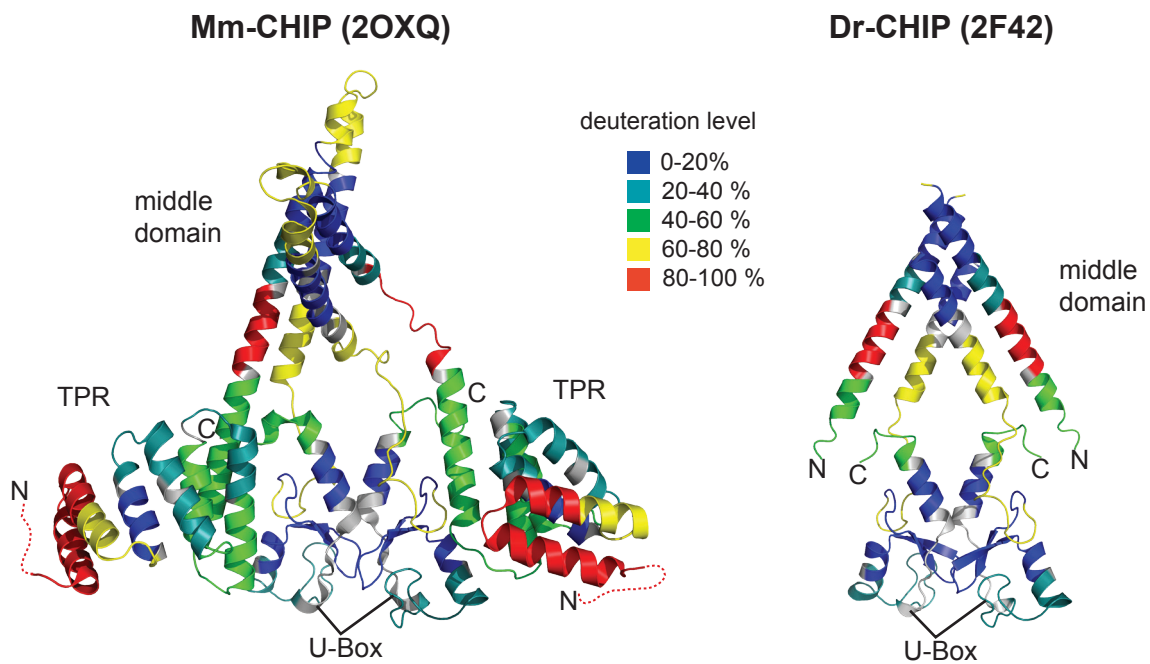




Suppl. Fig. 1: Conformational flexibility of CHIP. Deuteron incorporation shown on cartoon representation of the asymmetric mouse CHIP dimer structure in complex with MEEVD-peptides (residues 25-282, PDB code 2C2L) and for comparison the symmetric *Danio rerio* CHIP fragment (residues 127-304 corresponding to residues 142-301 of mouse CHIP; PDB code 2F42). The deuteration level of analyzed peptides after 10 s in D₂O is shown in the color code indicated. Gray color indicates regions for which no exchange data are available. The dashed line symbolizes the 24 N-terminal residues missing in the crystal structure.