



Figure 3 - figure supplement 1. Comparison of the MlaBDEF_{ab} and MlaBDEF_{ec} complex. The map of MlaBDEFab (this study) is shown in grey, overlaid to that of MlaBDEF_{ec} (ekiert et al.) in yellow, aligned in the cytosolic domains density. The overall architecture of both complexes are similar, however MlaD appears offset by $\sim 20^\circ$ in the *A. baumannii* complex, compared to the *E.coli* complex. It remains to be determined if this corresponds to a difference between species, or if the two structures were obtained in different states of substrate binding and/or nucleotide hydrolysis.