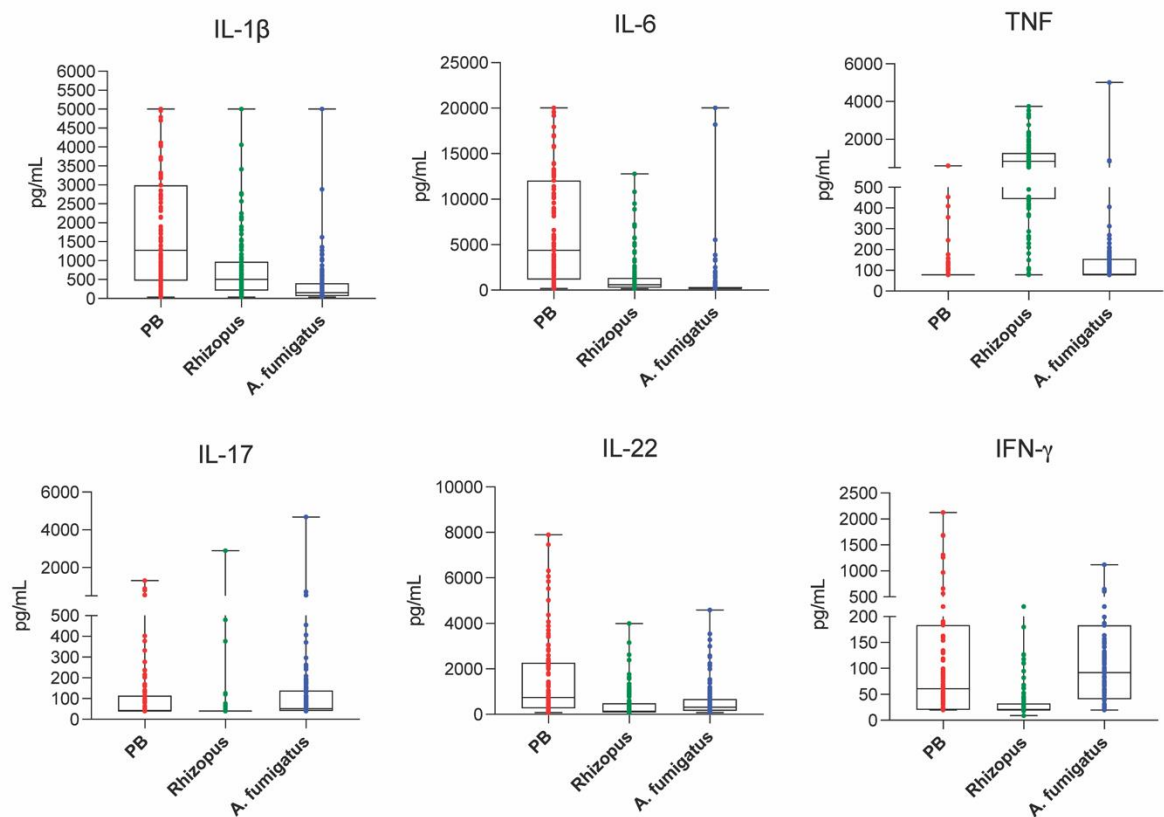
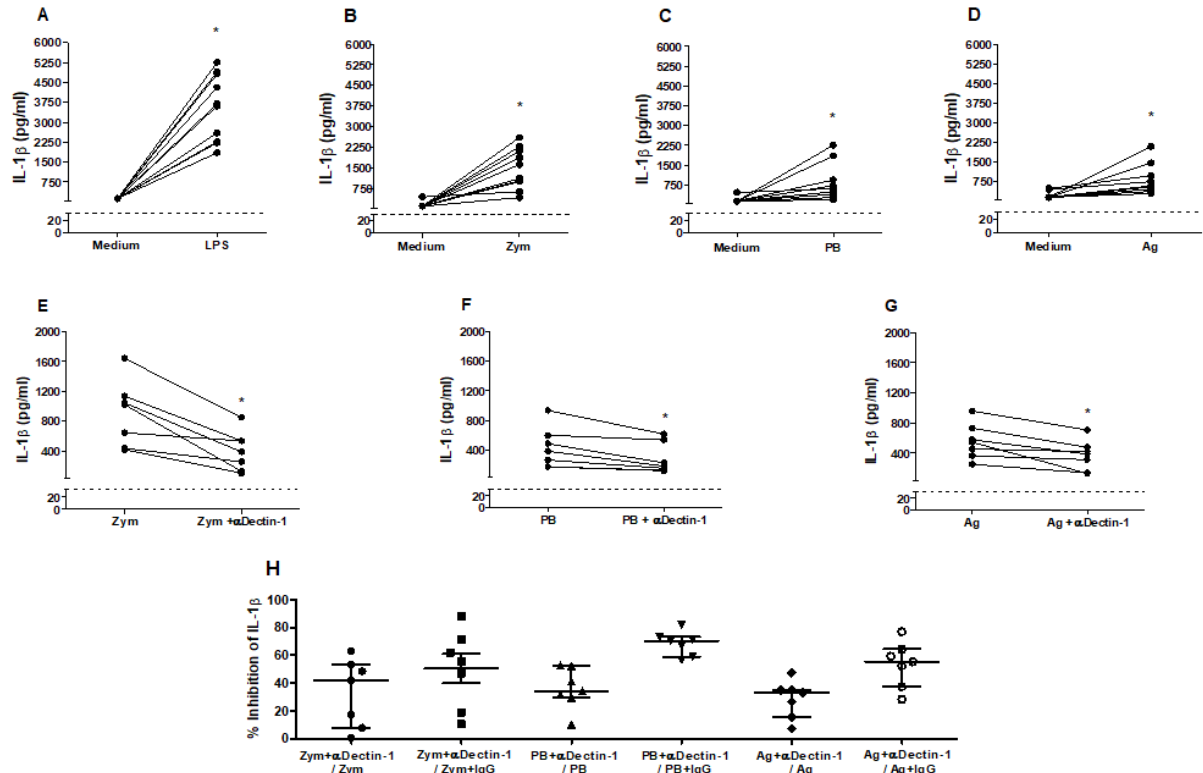


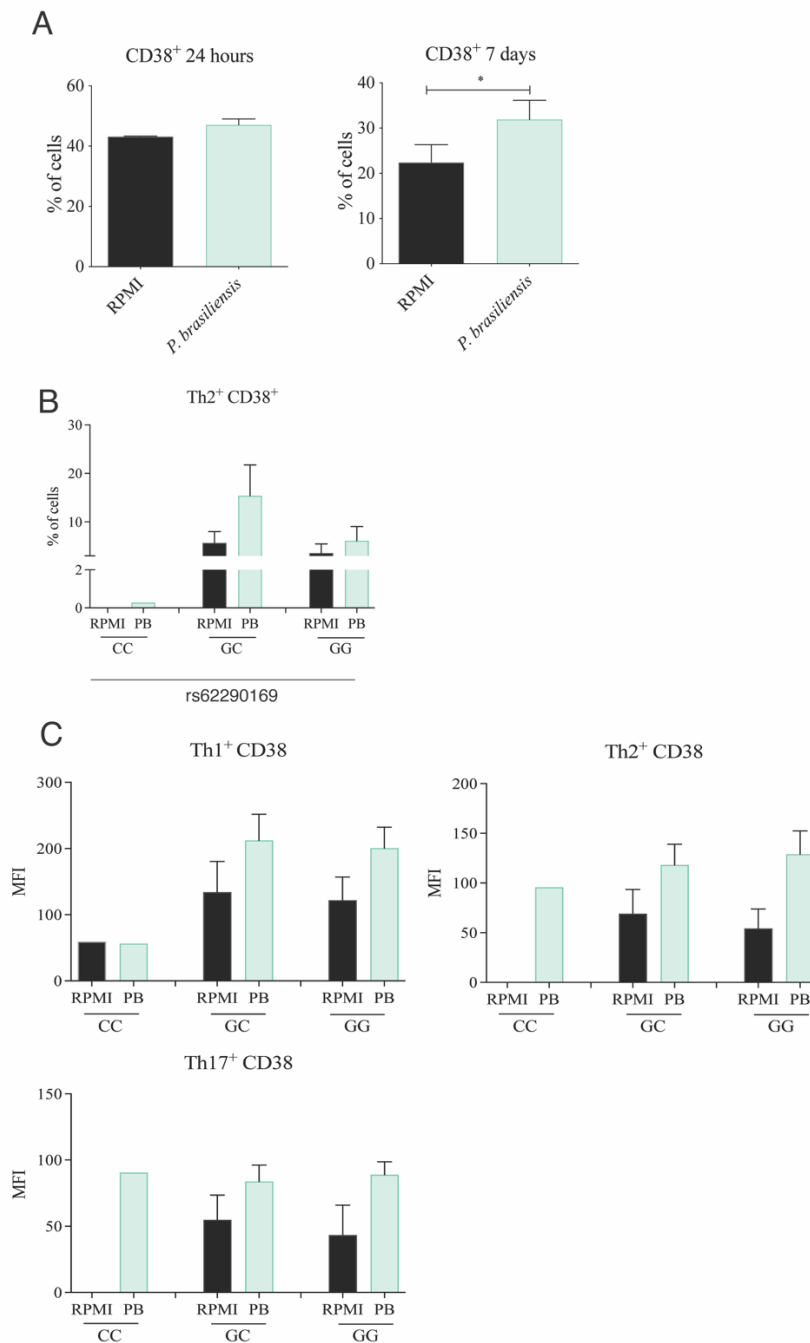
Supplementary Information



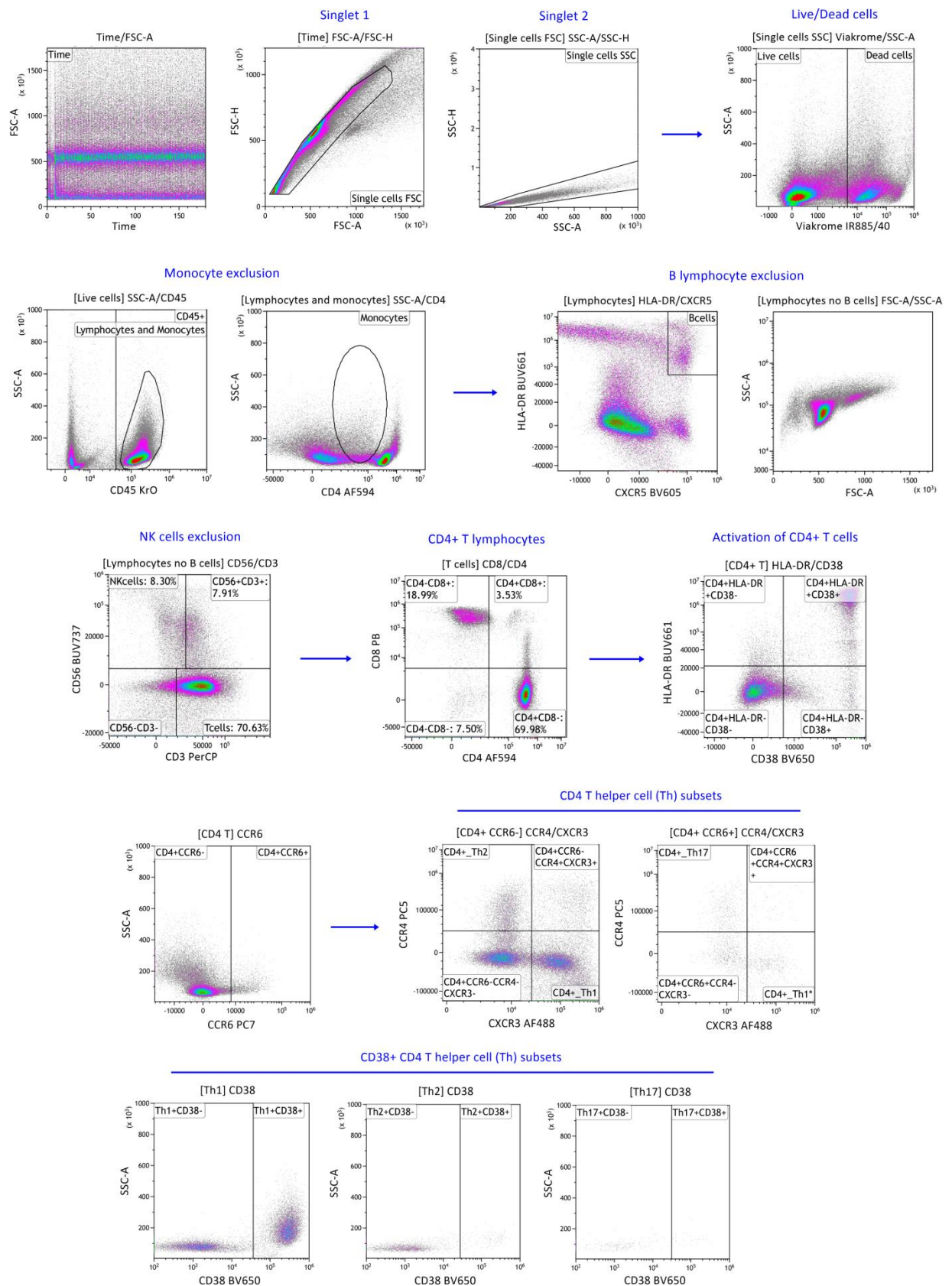
Supplementary Figure S1: Inter-individual variation upon stimulation. IL-1 β , TNF, IL-6 (24 h of incubation), IFN γ , IL-22 and IL-17 (seven days of incubation) levels after PBMC stimulation with heat-killed yeasts of *P. brasiliensis*, conidia of *A. fumigatus* and or *R. oryzae* (2.5 PBMCs:1 fungus; cytokines were measured by ELISA in pg/mL); The data is displayed as box-plots with median, quartiles with maximal and minimal values.



Supplementary Figure S2. Blocking of Dectin-1 decreases IL-1 β production induced by *P. brasiliensis*. PBMCs (2.5×10^5 cells/well) obtained from healthy donors were incubated in the absence or presence of neutralizing anti-Dectin-1 (α Dectin-1 at 10 μ g/mL) or control IgG antibodies (IgG at 10 μ g/mL) for 1 h. Then, the cells were stimulated with 100 ng/mL of lipopolysaccharide (LPS), 50 μ g/mL of Zymosan (Zym), 1×10^6 yeasts/mL (2.5 PBMCs:1 yeast) of heat-killed *P. brasiliensis* (PB) or 20 μ g/mL of particulate antigens from *P. brasiliensis* yeasts (Ag) for 24 h. The supernatant was collected to measure the amount of IL-1 β by ELISA. In A-D: The IL-1 β production induced by LPS, Zym, PB or Ag compared to the negative control, Medium (n = 10). In E-G: The amount of IL-1 β produced after stimulation of PBMCs with Zym, PB or Ag, in the absence or presence of neutralizing anti-Dectin-1 antibodies (n = 7) (Wilcoxon paired test, * p < 0.05). (H) Percentage of inhibition of IL-1 β due to blocking of Dectin-1 in relation of stimuli in the absence or presence of IgG control antibodies (n = 7). The data are presented as individual values, medians and interquartile ranges.



Supplementary Figure S3. Frequency of CD38 expressing cells. (A) Frequency of cells expressing CD38 in total PBMCs stimulated with *P. brasiliensis* for 24 hours and 7 days, by flow cytometry (n = 6, Wilcoxon paired test, * p < 0.05). (B) CD38 protein expression upon exposure to *P. brasiliensis* for seven days assessed by flow cytometry in Th2 cells of individuals part of the 200FG cohort carrying the different genotypes of the rs62290169 SNP (*PROM1* gene) CC (n = 1), GC (n = 4), GG (n = 6) and (D) CD38 mean fluorescence intensity (MFI) in Th1, Th2 and Th17 cells (Wilcoxon paired test, * p < 0.05).



Supplementary Figure S4. Representative dot plots of the indicated receptors in PBMCs upon *P. brasiliensis* exposure for 7 days. Related to figure 4 and supplementary 2. The cells were stained 4

with anti-HLA-DR, anti-CD56, anti-CD3, anti-CD4, anti-CD8, anti-CD45, anti-CXCR5, anti-CD38, anti-CXCR3, anti-CCR4, anti-CCR6 and ViaKrome as viability dye. Percentages of Th1, Th2, and Th17 expressing CD38 were obtained from the indicated CD4⁺ T helper cell subsets.