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Supporting Information

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**Bromodomain inhibitor I-BET151 suppresses immune responses during
fungal–immune interaction**

Compound Number	Cytokine	Mean pIC50	SD	Range	Number of curves (n)	Mean Hill Slope	Mean asymptote max
GSK1210151A	TNF α	6,5	0,43	5.9 - 7.8	68	1,3	105
GSK1210151A	IL-6	6,1	0,36	5.5 - 7.8	116	1,1	98

% Inhibition	[GSK1210151A] M	
	TNF α	IL-6
1,25	1,10E-08	1,50E-08
2,5	1,89E-08	2,84E-08
5	3,28E-08	5,46E-08
10	5,83E-08	1,08E-07
20	1,09E-07	2,25E-07
30	1,65E-07	3,68E-07
40	2,31E-07	5,49E-07
50	3,16E-07	7,94E-07
60	4,32E-07	1,15E-06
70	6,07E-07	1,72E-06
80	9,19E-07	2,80E-06
90	1,71E-06	5,85E-06
95	3,05E-06	1,15E-05
97,5	5,30E-06	2,22E-05

Figure S1. Target engagement plots for GSK1210151A (I-BET151).

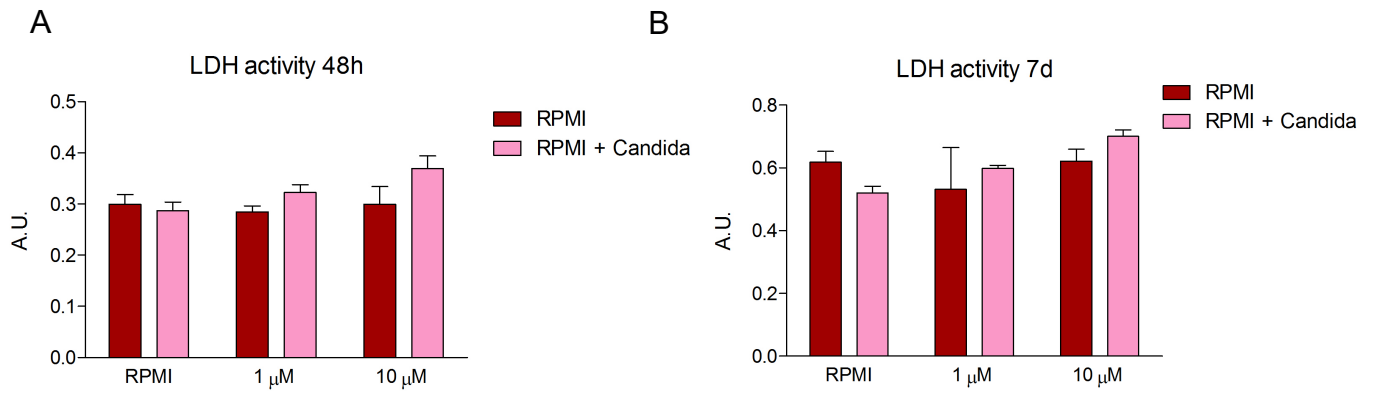


Figure S2. Cytotoxicity assay for I-BET151 treatment. Detection of cell death by lactate dehydrogenase (LDH) assay in PBMCs treated with control medium, 1 μ M or 10 μ M I-BET151 and stimulated with RPMI or 10^6 heat-killed *C. albicans* conidia during 48h (A) or 7 days (B) (mean \pm SEM, n =3 (from 3 different individual donors)).

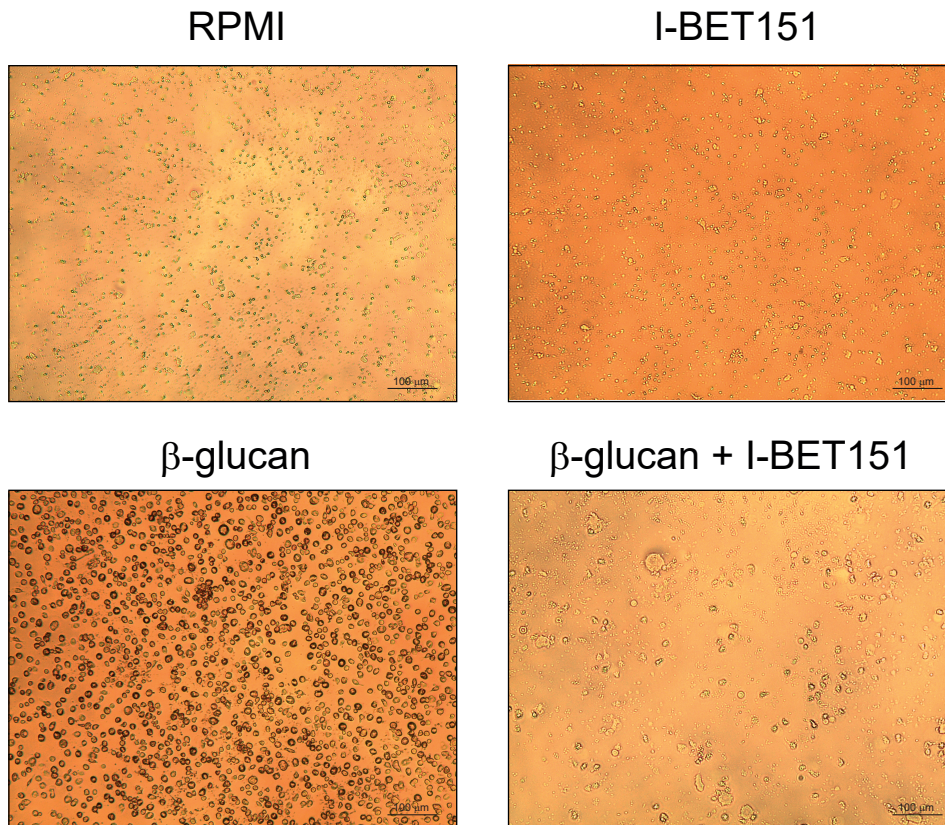


Figure S3. Morphological changes induced by trained immunity.

Morphology of cells after 24 h of training and 6 days of rest when the cells were trained with RPMI (negative control), 10 μM I-BET151, 1 mg/ml β-glucan or 10 μM I-BET151 + 1 mg/ml β-glucan. Pictures were taken at day 6, magnification x10.

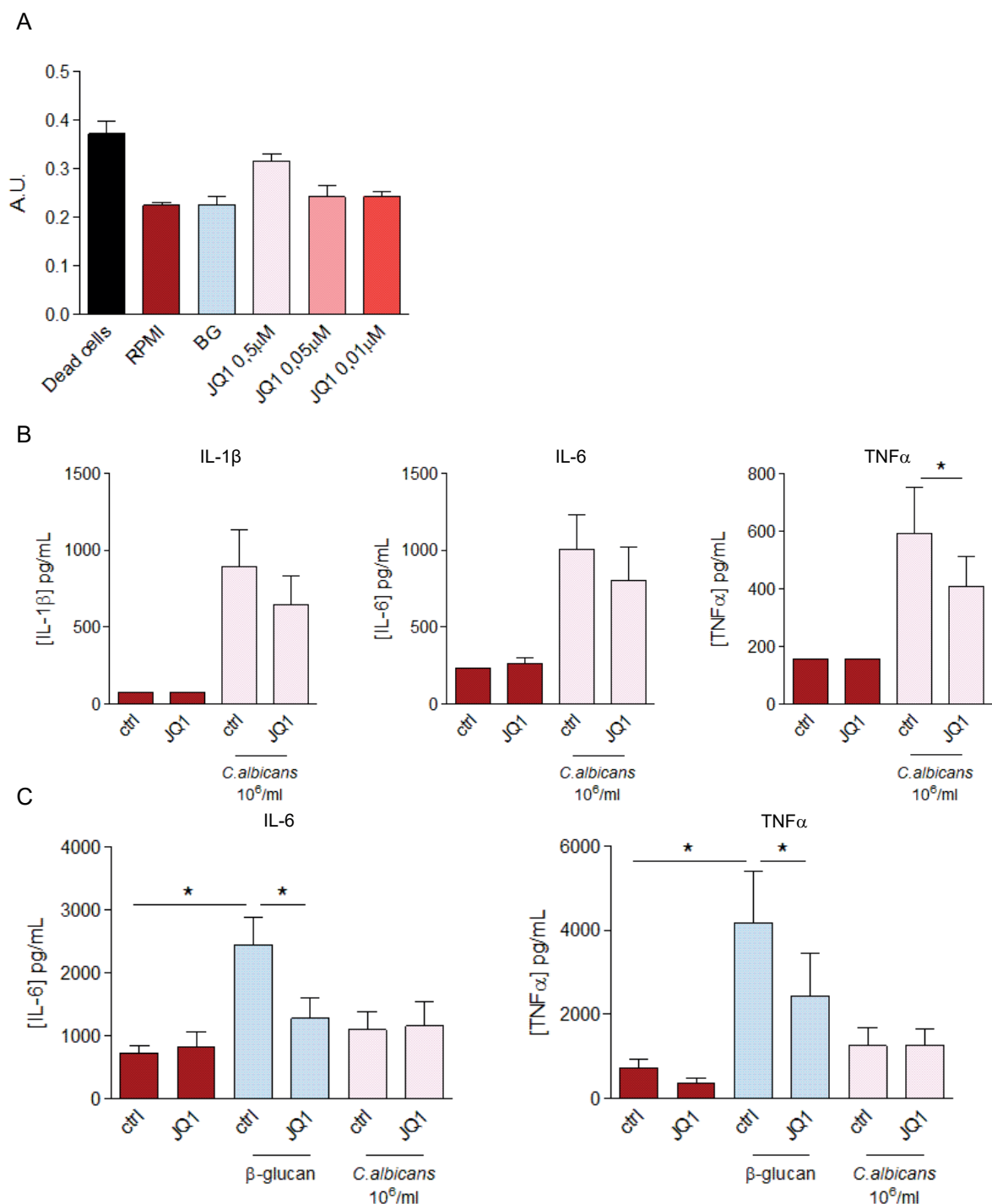


Figure S4. Effects of JQ1 in the induction of trained immunity.

(A) Cytotoxicity assay for the treatment of cells with three different concentrations of JQ1. Dead cells were used as a positive control. (B) IL-1 β , IL-6 and TNF α production by human monocytes treated or not with 0.05 μ M JQ1 and stimulated with 10⁶/ml heat-killed *C. albicans* conidia for 24 h. (C) IL-6 and TNF α production by human monocytes treated or not with 0.05 μ M JQ1 and stimulated with RPMI, 1 μ g/mL β -glucan or 10⁶/ml heat-killed *C. albicans* conidia, following a similar protocol displayed in Figures 3A and 3D. Cells were re-exposed to 10 ng/ml LPS after 6 days, and cytokines were measured 24 h later. Mean \pm SEM, n=6 (from 6 different individual donors); pooled from 2 independent experiments with 3 individual donors each. *p<0.05, Wilcoxon signed-rank test.

1 **Table 1. Primer pairs used for epigenetic study.**

<i>IL6.1</i>	FW	TCGTGCATGACTTCAGCTTT ²
<i>IL6.1</i>	RV	GCGCTAAGAAGCAGAACCAC ³
<i>IL6.2</i>	FW	AGGGAGAGCCAGAACACAGA
<i>IL6.2</i>	RV	GAGTTTCCTCTGACTCCATCG ⁴
<i>TNFA.1</i>	FW	AGAGGACCAGCTAAGAGGGA
<i>TNFA.1</i>	RV	AGCTTGTCAGGGGATGTGG ⁵
<i>TNFA.2</i>	FW	GTGCTTGTTCTCAGCCTCT
<i>TNFA.2</i>	RV	ATCACTCCAAAGTGCAGCAG ⁶
<i>TNFA.3</i>	FW	TGTCTGGCACACAGAAGACA ⁷
<i>TNFA.3</i>	RV	CCCTGAGGTGTCTGGTTTTTC
<i>TNFA.4</i>	FW	AGCCAGCTGTTCTCCTTTA ⁸
<i>TNFA.4</i>	RV	TTAGAGAGAGGTCCCTGGGG ⁹
<i>TNFA.5</i>	FW	TGATGGTAGGCAGAACTTGG
<i>TNFA.5</i>	RV	ACTAAGGCCTGTGCTGTTCC ¹⁰
<i>TNFA.6</i>	FW	CAGGCAGGTTCTCTTCTCT
<i>TNFA.6</i>	RV	GCTTTCAGTGCTCATGGTGT ¹¹

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14 **Supplementary Methods**

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16 **LPS stimulated human blood assay**

17 Blood was collected into 10 IU / mL sodium heparin and incubated for 30 min at 37°C in
 18 a concentration response (10 µM-0.51nM) of I-BET151, prior to the addition of 200 ng/mL
 19 LPS. Supernatants were removed after 22h incubation 37°C and analysed for IL-6 and
 20 TNFα using MesoScaleDiscovery (MSD) technology. ICxx calculations were determined
 21 using the formula $IC_{xx} = ((\% \text{ inhibition} / (100 - \% \text{ inhibition}))^{(1/\text{Hill Slope})}) * IC_{50}$.

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23 **Viability assays**

24 Cell viability was assessed using CytoTox 96® Non-Radioactive Cytotoxicity Assay
 25 (Promega). Released LDH in culture supernatants is measured with a 30-minute coupled
 26 enzymatic assay. The amount of color formed is proportional to the number of lysed cells.