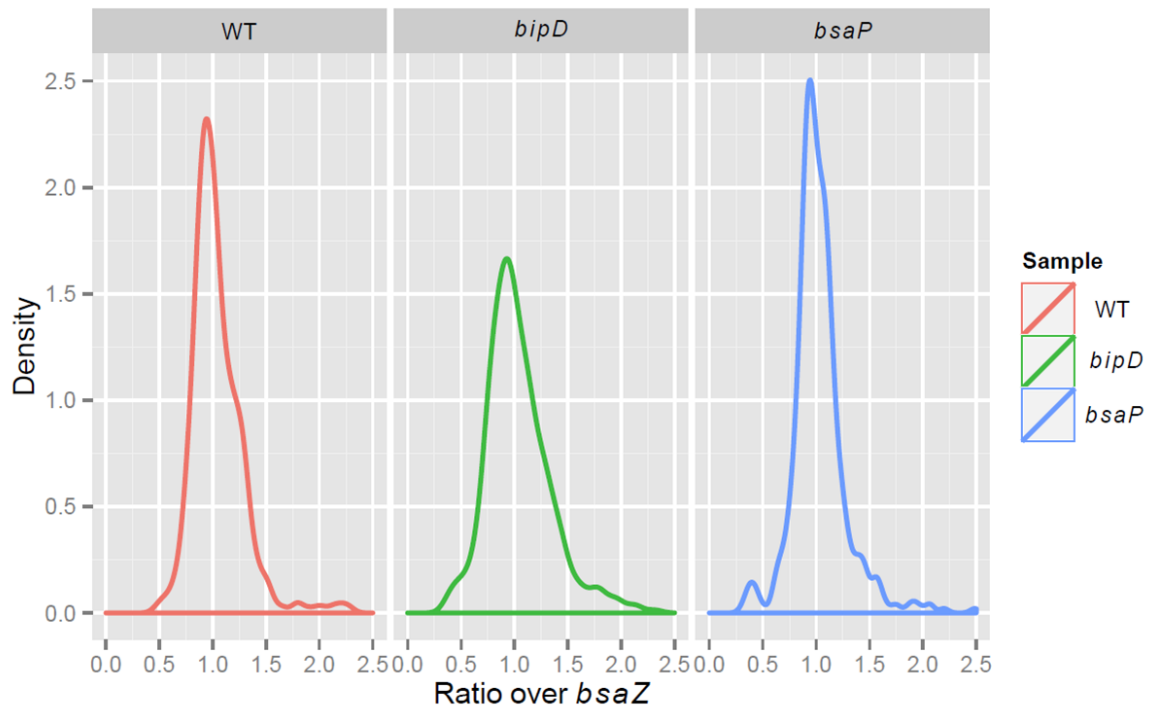**Figure S5.**

Figure S5. Distribution of iTRAQ ratios ranging between 0 and 2.5.

Density plot constructed with R software (www.r-project.org) illustrating the distribution of the median normalised iTRAQ ratios of proteins with a ratio between 0 and 2.5 in each of the three datasets where protein abundance in either 10276 WT (WT), 10276 *bipD*::pDM4 (*bipD*) or 10276 Δ*bsaP* (*bsaP*) samples were compared to the 10276 *bsaZ*::pDM4 (*bsaZ*) sample. Ratios close to 1 represent proteins that were present in similar quantities in the supernatants of a given strain when compared to the *bsa*-null mutant.