

## **Supplementary information**

### **N-cycle genes abundance determined N mineralization rate following afforestation in the Loess Plateau of China**

Yaping Zhao<sup>1,2</sup>, Yuqing Zhao<sup>1,2</sup>, Shuohong Zhang<sup>1,2</sup>, Yilin Xu<sup>1,2</sup>, Xinhui Han<sup>1,2</sup>, Gaihe Yang<sup>1,2</sup>, Chengjie Ren<sup>1,2\*</sup>

<sup>1</sup>College of Agronomy, Northwest A&F University, Yangling, 712100 Shaanxi, China;

<sup>2</sup>Shaanxi Engineering Research Center of Circular Agriculture, College of Agronomy, Northwest A&F University, Yangling, 712100 Shaanxi, China

\* Corresponding author: Chengjie Ren

Email: Rencj1991@nwsuaf.edu.cn; Tel: +8613488354909

**Table S1.** Results from the redundancy analysis (RDA) to identify the conditional effects of soil properties on soil nitrogen functional genes abundance.

Name	Explains %	Contribution %	pseudo-F	<i>P</i>
MBN	53.8	60.5	15.2	<b>0.002</b>
Clay	11	12.4	3.8	<b>0.004</b>
Silt	3.3	3.7	1.1	0.348
TIN	3	3.3	1	0.408
SOC	4.2	4.8	1.6	0.164
TN	3.7	4.1	1.4	0.206
C/N	2	2.3	0.7	0.61
SM	3	3.4	1.2	0.324
BD	2.2	2.5	0.8	0.528
NO <sub>3</sub> <sup>-</sup>	1.6	1.8	0.5	0.712
PH	1	1.1	0.3	0.868

The conditional effects represent the proportion of variation explained by the individual environmental variables (soil properties). Significant results are shown in bold.

BD, bulk density; SOC, total soil organic C; TN, total nitrogen; C/N, the ratio of SOC and TN; TIN, total inorganic nitrogen; MBN, microbial biomass nitrogen; SM, soil moisture.

**Table S2** Results of final plspm model presented in figure 4.

<b>Number</b>	<b>Relationships</b>	<b>Direct</b>	<b>Indirect</b>	<b>Total</b>
1	Soil physical properties →Soil chemical properties	-0.618	0.000	-0.618
2	Soil physical properties →Soil microbial community	0.061	-0.534	-0.474
3	Soil physical properties →Nitrification	-0.011	-0.561	-0.572
4	Soil physical properties →Denitrification	-0.486	-0.126	-0.611
5	Soil physical properties →Soil mineralization rate	0.144	-0.664	-0.520
6	Soil chemical properties →Soil microbial community	0.864	0.000	0.864
7	Soil chemical properties →Nitrification	0.580	0.370	0.950
8	Soil chemical properties →Denitrification	-0.932	1.199	0.267
9	Soil chemical properties →Soil mineralization rate	-0.136	0.874	0.738
10	Soil microbial community →Nitrification	0.428	0.000	0.428
11	Soil microbial community →Denitrification	0.419	0.377	0.796
12	Soil microbial community →Soil mineralization rate	-0.156	0.776	0.620
13	Nitrification →Denitrification	0.881	0.000	0.881
14	Nitrification→ Soil mineralization rate	0.928	0.420	1.348
15	Denitrification →Soil mineralization rate	0.477	0.000	0.476

**Table S3.** Results of final plspm model presented in figure 4.

	<b>Type</b>	<b>R<sup>2</sup></b>	<b>Block Communality</b>	<b>Mean Redundancy</b>	<b>AVE</b>
Soil physical properties	Exogenous	0.000	0.766	0.000	0.766
Soil chemical properties	Endogenous	0.382	0.791	0.302	0.791
Soil microbial community	Endogenous	0.685	0.817	0.560	0.817
Nitrification	Endogenous	0.942	0.549	0.517	0.549
Denitrification	Endogenous	0.662	0.636	0.421	0.636
Soil mineralization rate	Endogenous	0.910	1.000	0.910	1.000

**Table S4 Loading coefficient between observation variables and latent variables.** SPP: soil physical properties, SCP: soil chemical properties, MC: microbial community, Ni: nitrification, De: denitrification, R<sub>m</sub>: Soil net mineralization rate. \* indicates that there is a significant relationship between latent variables.

Latent variables	Observation variables	SPP	SCP	MC	Ni	De	R <sub>m</sub>
SPP	Bulk density	0.849	-0.599*	-0.458	-0.520	-0.302*	-0.312
	Silt	0.901	-0.496*	-0.380	-0.486	-0.729*	-0.574
SCP	Soil organic C	-0.609	0.944	0.858*	0.946*	0.675*	0.854
	Total N	-0.472	0.830	0.556*	0.682*	0.175*	0.441
MC	Shannon	-0.520	0.723	0.916	0.814*	0.712	0.789
	NMDS1	-0.325	0.775	0.892	0.837*	0.510	0.717
Ni	<i>amoB</i>	0.034	0.316	0.403	0.384	0.080	0.210*
	<i>narH,narY,nxrB</i>	-0.619	0.927	0.876	0.975	0.690	0.901*
De	<i>norC</i>	-0.169	0.285	0.340	0.310	0.615	0.303*
	<i>norB</i>	-0.663	0.533	0.678	0.669	0.946	0.864*
R <sub>m</sub>	net mineralization rate	-0.520	0.777	0.835	0.892	0.825	1.000

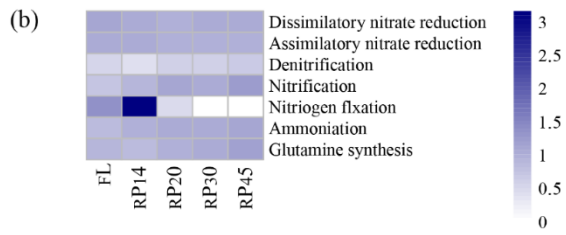
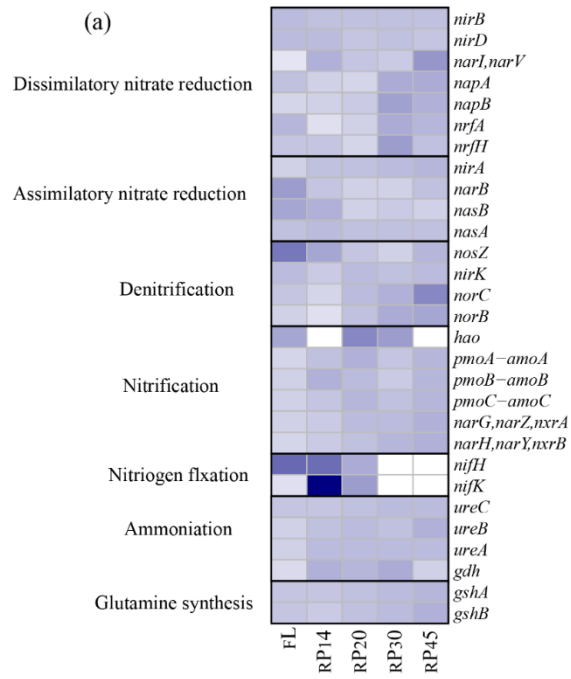
**Table S5. Changes of gene abundance (Nitrification genes and Denitrification genes) along with afforestation chronosequence.**

	Gene	FL	RP14	RP20	RP30	RP45	*P
Nitrification	<i>hao</i>	0.62±0.09a	0b	0.85±0.22a	0.68±0.03a	0b	<b>0.001</b>
	<i>pmoA-amoA</i>	8.54±1.22c	12.25±0.15ab	15.07±0.59a	11.33±1.24b	13.71±0.62ab	<b>0.003</b>
	<i>pmoB-amoB</i>	10.32±1.25c	16.17±1.37a	14.70±1.09ab	11.06±0.70b	14.97±1.61ab	<b>0.027</b>
	<i>pmoC-amoC</i>	13.02±1.15b	15.74±0.53ab	19.18±1.58a	16.31±1.82ab	18.85±1.54a	0.059
	<i>narG,narZ,nxrA</i>	34.80±2.27c	40.35±1.64bc	52.68±2.33a	51.77±5.49ab	59.71±4.83a	<b>0.004</b>
	<i>narH,narY,nxrB</i>	16.59±0.34c	19.82±1.17c	24.14±1.05b	27.68±1.82ab	29.90±1.26a	<b>0.000</b>
Denitrification	<i>nosZ</i>	1.49±0.57a	0.65±0.33ab	0.63±0.27ab	0.52±0.12ab	0b	0.090
	<i>nirk</i>	20.09±1.33a	16.74±1.00a	20.17±3.88a	18.89±1.98a	19.92±1.79a	0.792
	<i>norC</i>	0.66±0.52ab	0.53±0.23ab	0.80±0.26ab	0b	1.41±0.16a	0.062
	<i>norB</i>	7.41±0.86d	4.72±1.49cd	9.33±0.76bc	11.91±1.44ab	13.29±1.16a	<b>0.003</b>

**Table S6. Changes of gene abundance (Six nitrogen metabolism pathways) along with afforestation chronosequence.**

	<b>FL</b>	<b>RP14</b>	<b>RP20</b>	<b>RP30</b>	<b>RP45</b>	<b>*P</b>
Nitrification	83.89±1.81d	104.34±2.46c	126.60±4.07ab	118.82±8.18b	137.24±4.81a	<b>0.000</b>
Denitrification	29.65±1.14ab	22.64±1.79b	30.92±4.35ab	31.63±3.73ab	34.90±0.85b	0.094
Assimilatory nitrate reduction	239.39±2.73ab	245.36±11.96a	221.56±3.14b	227.60±2.80ab	231.17±4.85ab	0.127
Dissimilatory nitrate reduction	358.73±2.54a	330.55±11.06b	322.95±3.84b	340.43±4.80ab	342.53±9.31ab	<b>0.045</b>
Ammoniation	205.87±4.32c	233.64±10.58bc	250.78±11.77ab	249.21±10.26ab	266.57±5.95a	<b>0.008</b>
Glutamine synthesis	204.34±6.02cd	197.66±3.82d	223.85±4.01bc	233.75±9.91b	259.55±7.02a	<b>0.000</b>

**Fig.S1.**



**Fig.S2.**

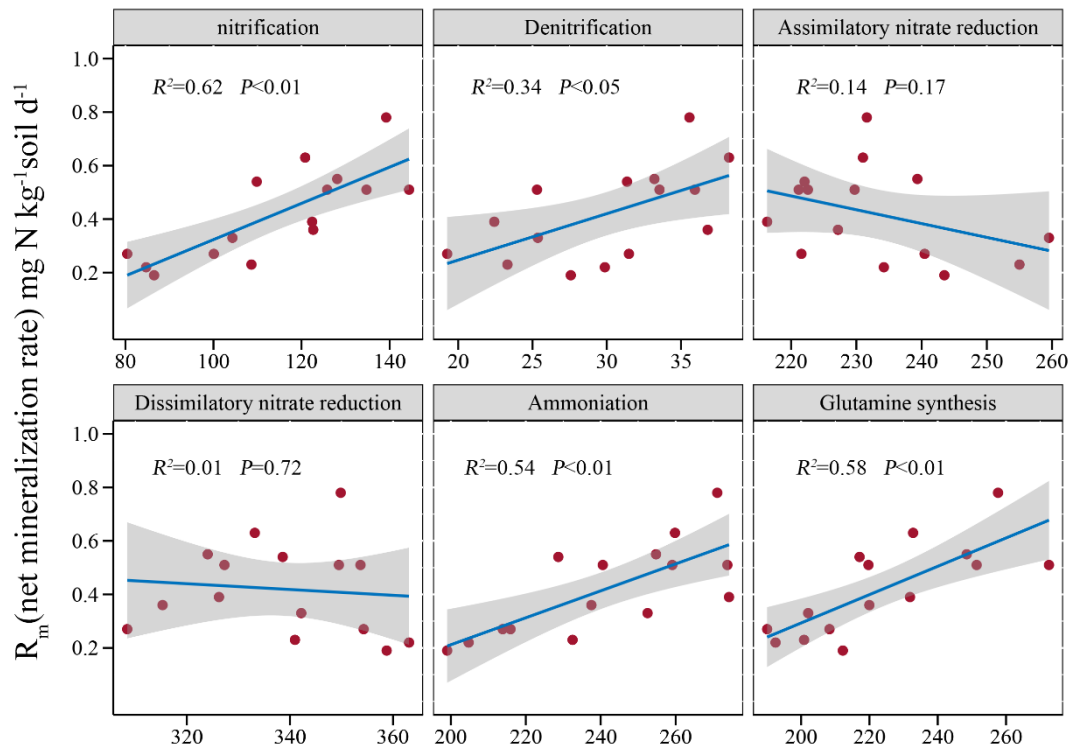
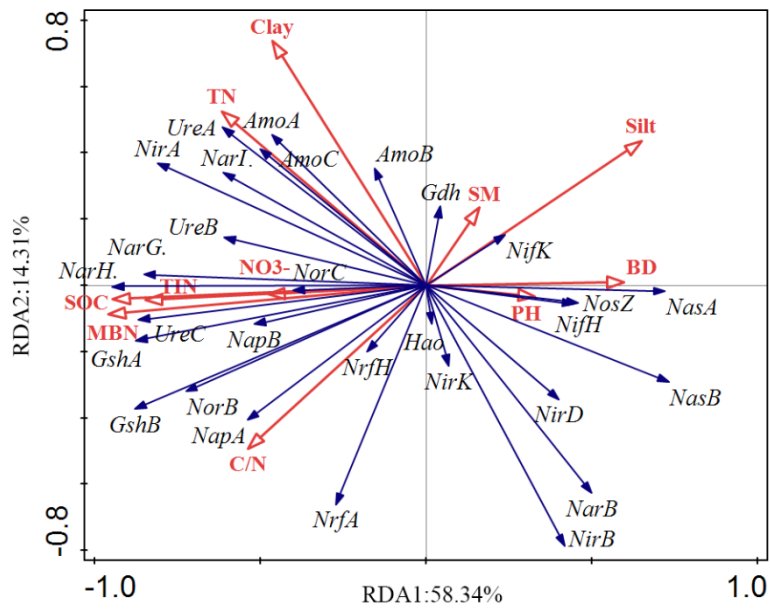


Fig.S3.



**Fig.S4.**

