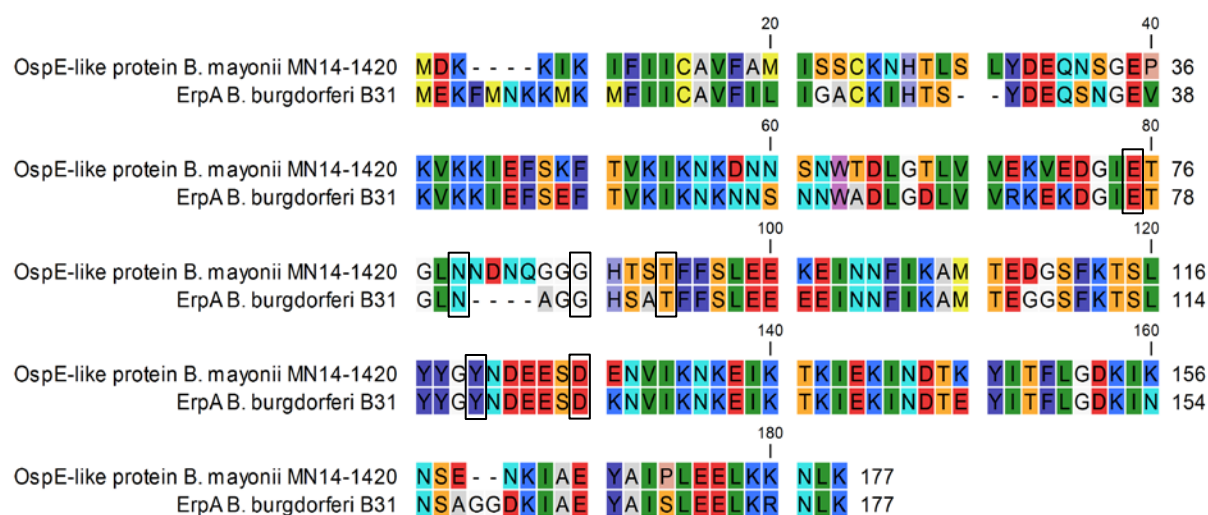


Supplementary Figure 3



Sequence alignment of ErpA of *B. burgdorferi* B31 and the OspE-like protein of *B. mayonii*.

Deduced amino acid sequences of ErpA (BBP38, NP_051199.2) and the OspE-like protein (WP_075552645.1) were used to create an alignment by CLC Sequence Viewer, version 8.0 (QIAGEN Aarhus A/S, Denmark). Residues involved in binding of ErpA to FH are boxed.