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 18: Data used to calculate unique *Listeria monocytogenes* high-confidence core genomes.

Strain	NCBI Database	NCBI Number
CC70B	Short-read archive	SRR835447
J1776	Nucleotide	CP006598.1
Scott-A	Nucleotide	NZ_CM001159.1
R2-502	Nucleotide	CP006594.1
J2-064	Nucleotide	CP006592.1
Lm19860	Nucleotide	KK211032.1
Lm26686	Nucleotide	KK211038.1
F6900	Nucleotide	NZ_AARU02000001.1 – NZ_AARU020000023.1
J2818	Nucleotide	NZ_AARX02000001.1 – NZ_AARX02000024.1
Lm25180	Nucleotide	KK211191.1
Lm21045	Nucleotide	KK211193.1
WSLC1001	Nucleotide	CP007160.1
C1-387	Nucleotide	CP006591.1
NCCP_15743	Nucleotide	NZ_APJT01000001.1 – NZ_APJT01000005.1
HPB1013	Short-read archive	SRR1639783
HPB1014	Short-read archive	SRR1640020
HPB1015	Short-read archive	SRR1640113
HPB1022	Short-read archive	SRR1640203
HPB1027	Short-read archive	SRR1640093
HPB1031	Short-read archive	SRR1640143
HPB1042	Short-read archive	SRR1639796
HPB1045	Short-read archive	SRR1640206
HPB1074	Short-read archive	SRR1639771
HPB1075	Short-read archive	SRR1640205
HPB1077	Short-read archive	SRR1640000
HPB1092	Short-read archive	SRR1640007
HPB1102	Short-read archive	SRR1639767
HPB1111	Short-read archive	SRR1640136
HPB1174	Short-read archive	SRR1639774
HPB1175	Short-read archive	SRR1639778
HPB1225	Short-read archive	SRR1640172
HPB1340	Short-read archive	SRR1640144

HPB1341	Short-read archive	SRR1640146
HPB1631	Short-read archive	SRR1640221
HPB1654	Short-read archive	SRR1640175
HPB1679	Short-read archive	SRR1639790
HPB1707	Short-read archive	SRR1640208
HPB1945	Short-read archive	SRR1640116
HPB2	Short-read archive	SRR1640183
HPB2088	Nucleotide	JOKU00000000.1
HPB2129	Short-read archive	SRR1640122
HPB2130	Short-read archive	SRR1640129
HPB2134	Short-read archive	SRR1640130
HPB2141	Short-read archive	SRR1640003
HPB2166	Short-read archive	SRR1640184
HPB2263	Short-read archive	SRR1640039
HPB2264	Short-read archive	SRR1640043
HPB23	Short-read archive	SRR1639677
HPB2386	Short-read archive	SRR1640193
HPB2412	Short-read archive	SRR1640211
HPB285	Short-read archive	SRR1640177
HPB2864	Short-read archive	SRR1640182
HPB2865	Short-read archive	SRR1640187
HPB2866	Short-read archive	SRR1640188
HPB2921	Short-read archive	SRR1640173
HPB2980	Short-read archive	SRR1640195
HPB330	Short-read archive	SRR1640223
HPB37	Short-read archive	SRR1639999
HPB394	Short-read archive	SRR1640131
HPB396	Short-read archive	SRR1639755
HPB4	Short-read archive	SRR1640179
HPB424	Short-read archive	SRR1640213
HPB518	Short-read archive	SRR1640164
HPB5209	Short-read archive	SRR1640224
HPB5252	Short-read archive	SRR1640125
HPB535	Short-read archive	SRR1640222
HPB5415	Nucleotide	JOKV00000000.1
HPB5767	Short-read archive	SRR1640123
HPB6355	Short-read archive	SRR1640120
HPB7	Short-read archive	SRR1639760

HPB7171 [58]	Nucleotide	JMMW01000000.1
HPB769	Short-read archive	SRR1640108
HPB770	Short-read archive	SRR1640109
HPB771	Short-read archive	SRR1640069
HPB772	Short-read archive	SRR1640111
HPB773	Short-read archive	SRR1640089
HPB774	Short-read archive	SRR1640090
HPB8	Short-read archive	SRR1639674
HPB862	Short-read archive	SRR1639793
HPB9	Short-read archive	SRR1639763
HPB913	Short-read archive	SRR1640166
HPB960	Short-read archive	SRR1640119
HPB962	Short-read archive	SRR1639780
HPB963	Short-read archive	SRR1639787
HPB976	Short-read archive	SRR1640169
HPB1024*	Short-read archive	SRR1640197
HPB1035*	Short-read archive	SRR1640198
HPB1043*	Short-read archive	SRR1640199
HPB1050*	Short-read archive	SRR1640115
HPB1053*	Short-read archive	SRR1639663
HPB1104*	Short-read archive	SRR1640024
HPB1110*	Short-read archive	SRR1640201
HPB1342*	Short-read archive	SRR1640152
HPB1380*	Short-read archive	SRR1640162
HPB2131*	Short-read archive	SRR1686399
HPB22*	Short-read archive	SRR1639669
HPB2863*	Short-read archive	SRR1640181
HPB5622*	Short-read archive	SRR1373535

Note: Asterisks indicate taxa with high-confidence core (HCC) profiles that exactly match a profile in the HCC database.