

Immunoproteomic Identification and Characterization of *Leishmania* Membrane Proteins as Non-Invasive Diagnostic Candidates for Clinical Visceral Leishmaniasis.

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Table S1. Molecular masses of urine reactive *Leishmania* peptides obtained from Imagelab.

Band No.	Mol. Wt. (kDa)	Relative Front (R _f)	Volume (Int)	Band %.	Lane %
1	120	0.198	5,005,392	14.0	1.5
2	97	0.253	4,432,848	12.4	1.3
3	91	0.271	4,268,157	12.0	1.2
4	72	0.292	3,883,404	10.8	1.1
5	63	0.345	7,100,688	19.8	2.1
6	55	0.386	9,989,280	27.9	2.9
7	51	0.410	8,012,105	19.7	2.0
8	45	0.467	1,141,140	3.2	0.3
9,	36	0.530	678,468	1.9	0.2
10	34	0.605	2,143,680	6.0	0.6
11	31	0.648	1,492,092	4.2	0.4

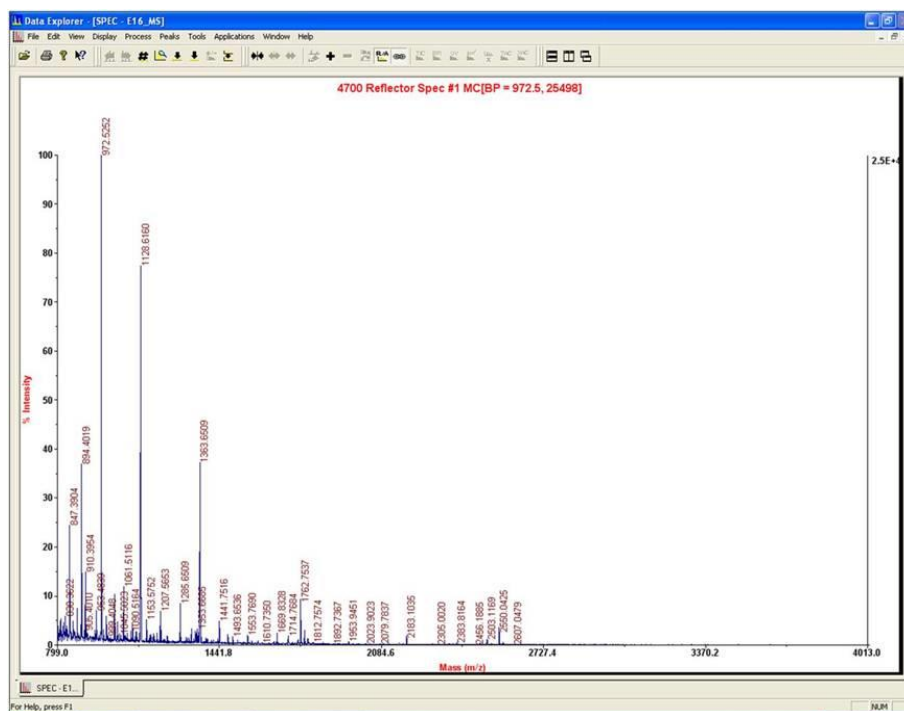


Figure S1. MALDI-TOF spectra of the tryptic fragments obtained from peptide 51 kDa.

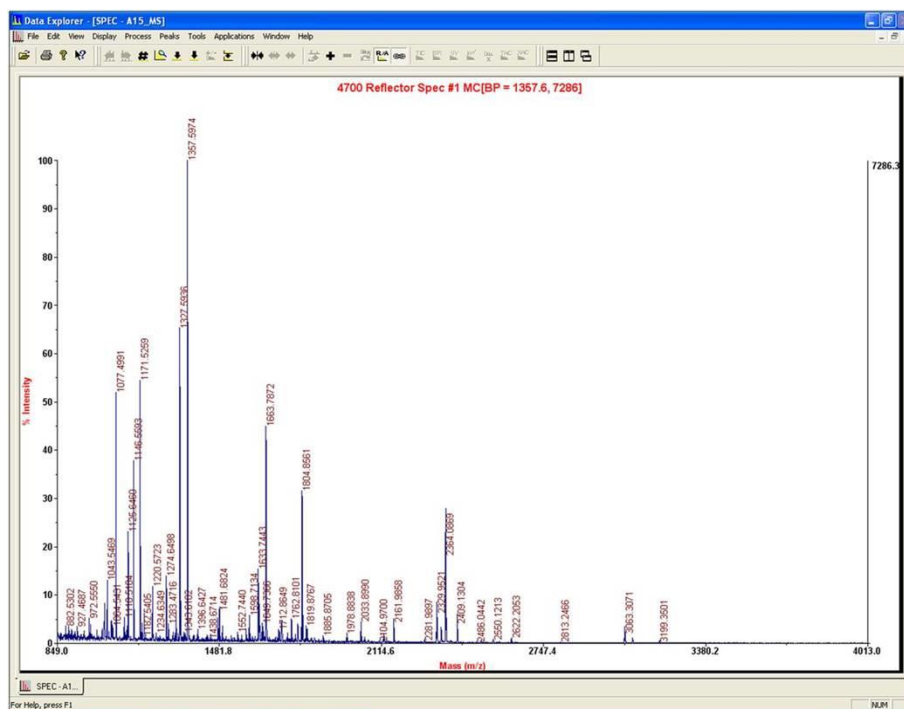


Figure S2. MALDI-TOF spectra of the tryptic fragments obtained from peptide 55 kDa.

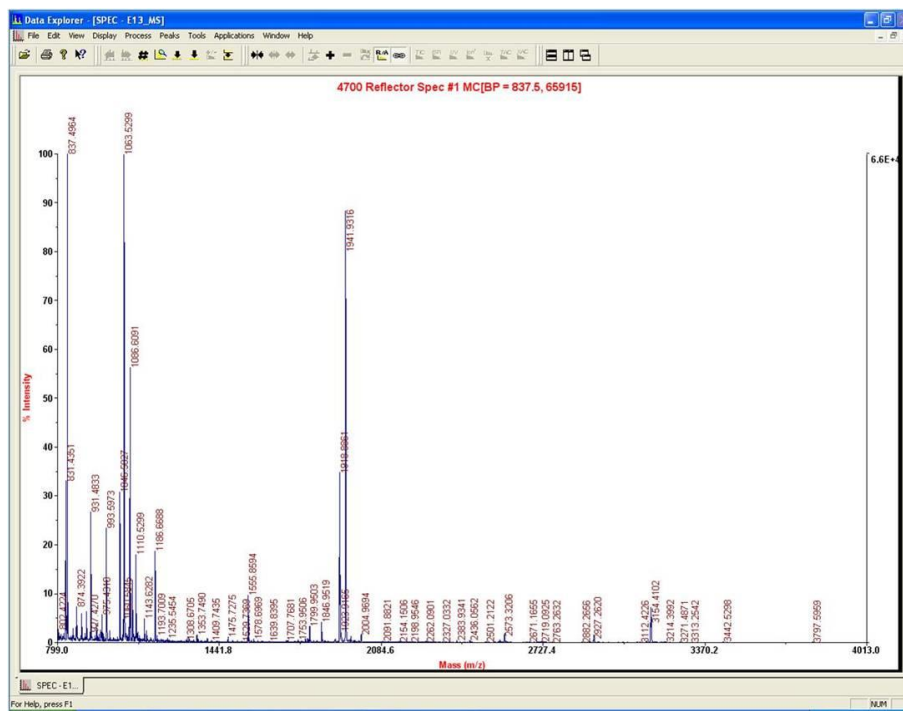


Figure S3. MALDI-TOF spectra of the tryptic fragments obtained from peptide 63 kDa.

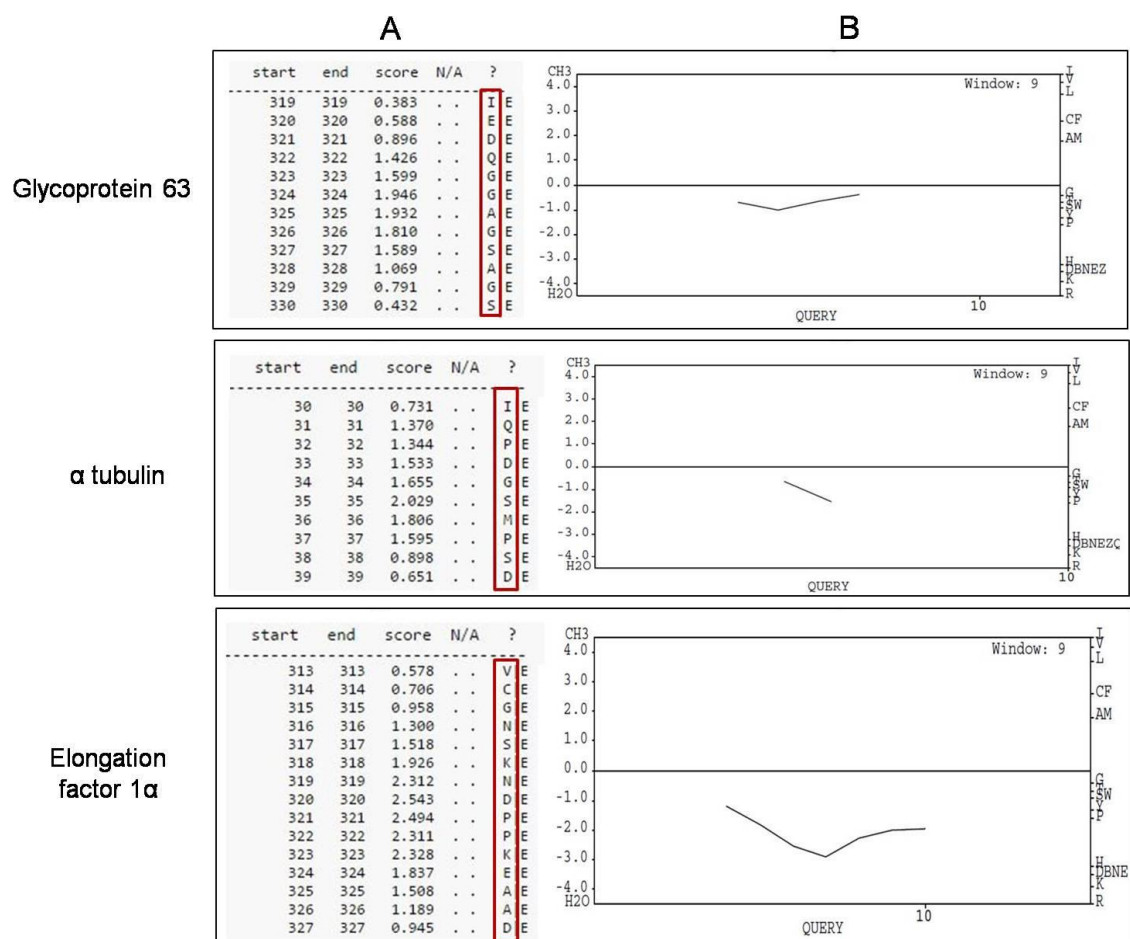


Figure S4. B cell epitope prediction of protein glycoprotein 63, α tubulin and elongation factor 1 α in BepiPred software (A) and Kyte & Doolittle hydropathy plot of selected epitopes (B).

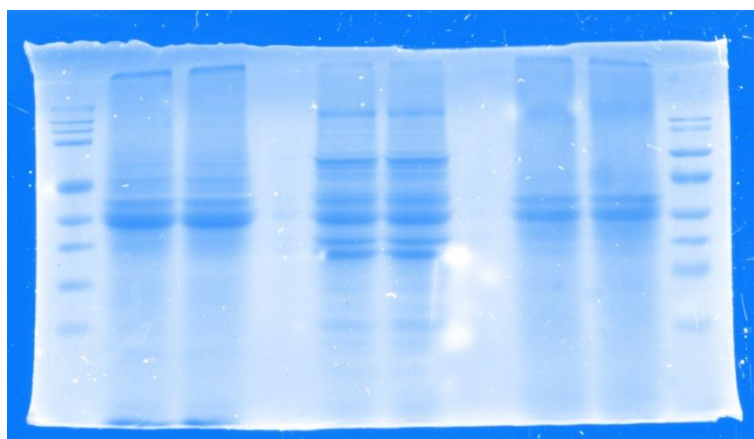


Figure S5. Original gel of figure 1

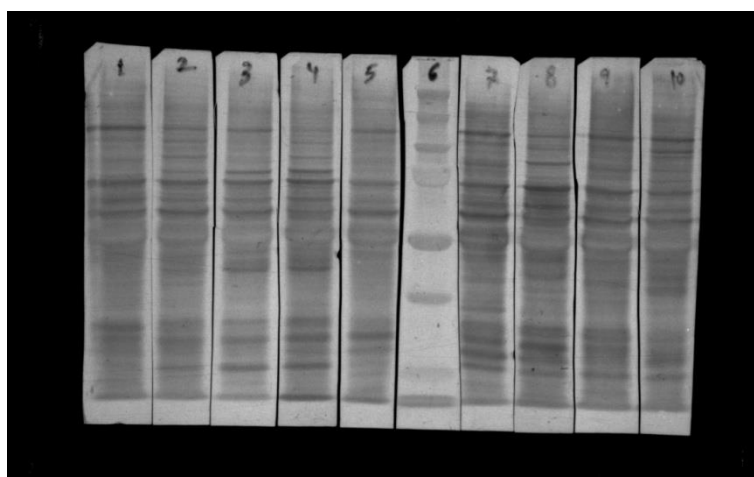


Figure S6. Original blot of figure 2A

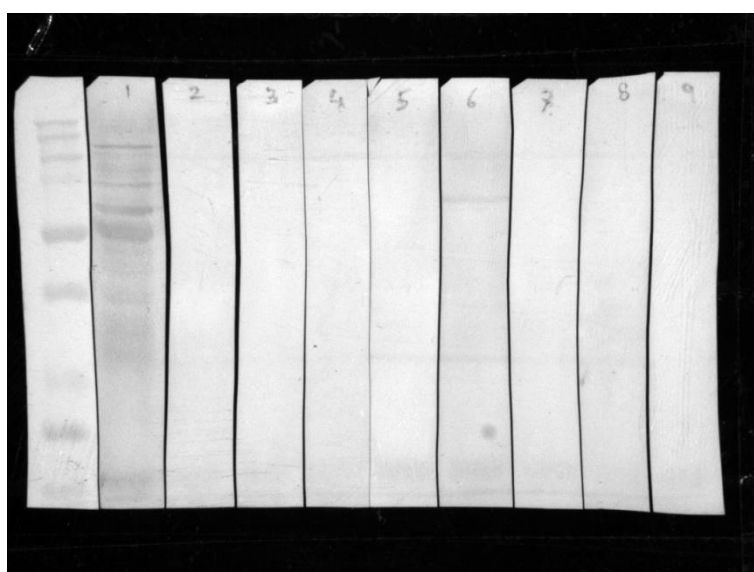


Figure S7. Original blot of figure 2B



Figure S8. Original blot of figure 3A

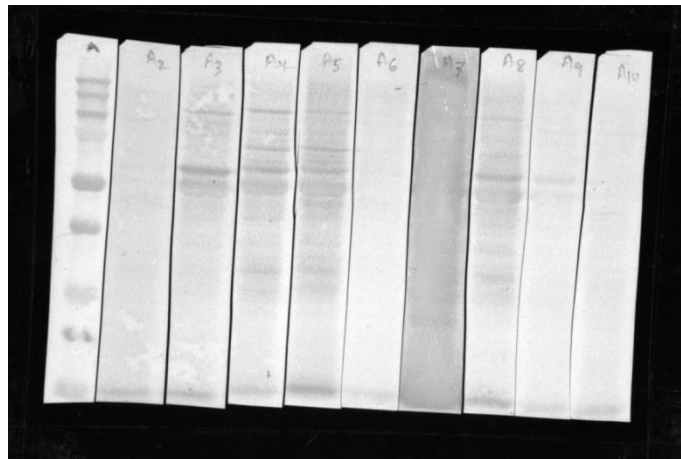


Figure S9. Original blot of figure 3B

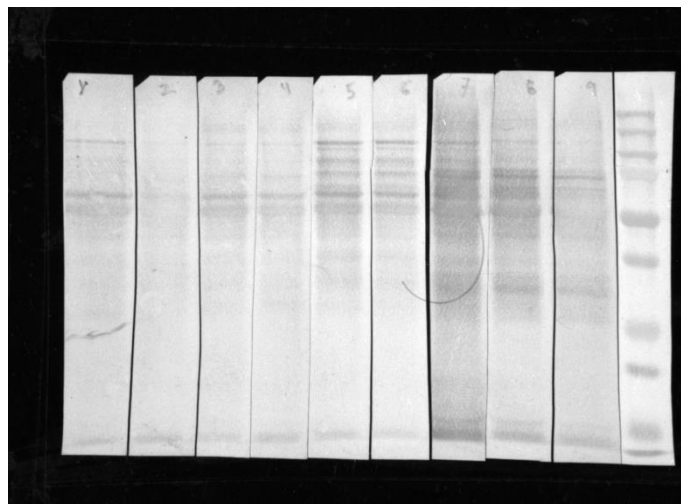


Figure S10. Original blot of figure 3C

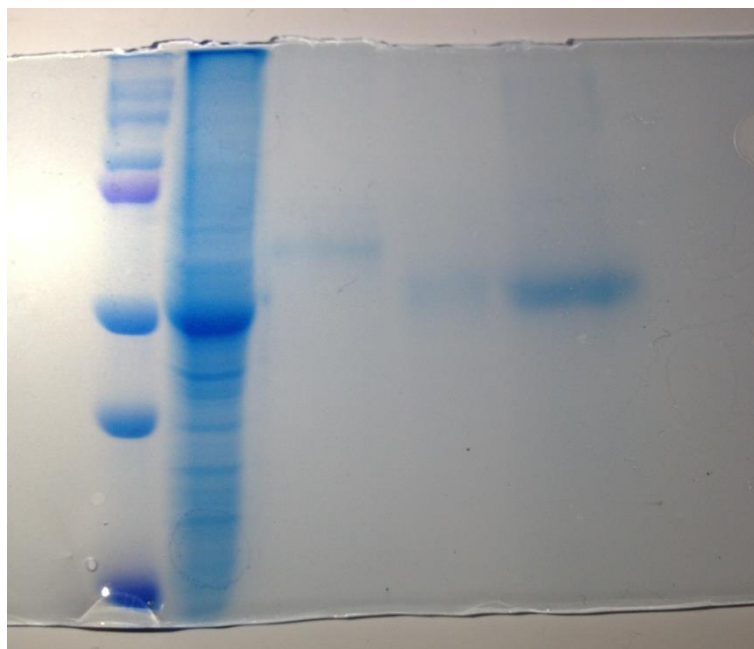


Figure S11. Original gel of figure 4B