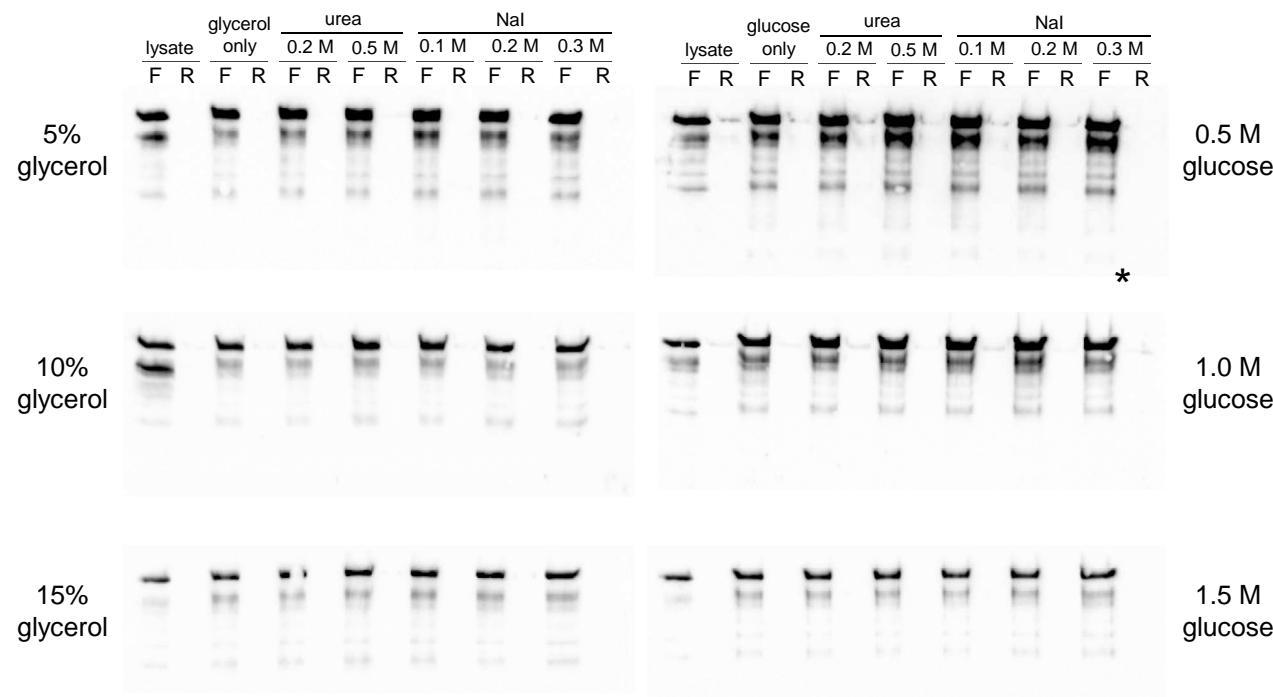


La motif →

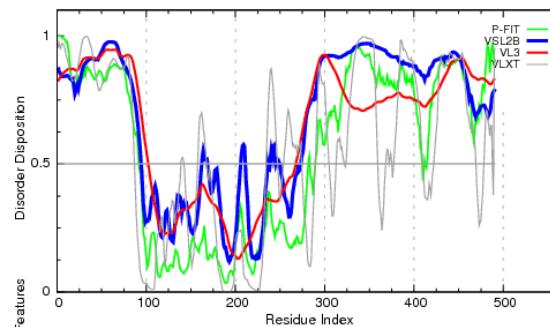
2° (NMR) 
 H.sapiens TASGGENE-REDLEQEWKPPDEELIKLVDQIEFYFSDENLEKDAFLLKHVRNNKLGYVSVKLLTSFKVKHLTRDWRTTAHALKYSVL
 F.catus TTSGGENE-HEDGEQEWKPPDQELIRLKLVQIEFYFSDENLEKDAFLLKHVRNNKLGYVSVKLLTSFKVKHLTRDWRTTAHALKYSMSL
 M.musculus TTSGGEND-REDLEPEWRPDEELIRLKLVQIEFYFSDENLEKDAFLLKHVRNNKLGYVSVKLLTSFKVKHLTRDWRTTAHALKYSVTL
 C.hircus TTSGGENE-RDDLEQEWPQDEELIKLVDQIEFYFSDENLEKDAFLLKHVRNNKLGYVSVKLLTSFKVKHLTRDWRTTAHALKYSVTL
 B.taurus TTSGGENE-RDDLEQEWPQDEELIKLVDQIEFYFSDENLEKDAFLLKHVRNNKLGYVSVKLLTSFKVKHLTRDWRTTAHALKYSVTL
 R.norvegicus TTSGGEND-REDLEPEWKPPENDLIRLKLVQIEFYFSDENLEKDAFLLKHVRNNKLGYVSVKLLTSFKVKHLTRDWRTTAHALKYSVTL
 G.gallus RSSGGEND--DDSDQNWKPPENDLIQKLVQIEYYFSDENLEKDAFLLKHVRNNKMGMGVSVKLLTSFKVKHLTRDWRTTTHALKYSDML
 M.gallopavo NSSGGEND--DDSDQNWKPPENDLIQKLIAQIEYYFSDENLEKDAFLLKHVRNNKMGMGVSVKLLTSFKVKHLTRDWRTTAHALKYSIDML
 C.cornix APCGGEND--DDSDQNWKPPENDLIQKLIAQIEYYFSDENLEKDAFLLKHVRNNKMGMGVSVKLLTSFKVKHLTRDWRTTAYALKYSNTL
 P.adeliae RRNGGEND--DDSDQWKWPPENDLIQKLIAQIEYYFSDENLEKDAFLLKHVRNNKMGMGVSVKLLTSFKVKHLTRDWRTTAYALKYSDTL
 C.pelagica VGGGEND--DDSDQNWKPPENDLIQKIIAQIEYYFSDENLEKDAFLLKHVRNNKMGMGVSVKLLTSFKVKHLTRDWRTTAYALKYSDTL
 O.hoazin KASGGEND--DDSDQNWKPPENDLIQKLIAQIEYYFSDENLEKDAFLLKHVRNNKMGMGVSVKLLTSFKVKHLTRDWRTTAYALKYSDIL
 X.tropicalis FDFTETLD-DSSYEGDSGIPDLKLVRKIVSQVEFYLSDENLSHDAFLLKHVQKNMGMGVSIKLTSFKVKYLTRDWRTKTLYALRFSEQL
 X.laevis ATSGGEND-GDELDQDWKPPDAELIQLKLIQIEYYLSDENLEKDAFLLKHVRNNKMGMGVSVKLLTSFKVKHLTRDWRTTAYALRYSNLL
 E.lucius GTSGGELE----EESWQPPDPPELTIQKLVAQIEYYLSDENLEHDAFLLKHVRNNKLGFVSVKLLTSFKVKHLTRDWRTTAYALRHSTIL
 O.latipes GTSGGELE----EESWQPPDPPELTIQKLVAQIEYYLSDENLEHDAFLLKHVRNNKLGFVSVKLLTSFKVKHLTRDWRTTAYALKHSKML
 T.rubripes GTSGGELE----DESWKPPDPPELTIQKLVAQIEYYLSDENLEHDAFLLKHVRNNKLGFVSVKLLTSFKVKHLTRDWRTTAFALARHSQIL
 D.rerioA GTSGGELE----EESWQPPDPPELTIQKLVAQIEYYLSDENLEHDAFLLKHVRNNKLGFVSVKLLTSFKVKHLTRDWRTTAYALRHSNLL
 D.rerioB TDLTDVFEEDFCEGELWIPPNDVTQQIATQLENYLSDENLSDDAFLLKHVQRNMGMGVSLKLLTSFKVKIRDLTTRDTAAARTSPQL
 X.maculatus GTSGGELE----EESWQPPDPPELTIQKLVTQIEFYLSDENLEHDAFLLKHVRNNKLGFVSVKLLTSFKVKHLTRDWRTTAYALKHSKIL
 Consensus_aa .s.ssp.-.pWbPP-.LhpK11sQ1E@yh\$DENLp+DAFLLKhvpRNkHG@vs1KLLTSFKK1cLTRDWRTThe@L+s..L

The figure displays a sequence logo for the DIRRISSRYSQVGTQECAIVEFEEVEAAIKAHEFMITES sequence across multiple species. The logo uses colored bars to represent the frequency of each amino acid at each position. A blue bar indicates positions 1-4, a grey bar indicates positions 5-8, and a green bar indicates positions 9-12. The x-axis represents the sequence positions, and the y-axis lists the species: H.sapiens, F.catus, M.musculus, C.hircus, B.taurus, R.norvegicus, G.gallus, M.gallopavo, C.cornix, P.adeliae, C.pelagica, O.hoazin, X.tropicalis, X.laevis, E.lucius, O.latipes, T.rubripes, D.rerioA, D.rerioB, X.maculatus, and Consensus aa. The logo shows high conservation of the sequence across all species.

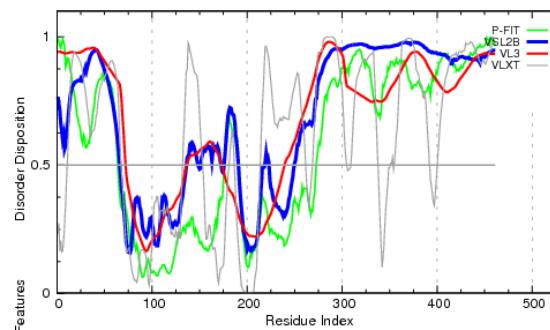
Supplemental Figure 1.



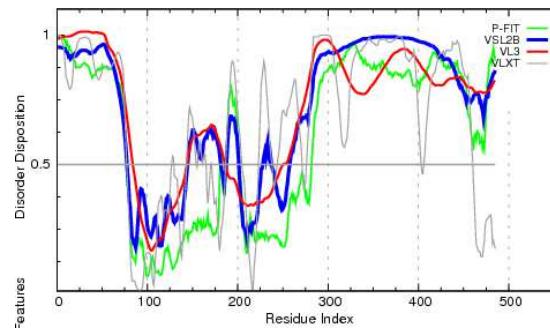
Supplemental Figure 2.



HsLARP6



XmLARP6



DrLARP6a

Supplemental Figure 3.