Supplementary Table S1. Proteins identified in *L.(V.) braziliensis* control strain (MHOM/BR/94/M15176) by analysis carried out with the "Protein Lynx" software using the *L.(V.) braziliensis* database, from data collected by mass spectrometry of the tryptic peptides.

Protein	Entry	mW (Da)	pI	Peptides	Theoretical Peptides	Coverage (%)
ATPase alpha subunit *L. braziliensis*	134059187	62576	9.9697	8	40	12.1951
putative heat shock 70 related protein 1 mitochondrial precursor *L. braziliensis*	134064193	70484	5.8301	9	59	16.4363

Supplementary Table S2. Proteins identified in *L.(V.) braziliensis* isolated from the patient sample by analysis carried out with the "Protein Lynx" software using the *L.(V.) braziliensis* database, from data collected by mass spectrometry of the tryptic peptides.

Protein	Entry	mW (Da)	pI	Peptides	Theoretical Peptides	Coverage (%)
ATPase alpha subunit *L. braziliensis*	134059187	62576	9.9697	4	40	7.3171
putative heat shock 70 related protein 1 mitochondrial precursor *L. braziliensis*	134064193	70484	5.8301	13	48	19.5719