The PyMOL session file (fig sup 1) was derived from a model of a SpoIVFB tetramer in complex with two Pro-K monomers reported previously (1). Only SpoIVFB chain A is shown in Figure 4-figure supplement 1A. In Figure 4-figure supplement 1B, predicted TMS2 of BofA is also shown, in an orientation to satisfy experimentally-observed cross-links (Figure 4, Figure 4-figure supplement 5, and Supplementary File 1).

Reference

1. Halder S, Parrell D, Whitten D, Feig M, & Kroos L (2017) Interaction of intramembrane metalloprotease SpoIVFB with substrate Pro-sK. *Proc. Natl. Acad. Sci. USA* 114:E10677-E10686.