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Protein structure prediction results for the protein sequence MGVSVETTQGLGRRVTITIAADSIETAVKSELVNVAKKVRIDGFRKGKVP MNI VAQRYGA SVRQDVLGDLMSRNFI DAI I KEKINPAGAPTYVPGEYKLGEDFTYSVEFEVYPEVELQGL EAI EVEKPIVEVT DADV DGM L DTLRKQQATWKEKDGAVEAEDRVTI DFTGSVDGE EFGG KASDFVLAMGQGRMIPGFEDGIKGHKAGEEFTIDVTFPEEYHAENLKGKAAKFALNKKV EERELPELTAEFIKRFGVEDGSVEGLRAEVRKKNMERELKSAIRNRVKSQAIEGLVKANDI DVPAALI DSEIDVLR RQA AQRFGGNEKQAL ELPREL FEEQAKRRVVVGLLLGEVIRTNEL KADEERVKGLIEEMASAYEDPKEVIEFY SKNKE LMDNMRNV ALEEQAVEAVLAKAKVTEK. The figure displays the sequence, secondary structure, SS confidence, disorder, and disorder confidence for each residue.

