

Supplementary Data

Identification of MicroRNAs as Potential Blood-Based Biomarkers for Diagnosis and Therapeutic Monitoring of Active Tuberculosis

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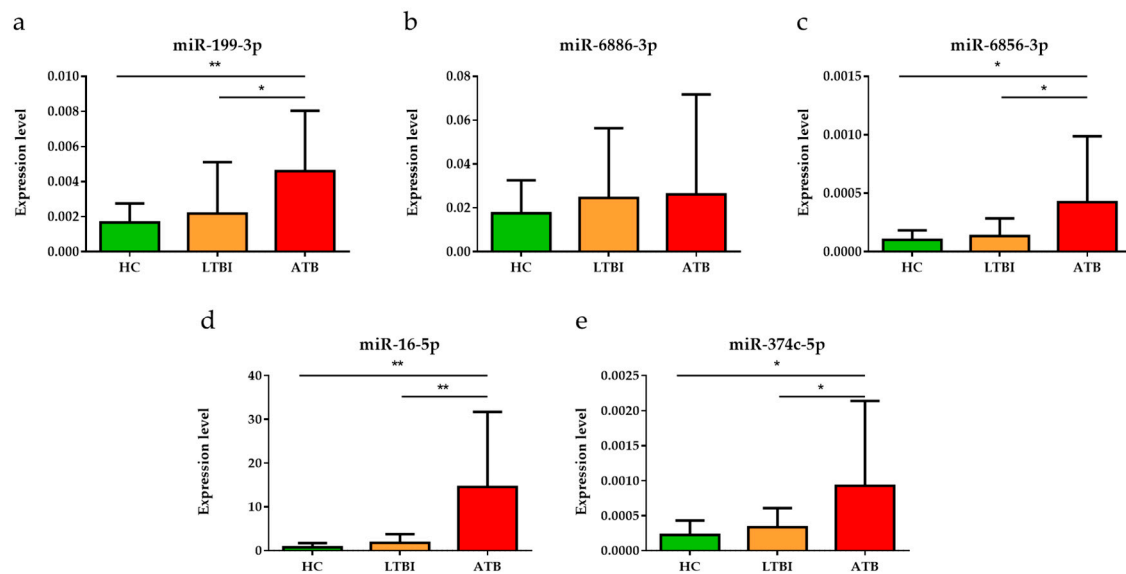


Figure S1. Validation of the expression levels of the selected miRNAs among study populations by qRT-PCR. The expression levels of (a) miR-199-3p (both miR-199a-3p and miR-199b-3p), (b) miR-6886-3p, (c) miR-6856-3p, (d) miR-16-5p, and (e) miR-374c-5p were measured in patients with active tuberculosis (ATB), individuals with latent tuberculosis infection (LTBI), and healthy controls (HCs). The data are shown as mean \pm standard deviation. * $p < 0.05$, ** $p < 0.01$.