Supplemental Figure 3

Name	Sequence
MLST_nifS-12	ΑΑΑΑΤΑGAAAAAACAAGAGAACTTGTAAAAAATTTCATTAATGCAGAATCTGCAAAAAAT
Assembled_MM1_nifS	ΑΑΑΑΤΑGAAAAAAAAAAGAGAACTTGTAAAAAATTT T ATTAATGCAGAATCTGCAAAAAAT
05e484a1-7b54-48fb	AAATAGAAAAAAAAGAGAAACTTGTAAAATT T TTATTAAT-GAATCTGCAAAAT
2e08a83a-bc8f-4976	AACCCGAAAAACAAGAGAACTTATAAAATT T ATTAATGCAGAATCTGCAAAATAT
88624a5a-f257-4af9	AAAGCCGAAAACAAGAGAACTTGTAAAATTT T ATTAATGCAGAATCTGCA-AATAT
ebeaba4b-a7f2-4c91	ΑΑΑΤΑGAAAAACAAGAGAACΤΤGTAAATTT T ATTAATGCAGAACTCTGCAAAATA
a753454d-707d-4b83	AAATAGAAAACAAGGGGACTGTAAATTT T ATTAATGCAGAAT
2cdd8d93-15cd-449d	CAAAATAGAGCTAAGAATTGACAATTT T GACTGCTTGGAGTCTGCAAAATAA
5a187c5f-d9c6-40d1	AAATAGAAAAACAAGAGAACTTAAAAATTT T ATTAATGTGCAGAATCTGCAAAA-
c5dc593f-c1fa-40f8	AAAGGCGAAAACAAGAGAACTTGAAAATTT T ATTAATGCAGAATCTGCAAAATAT
c3fed319-9ceb-4396	AAATAGTAGAAAAAACAGGGGACTTGTAAAATTTTC T CTTAATGCAGAATCTGCAAAATAT
e7fb0bcb-85cb-4a2a	AAAATAGAAAACAAGAGAACTGCTAAATTT T ATTAATTGCAGAATCTG
0177d47f-9cdd-4c4e	AAATAGAAAAAAAAGAGAAACTTGTAAAGTT T ATTAATGCAGAATCTGCAAAA
	*

Name	Sequence
MLST_pepX-1	TACAATGCTCA T GACAATTTAACAGTAATCAGTAGCACTAAAAAAGCAATAAAAGA
Assembled_MM1_pepX	TACAATGCTCA C GACAATTTAACAGTAATCAGTAGCACTAAAAAAAGCAATAAAAGA
817a604f-2391-44b1	ACAATGCGGCA C GACAATTTAACAGTAATCAGTAGCACTAAAAGCAATAAAGATAA
a216f821-fd7a-4ebb	ATACATGCTCA C GACAATTTAACGATGAAGATCAGCACTAAAAAGCAATAAAAGAT
aa375ec3-643a-4f7e	-TACAATTCAA <mark>G</mark> GACAATTTAACAGTAATGAGTAGCACTAAAGCAATAAAATAA
26deeb2d-2462-48ce	TACAATGCTCA C G-ACAATTAACAGCAATCAGTAGCACTAAAAGCAATAAAAGATA
D6a5329d-8245-4941	-ACCAATTCCA C GACATTTGAATAGGTACGTATTAGTAGCACTAAAGCAATAGACAGATA
6e1fdff6-2933-419d	TACAATGCTCA C GACAATTTAAATGATCAGTAGCACTAAAAGGGCAATAAAGAT
cfa9c429-2b73-4daf	TACAATGCTCA C GACAATTTAACAGTAATGAGTAGCACTAAAAGCAATAAAAGATA
b4a50839-4edc-485b	ACATCCTCTCA C GACAATTAACAGTAAACCAGTAGCACTCAAAAAGCAATAGAATA
64f1cd9f-45c5-47a7	TACAATGCTCA C G-ACAATTAACAGTAATCAGTAGCACTAAAACTAATAAAAGATA
	*

Nanopore sequencing reads confirm the novel SNPs in *nifS* and *pepX*. Top: The de novo assembly of the *nifS* gene matches the MLST *nifS* type 12 except for a C->T polymorphism. Eleven nanopore reads align to this region and all contain the T variant. Bottom: The de novo assembly of the *pepX* gene matches the MLST *pepX* type 1 except for a T->C polymorphism. Nine nanopore reads align to this region and eight contain the C variant. The two SNPs of interest are noted with a red asterisk (*).