



S3 Fig: Fluorescence imaging process overview to calculate neutrophil nuclei and neutrophil extracellular trap (chromatin) area. Scale bars: 100 μm (A-C) and 50 μm (D-I).

Isolated neutrophils were stimulated with *Mtb* and stained with Hoechst 33342 (A and boxed area D) and PL2-3 (B and boxed area E). C and the boxed area F show the overlap (blue, DNA; yellow, chromatin). The segmentation of fluorescent signals used to calculate the neutrophil nuclei area (G) and neutrophil extracellular trap area (H) with the overlap shown in (I). The scale bars represent 100 μm (A-C) and 50 μm (D-I).