

Supplementary Table 1. Bacterial strains and plasmids used in this study

Strains or plasmids	Characteristic(s)	Source or reference
<i>Escherichia coli</i>		
EC100D	For identifying Tn5-insertional site	Epicentre
DH5 α	For cloning	Promega
<i>Acidovorax citrulli</i>		
KACC17005	Wild-type, Rif ^r	Park et al. (2017)
<i>Ac</i> (EV)	Wild-type carrying the pBBR1-MCS5, Rif ^r , Gm ^r	This study
<i>Ac</i> Δ yppAc	Knockout mutant, Tn5 transposon inserted in <i>yppAc</i> gene, Rif ^r , Kan ^r	This study
<i>Ac</i> Δ yppAc(EV)	Knockout mutant carrying the pBBR1-MCS5, Rif ^r , Kan ^r , Gm ^r	This study
<i>Ac</i> Δ yppAc(YppAc)	Complemented strain, <i>Ac</i> Δ yppAc carrying the MCS5-YppAc, Rif ^r , Kan ^r , Gm ^r	This study
Plasmid		
pGem-T easy	T-A cloning vector, Amp ^r	Promega
pGem-YppAc	pGem-T easy vector ligating 1,628 bp of <i>yppAc</i> gene, Amp ^r	This study
pBBR1-MCS5	Broad-host-range vector, LacZ promoter, Gm ^r	Kovach et al. (1995)
pMCS5-YppAc	pBBR1-MCS5 carrying the <i>yppAc</i> gene from pGem-YppAc plasmid, Gm ^r	This study

Rif^r, Kan^r, Gm^r and Amp^r represent resistance to rifampicin, kanamycin, gentamycin, and ampicillin, respectively.

Supplementary Table 2. Protein and peptide spectral matches (PSM) between *Ac* and *Ac* Δ yppAc

Strain	1st		2nd		3rd		Shared proteins in 3 biological replicates
	Protein	PSM	Protein	PSM	Protein	PSM	
<i>Ac</i>	1,091	50,387	1,069	50,459	1,097	50,383	1,013
<i>Ac</i> Δ yppAc	1,156	55,911	1,147	55,863	1,127	55,981	1,047

References

- Park, H.-J., Seong, H. J., Sul, W. J., Oh, C.-S. and Han, S.-W. 2017. Complete genome sequence of *Acidovorax citrulli* strain KACC17005, a causal agent for bacterial fruit blotch on watermelon. *Korean J. Microbiol.* 53:340-341.
- Kovach, M. E., Elzer, P. H., Hill, D. S., Robertson, G. T., Farris, M. A., Roop, R. M. 2nd and Peterson, K. M. 1995. Four new derivatives of the broad-host-range cloning vector pBBR1MCS, carrying different antibiotic-resistance cassettes. *Gene* 166:175-176.