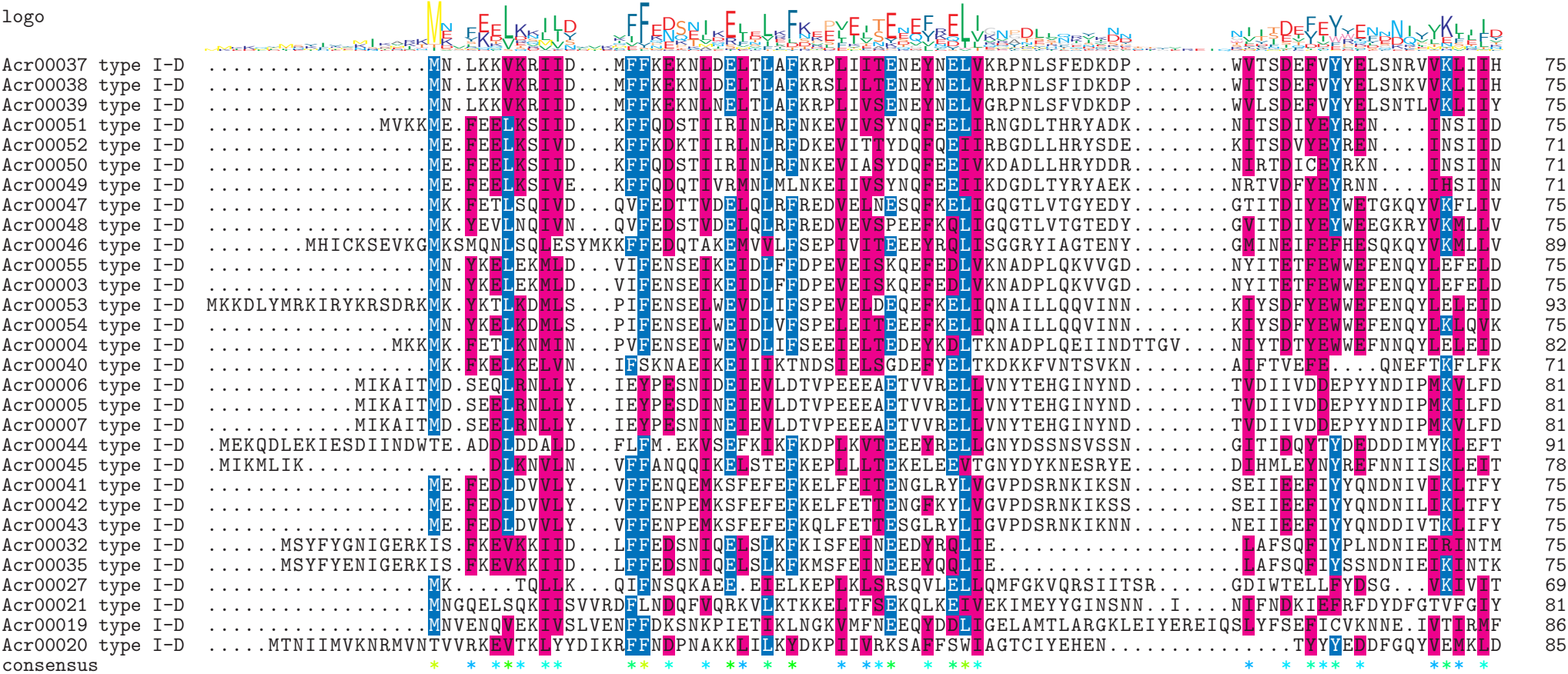


logo



A sequence logo representing the conservation of amino acids at each position of the protein. The y-axis indicates the information content or conservation score for each position. The x-axis shows the corresponding amino acid residues. Notable features include a highly conserved 'Y' at the start, followed by 'K', 'D', 'N', 'K', 'L', 'E', 'I', 'F', 'W', 'R', 'E', 'K', 'K'. The C-terminal region shows more variability with residues like 'S', 'E', 'E', 'D', 'T', 'L', 'S', 'A', 'N', 'D', 'I', 'T', 'K', 'T', 'I', 'K'.

- X non conserved
- X similar
- X >50% conserved