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The figure displays protein structure prediction results for a protein of 370 amino acids. The results are presented in a tabular format for the full-length protein and a truncated version (1-100 aa).

Segment	Sequence	Secondary structure	SS confidence	Disorder	Disorder confidence
1-60	M N Y S H D N W S A I L A H I G K P E E L D T S A R N A G A L T R R R E I R D A A T L L R L G L A Y G P G G M S L R E V	Alpha-helix, Alpha-helix, Alpha-helix, Alpha-helix, Alpha-helix, Alpha-helix, Alpha-helix, Alpha-helix, Alpha-helix, Alpha-helix	High	Low	High
61-120	T A W A Q L H D V A T L S D V A L L K R L R N A A D W F G I L A A Q T L A V R A A V T G C T S G K R L R L V D G T A I S	Alpha-helix, Alpha-helix, Alpha-helix, Alpha-helix, Alpha-helix, Alpha-helix, Alpha-helix, Alpha-helix, Alpha-helix, Alpha-helix	High	Low	High
121-180	A P G G G S A E W R L H M G Y D P H T C Q F T D F E L T D S R D A E R L D R F A Q T A D E I R I A D R G F G S R P E C I	Alpha-helix, Alpha-helix, Alpha-helix, Alpha-helix, Alpha-helix, Alpha-helix, Alpha-helix, Alpha-helix, Alpha-helix, Alpha-helix	High	Low	High
181-240	R S L A F G E A D Y I V R V H W R G L R W L T A E G M R F D M M G F L R G L D C G K N G E T T V M I G N S G N K K A G A	Alpha-helix, Alpha-helix, Alpha-helix, Alpha-helix, Alpha-helix, Alpha-helix, Alpha-helix, Alpha-helix, Alpha-helix, Alpha-helix	High	Low	High
241-300	P F P A R L I A V S L P P E K A L I S K T R L L S E N R R K G R V V Q A E T L E A A G H V L L L T S L P E D E Y S A E Q	Alpha-helix, Alpha-helix, Alpha-helix, Alpha-helix, Alpha-helix, Alpha-helix, Alpha-helix, Alpha-helix, Alpha-helix, Alpha-helix	High	Low	High
301-360	V A D C Y R L R W Q I E L A F K R L K S L L H L D A L R A K E P E L A K A W I F A N L L A A F L I D D I I Q P S L D F P	Alpha-helix, Alpha-helix, Alpha-helix, Alpha-helix, Alpha-helix, Alpha-helix, Alpha-helix, Alpha-helix, Alpha-helix, Alpha-helix	High	Low	High
361-370	P R S A G S E K K N	Alpha-helix	High	Low	High

Confidence Key
High(9)  Low (0)
? Disordered
 Alpha helix
 Beta strand