Supporting information

Relative contributions of denitrification and anammox to nitrogen removal in riverine

wetlands across China

Danli Deng 1, 2, 3, Di Xu 3, 4, Gang He 3, 4, Bangjing Ding 3, 4, Wenzhi Liu 3, 4*

¹ Post Doctoral Research Station of Hydraulic Engineering of Three Gorges University, Yichang 443002,

China

² Hubei Field Observation and Scientific Research Stations for Water Ecosystem in Three Gorges Reservoir,

China Three Gorges University, Yichang 443002, China

³ Hubei Key Laboratory of Wetland Evolution & Ecological Restoration, Wuhan Botanical Garden, Chinese

Academy of Sciences, Wuhan 430074, China

⁴ Danjiangkou Wetland Ecosystem Field Scientific Observation and Research Station, the Chinese Academy

of Sciences & Hubei Province, Wuhan 430074, China

*Corresponding author: Wenzhi Liu

Phone: +86 27 87700866;

Fax: +86 27 87700877;

Email: liuwz@wbgcas.cn

Determination of nitrogen removal potential

The denitrification and anammox rates of channel sediments, riparian rhizosphere soils, and

depth-selected riparian bulk soils (0-20, 60-80, and 160-180 cm in depth) were quantified using the ¹⁵N

isotope pairing method [1,2]. Briefly, three 5 g replicates of homogenized sediment or soil samples were

weighed into 12-mL cylindrical glass vials containing helium-purged distilled water [3]. Subsequently, all

vials were pre-incubated in a horizontal shaker (180 rpm) at 24°C for 3 days to remove the background NO₃-

and NO₂-, as well as residual oxygen [4]. All vials were then divided into three groups and injected with 100

μL helium-purged stock solution: ¹⁵NH₄Cl (98 at.% ¹⁵N), ¹⁵NH₄Cl + K¹⁴NO₃, or ¹⁴NH₄Cl + K¹⁵NO₃ (99.99

at.% ¹⁵N). This resulted in a final total N concentration of 100 μmol in each vial ^[5]. Additions of ¹⁵NH₄⁺ solutions were used to verify that background NO₃⁻-N and NO₂⁻-N have been consumed, while additions of ¹⁵NH₄⁺-N + ¹⁴NO₃⁻-N solutions were used to confirm the presence of anammox process. Additions of ¹⁴NH₄⁺-N + ¹⁵NO₃⁻-N solutions were used to determine denitrification and anammox rates. The incubation was terminated at five-time intervals (0, 1, 2, 4, and 8 h) by injecting 200 μL of 7 M ZnCl₂ solution. The isotopic abundance of ²⁸N₂, ²⁹N₂, and ³⁰N₂ in the glass vials were measured using isotope ratio mass spectrometry (Gasbench II and Delta V Advantage, Thermo Finnigan, Germany).

Analysis of abundance of denitrifying and anammox bacteria

DNA was extracted from replicate soil and sediment samples using a PowerSoil DNA Isolation Kit (MoBio Laboratories Inc., CA, USA), and DNA concentrations were determined using a NanoDrop 2000 Spectrophotometer (Thermo Fisher Scientific, MA, USA). The abundance (i.e., copy number) of nirK, nirS, and hzsB genes was measured in triplicate by using a Roche LightCycler480 real-time PCR system (Roche Diagnostics, Mannheim, Germany) with the fluorescent dye SYBR green qPCR method. The reduction of NO₂-N to nitric oxide (NO) is the rate-limiting step of denitrification, which is catalyzed by either a copper-containing enzyme encoded by the nirK gene or a cytochrome-cd1 enzyme encoded by the nirS gene^[6]. The hzsB gene encoded for hydrazine synthase β -subunit which catalyzes the production of hydrazine from NH₄⁺-N and NO₂⁻-N during the anammox process ^[4]. Detailed procedures of qPCR were described previously^[7]. In brief, the abundance (i.e., copy number) of nirS, nirK, and hzsB genes was determined in triplicate using a Roche LightCycler480 real-time PCR System (version 1.5.0; Roche Diagnostics, Mannheim, Germany) with the fluorescent dye SYBR green quantitative PCR method. Primer sets of nirSCd3aF/nirSR3cdR, nirKF1aCu/nirKR3Cu, and hzsB396F/hzsB742R were applied for the nirS, nirK, and hzsB genes, respectively (Table S2). The 20-µL quantitative PCR mixture contained 10 µL of SyberGreen qPCR Master Mix (2×), 1 μL of primers (10 μM), and 2 μL of DNA template. The primers and qPCR protocol are listed in Supplementary Table S2. Standard curves were constructed with serial plasmid dilutions of a known amount of plasmid DNA containing a fragment of the nirS, nirK, and hzsB genes.

Measurements of soil, sediment, and water physicochemical properties

Soil pH was measured at a soil-to-water ratio of 1:5 (v/v) using a pH meter (Hanna Instruments, Padova, Italy), whereas soil moisture content was determined gravimetrically by drying 20 g of fresh soil at 105°C

for 24 hours. T The soil TN, TC, and TOC concentrations of air-dried soil samples were determined by an elemental analyzer. (Vario TOC cube, Hanau, Germany). Soil or sediment physicochemical properties, including moisture, pH, total N (STN), NH₄⁺-N, NO₃⁻-N, total carbon (STC) and organic carbon (SOC) were determined following published analytical methods ^[8,9]. Soil Fe²⁺ and AFe concentrations were measured using the phenanthroline spectrophotometric method ^[10]. The concentrations of water TN, NH₄⁺-N, NO₃⁻-N, TC, and TOC were measured in the laboratory, as described previously^[11]. The concentrations of NH₄⁺ and NO₃⁻ in water were analyzed by a continuous flow analyzer (EasyChem plus, Systea, Italy). The water TN, TC, and TOC concentrations of samples were determined by a Total Organic Carbon Analyzer Analyzer (Liquid). (Vario TOC cube, Hanau, Germany).

References

- [1] Thamdrup B, Dalsgaard T. 2002. Production of N₂ through anaerobic ammonium oxidation coupled to nitrate reduction in marine sediments. Applied and Environmental Microbiology 68: 1312-1318 https://doi.org/10.1128/aem.68.3.1312-1318.2002
- [2] Risgaard-Petersen N, Nielsen L, Rysgaard S, Dalsgaard T, Meyer R. 2003. Application of the isotope pairing technique in sediments where anammox and denitrification coexist. Limnology and Oceanography-methods 1: 63-73 https://doi.org/10.4319/lom.2003.1.63
- [3] Deng D, Pan Y, Liu G, Liu W, Ma L. 2020. Seeking the hotspots of nitrogen removal: A comparison of sediment denitrification rate and denitrifier abundance among wetland types with different hydrological conditions. Science of the Total Environment 737: 140253 https://doi.org/10.1016/j.scitotenv.2020.140253
- [4] Deng D, Ding B, He G, Ji M, Yang Y, et al. 2023. The contribution of anammox to nitrogen removal is greater in bulk soils than in rhizosphere soils in riparian wetlands along the Yangtze River. Global Biogeochemical Cycles 37(5): e2022GB007576 https://doi.org/10.1029/2022gb007576
- [5] Deng D, Yang Z, Yang Y, Wan W, Liu G, et al. 2024. Metagenomic insights into nitrogen-cycling microbial communities and their relationships with nitrogen removal potential in the Yangtze River. Water Research 265: 122229 https://doi.org/10.1016/j.watres.2024.122229
- [6] Wallenstein M, Myrold D, Firestone M, Voyteck M. 2006. Environmental controls on denitrifying communities and denitrification rates: insights from molecular methods. Ecological Applications 16(6): 2143-2152 https://doi.org/10.1890/1051-0761(2006)016[2143:ecodca]2.0.co;2
- [7] Jiang X, Yao L, Guo L, Liu G, Liu W. 2017. Multi-scale factors affecting composition, diversity, and abundance of sediment denitrifying microorganisms in Yangtze lakes. Applied Microbiology and Biotechnology 101: 8015-8027 https://doi.org/10.1007/s00253-017-8537-5
- [8] Liu W, Wang Z, Zhang Q, Cheng X, Lu J, et al. 2015. Sediment denitrification and nitrous oxide

- production in Chinese plateau lakes with varying watershed land uses. Biogeochemistry 123: 379-390 https://doi.org/10.1007/s10533-015-0072-9
- [9] Jiang X, Liu W, Yao L, Liu G, Yang Y. 2020. The roles of environmental variation and spatial distance in explaining diversity and biogeography of soil denitrifying communities in remote Tibetan wetlands. FEMS Microbiology Ecology 96(5): fiaa063 https://doi.org/10.1093/femsec/fiaa063
- [10] Herrera L, Ruiz P, Aguillon J, Fehrmann. 1989. A new spectrophotometric method for the determination of ferrous iron in the presence of ferric iron. Journal of Chemical Technology and Biotechnology 44(3): 171-181 https://doi.org/10.1002/jctb.280440302
- [11] Liu W, Jiang X, Zhang Q, Li F, Liu G. 2018. Has submerged vegetation loss altered sediment denitrification, N₂O production and denitrifying microbial communities in subtropical lakes? Global Biogeochemical Cycles 32: 1195-1207 https://doi.org/10.1029/2018gb005978