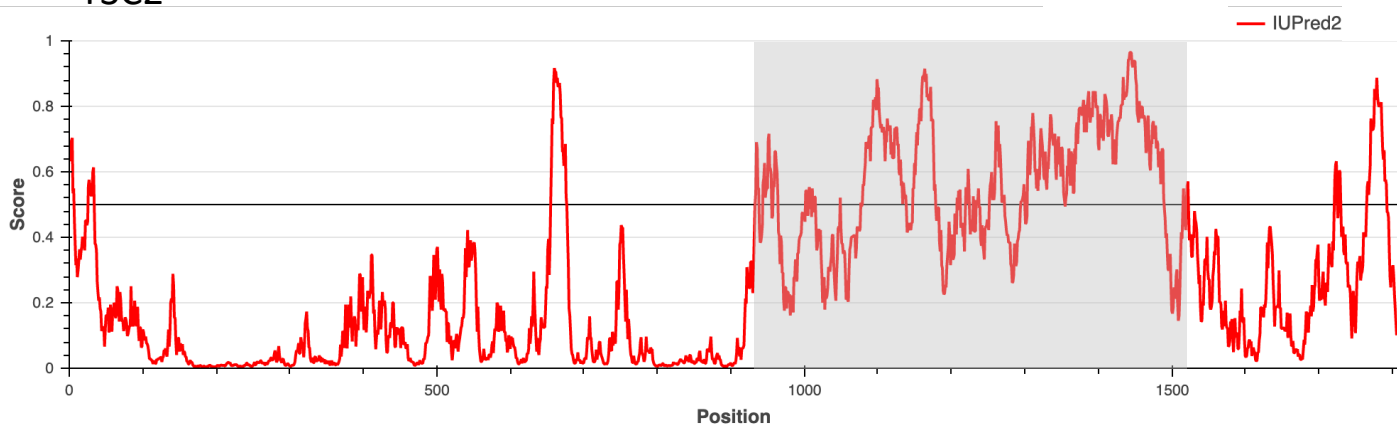


## TSC2



### Supplementary figure S2.

**Supplementary figure S2.** Intrinsically disordered regions in TSC2. Prediction of protein disorder using the IUPred web server for the human TSC2 protein. The horizontal axis represents the protein chain and the vertical axis represents the probability of each residue to be disordered. Residues with values above 0.5 are predicted to be disordered and values below 0.5 indicate an ordered structure. The broad intrinsically disordered region (ID) consistent with stress granule localization is shaded.