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Supplementary Figure 1. Percentage of IFNγ-producing CD4+ splenocytes from mock-, BCG- and MTBVAC- vaccinated C57BL/6 (a), BALB/c (b) and C3H/HeNRj (c) mice following stimulation with PPD (5 µg/ml), ESAT6 (2 µg/ml), CFP10 (2 µg/ml) and Ag85B (2 µg/ml). Data are representative from one of two independent experiments (n=5 mice/group/experiment). All data are mean ± SEM. ns, non-significant; *, p < 0.05; **, p < 0.01 ***, p < 0.001 by unpaired t-student test. (d), (e), CD4+, CD8+ and CD4-CD8- cells were analyzed from an IFNγ-positive gated population of splenocytes harvested from MTBVAC-vaccinated C3H mice. (d) Representative dot-plot is shown in the figure. (e) Data from the graph is the mean from one experiment (n=5).



Supplementary Figure 2. Culture-filtrate protein samples from BCG and MTBVAC separated by SDS-PAGE and stained with Colloidal blue. Three independent extractions from each vaccine strain are represented in the image. A differential band found in MTBVAC (pointed with an arrow) was identified as Ag85B by MALDI-TOF MS.



Supplementary Figure 3. Dissemination and replication of MTBVAC and BCG in lymphoid organs from C3H mice. MTBVAC and BCG CFUs in draining lymph nodes (inguinal and axillary) (**a**) and spleen (**b**) at 7, 14 and 28 days following subcutaneous vaccination of C3H in the right rear flank. Data are derived from n=6 mice. All data are mean ± SD. ns, non-significative; *, p < 0.05; by two-way ANOVA and Bonferroni post-test.



b



Supplementary Figure 4. Construction of MTBVAC mutant substrains. (a) General scheme of the method used to construct the MTBVAC mutants tested in this study. (b) Analysis of recombinant MTBVAC colonies after recombineering. Agarose electrophoresis shows CFP10/ESAT6 region amplified with flanking primers. Wild-type amplicon: 0.8 kbp; mutant amplicon: 1.7 kbp.

а



Supplementary Figure 5. *In vitro* CFP10 presentation assay. IFN γ production by purified CD4+ splenocytes from BCG-, MTBVAC- and MTBVAC Δ E6C10- vaccinated, or unvaccinated C3H/HeNRj mice following incubation with syngeneic BMDM previously pulsed with CFP10 increasing concentrations. Data represent the mean of three replicates ± SEM.

H37Rv glycerol stocks



Supplementary Figure 6. Lung bacterial load one day post low-dose H37Rv intranasal challenge. Results correspond to the two H37Rv glycerol stocks used in the present study. Data are derived from n=10 mice (lot 070513) and n=6 mice (lot 200814). All data are mean ± SD.



Supplementary Figure 7. Full western blots performed in this study. (a) Full blot corresponding to intracellular protein analysis from figure 1b. Blot was sequentially stained with anti-GroEL and anti-CFP10. Then membrane was incubated with a stripping solution and re-probed with anti-ESAT6. (b) Full blot corresponding to culture-filtrate protein fraction analysis from figure 1b. Blot was sequentially probed with anti-GroEL and anti-CFP10, and then re-probed with anti-Ag85A and anti-ESAT6 after stripping treatment. (c) Full blot corresponding to figure 1c. Blot was sequentially stained with anti-GroEL, anti-Ag85A and anti-CFP10, and then re-probed with anti-CFP10, and then re-probed with anti- anti-ESAT6 after stripping treatment. (d) Full blot corresponding to figure 3a. Membrane was divided in two fragments. The upper one was probed with anti-GroEL, and the lower with anti-CFP10. Then, this was re-probed with anti-ESAT6 following stripping treatment. Red rectangles correspond to the blot fragments shown in the main figures.