

**Supporting Information for:**

The *Listeria monocytogenes* Core-Genome Sequence Typer (LmCGST): a bioinformatic pipeline for molecular characterization with next-generation sequence data

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**Additional file 11: Summary statistics describing data obtained from Illumina sequencing runs of *Listeria monocytogenes* strain 08-5578 genomic DNA samples.**

<b>Run</b>	<b>Yield (Gb)</b>	<b>Average Cluster Density (k/mm<sup>2</sup>)</b>	<b>Total Reads</b>	<b>Reads Passing Filter</b>
1	3.96	478.5	9357411	8506127
2	1.16	178.5	5077244	2302912
3	7.76	1075.0	19068628	16320666

Note: Genomic DNA was extracted and divided into four subsamples. Each subsample was sequenced during three separate runs on an Illumina MiSeq benchtop sequencer.