The PyMOL session file (Figure 7C) was derived from the model ‘fb.sigk.bofa.fa’ of a SpoIVFB tetramer in complex with one molecule each of full-length BofA and parts of Pro-K (residues 38-114) and SpoIVFA (residues 65-111), by hiding the B, C, and D chains of SpoIVFB. The images shown in Figure 7 A and B were then derived by hiding additional parts of the model.