

Supplementary Information

Wastewater-influenced estuaries are characterized by disproportionately high nitrous oxide emissions but overestimated IPCC emission factor

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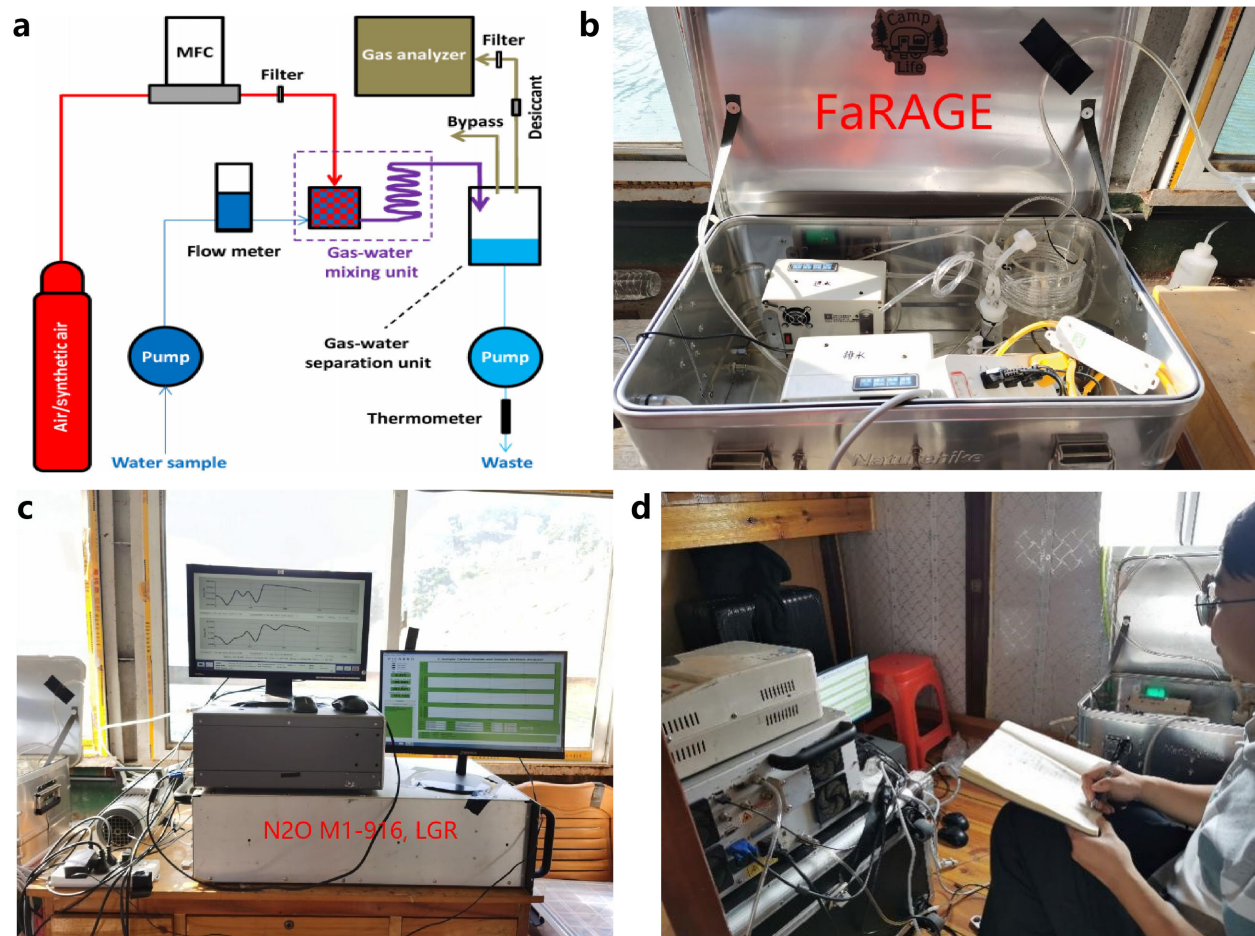
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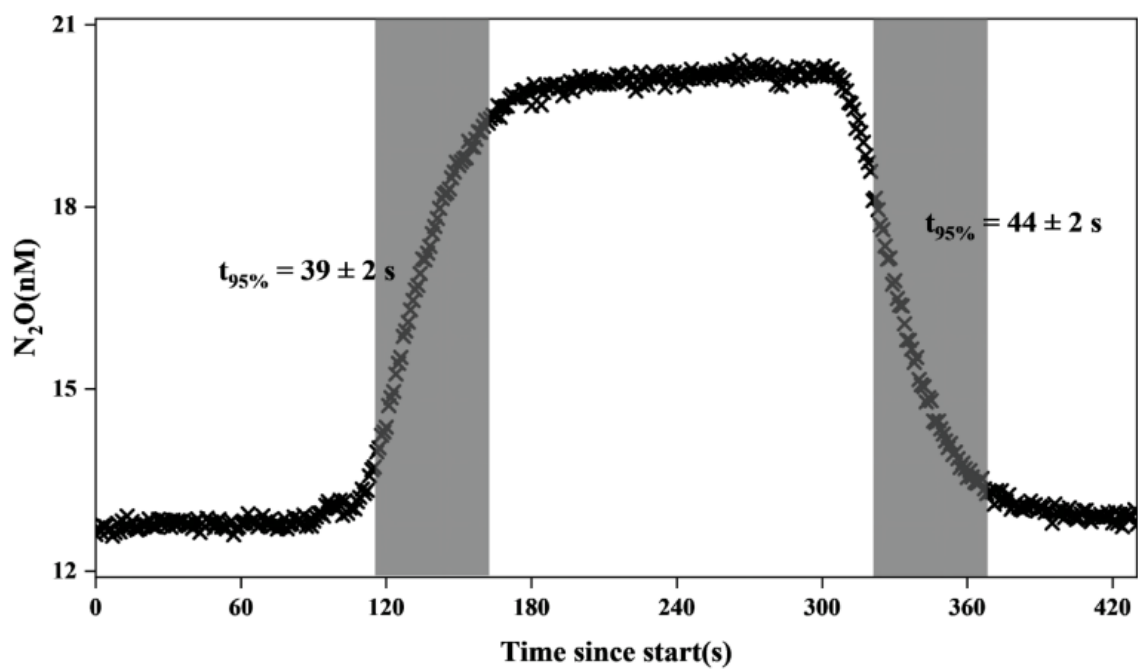
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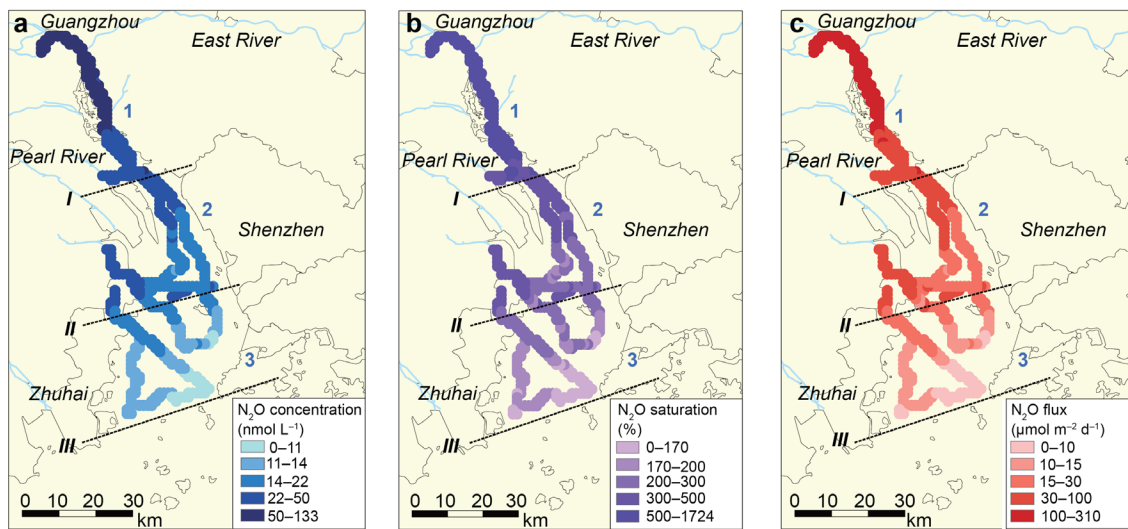
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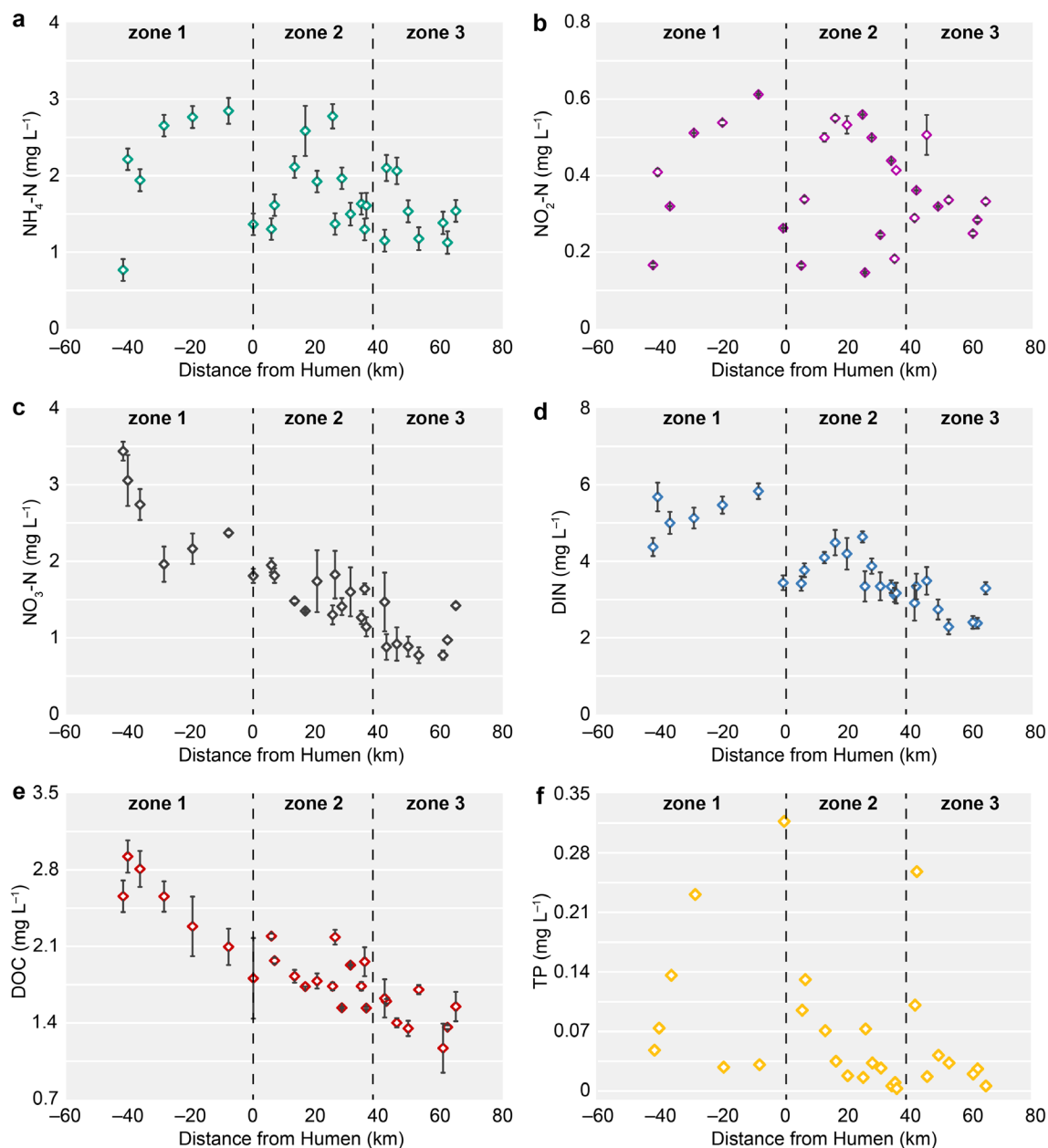
Supplementary Figure 1. High-resolution N₂O monitoring platform used in this study. a, Schematic the FaRAGE system¹. **b,** Photo of the FaRAGE system used in the field campaign. **c,** The FaRAGE system was coupled with a gas analyzer (N₂O M1-916, LGR). **d,** A researcher monitoring the system.



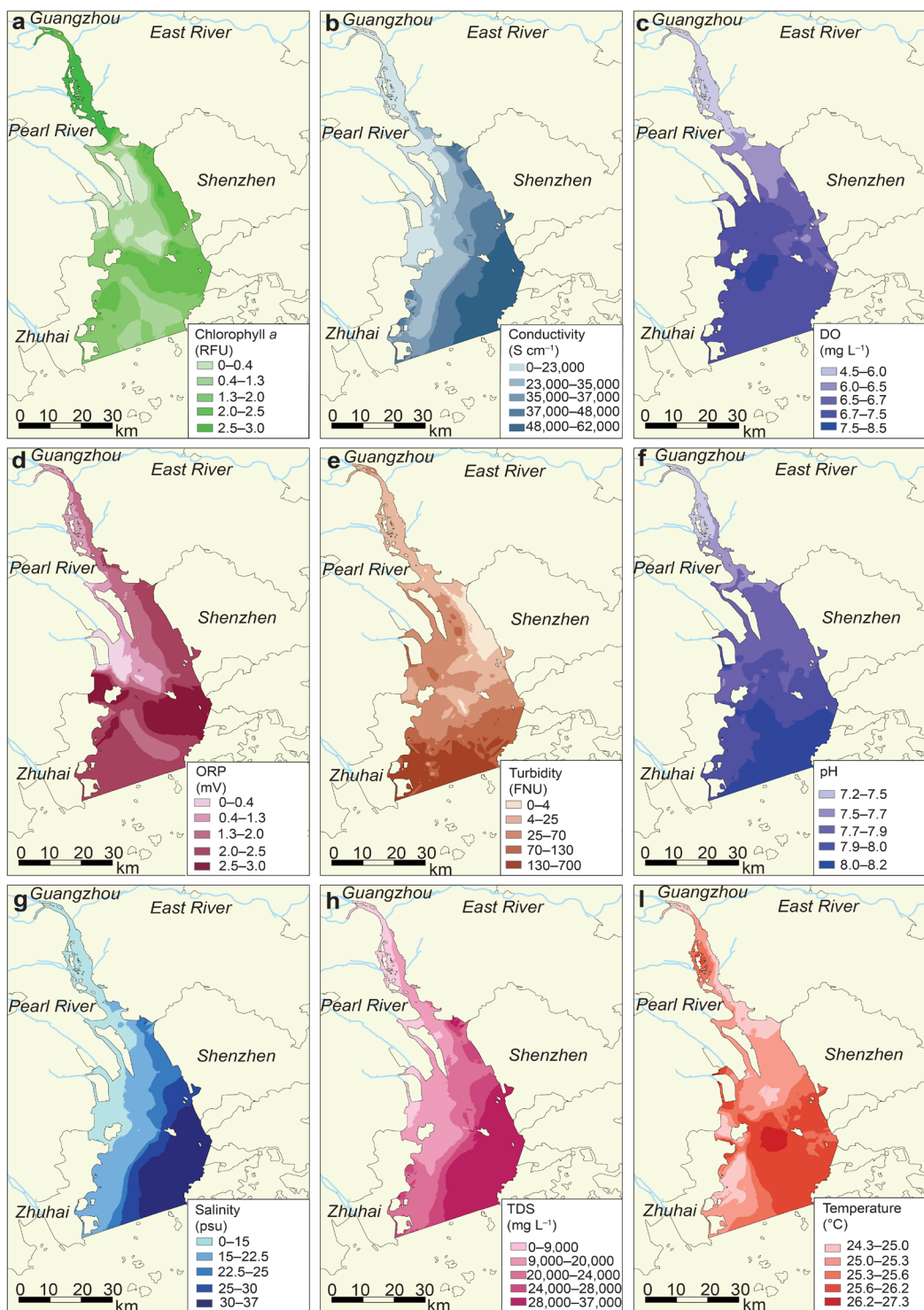
Supplementary Figure 2. Response process of the sample entering into the plateaus of measurement with a response time of 41 ± 2 s using the FaRAGE-based N₂O measurements.



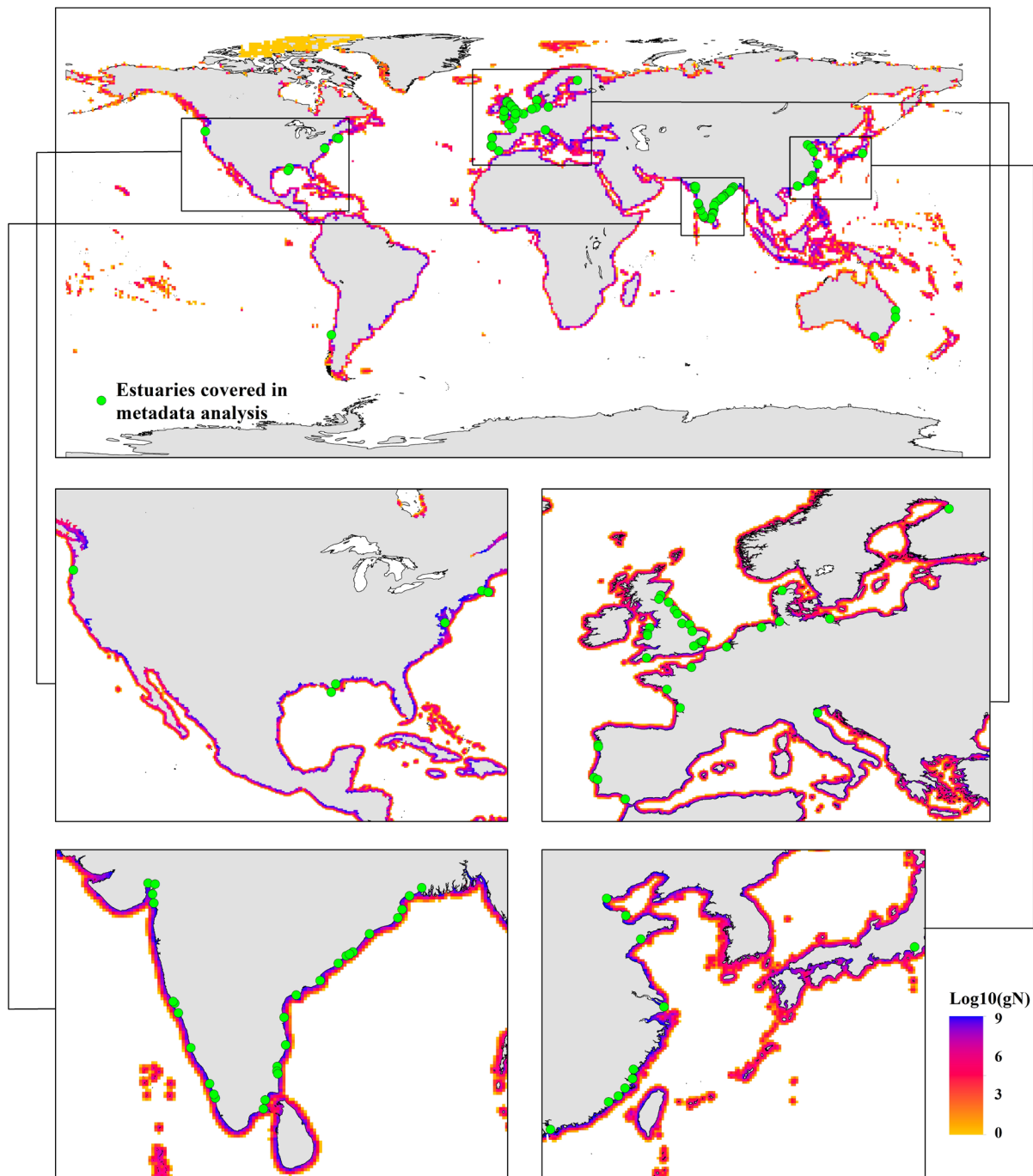
Supplementary Figure 3. Spatial distributions of N_2O concentration, saturation, and flux in PRE from the real-time, high-resolution monitoring. a, Dissolved N_2O concentration. b, N_2O saturation. c, N_2O flux. The base map is from RESDC data (<http://www.resdc.cn/>).



Supplementary Figure 4. Distribution of physicochemical parameters in PRE as a function of distance from Humen. a, $\text{NH}_4\text{-N}$. b, $\text{NO}_2\text{-N}$. c, $\text{NO}_3\text{-N}$. d, DIN. e, DOC. f, TP. The distance is positive downstream of Humen Outlet. Dashed lines in black delineate three zones of PRE: zone 1—upper Estuary (Humen upstream), zone 2—middle Estuary (Inner Lingdingyang), and zone 3—lower Estuary (Outer Lingdingyang).



Supplementary Figure 5. Spatial distribution of physicochemical parameters. a, Chlorophyll a. b, Conductivity. c, DO. d, ORP. e, Turbidity, f, pH. g, Salinity. h, Total dissolved solids (TDS). i, Temperature. The base map is from RESDC data (<http://www.resdc.cn/>).



Supplementary Figure 6. The distribution of wastewater nitrogen input and estuarine locations covered in the meta-analysis.

Supplementary Table 1. Goodness of fit for statistics of linear, efficiency loss and Michaelis-Menten model in global, Asian, European, American and Oceania estuaries.

		RSE	Adjusted R ²	p-value	RMSE	AIC
Global	Michaelis–Menten	0.304	0.254	2.547e–41	0.298	32.20
	Linear	0.299	0.274	6.480e–06	0.294	30.50
	Efficiency loss	0.250	0.489	1.078e–10	0.246	8.28
Asian	Michaelis–Menten	0.269	0.281	3.367e–19	0.258	9.43
	Linear	0.306	0.072	0.100	0.294	16.10
	Efficiency loss	0.277	0.238	0.007	0.266	11.00
European	Michaelis–Menten	0.232	0.423	5.397e–13	0.218	2.46
	Linear	0.256	0.299	0.014	0.240	5.77
	Efficiency loss	0.233	0.419	0.003	0.219	2.57
American and Oceania	Michaelis–Menten	0.248	0.149	1.216e–10	0.233	4.65
	Linear	0.1898	0.504	0.001	0.178	–4.50
	Efficiency loss	0.2168	0.351	0.007	0.2038	0.042

Supplementary Table 2. Primer pairs used in qPCR assays and the corresponding amplification conditions.

Gene	Primer	Sequence (5'-3')	Thermal profile	Reaction mixture	Amplification efficiency (%)	Ref.
AOA	arch-amoA-23F	ATGGTCTGGCTWAGACG	95°C for 3 min, 40 cycles of 5 s at 95°C, 30 s at 58°C, and 1 min at 72°C	10 µL 2X ChamQ SYBR	89	1
	arch-amoA-616R	GCCATCCATCTGTATGTCCA		Color qPCR		
AOB	CTO189f	GGAGRAAAGCAGGGGATCG	95°C for 3 min, 40 cycles of 5 s at 95°C, 30 s at 58°C, and 1 min at 72°C	Master Mix, 0.8 µL of each primer (5 µM),	105	2
amoA	CTO654r	CTAGCYTTGTAGTTTCAAAC GC		0.4 µL 50 X ROX		
nirS	cd3aF	GTSAACGTAAGGARACSGG	95°C for 3 min, 40 cycles of 5 s at 95°C, 30 s at 58°C, and 1 min at 72°C	0.4 µL 50 X	104	3
	R3cd	GASTTCGGRTGSGTCTTGA				
nirK	FlaCu	ATCATGGTSGTGCCGCG	95°C for 3 min, 40 cycles of 5 s at 95°C, 30 s at 58°C, and 1 min at 72°C	Reference Dye	101	4
	R3cu	GCCTCGATCAGRTTGTGGTT				
narG	narG-f	TCGCCSATYCCGGCSATGTC	95°C for 3 min, 40 cycles of 5 s at 95°C, 30 s at 58°C, and 1 min at 72°C	1, and 2 µL Template (DNA)	99	5
	narG-r	GAGTTGTACCAGTCRGC SGAYTCSG				
nosZ	nosZF	CGYTGTTCMTCGACAGCCAG	95°C for 3 min, 40 cycles of 5 s at 95°C, 30 s at 58°C, and 1 min at 72°C		96	6
	nosZR	CATGTGCAGNGC RTGGCAGAA				

Supplementary References

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