

Figure S2 a) B-factor distributions of the backbone atoms of the B-factors normalized as z-scores.

Top row: 165 structures with resolution better than 1.9 Å (bin R1).

Middle row: 357 structures with resolution 1.9-2.5 Å (bin R2).

Bottom row: 187 structures with resolution 2.5-3.0 Å (bin R3).

Water bridges considered between amino acids and DNA phosphates.

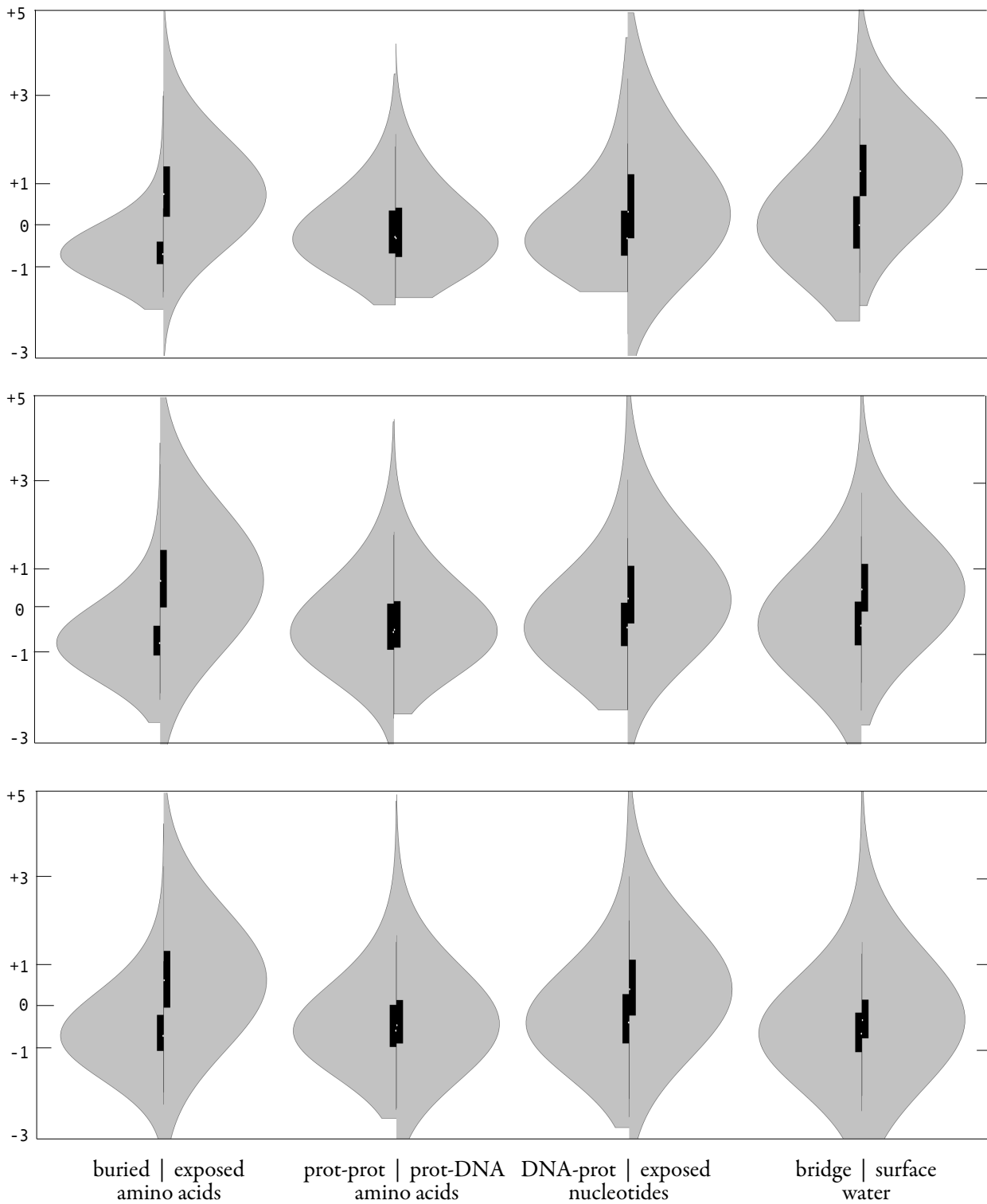


Figure S2 b) B-factor distributions of the side chain atoms of the B-factors normalized as z-scores.

Top row: 165 structures with resolution better than 1.9 Å (bin R1).

Middle row: 357 structures with resolution 1.9-2.5 Å (bin R2).

Bottom row: 187 structures with resolution 2.5-3.0 Å (bin R3).

Water bridges considered between amino acids and DNA bases.

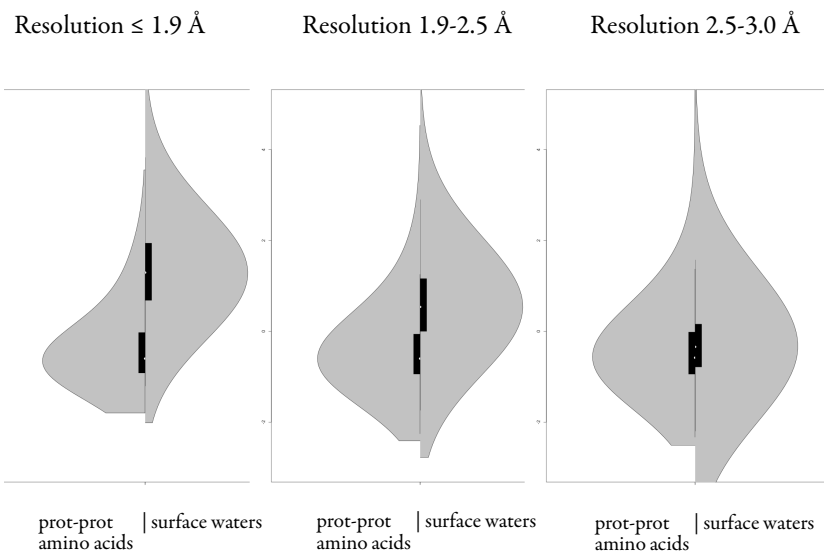


Figure S2 c) Comparison of normalized B-factor distributions (normalized as z-scores) in the three resolution bins for the backbone atoms of amino acids in contact with another protein (group of residues *protein-protein aa*) and the first-shell water molecules (group *surface w*).