

SUPPLEMENTARY FIGURE 2: PFGE diversity of 110 *Listeria monocytogenes* isolates and correspondence with MLST and MvLST data.

The UPGMA tree was constructed based on the number of different bands observed on *ApaI* and *AscI* patterns using the ‘average of experiments function’ in BioNumerics v6.6. Lin., lineage; CC: clonal complex; ST: sequence type. The figure is spread on two pages, with two strains being repeated in order to mark continuity.



