

IDP spot	Organism	Protein ID	Function	Score	Best Combination
					REM465_GLOBPIPE_IUPRED_VSL2B
IUP1	<i>L. major</i>	LmjF27.0240	kinetoplast-associated protein-like protein	285	X
IUP4	<i>L. major</i>	LmjF32.2520	hypothetical protein, unknown function	31	X
IUP5	<i>L. major</i>	LmjF09.0910	calmodulin, putative	505	X
IUP6	<i>L. major</i>	LmjF27.0240	kinetoplast-associated protein-like protein	435	X
IUP8	<i>L. major</i>	LmjF05.0380	microtubule-associated protein, putative	188	X
IUP10	<i>L. major</i>	LmjF05.0380	microtubule-associated protein, putative	449	X
IUP11	<i>L. major</i>	LmjF05.0380	microtubule-associated protein, putative	228	X
IUP25	<i>L. major</i>	LmjF23.1020	hypothetical protein, unknown function	1973	X
IUP26a	<i>L. major</i>	LmjF04.0770	nascent polypeptide associated complex subunit-like protein	1093	X
IUP26b	<i>L. major</i>	LmjF19.1160	hypothetical protein, conserved	1054	X
IUP27	<i>L. major</i>	LmjF04.0770	nascent polypeptide associated complex subunit-like protein	1854	X
IUP32	<i>L. major</i>	LmjF28.2770	heat-shock protein hsp70, putative	608	X
IUP34	<i>L. major</i>	LmjF30.2460	heat shock 70-related protein 1, mitochondrial precursor, putative	1591	X
IUP37	<i>L. braziliensis</i>	no hits	-	-	
IUP40	<i>L. braziliensis</i>	no hits	-	-	
IUP42	<i>L. braziliensis</i>	LbrM25_V2.0580	eukaryotic initiation factor 5a, putative	2160	X
IUP43	<i>L. braziliensis</i>	LbrM28_V2.1300	glucose-regulated protein 78, putative	363	X