

**Table S1** Major topological properties of the co-occurrence network of microbiome in rhizosphere of cucumber.

Groups	Treatments	Nodes	Nodes connected with <i>Strep.</i>	Number of Links	ND (Network diameter)	MD (modularity)	CC (Clustering coefficient)	GD (Graph density)	AD (Average degree)	APL (Average path length)
Bacteria	Control	296	20	2342	4	0.484	0.318	0.054	15.824	2.641
	With <i>Strep.</i>	313	41	3536	4	0.441	0.350	0.066	15.525	2.525
Fungi	Control	62	1	405	5	0.218	0.526	0.214	13.065	2.26
	With <i>Strep.</i>	69	4	399	6	0.285	0.462	0.170	11.565	2.498

**Table S2** Biomass of cucumber after the inoculation with *Streptomyces* Act12<sup>1</sup>

Inoculation	Concentration g kg <sup>-1</sup> soil	Biomass in 20 <sup>th</sup> d g plant <sup>-1</sup>		Biomass in 45 <sup>th</sup> d g plant <sup>-1</sup>	
		Mean	ΔCtrl%	Mean	ΔCtrl%
CK		13.91±0.94ab	-	19.28±0.73ab	-
Act12 <sup>2</sup>	1.0	16.87±2.62a	21.2	21.79±3.58a	13.0
	1.5	16.53±3.04a	18.8	22.47±1.67a	16.5
	2.0	15.24±1.93a	9.6	19.69±1.65ab	2.1
HK <sup>3</sup>		10.13±1.31c	-	19.90±1.19bc	-
Act12+HK	1.0	15.38±2.47b	51.9	19.77±0.67c	-0.6
	1.5	18.12±1.37ab	78.9	25.60±4.32a	28.6
	2.0	18.95±1.52a	87.0	23.71±0.66ab	19.2

<sup>1</sup>Three duplicates were set up in a pot experiment planting with cucumber, control was treated without *Streptomyces*. Values are means ± standard deviation ( $n = 3$ ). Different letters indicate statistically significant difference between the control and *Streptomyces* treatments, at the level of  $P < 0.05$ .

<sup>2</sup>The *Streptomyces* bio-agent was  $2.59 \times 10^{11}$  CFU·g<sup>-1</sup>.

<sup>3</sup>HK represents inoculated with pathogens *F. oxysporum* with concentration of  $10^2$  CFU·g<sup>-1</sup> soil.

**Table S3** Root weight and length of cucumber after the inoculation with *Streptomyces* Act12<sup>1</sup>

Inoculation	Concentration g kg <sup>-1</sup> soil	Root weight in 20 <sup>th</sup> d g plant <sup>-1</sup>		Root weight in 45 <sup>th</sup> d g plant <sup>-1</sup>		Root length in 45 <sup>th</sup> d cm	
		Mean	ΔCtrl%	Mean	Δ Ctrl	Mean	Δ Ctrl
CK		0.55±0.15b	-	1.70±0.17b	-	7.94±0.55b	-
Act12 <sup>2</sup>	1.0	1.42±0.22a	157.7	2.09±0.33a	22.9	12.45±0.56a	56.8
	1.5	1.79±0.35a	225.0	1.95±0.09ab	14.4	12.49±0.99a	57.3
	2.0	0.91±0.07b	66.1	1.78±0.14ab	4.4	13.02±0.77a	63.9
HK <sup>3</sup>		0.79±0.28c	-	1.59±0.17a	-	6.90±0.97c	-
Act12+HK	1.0	1.34±0.37b	69.6	1.85±0.02a	16.6	12.04±1.93bc	74.6
	1.5	1.70±0.3ab	114.6	2.08±0.3a	30.6	18.12±0.29a	162.
	2.0	1.96±0.36a	147.5	1.73±0.43a	9.0	14.49±2.32b	110.

<sup>1</sup>Three duplicates were set up in a pot experiment planting with cucumber, control was treated without *Streptomyces*. Values are means ± standard deviation ( $n = 3$ ). Different letters indicate statistically significant difference between the control and *Streptomyces* treatments, at the level of  $P < 0.05$ .

<sup>2</sup>The *Streptomyces* bio-agent was  $2.59 \times 10^{11}$  CFU·g<sup>-1</sup>.

<sup>3</sup>HK represents inoculated with pathogens *F. oxysporum* with concentration of  $10^2$  CFU·g<sup>-1</sup> soil.

**Table S4** Root activity after the inoculation with *Streptomyces* Act12<sup>1</sup>

Inoculation	Concentration g kg <sup>-1</sup> soil	Root activity in 20 <sup>th</sup> d g·g <sup>-1</sup> ·h <sup>-1</sup>		Root activity in 45 <sup>th</sup> d g·g <sup>-1</sup> ·h <sup>-1</sup>	
		Mean	ΔCtrl%	Mean	ΔCtrl%
CK		0.12±0.03d	-	0.07±0.03a	-
Act12 <sup>2</sup>	1.0	1.01±0.13a	741.6	0.08±0.03a	17.1
	1.5	0.70±0.02b	482.0	0.07±0.03a	-5.9
	2.0	0.26±0.01c	114.3	0.05±0.02a	-23.1
HK <sup>3</sup>		0.15±0.06d	-	0.07±0.02b	-
Act12+HK	1.0	0.69±0.00ab	362.0	0.18±0.01a	161.0
	1.5	0.80±0.04a	433.4	0.09±0.03b	32.3
	2.0	0.50±0.00bc	232.8	0.07±0.01b	0.5

<sup>1</sup>Three duplicates were set up in a pot experiment planting with cucumber, control was treated without *Streptomyces*. Values are means ± standard deviation ( $n = 3$ ). Different letters indicate statistically significant difference between the control and *Streptomyces* treatments, at the level of  $P < 0.05$ .

<sup>2</sup>The *Streptomyces* bio-agent was  $2.59 \times 10^{11}$  CFU·g<sup>-1</sup>.

<sup>3</sup>HK represents inoculated with pathogens *F. oxysporum* with concentration of  $10^2$  CFU·g<sup>-1</sup> soil.

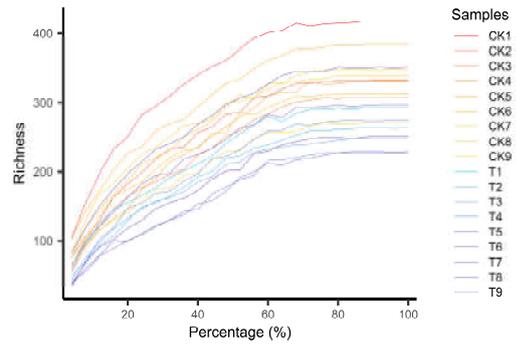
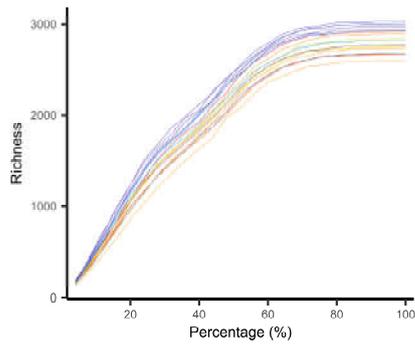
**Table S5** Leaf area and SPAD after the inoculation with *Streptomyces* Act12<sup>1</sup> in the 45<sup>th</sup> day.

Inoculation	Concentration g kg <sup>-1</sup> soil	Leaf area cm <sup>2</sup>		SPAD	
		Mean	ΔCtrl%	Mean	ΔCtrl%
	CK	53.08±9.15b	-	39.74±1.96a	-
Act12 <sup>2</sup>	1.0	77.33±7.97a	45.7	43.38±5.02a	9.2
	1.5	77.13±12.34a	45.3	41.58±4.11a	4.6
	2.0	62.18±4.10b	17.1	41.01±2.54a	3.2
	HK <sup>3</sup>	54.97±4.15c	-	38.37±3.85a	-
Act12+HK	1.0	64.54±5.29bc	17.4	40.34±2.51a	5.1
	1.5	82.45±16.29a	50.0	41.34±2.28a	7.7
	2.0	73.67±3.09ab	34.0	40.96±2.17a	6.8

<sup>1</sup> Three duplicates were set up in a pot experiment planting with cucumber, control was treated without *Streptomyces*. Values are means ± standard deviation ( $n = 3$ ). Different letters indicate statistically significant difference between the control and *Streptomyces* treatments, at the level of  $P < 0.05$ .

<sup>2</sup> The *Streptomyces* bio-agent was  $2.59 \times 10^{11}$  CFU·g<sup>-1</sup>.

<sup>3</sup> HK represents inoculated with pathogens *F. oxysporum* with concentration of  $10^2$  CFU·g<sup>-1</sup> soil.



(a) (b)  
**Figure S1.** The rarefaction curve by each samples of the rhizosphere microbiome associated with cucumber plants in *Streptomyces*-inoculated soil (T) and uninoculated control soil (CK) *Streptomyces*. (a) The rarefaction curves of bacteria; (b) the rarefaction curves of fungi.