

1 ***Supplementary Information for Scientific Reports***

2 **Soil microbial quantification approaches coupling with relative abundances reflecting the changes of**
3 **taxa in different locations**

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24 **Supplementary Table S1. Summary of soil characteristics of the two sampling sites in this study.**

Location	Elevation (m)	Latitude	Soil pH ^a	Soil moisture (%)	Total carbon content (TOC, mg/g)
Beijing	810-1349	40°31'10"N	6.17±0.48	21.49±7.35	46.69±27.96
		115°49'32"E			
Tibet	5088-5116	29°36'21"N	6.53±0.22	1.50±0.35	26.28±5.27
		85°45'09"E			

25 ^a Average individual soil variables (pH, moisture, and TOC) over 10 samples.

26 **Supplementary Table S2. Dissimilarity tests of microbial communities from two sampling sites^a.**

Dissimilarity Beijing vs. Tibet	MRPP ^b		ANOSIM ^c		ADONIS ^d	
	δ	<i>P</i>	<i>R</i>	<i>P</i>	<i>F</i>	<i>P</i>
Total	0.715	0.002	0.288	0.002	0.14	0.001
Proteobacteria	0.708	0.003	0.269	0.005	0.139	0.001
Actinobacteria	0.693	0.003	0.233	0.007	0.122	0.003
Acidobacteria	0.713	0.002	0.256	0.002	0.139	0.002
Bacteroidetes	0.755	<0.001	0.347	0.001	0.146	0.002
Verrucomicrobia	0.698	<0.001	0.318	0.003	0.16	0.003
Planctomycetes	0.798	0.002	0.379	0.001	0.125	0.001
Chloroflexi	0.72	0.005	0.271	0.004	0.139	0.007
Gemmatimonadetes	0.715	<0.001	0.269	0.003	0.148	0.001
Crenarchaeota	0.699	0.002	0.223	0.01	0.142	0.004
Cyanobacteria	0.866	0.003	0.213	0.015	0.109	0.004

27 ^aDifferent statistical approaches were used with Bray-Curtis distances, and *P* values were of corresponding significance

28 tests.

29 ^bMRPP, multiresponse permutation procedure.

30 ^cANOSIM, analysis of similarity.

31 ^dADONIS, permutational multivariate analysis of variance with the Adonis function.

32 **Supplementary Table S3. Spearman and Kendall rank correlation (r values, n=20) between different**
 33 **measurements.**

Spearman correlation	ATP	FCM	qPCR	PLFA	MBC
ATP	1	0.839**	0.793**	0.496**	-0.179
FCM		1	0.854**	0.518**	-0.245
qPCR			1	0.504**	-0.181
PLFA				1	-0.142
MBC					1
Kendall correlation	ATP	FCM	qPCR	PLFA	MBC
ATP	1	0.642**	0.586**	0.311*	-0.132
FCM		1	0.670**	0.364*	-0.111
qPCR			1	0.402**	-0.085
PLFA				1	-0.085
MBC					1

34 *: $P < 0.05$; **: $P < 0.01$

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36 **Supplementary Table S4. Correlation tests (r values, n=20) between different measurements and the**
37 **environmental factors.**

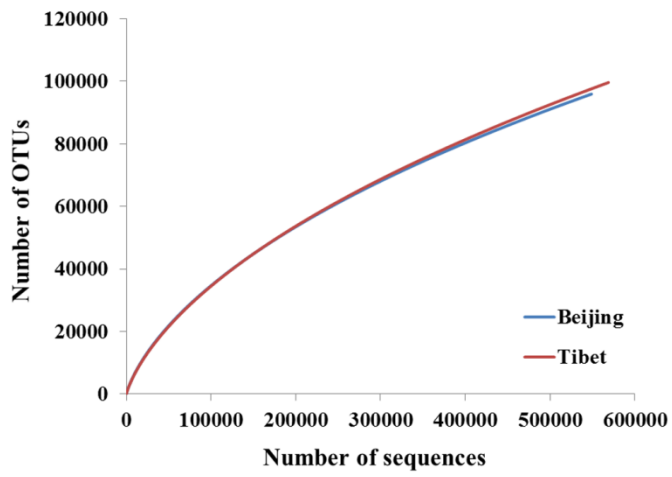
Pearson correlation	Moisture	TOC
ATP	0.806***	0.708***
FCM	0.803***	0.697**
qPCR	0.679***	0.748***
PLFA	0.650**	0.626**
MBC	-0.245	0.041

38 **: $P < 0.01$; *** : $P < 0.001$

39 **Supplementary Table S5. Correlation tests (r values, n=20) between the environmental factors and the EAA of**
 40 **the major phyla based on the ATP measurement.**

Pearson correlation	Moisture	TOC
Proteobacteria	0.784**	0.585**
Acidobacteria	0.698**	0.511*
Actinobacteria	0.741**	0.608**
Bacteroidetes	0.575**	0.528*
Verrucomicrobia	0.595**	0.508*
Chloroflexi	0.665**	0.512*
Gemmatimonadetes	0.651**	0.433
Planctomycetes	0.706**	0.577**
Cyanobacteria	-0.596**	-0.227

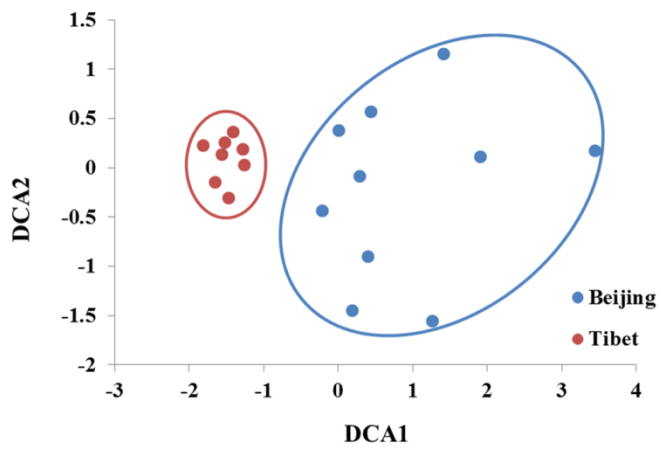
41 *: $P < 0.05$; **: $P < 0.01$



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43 **Supplementary Figure S1. Rarefaction curves based on 16S rRNA gene amplicon sequencing of soil microbial**

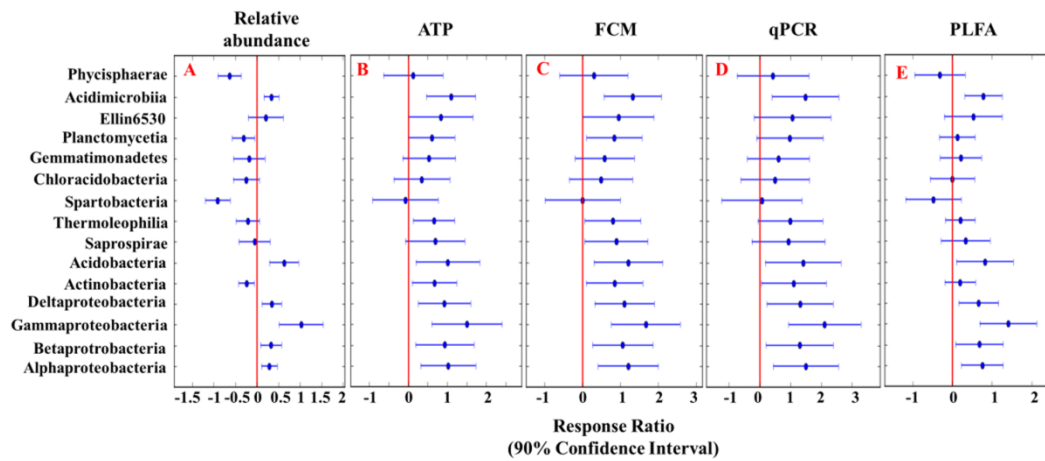
44 **communities.**



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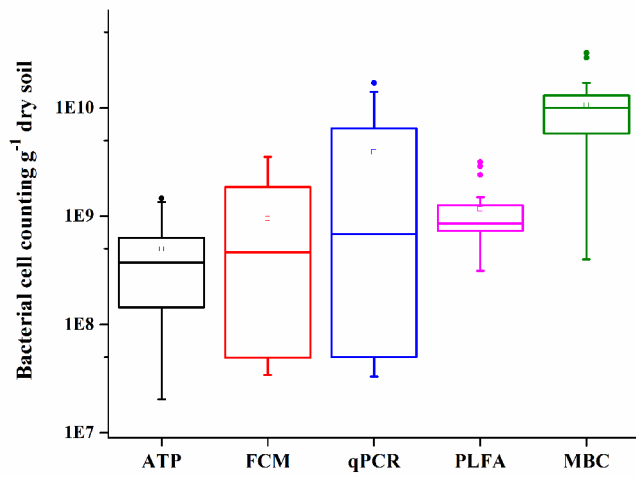
46 **Supplementary Figure S2. Detrended correspondence analysis (DCA) of microbial community composition in**

47 **Beijing and Tibet sampling sites based on 16S rRNA gene sequencing.**



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49 **Supplementary Figure S3. Response ratio for the differences between the relative abundances and absolute**
50 **abundances of the major class in the two sampling sites.** Significance was determined using the response ratio
51 analysis at a 95% confidence level. The 95% CI of a response variable without overlapping with zero represent a
52 significant result, otherwise, with non-significance. The vertical solid line was drawn when the variables of Beijing
53 samples equaled with those of Tibet samples.



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55 **Supplementary Figure S4. Bacterial cell numbers assessed by five different methods in Beijing and Tibet sites.**

56 Horizontal lines show median values, □ indicate mean values, boxes denote values comprised within the lower and

57 upper quartile of the data, the vertical lines represent ranges, and • indicate outliers.