Supporting Information Text S17: Protein A-B domain supplementary figures

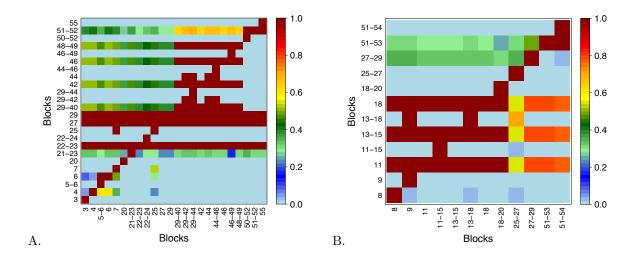


Figure 1: Protein A-B domain: BIS unclustered correlated distribution score matrices. BIS analysis realized on the dataset of 452 sequences. A: matrix computed for d = 0. The clustered matrix is reported in Fig. 2A-Text S17 (clusters have symmetric and environmental scores ≥ 0). B: matrix computed for d = 1.

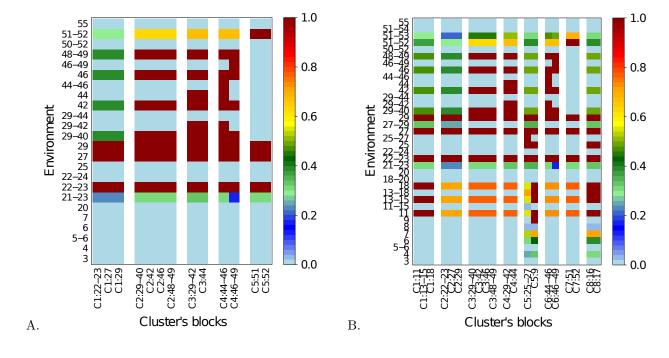


Figure 2: Protein A-B domain: matrices computed for d = 0 and $d \le 1$.

BIS analysis realized on the dataset of 452 sequences. Clustered correlated distribution score matrices for d=0 (**A**) and $d \le 1$ (**B**), where symmetric and environmental scores used for selecting clusters are ≥ 0 . Notice that Fig. 4A is made of clusters of this matrix with scores = 1.