	10 20	420 430 440
*Accession N° Lmip	F GGCAATGTCAACGACAATTGC	ACTGGCAAGTTAATTGATGGCCA Lmip_I
U92205 L. feeleii sg. 1	TGCG	TTGAATAC
AF022341 L. feeleii sg. 2	TGCG	TTGAATAC
JN191172 L. tunisiensis	TGCA	TTGAATACT
U92209 L. jordanis	AT.CGTGCG	AAGGAAATGGC.TTCCTTC.A.G.ATG.
X83036 L. longbeachae	TAGGTGCA	CCGGCACTGGC.CAACC.GG.A.ATCAG.
S62141 L. micdadei	ATGCA	GAGGCGATGGCTACCG.CCA.G.ATG.
U91637 L. dumoffii	TTGGTGCT	CAGGTTCAGGC.CTCC.GG.A.ATCTG.
U91607 L. anisa	TTGTGAGCT	CAGGTACAGGAAATACCCGG.A.A.CGG.
U91609 L. bonzemanii	TTGTTGCT	CAGGCACAGGAAATACCCGGCA.ATCGG.
AF095229 L. pneumophila sg. 1	TAGGTGCA	CTGGAAATGGTTACCTGG.A.ATCGG.

Supplementary Figure 1 Alignment of Lmip primers and DNA sequences of *mip* gene of *Legionella* species (ClustalW, Bioedit 7.2). These sequences were taken from the GenBank database. (A) Lmip_F primer. (B) Lmip_R primer. sg., serogroup. *Genbank accession number.