

Table 1 | Oligonucleotide primers used for qRT-PCR in this study.

Name	Sequence (5'-3')
Actin-F	TGGGATGATATGGAGAAGATATGG
Actin-R	GGCTTCAGT TAGGAGGACAGGA
lea-F	GTCCTAACATCTCCTCCAA
lea-R	CAGGGTAAT CGCATCAG
tdi65-F	CGTCCTCTGTTGTCCC
tdi65-R	AGTGAAGGCAATGAAGC
ltpg2-F	TACTGGACC GTTGAGCA
ltpg2-R	GGTGGTGTGGTGGTGTAA
nced1-F	AGGCAACAGTGAAACTTCCATCAAG
nced1-R	TCCATTAAAGAGGATATTACCGGGAC

Table S2 | Drought tolerance efficacy of wild isolates under greenhouse conditions.

Strains	Possible Species	Drought Tolerance Efficacy*
-	-	+
Blt-11	<i>Acinetobacter</i> sp.	++
HS1-1	<i>Acinetobacter</i> sp.	+++
DS19	<i>Arthrobacter globiformis</i>	+
HS23-4	<i>Arthrobacter globiformis</i>	++
54	<i>Bacillus amyloliquefaciens</i>	++++
2YN11	<i>Bacillus amyloliquefaciens</i>	++
5YN11	<i>Bacillus amyloliquefaciens</i>	++
HSSN09	<i>Bacillus amyloliquefaciens</i>	++
W13-3	<i>Bacillus cereus</i>	++
Blgb-12	<i>Bacillus cereus</i>	++
DS38	<i>Bacillus cereus</i>	++
HYN1	<i>Bacillus cereus</i>	+
HS10	<i>Bacillus lincheniformis</i>	+
31	<i>Bacillus megaterium</i>	++
3BS3	<i>Bacillus megaterium</i>	++
HS15	<i>Bacillus megaterium</i>	++
DS58	<i>Bacillus pocheonensis</i>	+
Ljb-10	<i>Bacillus pumilus</i>	++
DS22	<i>Bacillus pumilus</i>	+
YF31	<i>Bacillus</i> sp.	++
Lgn-13	<i>Bacillus</i> sp.	+++
Lgn-3	<i>Bacillus</i> sp.	++
Lgn-4	<i>Bacillus</i> sp.	+
DSSN12	<i>Bacillus subtilis</i>	+
Blgb-17	<i>Bacillus subtilis</i>	++
1BQN14	<i>Enterobacter asburiae</i>	++
YT8	<i>Enterobacter ludwigii</i>	++
3BJN7	<i>Enterobacter</i> sp.	++

Lgn-6

Enterobacter sp.

+ +

*Drought tolerance efficacy: Classification of isolates according to survival rate of tomato plant after 15 days of drought treatment and then 5 days of rewatering, where (+) 0-20%; (++) 21%-40%; (+++) 41%-60%; (++++) \geq 61%.

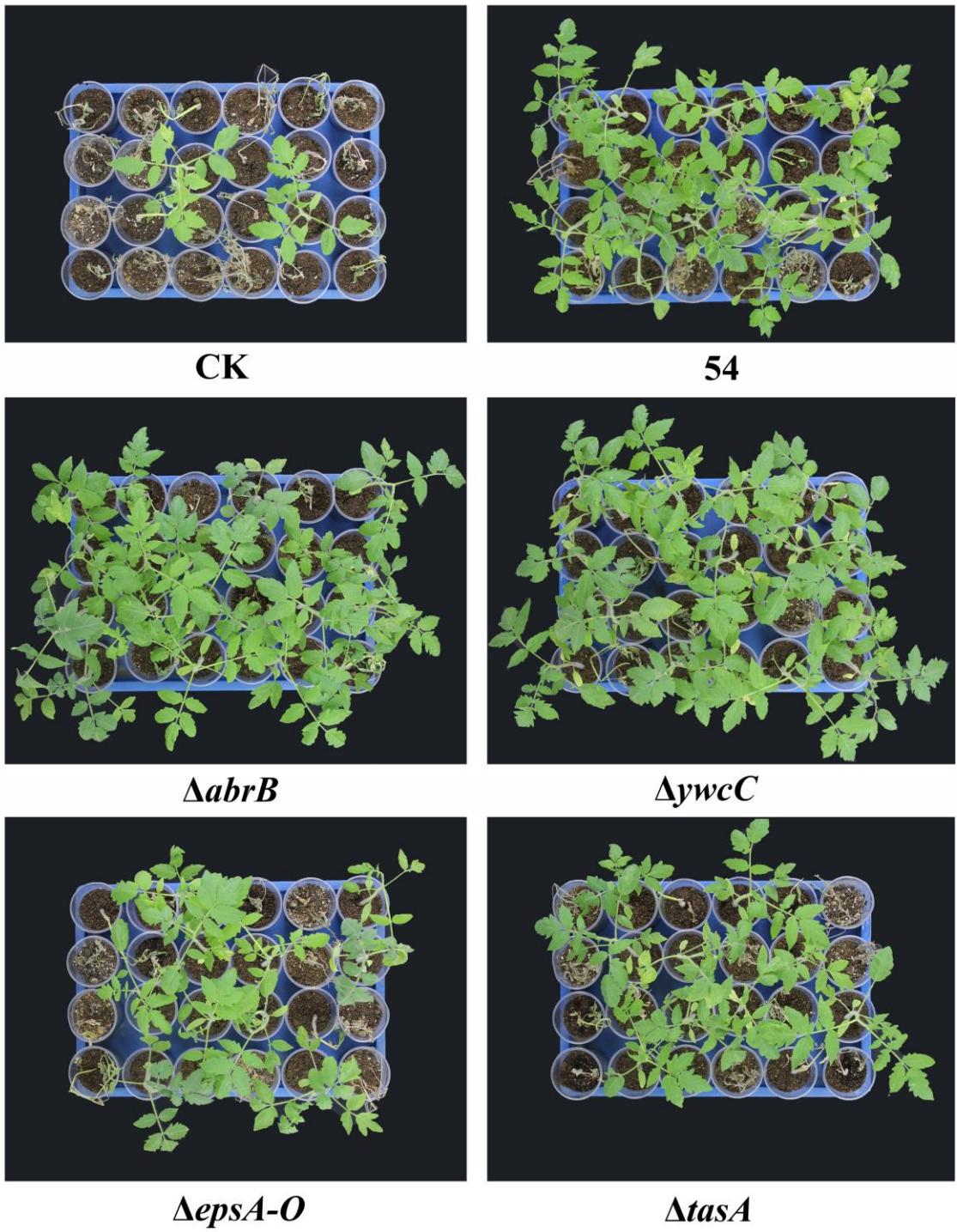


Figure S1. Biofilm formation is positively involved in *B. amyloliquefaciens* 54-mediated drought tolerance of tomato plants. Photographs of each treatment were taken after 15 days of drought treatment and then 5 days of rewatering.