

Figure S4

Protein-Protein binding domain

>query: stretch=5 crowd_predictions=7 gap=20 itr=0

MGFHSFDVYFFFLSCPNNFCFSLSLSNFQKKS~~RD~~KERE

PP-PP-----P-----P-----

IQTNYPTFKKSKKNTKPRKKKARKKMGKYSELIDAGVRIA

----PP-----P-----

ARFHS~~HC~~PQTARMYYHPPPPTTAESGPTQRYPPQDGGVLG

---P-----P-----PPPPPPPPPPPPPPPPPPPP--PP

CKGSSSGVDITKDLILHSIC

PPP-P-----P