

Supplementary Figure 1: Interactions of the cholesterol with selected cytokines possess diagnostic value in the pathogenesis of cutaneous leishmaniosis. Proteins are demonstrated as spheres while cholesterol is shown as capsule-shaped node. Stronger associations are represented by thicker lines. Protein-protein interactions are shown in grey, chemical-protein interactions in green and interactions between chemicals in red.

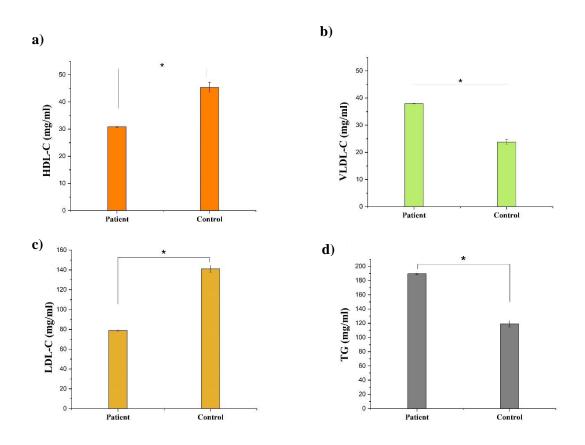
Protein-Protein Network Stats

- number of nodes: 22
- number of edges: 83
- average node degree: 7.55

clustering coefficient: 0.88

expected number of edges: 24

PPI enrichment p-value: 0



Supplementary Figure 2: Average total of (a) HDL-C, (b) VLDL-C, (c) LDL-C and (d) TG.

HDL-C = high-density lipoprotein-cholesterol; VLDL-C = very-low-density lipoprotein-cholesterol;LDL-C = low density lipoprotein-cholesterol; <math>TG = triglycerides. $*P \leq 0.05$.

	Cholesterol in mg/dL	IL-10 in ng/L	IL-12 in ng/L	TNF-α in ng/L
Patient (n = 50)	143.23 ± 1.27	523.77 ± 8.29	202.48 ± 4.20	173.57 ± 18.96
Control $(n = 25)$	156.40 ± 1.72	14.08 ± 1.72	144.09 ± 5.95	15.98 ± 0.92
T-test	4.335*	21.349*	15.557*	61.570*
P value	0.0001	0.0001	0.0001	0.0001

Supplementary Table 1: The serum cholesterol and cytokine profile in cutaneous leishmaniosis

 $IL = interleukin; TNF-\alpha = tumor necrosis factor alpha.$

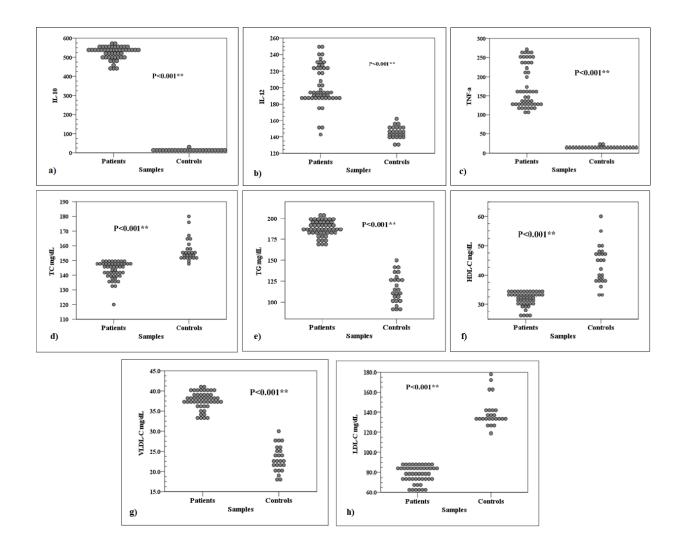
**P* ≤0.01.

Supplementary Table 2: Comparison between patients and control in lipid profile

	Mean ± SE in mg/dL						
	Cholesterol	Triglyceride	HDL	VLDL	LDL		
Patients	141.47 ± 0.71	189.44 ± 1.01	$\begin{array}{c} 30.80 \pm \\ 0.32 \end{array}$	37.90 ± 0.22	78.85 ± 1.02		
Control	161.65 ± 2.94	119.05 ± 4.48	45.35 ± 1.89	$\begin{array}{c} 23.81 \pm \\ 0.90 \end{array}$	141.01 ± 3.71		
T-test	4.079*	6.011*	2.321*	1.247*	5.445*		
P value	0.0001	0.0001	0.0001	0.0001	0.0001		

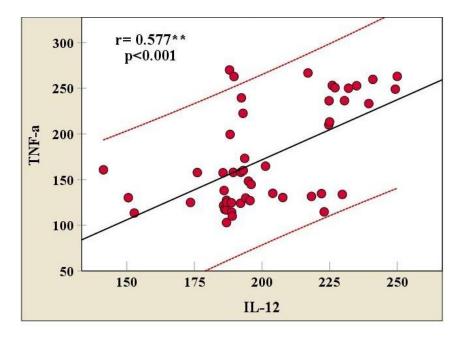
SE = standard error; *HDL* = high-density lipoprotein; *VLDL* = very-low-density lipoprotein; *LD* = low density lipoprotein.

**P* ≤0.01.



Supplementary Figure 3: The dot plots of differences in the distribution of cytokines and lipid profiles levels between cases and controls of (a) IL-10, (b) IL-12, (c) TNF- α , (d) TC, (e) TG, (f) HDL-C, (g) VLDL-C and (h) LDL-C.

 $IL = interleukin; TNF-\alpha = tumor necrosis factor alpha; TC = total cholesterol; TG = triglycerides; HDL-C = high-density lipoprotein-cholesterol; VLDL-C = very-low-density lipoprotein-cholesterol; LDL-C = low density lipoprotein-cholesterol.$



Supplementary Figure 4: The dot plot of significant correlation of IL-12 and TNF- α . The separate red line expressed as 95% CI.

 $IL = interleukin; TNF-\alpha = tumor necrosis factor alpha; CI = confidence interval.$