

Figure S3. Homologous analysis of hMLKL (1-178) and mMLKL (1-168) amino acid sequences.

The helices were labeled based on the solution structures of full-length mouse MLKL (PDB accession code 4BTF) and human MLKL2-178 (PDB accession code 2MSV). 'TM' indicates 'transmembrane'. Residues that are conserved across all sequences are highlighted in yellow.