

CORRECTION

Correction: Temporal Expression of Peripheral Blood Leukocyte Biomarkers in a *Macaca fascicularis* Infection Model of Tuberculosis; Comparison with Human Datasets and Analysis with Parametric/Non-parametric Tools for Improved Diagnostic Biomarker Identification

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The text in Fig 1 highlighting rows of expression data by week is incorrectly reversed. The authors have provided a corrected version here.



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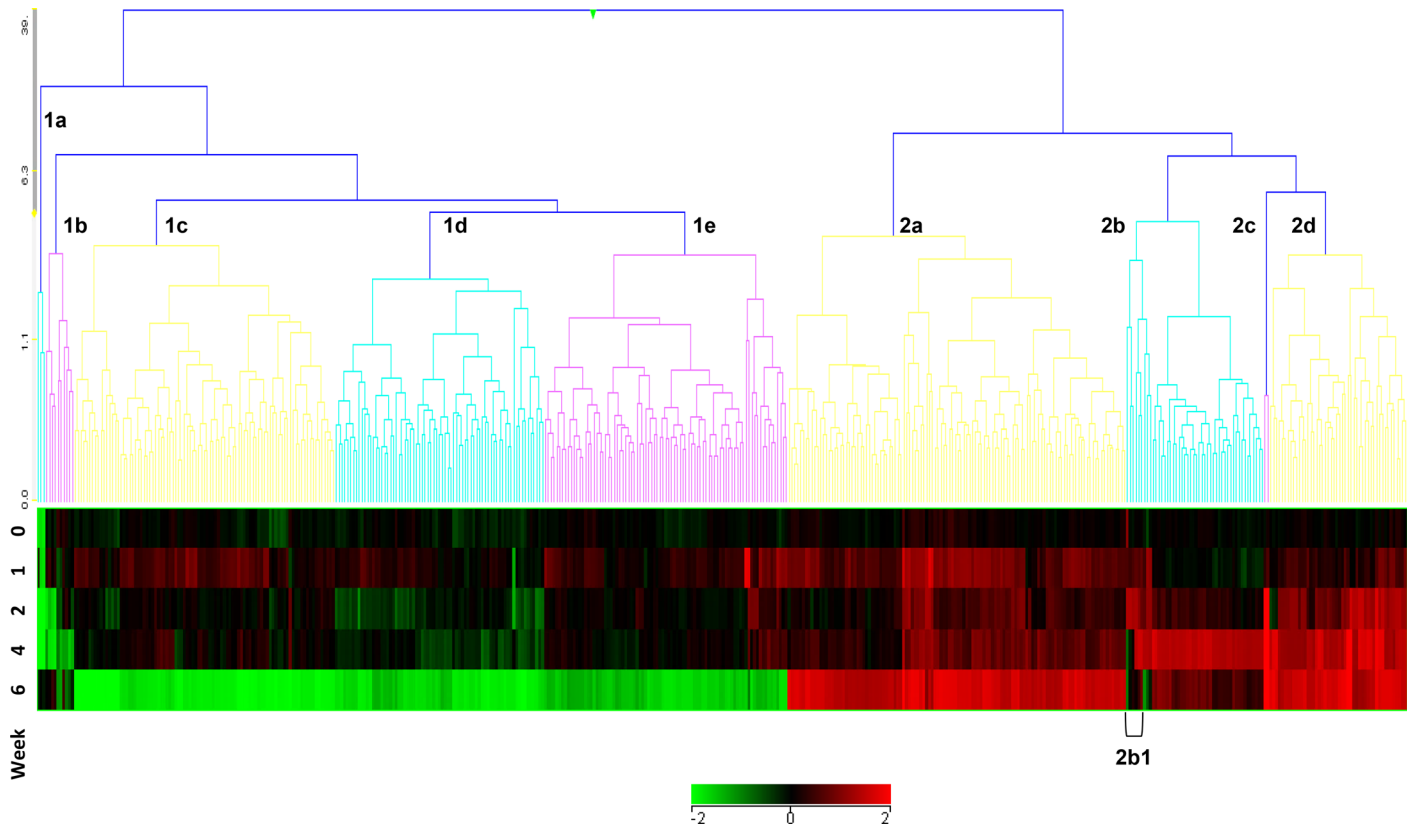


Fig 1. Cluster analysis of temporally expressed entities in peripheral blood leukocytes of Cynomolgus Macaques (all animals) pre-(week 0) or post (weeks 1–6) aerosol-challenge with *M. tuberculosis*. These exhibit patterns of up- (cluster 2) or down-regulation (cluster 1) across the six week time course of the experiment. Cluster 2b1 (highlighted) contains co-expressed entities, FOS, IL8 and KLF2.

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Reference

1. Javed S, Marsay L, Wareham A, Lewandowski KS, Williams A, Dennis MJ, et al. (2016) Temporal Expression of Peripheral Blood Leukocyte Biomarkers in a *Macaca fascicularis* Infection Model of Tuberculosis; Comparison with Human Datasets and Analysis with Parametric/Non-parametric Tools for Improved Diagnostic Biomarker Identification. PLoS ONE 11(5): e0154320. doi:[10.1371/journal.pone.0154320](https://doi.org/10.1371/journal.pone.0154320) PMID: [27228113](https://pubmed.ncbi.nlm.nih.gov/27228113/)