

Supplementary Figure S3: SNP cluster PDS000033428.133 has highly variable AMR gene content driven partially by ISs in family IS6. The rings, from interior to exterior, represent pMHMC-004 coverage, pMHMC-004 presence, and then in larger groups of rings, pMHMC-004 AMR gene content and pMHMC-012 AMR gene content. Clades of interest within this SNP cluster are color coded directly on the tree and are the same as in Figure 6.