

Fig. 1a

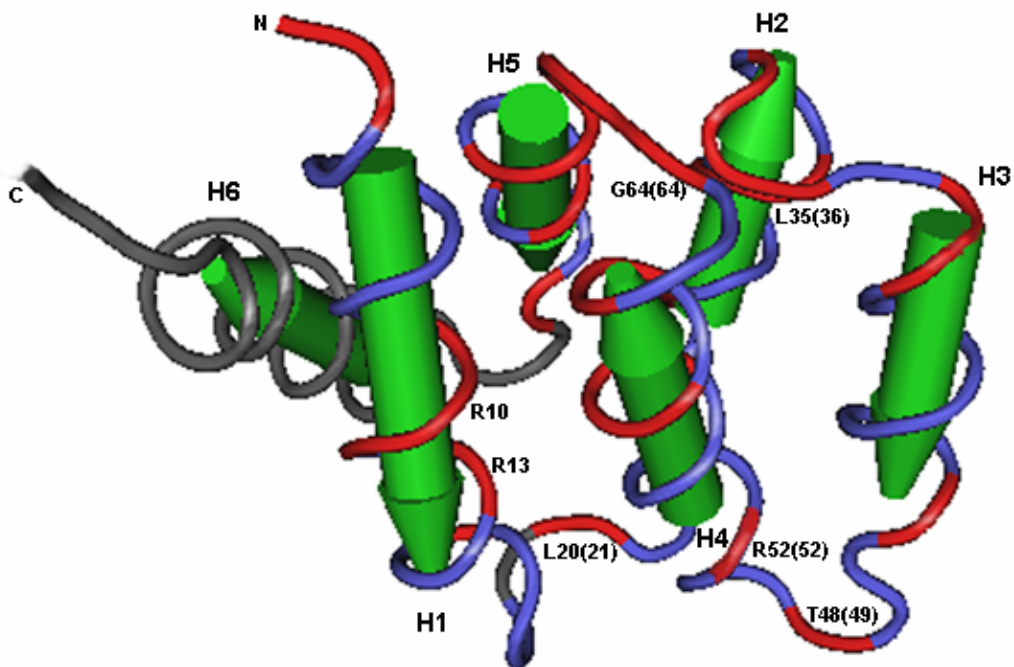


Fig. 1b

Supplementary Fig. 1: Structural comparison of *acasp* CARD predicted amino acid sequence with known crystal structures from humans. Sequence conserved between *acasp* and (a) MMDB Apaf-1 CARD when in complex with procaspase 9 (PDB: 3ygs gi: 7767041) and (b) RAIDD (PDB: 3crd, gi: 4388927). Identical sequence is shown in red, non-identical in blue and unaligned sequence in gray. Helical (H) secondary structures are depicted as green arrows and numbered in structure order. Selected *acasp* amino acids positions are depicted with the corresponding known human structure residue positions in parentheses.