



Fig. S5. Genetic organization of the QS regulatory genes in *B. thailandensis* E264. *btaI1* and *btaR1* are not located next to each other and are divergently transcribed in *B. thailandensis* E264. The promoter region of *btaI1* contains a putative *lux-box* sequence centered 73.5 bp upstream of the *btaI1* translation start site (CCCTGTAAGGGTTAACAGTT). *btaI2* and *btaR2* are also not located next to each other and are transcribed in the same direction on the genome of *B. thailandensis* E264. The promoter region of *btaI2* contains a putative *lux-box* sequence centered 65.0 bp upstream of the *btaI2* translation start site (ACCTGTAGAAATCGTAGT). *btaI3* and *btaR3* are also transcribed in the same direction and are located next to each other in *B. thailandensis* E264.