

## Supplementary File 1

Table S1. BLASTn homology analysis of 16S rRNA gene fragment of the SRS89 strain.

	Accession No.	Query Cover	E value	Per. Ident.
<i>Pantoea agglomerans</i> strain DSM 3493	NR_041978	100%	0.0	99.54%
<i>Pantoea agglomerans</i> strain NCTC9381	NR_114735	100%	0.0	99.23%
<i>Pantoea agglomerans</i> strain NBRC102470	NR_114111	100%	0.0	99.31%
<i>Pantoea vagans</i> strain LMG24199	NR_116115	100%	0.0	99.08%

Figure S1. Alignment of 16S rRNA gene fragment of the SRS89 strain to *P. agglomerans* (NR\_041978, NR\_114735, NR\_114111) and *P. vagans* (NR\_116115) strains.