

Table S1. Preliminary screening of bacteria against *A. flavus*

Strain	Identification	Antagonistic ability
2-1	<i>Lactobacillus plantarum</i>	-
3-2	<i>Lactobacillus alimentarius</i>	-
5-1	<i>Lactobacillus plantarum</i>	-
5-4	<i>Lactobacillus hilgardii</i>	-
7-2	<i>Lactobacillus pentosus</i>	-
7-4	<i>Lactobacillus alimentarius</i>	-
8-1	<i>Lactobacillus plantarum</i>	-
9-1	<i>Lactobacillus namurensis</i>	-
9-3	<i>Lactobacillus paracasei</i>	-
10-1	<i>Lactobacillus alimentarius</i>	-
13-4	<i>Lactobacillus alimentarius</i>	-
16-1	<i>Lactobacillus plantarum</i>	-
16-4	<i>Lactobacillus pentosus</i>	-
17-1	<i>Weissella hellenica</i>	-
19-1	<i>Lactobacillus pentosus</i>	-
19-2	<i>Lactobacillus alimentarius</i>	-
21-1	<i>Lactobacillus plantarum</i>	-
21-3	<i>Lactobacillus pentosus</i>	-
21-4	<i>Lactobacillus harbinensis</i>	-
21-5	<i>Pediococcus ethanolidurans</i>	-

24-1	<i>Lactobacillus plantarum</i>	-
25-1	<i>Lactobacillus alimentarius</i>	-
28-1	<i>Lactobacillus alimentarius</i>	-
29-1	<i>Lactobacillus alimentarius</i>	-
33-3	<i>Lactobacillus plantarum</i>	-
CICC6239	<i>Lactobacillus brevis</i>	-
CICC24450	<i>Lactobacillus brevis</i>	-
CGMCC1.2427	<i>Lactobacillus fructivorans</i>	-
4		-
E11		+
V1J		+
V2		+
J0		-
J1		-
Y1①	<i>Bacillus subtilis</i>	-
D2④		-
V1J2		+
V2②		-
VIJ1		+
2×1		+
2399		-

Fy1q	-
67	-
10081	-
y×10②	-
y×10②'	-
GQy8	-
1#	+
H6	+
B1⑥	-
Y1③	-
VIJ3	-
C1②	-
10063	+
H1	-
9932	+
B1-3	-
Z4-5	-
BJ3-2	-
KCKK	-
KCFZ	-
KCW	-

KCB

-

(Note: "+" denoted the strain had obvious bacteriostasis circle, "-" denoted the strain didn't have obvious bacteriostasis circle)

Table S2. The abilities of *B. subtilis* E11, *B. subtilis* V1J1, and *B. subtilis* 9932 to degrade AFB₁ at different time points.

Strains	Time (h)				
	24	48	72	96	120
<i>B. subtilis</i> E11	81.34 ±3.25%	88.86 ±1.92%	92.85 ±0.48%	95.38 ±0.69%	96.28 ±0.62%
<i>B. subtilis</i> 9932	82.72 ±4.95%	86.89 ±0.57%	88.68 ±1.62%	89.31 ±0.41%	95.17 ±0.41%
<i>B. subtilis</i> V1J1	84.09 ±4.36%	89.03 ±1.00%	90.81 ±2.06%	94.73 ±1.00%	95.58 ±0.71%

The values were reported as average ± SD and were the mean of three independent analyses.

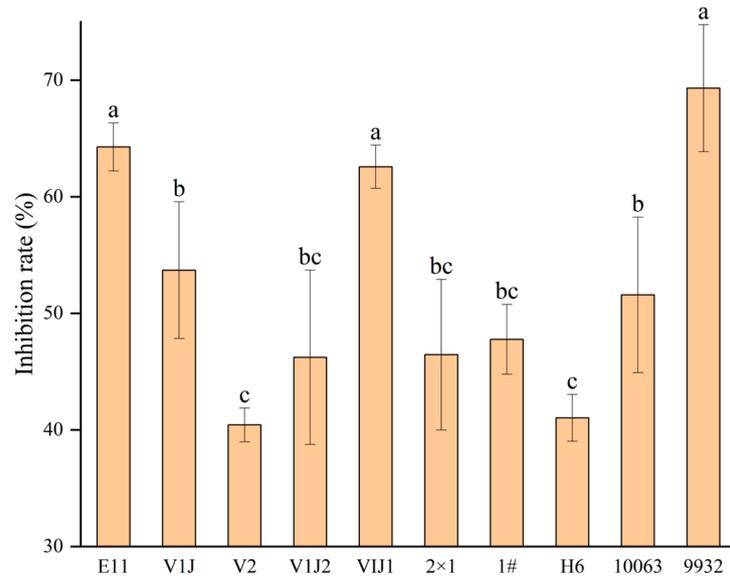


Figure S1. Rescreening of antagonistic bacteria against *A. flavus*. Values were expressed as means \pm SD;

a, b, c indicated significant differences of inhibition rate of different strains to *A. flavus* ($p < 0.05$).

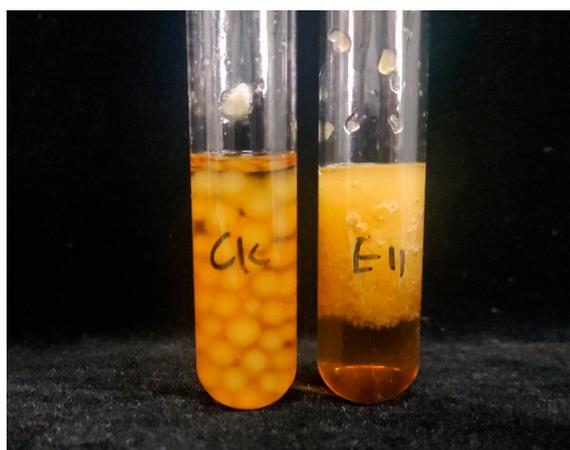


Figure S2. The effect of the fermentation supernatant of *B. subtilis* E11 on *A. flavus* mycelia. The tube on the left was the morphology of *A. flavus* mycelia after 72h culture without the fermentation supernatant of *B. subtilis* E11; The tube on the right showed the mycelia morphology of *A. flavus* co-cultured with *B. subtilis* E11 fermentation supernatant after 72h.