

## *Supplementary Material*

### **High levels of PF4, VEGF-A, and classical monocytes correlate with the platelets count and inflammation during active tuberculosis**

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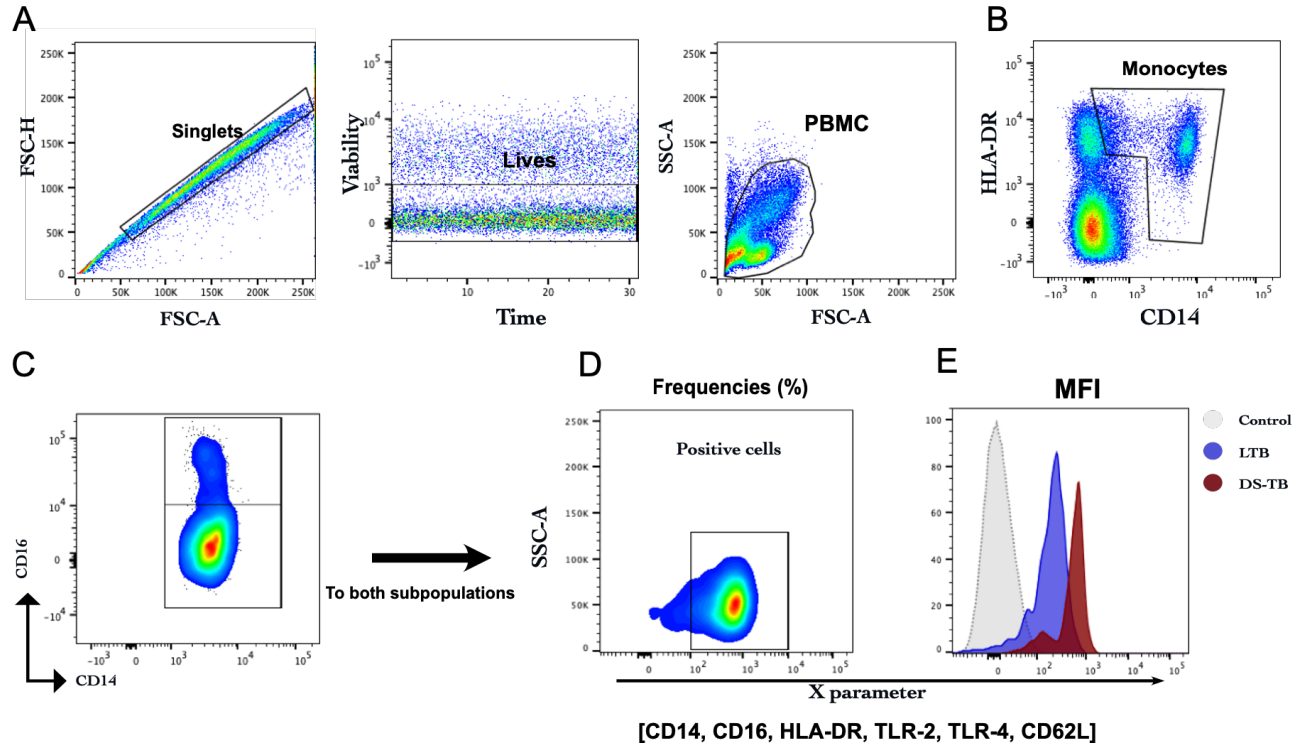
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## Figures

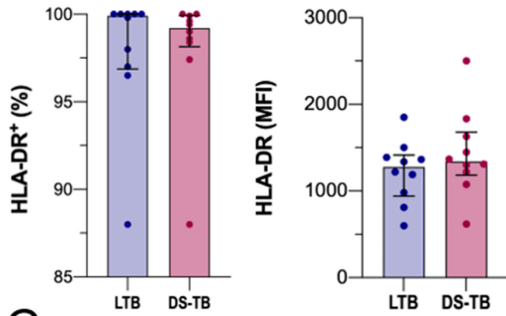


**Figure S1. Gating strategies for flow cytometrical analysis.** Representative example of the results obtained in cell subpopulation monocytes subsets from peripheral blood analysis (5,000 threshold): singlets cells (**A**) were identified through forward scatter (FSC-A versus FSC-H). Then, viable cells were selected, followed by forward scatter (FSC) and side scatter (SSC) dot plot to selected live monocytes based on HLA-DR and CD14 expression (**B**). Two subpopulations of monocytes were identified based on CD14 and CD16 expression as Classical (CD14<sup>+</sup>) and Non-classical (CD14<sup>+</sup>CD16<sup>+</sup>) (**C**). The frequency (**D**) and mean fluorescent intensity (MFI) (**E**) of indicated molecules were calculated. All MFI analyzed included a latent (blue), active (red) tuberculosis patients and a negative control (gray).

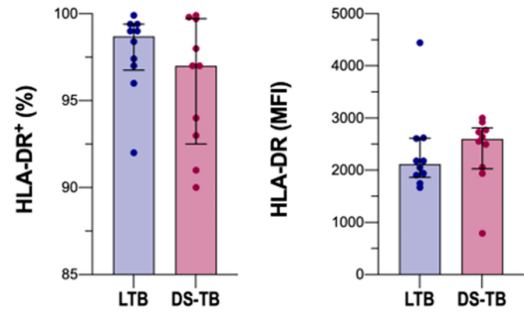
## Classical

## Non-classical

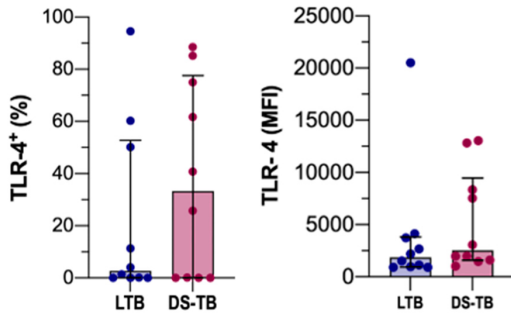
**A**



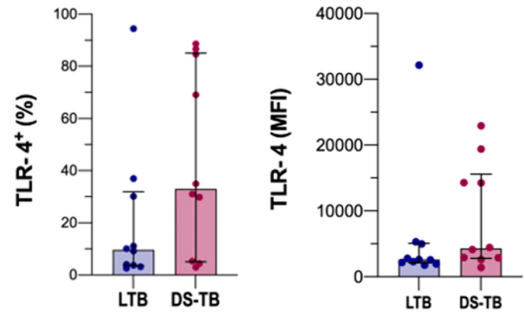
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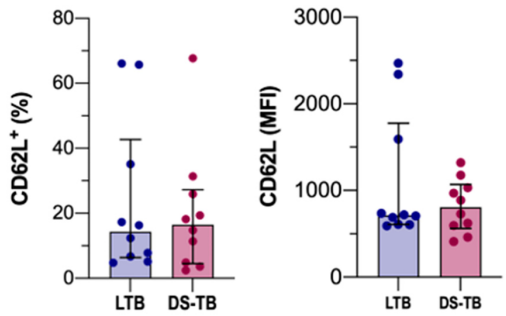
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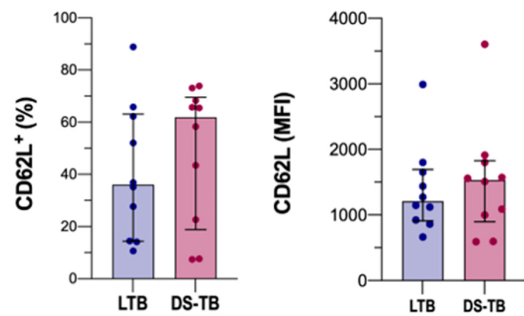
**D**



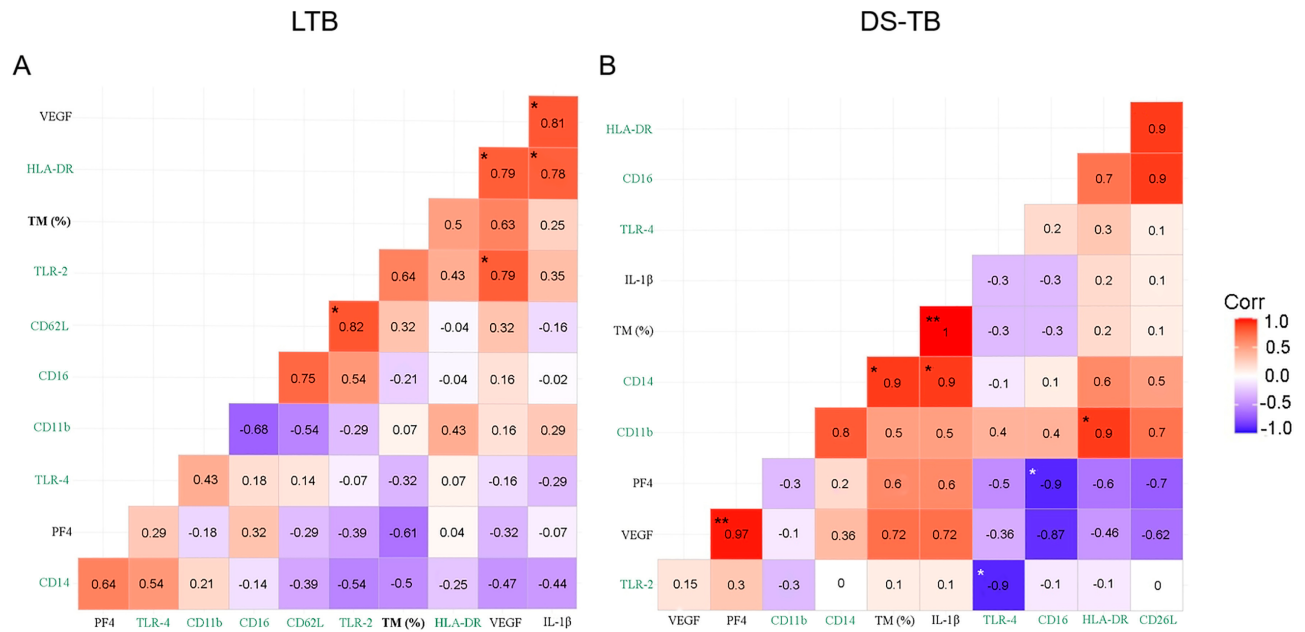
**E**



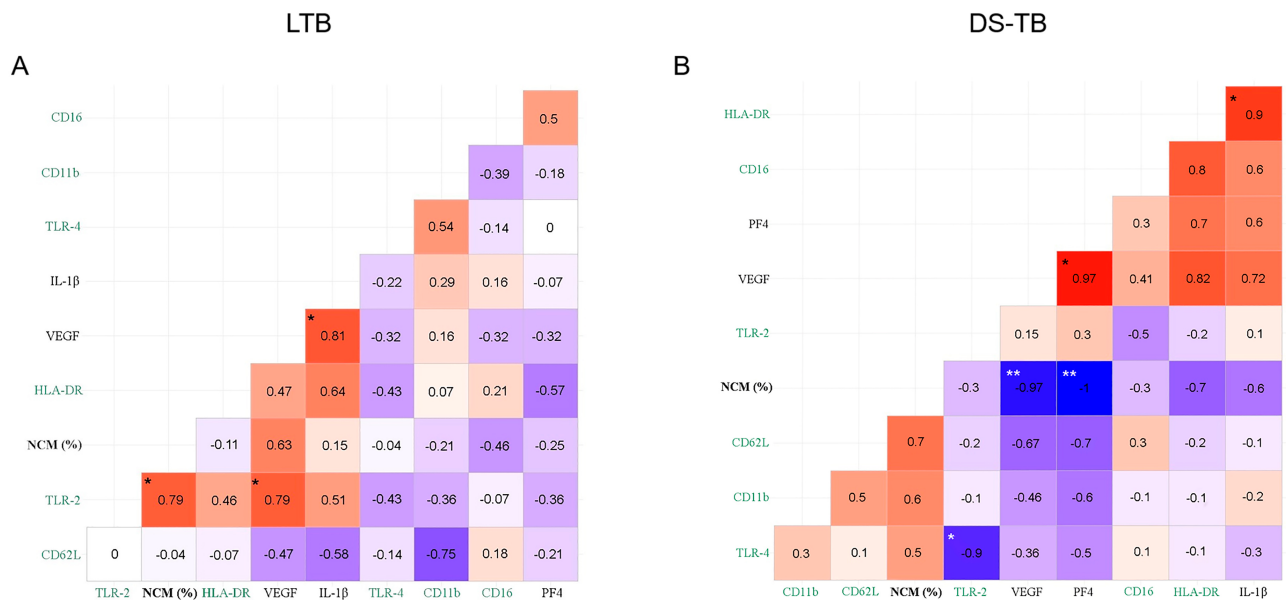
**F**



**Figure S2. Frequency of subsets monocytes cells.** Frequency and mean fluorescent intensity (MFI) of total monocytes cells in latent (LTB) and active (DS-TB) TB patients according to HLA-DR (**A, B**), TLR-4 (**C, D**) and CD62L (**E, F**) on classical (**A, C** and **E**) and non-classical (**B, D** and **F**) monocytes subsets. Data are shown as median with interquartile range (IQR, 25–75). The statistical comparison was performed using Mann-Whitney U Test.



**Figure S3. Correlation between Total monocytes (TM) and plasmatic levels of soluble molecules in latent (A) and active (B) TB patients.** Correlations are presented with the value of Spearman's Rho in the corresponding box. Significant correlations are represented with asterisks (\*  $p < 0.05$ , \*\*  $p < 0.01$ ). The red color indicates a strong positive correlation and the blue color indicates a strong negative correlation. Green labels are mean fluorescent intensity values.



**Figure S4. Correlation between non-classical monocytes (NCM) and plasmatic levels of soluble molecules in latent (A) and active (B) TB patients.** Correlations are presented with the value of Spearman's Rho in the corresponding box. Significant correlations are represented with asterisks (\*  $p < 0.05$ , \*\*  $p < 0.01$ ).

< 0.05, \*\*p < 0.01). The red color indicates a strong positive correlation and the blue color indicates a strong negative correlation. Green labels are mean fluorescent intensity values.

## Tables

**Table S1.** Antibodies used for flow cytometry, ELISA and Bio-Plex.

Molecule/Product	Conjugate to	Clone	Catalog	Company
CD16	PE-Cy7	3G8	302016	Biolegend
CD11b	PerCP-Cy5.5	ICRF44	301328	Biolegend
CD284	PE	HTA125	312806	Biolegend
CD282	FITC	TL2.1	309706	Biolegend
HLA-DR	APC-Cy7	L243	307617	Biolegend
CD14	BV510	63D3	367123	Biolegend
CD62L	BV421	DREG-56	304828	Biolegend
Viability	PE TexRed	NA <sup>‡</sup>	423109	Biolegend
CXCL4/PF4	NA <sup>‡</sup>	NA <sup>‡</sup>	DPF40	R&D System
MCP-1	NA <sup>‡</sup>	NA <sup>‡</sup>	438804	BioLegend
IL-17A	NA <sup>‡</sup>	NA <sup>‡</sup>	433914	Biolegend
MUC5B	NA <sup>‡</sup>	NA <sup>‡</sup>	MBS2024599	MyBioSource
IL-6	NA <sup>‡</sup>	NA <sup>‡</sup>	171B0006M	Bio-Rad Laboratories
IP-10	NA <sup>‡</sup>	NA <sup>‡</sup>	171B5020M	Bio-Rad Laboratories
VEGF-A	NA <sup>‡</sup>	NA <sup>‡</sup>	171B5027M	Bio-Rad Laboratories

PDGF-AB	NA <sup>‡</sup>	NA <sup>‡</sup>	171B5024M	Bio-Rad Laboratories
Reagent Kit III with Flat Bottom Plate	NA <sup>‡</sup>	NA <sup>‡</sup>	171-304090M	Bio-Rad Laboratories
Human Cytokine Standards	NA <sup>‡</sup>	NA <sup>‡</sup>	171D50001	Bio-Rad Laboratories

<sup>‡</sup>NA, not applicable.

**Table S2.** Comparison of parameters evaluated with the healthy donors group.

Biomolecule	HD	LTB	DS-TB	<i>p</i> value HD vs LTB	<i>p</i> value HD vs DS-TB
PF4 (ng/mL)	2250 (1929- 2816)	2012 (1213- 2727)	3108 (2578- 3750)	ns	ns
PDGF-BB (pg/mL)	420.2 (30.81- 635.9)	65.98 (28.46- 199.1)	186 (19.39- 257.9)	ns	ns
VEGF-A (pg/mL)	0 (0-0)	0 (0-0.55)	9.64 (0- 22.69)	ns	0.0009
MCP-1 (pg/mL)	79.11 (61.22- 94.48)	99.04 (88.15- 118.3)	91.16 (71.91- 123.0)	ns	ns
IL-6 (pg/mL)	0.22 (0-	0.73 (0.4425-	14.78 (4.428-	ns	0.0002

	1.275)	1.0)	31.60)		
IL-1 $\beta$ (pg/mL)	0 (0-0)	0.035 (0- 0.1425)	0.465 (0.33- 0.7625)	ns	<0.0001
IP-10 (pg/mL)	118.3 (103.7- 332.7)	282 (161.7- 473.6)	2718 (1033- 5658)	ns	0.0003

Data are shown as median with interquartile range (IQR, 25–75). The statistical comparison was performed using Mann-Whitney U Test. HD: healthy donor, LTB: latent tuberculosis, DS-TB: drug-sensitive tuberculosis, PF4: platelet factor-4, PDGF-BB: platelet-derived growing factor, VEGF-A: vascular endothelial growing factor, MCP-1: monocyte chemoattractant protein-1, IL-6: interleukin-6, IL-1b: interleukin-1 beta, IP-10: Interferon- $\gamma$  inducible protein 10, ns: not significant.