



Figure S2. Distribution of distances between coevolving residues with minimal primary sequence separations

(A) The fraction of coevolving (black bars) or all (white bars) residue pairs with a primary distance of at least 10 amino acids that lie within the specified interval of physical distance from each other is depicted. (B) The fraction of coevolving (black bars) or all (white bars) residue pairs with a primary distance of at least 30 amino acids that lie within the specified interval of physical distance from each other is depicted.