

**Supplementary Table S1. Bacterial strains and plasmids used in this study**

Name	Description	Reference or source
<b>Strains</b>		
<i>S. suis</i> SC19	<i>S. suis</i> serotype 2	[1]
<i>S. suis</i> SS2041	Multiple-drug resistant strain	[2]
<i>E. coli</i> ATCC25922	<i>E. coli</i> standard strain	Laboratory stock
<i>S. aureus</i> ATCC29213	<i>S. aureus</i> standard strain	Laboratory stock
<i>S. aureus</i> 1213M4A	Multiple-drug resistant strain	Laboratory stock
<i>E. rhusiopathiae</i> 13013	Multiple-drug resistant strain	[3]
<i>S. mutans</i> ATCC25175	<i>S. mutans</i> standard strain	Laboratory stock
<i>B. subtilis</i> WB800N	Derivative of <i>B. subtilis</i> 168	[4]
<b>Plasmids</b>		
pET28a-dacA <sub>cyto</sub>	Plasmid for the expression of ssDacA, Kan <sup>r</sup>	This study

**Supplementary Table S2. Primers used in this study**

Strains	Sequence (5'-3')
ssDacA <sub>cyto</sub> -F	TAAGAAGGAGATATACCATGGCGCTATGCTTGAAAAACTC
ssDacA <sub>cyto</sub> -R	GTGGTGGTGGTGCTCGAGTTCATGTTGCCACCTCC

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- Ding, Y.; Zhu, D.; Zhang, J.; Yang, L.; Wang, X.; Chen, H.; Tan, C. Virulence determinants, antimicrobial susceptibility, and molecular profiles of *Erysipelothrix rhusiopathiae* strains isolated from China. *Emerg Microbes Infect* **2015**, *4*, e69, doi:10.1038/emi.2015.69.
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