



Figure S10 Heatmaps of M-3D Hi-C links aligned to prokaryotic reference genomes. Reads from the M-3D *Hind*III non-resuspended library (Table S3) were aligned to the draft assemblies of four prokaryotic species present in the M-3D sample. Each heatmap has a resolution of 10 Kb, and the legend indicates the \log_{10} of link density.