Supporting Information - Graebsch et al.

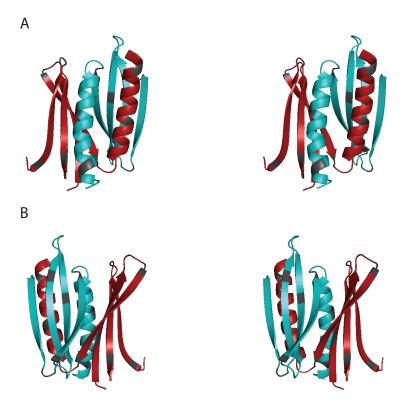


Figure S1 Stereoview of the crystal structure of *B. burgdorferi* Pur-alpha. (A) Ribbon backbone model with one monomer shown in red, the other in cyan. Every 10th residue is highlighted in grey (starting from residue 10). (B) Stereoview of (A), rotated 180° around the vertical axis.