



Supplementary Figure 5. Depiction of the genetic rearrangements (gene losses and inversions) among the Indian strains compared with the closely related *L. monocytogenes* 4b serotype strain. All of the genomes were analyzed and comparatively visualized with the Genome-Comparisons (GECO) comparative genomic tool. (Ref.1) The regions are pegged at positions F2365_0490 and F2365_2055 of *L. monocytogenes* F2365. HP: hypothetical protein.

1. Kuenne CT, Ghai R, Chakraborty T, Hain T. GECO--linear visualization for comparative genomics. *Bioinformatics* 2007; 23: 125–126.