**Figure S1.** ***In silico* prediction of N-glycosylation of FliC amino acid residues.** Graph indicating predicted N-glycosylation sites across the protein chain from the N- to C-terminus, with potential sites (green vertical lines) crossing the threshold (blue horizontal line at 0.5) as determined using NetNGlyc 1.0 (<http://www.cbs.dtu.dk/services/NetNGlyc/)>.