**S2 Fig. MST analysis on the interaction between *E. coli* SecA and *M. tuberculosis* SecA1 and SecA2.** MST measurement of *E. coli* SecA-Cy5 (open circles), *M. tuberculosis* SecA1-Cy5 (open triangles), and SecA2-Cy5 (open square) titrated with increasing concentrations of unlabeled *E. coli* SecA, The curves were fitted using the Hill-equation and the apparent Kd values were determined with a standard error of 3 measurements. The formation of the SecA homodimer showed an apparent Kd of 31 ± 1.5 nM, whereas for the *M. tuberculosis* SecA proteins, no interaction with *E. coli* SecA was observed.