Supplementary information

Aquatic macrophytes in morphological and physiological responses to the nanobubble technology application for water restoration

Shuo Wang[†], Yunsi Liu[†], Tao Lyu[‡], Gang Pan^{*, §}, Pan Li^{*,†,"}

[†]School of Environmental Science and Engineering, Tongji University, 1239 Siping Road, Shanghai, PR China.

[‡]Cranfield Water Science Institute, Cranfield University, College Road, Cranfield, Bedfordshire, MK43 0AL, UK

[§]Integrated Water-Energy-Food Facility (iWEF), School of Animal, Rural, and Environmental Sciences, Nottingham Trent University, Nottinghamshire NG25 0QF, UK

[#]School of Environmental Science and Engineering, State Key Laboratory of Control and Resource Reuse, Tongji University, 1239 Siping Road, Shanghai, PR China. *Corresponding authors: lipan@tongji.edu.cn (P. Li); gang.pan@ntu.ac.uk (G. Pan).

The SI contains 5 pages, which has 3 Figures and 2 Tables.

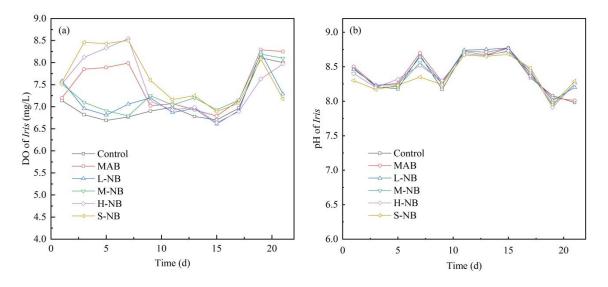


Figure S1. Changes in (a) DO level and (b) pH in water over the 21 days of cultivation of *Iris.* MAB, L-NB, M-NB, H-MB, S-NB represent macrobubble aeration, low, medium, high and super-high nanobubble aeration groups, respectively.

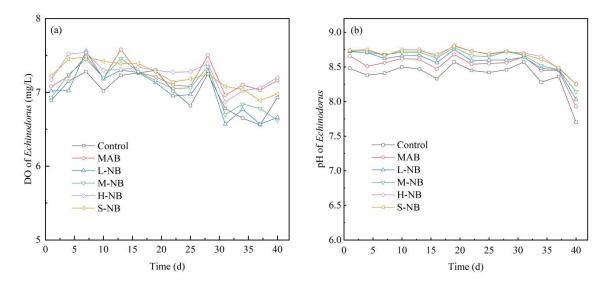


Figure S2. Changes in (a) DO level and (b) pH in the nutrient solution during the 40 days of cultivation of *Echinodorus*. MAB, L-NB, M-NB, H-MB, S-NB represent macrobubble aeration, low, medium, high and super-high nanobubble aeration groups, respectively.

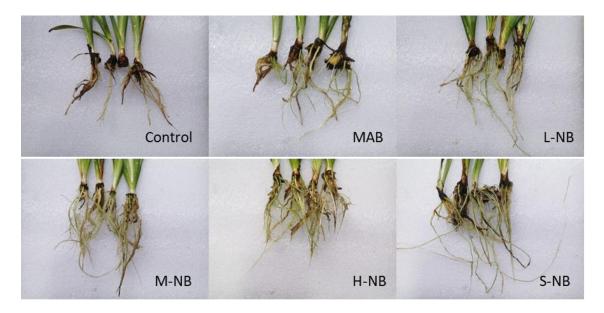


Figure S3. The appearance of *Iris* root at the end of the experiment. MAB, L-NB, M-NB, H-MB, S-NB represent macrobubble aeration, low, medium, high and super-high nanobubble aeration groups, respectively.

Table S1

Significantly enriched Gene Ontology (GO) classification of DEGs in *Iris* for MAB group vs. H-NB group (P-value < 0.05).

	-	-
Up	Down	Term_type
33	12	molecular_function
33	12	molecular_function
28	14	molecular_function
28	11	molecular_function
6	0	molecular_function
5	4	molecular_function
6	1	molecular_function
5	8	molecular_function
12	10	molecular_function
5	0	molecular_function
12	2	molecular_function
4	6	molecular_function
4	4	molecular_function
103	46	molecular_function
2	1	biological_process
7	0	biological_process
7	0	biological_process
7	0	biological_process
	0	biological_process
5	0	biological_process
2	3	biological_process
7	0	biological_process
7	7 0	biological_process
6	0	biological_process
46	29	biological_process
18	5	biological_process
103	38	biological_process
	$ \begin{array}{c} 33\\33\\28\\28\\28\\6\\5\\5\\5\\5\\5\\12\\5\\12\\4\\4\\103\\2\\7\\7\\7\\5\\2\\7\\7\\7\\5\\2\\7\\7\\7\\5\\2\\7\\7\\1\\8\end{array} $	$\begin{array}{cccccccccccccccccccccccccccccccccccc$

Table S2

Significantly enriched Gene Ontology (GO) classification of DEGs in *Echinodorus* for MAB group vs. S-NB group (P-value < 0.05).

Description	Up	Down	Term_type
structural constituent of ribosome	159	127	molecular_function
structural molecule activity	178	197	molecular_function
oxidoreductase activity	133	378	molecular_function
peptidase inhibitor activity	11	35	molecular_function
peptidase regulator activity	11	35	molecular_function
ribosome biogenesis	173	142	biological_process
ribonucleoprotein complex biogenesis	173	142	biological_process
cellular component biogenesis	216	216	biological_process
peptide biosynthetic process	169	164	biological_process
translation	169	158	biological_process
amide biosynthetic process	179	184	biological_process
peptide metabolic process	172	168	biological_process
oxidation-reduction process	131	359	biological_process
organonitrogen compound biosynthetic process	228	282	biological_process
cellular amide metabolic process	183	195	biological_process
cellular component organization or biogenesis	263	336	biological_process
organonitrogen compound metabolic process	314	467	biological_process
metabolic process	794	1580	biological_process
photosynthesis	12	93	biological_process
ribonucleoprotein complex	173	169	cellular_component
ribosome	168	141	cellular_component
macromolecular complex	352	672	cellular_component
thylakoid	14	97	cellular_component
thylakoid part	14	97	cellular_component
photosynthetic membrane	12	92	cellular_component
photosystem	11	86	cellular_component
thylakoid membrane	5	47	cellular_component
photosystem II oxygen evolving complex	3	40	cellular_component
photosystem II	7	63	cellular_component
intracellular non-membrane-bounded organelle	229	249	cellular_component