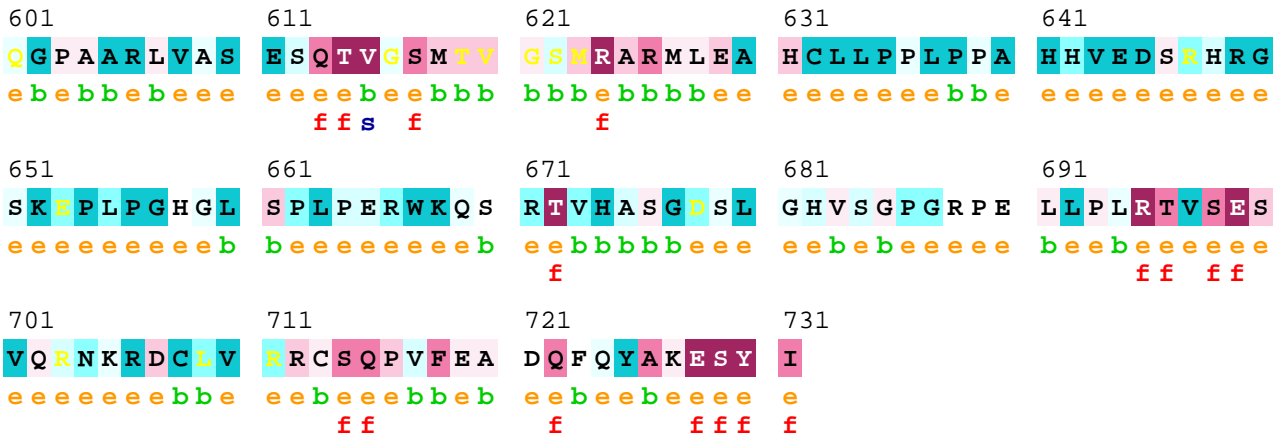


ConSurf Results

Supplementary Figure 1. Consurf Prediction showing Conservation Profile of Amino Acids in TAGAP protein





The conservation scale:



Variable Average Conserved

- e - An exposed residue according to the .
- b - A buried residue according to the .
- f - A predicted functional residue (highly conserved and exposed).
- s - A predicted structural residue (highly conserved and buried).
- x - Insufficient data - the calculation for this site was performed on less than 10% of the sequences.