

Supplementary Materials

Table S1 Oligonucleotides used for PCR amplification and sequencing of denitrification function gene fragments

gene	Primer	Primer fragments (5-3)	Length (bp)	annealing temperatures (°C)	Reference
<i>NosZ</i>	<i>NosZ</i> -1F	GGTAACCTTGACAACACCGA	1100	56	(Scala and Kerkhof, 1999)
	<i>NosZ</i> -2R	ATGACGAAGCCGTGAGACA			
<i>NirK</i>	<i>NirK</i> -1F	GG(A/C)ATGGT(G/T)CC(C/G)TGGCA	514	56	(Braker et al., 1998)
	<i>NirK</i> -2R	GCCTCGATCAG(A/G)TT(A/G)TGG			
<i>NapA</i>	<i>NapA</i> -1F	TCTGGACCATGGGCTTCAACCA	877	59	(Huang et al., 2013)
	<i>NapA</i> -2R	ACGACGACCGGCCAGCGCAG			
<i>NirS</i>	<i>NirS</i> -1F	CCTA(C/T)TGGCCGCC(A/G)CA(A/G)T	890	56	(Braker et al., 1998)
	<i>NirS</i> -2R	CGTTGAACTT(A/G)CCGGT			
<i>NorB</i>	<i>NorB</i> F	CGNGARTYCTSGARCARCC	669	55	(Garbeva et al., 2007)
	<i>NorB</i> R	CRTADGCVCCRWAGAAVGC			

Table S2 Oligonucleotides used for qPCR amplification and sequencing of denitrification function gene fragments

gene	Primer	Primer fragments (5-3)	Length (bp)	annealing temperatures (°C)	Reference
<i>NosZ</i>	<i>NosZ</i> -1527F	CGCTGTTCHTCGACAGYCA	250	56	(Wang et al., 2015)
	<i>NosZ</i> -1773R	ATRTCGATCARCTGBTCGTT			
<i>NirK</i>	<i>NirK</i> -1F	TCATGGTGCTGCCGCGKGCACGG	326	64	(Wang et al., 2015)
	<i>NirK</i> -2R	GAACCTTGCCGGTKGCCACAG			
<i>NapA</i>	<i>NapA</i> -1F	CCCAATGCTCGCCACTG	130	60	(Cheng et al., 2019)
	<i>NapA</i> -2R	CATGTTKGAGCCCCACAG			
<i>NorB</i>	<i>NorB</i> -1F	GCAGCCGATTTTACTCAGCG	171	55	(Pal et al., 2015)
	<i>NorB</i> -2R	TCTTTGGCGGCCACGATATT			
16S rRNA	16S rRNA-F	CGGTGAATACGTTTCYCGG	142	55	(He et al., 2017)
	16S rRNA-R	GGHTACCTTGTTACGACTT			

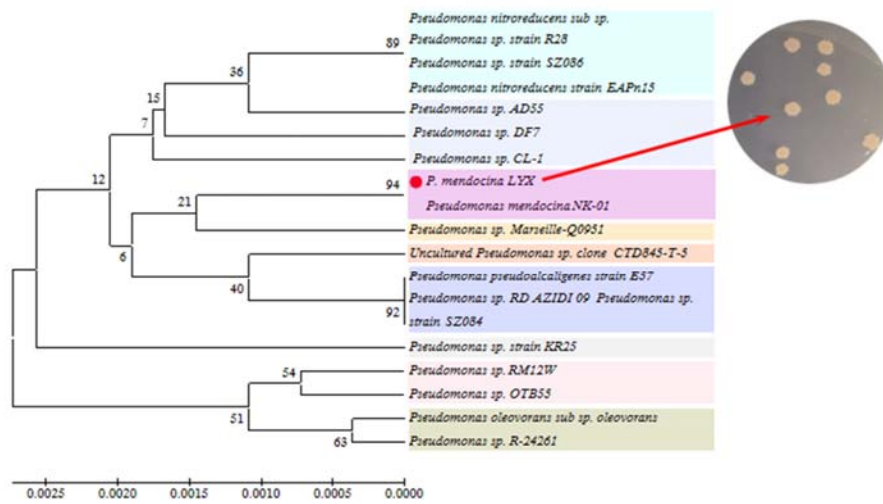


Fig. S1 Phylogenetic tree based on the 16S rRNA gene sequence of *Pseudomonas mendocina* LYX.

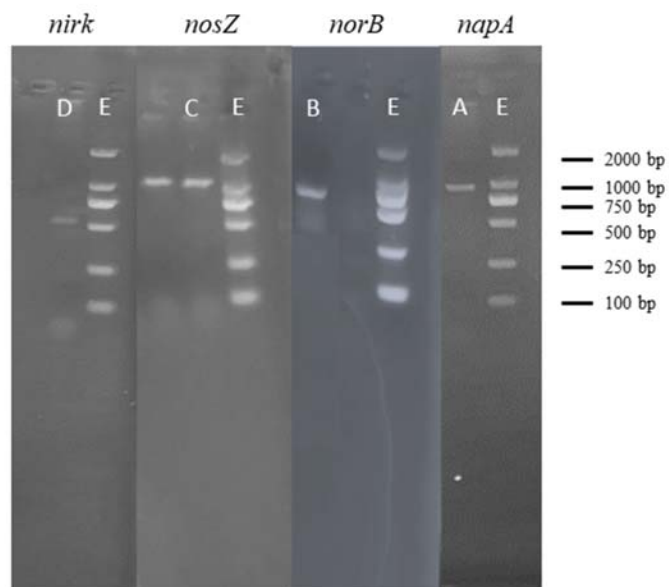


Fig. S2 Agarose gel electrophoresis of PCR amplified *napA*, *nirK*, *nosZ* and *norB* gene fragments from bacteria DNA, *napA*-1F/*napA*-2R(lanes A), using *norBF*/*norBR*(lanes B), *nosZ*-1F/*nosZ*-2R(lanes C), *nirK*-1F/*nirK*-2R(lanes D). Lanes E, DNA marker

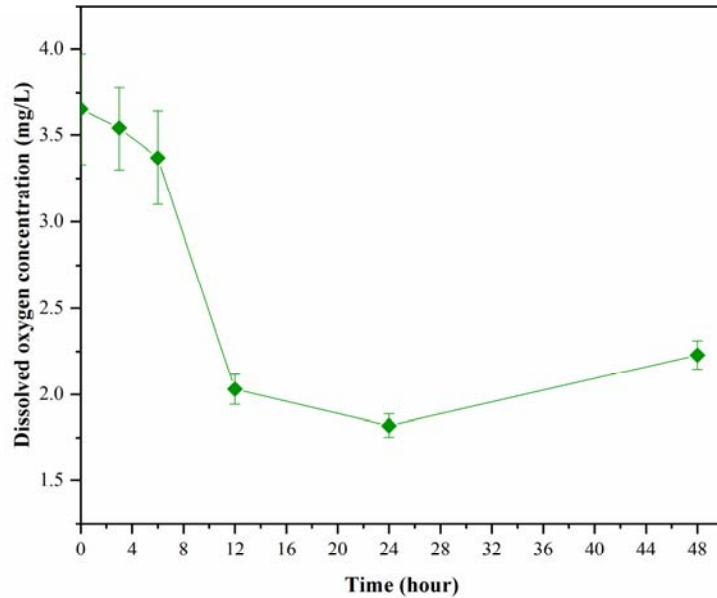


Fig. S3 Changes of dissolved oxygen concentration during nitrogen balance experiments

16SrRNA sequencing fragments:

AAATGCCGAGCCTACACATGCAAGTCGAGCGGATGAAAAGAGCTTGCTCCCTGATTTAGCGGCGGAC
GGGTGAGTAATGCCTAGGAATCTGCCTGGTAGTGGGGGATAACGTTCCGAAAGGAACGCTAATACCG
CGTACGTCCTACGGGAGAAAGCAGGGGACCTTCGGGCCTTGCCTATCAGATGAGCCTAGGTTCGGATT
AGCTAGTTGGTGAGGTAATGGCTCACCAAGGCGACGATCCGTAAGTGGTCTGAGAGGATGATCAGTCA
CACTGGAAGTACGACACGGTCCAGACTCTACGGGAGGCAGCAGTGGGGAATATTGGACAATGGGCG
AAAGCCTGATCCAGCCATGCCGCGTGTGTGAAGAAGGTCTTCGGATTGTAAAGCACTTTAAGTTGGGA
GGAAGGGCAGTAAGTTAATACCTTGCTGTTTTGACGTTACCGACAGAATAAGCACCGGCTAACTTCGT
GCCAGCAGCCGCGTAATACGAAGGGTGCAAGCGTTAATCGGAATTACTGGGCGTAAAGCGCGCGTA
GGTGGTTTCGTTAAGTTGGATGTGAAAGCCCCGGGCTCAACCTGGGAACTGCATCCAAAACCTGGCGAGC
TAGAGTACGGTAGAGGGTGGTGAATTTCTGTGTAGCGGTGAAATGCGTAGATATAGGAAGGAACA
CCAGTGGCGAAGGCGACCACCTGGACTGATACTGACACTGAGGTGCGAAAGCGTGGGGAGCAAACAG
GATTAGATACCCTGGTAGTCCACGCCGTAACGATGTCAACTAGCCGTTGGAATCCTTGAGATTTTAGT
GGCGCAGCTAACGCATTAAGTTGACCGCTGGGGAGTACGGCCGCAAGGTTAAAACCTCAAATGAATT
GACGGGGGCCCCGACAAGCGGTGGAGCATGTGGTTTAATTTGAAGCAACGCGAAGAACCCTTACCTGG
CCTTGACATGCTGAGAACTTTCCAGAGATGGATTGGTGCCTTCGGGAACTCAGACACAGGTGCTGCAT
GGCTGTTGTCAGCTCGTGTGCTGAGATGTTGGGTTAAGTCCCCTAACGAGCGCAACCCTTGTCCCTAGT
TACCAGCACGTTATGGTGGGCACTCTAAGGAGACTGCCGGTGACAAACCGGAGGAAGGTGGGGATGA
CGTCAAGTCATCATGGCCCTTACGGCCAGGGCTACACACGTGCTACAATGGTCGGTACAAAGGGTTGC
CAAGCCGCGAGGTGGAGCTAATCCATAAAAACCGATCGTAGTCCGGATCGCAGTCTGCAACTCGACTG
CGTGAAGTCGGAATCGCTAGTAATCGTGAATCAGAATGTCACGGTGAATACGTTCCCGGGCCTTGAC
ACACCGCCCGTACACCATGGGAGTGGGTTGCTCCAGAAGTAGCTAGTCTAACCTTCGGGGGGACGGT
ACCACGGAAGGATCCAGG

Reference

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