

Sequence alignment of lcsn_CK1 across various accessions. The alignment shows the sequence from position 1 to 450. A color-coded scheme highlights specific amino acid residues: blue for hydrophobic, red for polar, green for acidic, purple for basic, and yellow for aromatic. Conserved residues are shown in black.

Accessions: 17538560.w, 17544600.w, 17543206.w, 17539102.w, 17540210.w, 17541944.w, 17558800.w, 17562500.w, 17532057.w, 17534577.w, 17540212.w, 17541946.w, 17538256.w, 17556390.w, lcsn_CK1

Position 1 to 450:

17538560.w, 17544600.w, 17543206.w, 17539102.w, 17540210.w, 17541944.w, 17558800.w, 17562500.w, 17532057.w, 17534577.w, 17540212.w, 17541946.w, 17538256.w, 17556390.w, lcsn_CK1