Effects of continuous cropping of *Codonopsis tangshen* on soil bacterial community as determined by pyrosequencing

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Supplementary Table 1 The relative abundances (%) of the bacteria at the Phylum level

	FLZ1	**LZ1**	**FLZ2**	**LZ2**	**FLZ3**	**LZ3**
Proteobacteria	41.38	39.56	41.42	41.50	37.85	34.55
Acidobacteria	15.75	21.14	16.52	23.80	19.33	25.49
Gemmatimonadetes	7.89	8.66	6.48	6.84	12.50	13.08
Chloroflexi	10.57	5.27	10.42	5.23	10.11	7.39
Actinobacteria	8.00	7.80	8.31	7.23	3.77	7.83
Bacteroidetes	4.67	4.62	5.02	3.49	4.31	1.70
Nitrospirae	4.27	1.56	4.78	1.10	5.21	1.16
Verrucomicrobia	1.51	2.89	0.96	0.98	0.94	0.63
Planctomycetes	2.03	0.73	1.57	0.87	1.85	0.67
TM7	0.28	0.75	0.48	2.43	0.20	2.24
WS3	1.70	0.42	1.18	0.07	1.94	0.08
AD3	0.01	1.05	0.01	2.04	0.01	2.00
Others	1.93	5.56	2.85	4.41	1.99	3.20

LZ1, FLZ, LZ2, FLZ2, LZ3 and FLZ3 represent the soil samples collected from continuous cropping (LZ) and main crops (FLZ) at the May 4th (LZ1 and FLZ1), July 14th (LZ2 and FLZ2) and September 19th (LZ3 and FLZ3), 2016.

Supplementary Table 2 The relative abundances (%) of the bacteria at the class level

	FLZ1	LZ1	FLZ2	LZ2	FLZ3	LZ3
Alphaproteobacteria	16.05	19.62	21.78	29.29	16.93	16.05
Betaproteobacteria	10.38	7.91	7.24	2.90	9.67	10.38
Gammaproteobacteria	6.05	8.09	4.68	5.67	4.33	6.05
Deltaproteobacteria	8.79	3.94	7.57	3.63	6.85	8.79
Gemm-1	4.04	3.77	3.75	4.47	6.29	4.04
Acidobacteria-6	9.56	1.46	9.12	0.89	8.07	9.56
Acidobacteriia	0.25	7.79	0.39	9.00	1.33	0.25
Gemmatimonadetes	3.40	4.87	2.52	2.35	5.87	3.40
Actinobacteria	4.28	4.27	5.52	4.73	1.25	4.28
Solibacteres	0.89	5.19	1.57	4.60	2.68	0.89
Anaerolineae	6.07	1.25	5.33	0.41	5.46	6.07
DA052	0.03	4.36	0.11	7.66	0.20	0.03
Nitrospira	4.27	1.56	4.78	1.10	5.19	4.27
Acidimicrobiia	1.70	1.27	1.70	1.19	1.58	1.70
Saprospirae	1.97	1.46	2.26	1.24	2.46	1.97
Thermoleophilia	1.78	2.24	0.96	1.30	0.88	1.78
Ellin6529	1.86	0.63	2.26	0.57	2.63	1.86
Ktedonobacteria	0.04	2.11	0.02	2.91	0.09	0.04
Chloracidobacteria	1.66	0.46	1.74	0.37	3.12	1.66
Sphingobacteriia	0.54	2.24	0.70	1.51	0.41	0.54
Others	16.38	15.53	16.00	14.21	14.69	16.38

LZ1, FLZ, LZ2, FLZ2, LZ3 and FLZ3 represent the soil samples collected from continuous cropping (LZ) and main crops (FLZ) at the May 4th (LZ1 and FLZ1), July 14th (LZ2 and FLZ2) and September 19th (LZ3 and FLZ3), 2016.

Supplementary Table 3 The relative abundances (%) of the bacteria at the genus level

	FLZ1	**LZ1**	**FLZ2**	**LZ2**	**FLZ3**	**LZ3**
Rhodoplanes	3.16	2.64	4.52	2.49	2.04	1.56
Candidatus Koribacter	0.15	3.13	0.17	4.48	0.57	3.16
Candidatus Solibacter	0.42	2.69	0.87	2.13	1.46	3.82
Rhodanobacter	0.64	3.42	0.25	1.84	0.54	4.03
Nitrospira	2.19	1.11	2.71	1.07	2.41	1.02
Sphingobium	1.16	1.19	1.36	0.71	2.52	1.98
Rhodococcus	0.02	0.09	1.54	1.11	0.23	0.02
Unclassified genus	80.41	74.86	76.20	79.09	83.80	81.17
Others	11.85	10.87	12.38	7.09	6.43	3.24

LZ1, FLZ, LZ2, FLZ2, LZ3 and FLZ3 represent the soil samples collected from continuous cropping (LZ) and main crops (FLZ) at the May 4th (LZ1 and FLZ1), July 14th (LZ2 and FLZ2) and September 19th (LZ3 and FLZ3), 2016.

Supplementary Figure 1 Sampling plan design drawing