Table S1. Plasmids

Plasmid	Description	Source
pHSS6-GCU	Vector containing gonococcal uptake sequence (Kan <sup>r</sup> )	(1)
pUNCH412	FA19 tbpA in pET-11	(2)
pUNCH755	Vector containing truncated and nonfunctional <i>tbpB</i> gene, full length <i>tbpA</i> gene, and <i>tbpA</i> downstream region with an mTn3Cm insertion	(3)
pVCU150	pHSS6-GCU containing <i>tbpA</i> gene with K351A mutation, novel silent BamHI site at <i>tbpA</i> bp 1488-1493, and no stop codon	This study
pVCU151	pHSS6-GCU containing <i>tbpA</i> gene with D355A mutation, novel silent BamHI site at <i>tbpA</i> bp 1488-1493, and no stop codon	This study
pVCU152	pHSS6-GCU containing <i>tbpA</i> gene with D355K mutation, novel silent BamHI site at <i>tbpA</i> bp 1488-1493, and no stop codon	This study
pVCU153	pHSS6-GCU containing <i>tbpA</i> gene with N357A mutation, novel silent BamHI site at <i>tbpA</i> bp 1488-1493, and no stop codon	This study
pVCU154	pHSS6-GCU containing <i>tbpA</i> gene with Q358A mutation, novel silent BamHI site at <i>tbpA</i> bp 1488-1493, and no stop codon	This study
pVCU155	pHSS6-GCU containing <i>tbpA</i> gene with K359A mutation, novel silent BamHI site at <i>tbpA</i> bp 1488-1493, and no stop codon	This study
pVCU156	pHSS6-GCU containing <i>tbpA</i> gene with K359E mutation, novel silent BamHI site at <i>tbpA</i> bp 1488-1493, and no stop codon	This study
pVCU157	pHSS6-GCU containing <i>tbpA</i> gene with K359R mutation, novel silent BamHI site at <i>tbpA</i> bp 1488-1493, and no stop codon	This study
pVCU158	pHSS6-GCU containing <i>tbpA</i> gene with Q360A mutation, novel silent BamHI site at <i>tbpA</i> bp 1488-1493, and no stop codon	This study
pVCU159	pHSS6-GCU containing <i>tbpA</i> gene with Q360E mutation, novel silent BamHI site at <i>tbpA</i> bp 1488-1493, and no stop codon	This study
pVCU160	pHSS6-GCU containing <i>tbpA</i> gene with Q360K mutation, novel silent BamHI site at <i>tbpA</i> bp 1488-1493, and no stop codon	This study

pVCU161	pUNCH755 containing tbpA gene from pVCU150	This study
pVCU162	pUNCH755 containing tbpA gene from pVCU151	This study
pVCU163	pUNCH755 containing tbpA gene from pVCU152	This study
pVCU164	pUNCH755 containing tbpA gene from pVCU153	This study
pVCU165	pUNCH755 containing tbpA gene from pVCU154	This study
pVCU166	pUNCH755 containing tbpA gene from pVCU155	This study
pVCU167	pUNCH755 containing tbpA gene from pVCU156	This study
pVCU168	pUNCH755 containing tbpA gene from pVCU157	This study
pVCU169	pUNCH755 containing tbpA gene from pVCU158	This study
pVCU170	pUNCH755 containing tbpA gene from pVCU159	This study
pVCU171	pUNCH755 containing tbpA gene from pVCU160	This study
pVCU172	pUNCH755 containing <i>tbpA</i> gene with deletion from bp	This study
pVCU757	1453-1488 (loop 3 helix region) FA 1090 <i>tbpA</i> in pET22b	This study

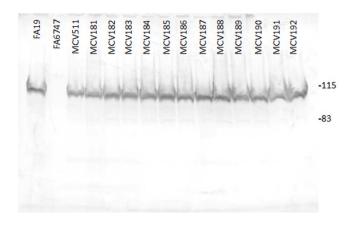


Fig S1. TbpA expression determination by western blot. WT and mutant gonococci were iron stressed in liquid CDM for 4 hours, standardized to cell density, pelleted, and lysed. Lysates were subjected to a bicinchoninic acid assay (Thermo) to assess protein levels, and then evenly loaded for SDS-PAGE and subsequent transfer to nitrocellulose. Equivalent protein loading was confirmed by ponceau staining. The western blot was probed with polyclonal TbpA antibody, and developed with the NBT/BCIP development system (Sigma). TbpA bands are present at 100 kDa.

## 11 References

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