

REFERENCES

- Abdelahad, N., Bolpagni, R., Lasinio, G. J., Vis, M. L., Amadio, C., Laini, A. & Keil, E. J. 2015. Distribution, morphology and ecological niche of *Batrachospermum* and *Sheathia* species (Batrachospermales, Rhodophyta) in the fontanili of the Po plain (northern Italy). *Eur. J. Phycol.* 50:318-329.
- Carlile, A. L. & Sherwood, A. R. 2013. Phylogenetic affinities and distribution of the Hawaiian freshwater red algae (Rhodophyta). *Phycologia* 52:309-319.
- Chapuis, I. S. 2016. Batrachospermales (Rhodophyta) from the Iberian Peninsula and the Balearic Islands: diversity and phylogeny. Ph.D. dissertation, Universidad de Granada, Granada, 175 pp.
- Chiasson, W. B., Johanson, K. G., Sherwood, A. R. & Vis, M. L. 2007. Phylogenetic affinities of the form taxon *Chantransia pygmaea* (Rhodophyta) specimens from the Hawaiian Islands. *Phycologia* 46:257-262.
- Chou, J. -Y. & Wang, W. -L. 2006. *Batrachospermum arcuatum* Kylin (Batrachospermales, Rhodophyta), a freshwater red alga newly recorded in Taiwan. *Taiwania* 51:58-63.
- Han, J. -F., Nan, F. -R., Feng, J., Lv, J. -P., Liu, Q., Kociolek, J. P. & Xie, S. -L. 2018. *Sheathia jinchengensis* (Batrachospermales, Rhodophyta), a new freshwater red algal species described from North China. *Phytotaxa* 367:63-70.
- Han, J. -F., Nan, F. -R., Feng, J., Lv, J. -P., Liu, Q., Lui, X. -D. & Xie, S. -L. 2019. *Sheathia matouensis* (Batrachospermales, Rhodophyta), a new freshwater red algal species from North China. *Phytotaxa* 415:255-263.
- Ji, L., Feng, J., Chen, L. & Xie, S. 2011. A molecular and morphological investigation of *Batrachospermum arcuatum* (Batrachospermales, Rhodophyta) in China. *Aquat. Bot.* 95:254-257.
- Ji, L., Xie, S. -L., Feng, J., Chien, L. & Wang, J. 2014. Molecular systematics of four endemic Batrachospermaceae (Rhodophyta) species in China with multilocus data. *J. Syst. Evol.* 52:92-100.
- Johnston, E. T., Lim, P. -E., Buhari, N., Keil, E. J., Djawad, M. I. & Vis, M. L. 2014. Diversity of freshwater red algae (Rhodophyta) in Malaysia and Indonesia from morphological and molecular data. *Phycologia* 53:329-341.
- Li, Q., Ji, L. & Xie, S. -L. 2010. Phylogenetic analysis of Batrachospermales (Florideophyceae, Rhodophyta) based on chloroplast *rbcL* sequences. *Acta Hydrobiol. Sin.* 34:20-28.
- Necchi, O. Jr., West, J. A., Ganesan, E. K., Yasmin, F., Rai, S. K. & Rossignolo, N. L. 2019. Diversity of the genus *Sheathia* (Batrachospermales, Rhodophyta) in northeast India and east Nepal. *Algae* 34:277-288.
- Salomaki, E. D., Kwadrans, J., Eloranta, P. & Vis, M. L. 2014. Molecular and morphological evidence for *Sheathia* gen. nov. (Batrachospermales, Rhodophyta) and three new species. *J. Phycol.* 50:526-542.
- Szinte, A. L., Taylor, J. C., Abosedo, A. T. & Vis, M. L. 2020. Current status of the freshwater red algal diversity (Rhodophyta) of the African continent including description of new taxa (Batrachospermales). *Phycologia* 59:187-199.
- Vis, M. L., Feng, J., Chiasson, W. B., Xie, S. -L., Stancheva, R., Entwisle, T. J., Chou, J. -Y. & Wang, W. -L. 2010. Investigation of the molecular and morphological variability in *Batrachospermum arcuatum* (Batrachospermales, Rhodophyta) from geographically distant locations. *Phycologia* 49:545-553.

Supplementary Table S2. Outgroup sequences used for phylogenetic analyses

Taxon	GenBank No.	
	<i>rbcL</i>	COI-5P
Batrachospermales		
<i>Acarposporophycos brasiliensis</i>	FJ386458	KU672388
<i>Batrachospermum gelatinosum</i>	GU810833	KM055327
<i>Kumanoa holtonii</i>	JN590004	JN604921
<i>Kumanoa louisianae</i>	JN590005	JN604924
<i>Kumanoa procarpa</i>	FJ386464	KM055332
<i>Lemanea fluviatilis</i>	KM055243	KM055333
<i>Lemanea fucina</i>	KJ825958	KU672391
<i>Lympha mucosa</i>	KM593865	KM593873
<i>Montagnia australis</i>	EU106056	KT802766
<i>Montagnia macrospora</i>	EU106060	KU672389
<i>Nocturama antipodites</i>	FJ386456	KT802754
<i>Nothocladus diatyches</i>	KT802848	KT802758
<i>Nothocladus theaquus</i>	KT802863	KT802763
<i>Paludicola keratophyta</i>	KJ825960	MN943976
<i>Paludicola leafensis</i>	MN943934	MN943981
<i>Paludicola johnhallii</i>	MN943932	MN943975
<i>Paludicola turfosa</i>	MN943940	MN943985
<i>Paralemanea</i> sp.	GQ285124	KM055335
<i>Petrohua bernabei</i>	AY960690	KM055336
<i>Psilosiphon scoparius</i>	AF029155	KU672392
<i>Sirodotia delicatula</i>	DQ646474	KU672394
<i>Sirodotia huillensis</i>	AF029157	EU636739
<i>Sirodotia suecica</i>	JF344718	KM055337
<i>Torularia atra</i>	KT802842	KT802757
<i>Torularia puiggariana</i>	FJ386462	KT802762
<i>Tuomeya americana</i>	KM055244	KM055330
<i>Virescentia viride-americana</i>	AF244112	KM055328
<i>Virescentia viride-brasiliensis</i>	KM078048	KM259993
<i>Visia cayennensis</i>	AY423393	KM055326
<i>Visia</i> sp.	KM055245	KM055325
<i>Volatus carrioni</i>	KM593861	KM593866
Thoreales		
<i>Nemalionopsis shawii</i>	KM005141	KM055241
<i>Thorea reikei</i>	KM005140	KM055239

COI-5P, 5' region of the cytochrome c oxidase subunit I.

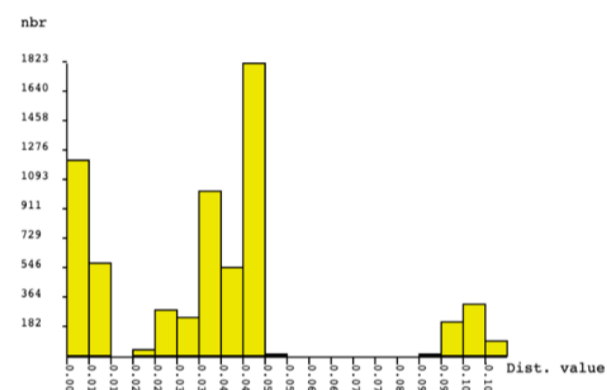
Partition with prior maximal distance $p = 7.74e-03$ Distance Simple Dist MinSlope = 1.500000

Group	No. of specimens	Species name	GenBank accessions
1	3	<i>S. abscondita</i>	MT441833, DQ393129, KC596161
2	6	<i>S. californica</i>	MT441834, MT441835, MT441836, MT441837, MT441838, MT441839
3	56	<i>S. dispersa</i> / <i>S. murphyi</i>	MN487058, MN487059, KF557553, KF557554, KC596123, KC596124, KC596125, KC596126, KC596127, KC596128, KC596129, KC596130, KC596131, KC596132, KC596133, KC596134, KC596135, KC596136, KC596137, KC596138, KC596139, KC596140, KC596141, KC596142, KC596143, KC596144, KC596145, KC596146, KC596147, KC596148, KC596149, KC596150, KC596151, KC596152, KC596153, KC596154, KC596155, KC596156, KC596157, KC596158, KC596159, KC596160, EF116869, EF116870, EF116871, EF116872, EF116873, EF116874, EF116875, EF116876, EF116877, DQ141320, GU457350, GU457349, GU457348, MN974517
4	18	<i>S. arcuata</i>	DQ393131, JX669741, JX669779, KF186240, KM077035, KM077036, KM077037, KM077038, KM077039, KM077040, KM077041, KM077042, KM593804, KM593810, KM593811, KM593812, KM593818, KM593824
5	13	<i>S. longipedicellata</i>	GU457346, GU457347, HQ677186, JN086517, JN086518, JN086519, JN086520, JN086521, JN086522, HQ677187, MH356749, HQ677188, MK746107
6	2	<i>S. assamica</i>	MN481450, MN481451
7	2	<i>S. indonepalensis</i>	MN481449, MN487060
8	12	<i>S. plantuloides</i>	MT441840, MT441841, MT441842, MT441843, MT441844, MT441845, MT441846, MT441847, MT441848, MT441849, MT441850, MT441851
9	2	<i>S. transpacific</i>	AY297054, DQ393130

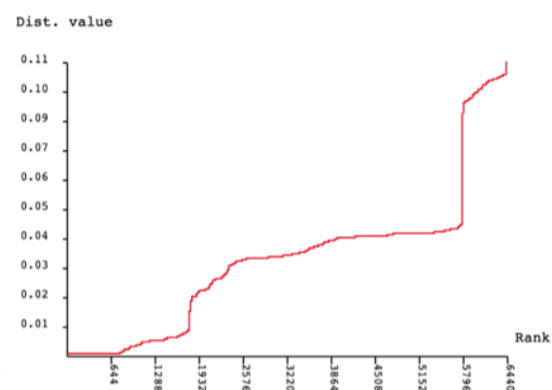
ABGD Web results using Simple Dist mesure of distance Left click **here** to save matrix distance file

Data: Sheathia_rbcL_022520.fasta

Histogram of distances **[save]**



Ranked distances **[save]**



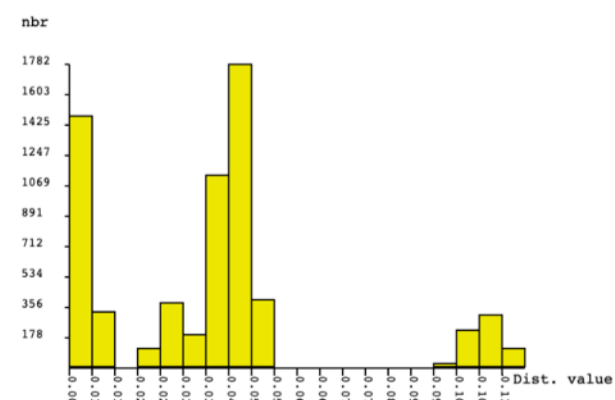
Partition with prior maximal distance $p = 2.78e-03$ Distance JC69 Jukes-Cantor MinSlope = 1.500000

Group	No. of specimens	Species name	GenBank accessions
1	3	<i>S. abscondita</i>	MT441833, DQ393129, KC596161
2	18	<i>S. arcuata</i>	DQ393131, JX669741, JX669779, KF186240, KM077035, KM077036, KM077037, KM077038, KM077039, KM077040, KM077041, KM077042, KM593804, KM593810, KM593811, KM593812, KM593818, KM593824
3	13	<i>S. longipedicellata</i>	GU457346, GU457347, HQ677186, JN086517, JN086518, JN086519, JN086520, JN086521, JN086522, HQ677187, MH356749, HQ677188, MK746107
4	2	<i>S. assamica</i>	MN481450, MN481451
5	6	<i>S. californica</i>	MT441834, MT441835, MT441836, MT441837, MT441838, MT441839
6	4	<i>S. dispersa</i>	DQ141320, GU457348, GU457349, GU457350
7	2	<i>S. indonepalensis</i>	MN481449, MN487060
8	1	<i>S. murphyi</i>	MN974517
9	12	<i>S. plantuloides</i>	MT441840, MT441841, MT441842, MT441843, MT441844, MT441845, MT441846, MT441847, MT441848, MT441849, MT441850, MT441851
10	2	<i>S. transpacific</i>	AY297054, DQ393130
11	35	<i>S. dispersa</i>	EF116869, EF116870, EF116871, EF116872, EF116874, EF116875, KC596123, KC596124, KC596125, KC596126, KC596127, KC596128, KC596129, KC596130, KC596133, KC596136, KC596139, KC596140, KC596141, KC596142, KC596143, KC596144, KC596146, KC596147, KC596148, KC596149, KC596150, KC596151, KC596152, KC596153, KC596154, KC596155, KC596156, KC596158, KC596159
12	11	-	EF116873, KC596131, KC596132, KC596134, KC596135, KC596137, KC596138, KC596145, KC596157, KF557553, KF557554
13	3	-	EF116876, EF116877, KC596160
14	2	-	MN487058, MN487059

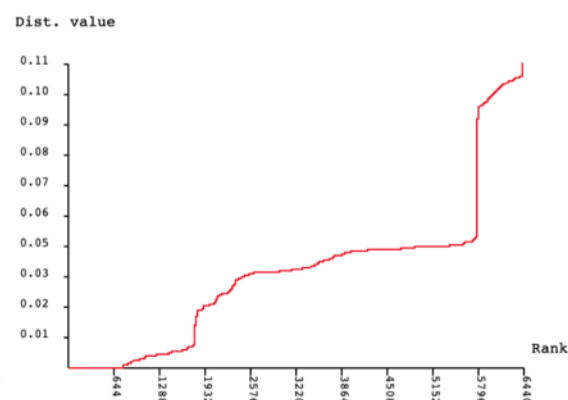
ABGD Web results using JC69 Jukes-Cantor mesure of distance Left click **here** to save matrix distance file

Data: Sheathia_rbcL_022520.fasta

Histogram of distances **[save]**



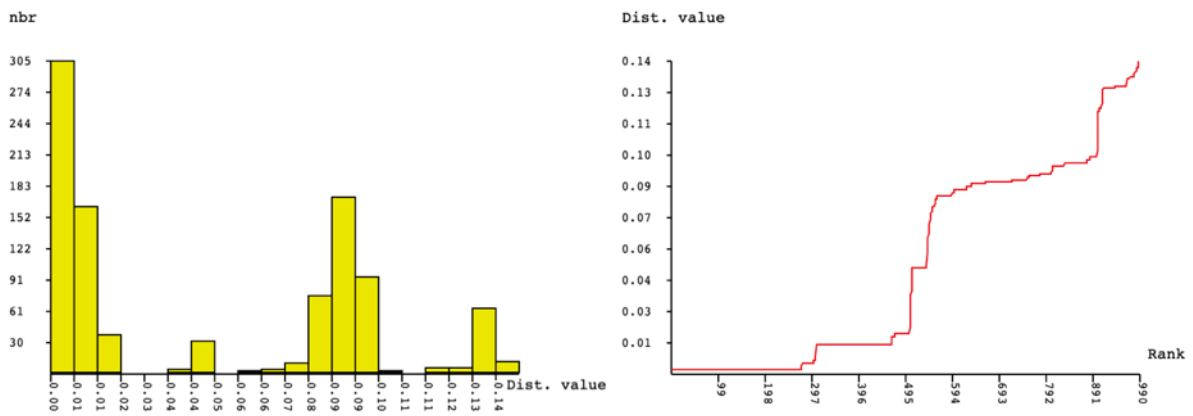
Ranked distances **[save]**



Supplementary Fig. S1. Results from Automatic Barcode Gap Discovery (ABGD) using all *rbcL* sequences for *Sheathia* species with homocortication in Supplementary Table S1.

Initial Partition with prior maximal distance $p = 1.00e-03$ Distance Simple Dist MinSlope = 1.500000

Group	No. of specimens	Species name	GenBank accessions
1	4	<i>S. arcuata</i>	KM592945, KM592946, KM592947
2	2	<i>S. californica</i>	MT441854, MT441855
3	24	<i>S. dispersa</i>	JX669681, KC596286, KC596287, KC596288, KC596289, KC596290, KC596291, KC596292, KC596293, KC596294, KC596297, KC596300, KC596303, KC596304, KC596305, KC596307, KC596308, KC596309, KC596310, KC596311, KC596312, KC596313, KC596314, KC596315
4	7	<i>S. dispersa</i>	KC596295, KC596296, KC596298, KC596299, KC596301, KC596302, KC596306
5	1	<i>S. dispersa</i>	KC596316
6	1	<i>S. abscondita</i>	MT441852
7	1	<i>S. transpacifica</i>	MT441857
8	1	<i>S. plantuloides</i>	MT441856
9	3	<i>S. longipedicellata</i>	KC511071, KC511072, KC511075
10	1	<i>S. murphyi</i>	MN974522

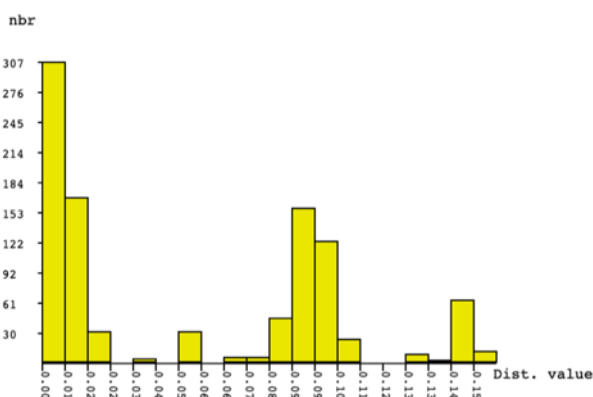


Partition with prior maximal distance $p = 2.78e-03$ Distance JC69 Jukes-Cantor MinSlope = 1.500000

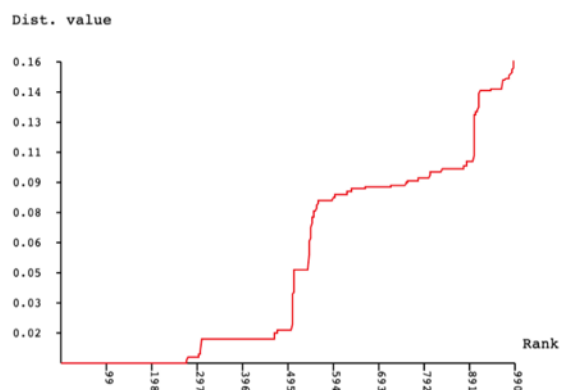
Group	No. of specimens	Species name	GenBank accessions
1	4	<i>S. arcuata</i>	KM592945, KM592946, KM592947
2	2	<i>S. californica</i>	MT441854, MT441855
3	1	<i>S. dispersa</i>	JX669681
4	7	<i>S. dispersa</i>	KC596295, KC596296, KC596298, KC596299, KC596301, KC596302, KC596306
5	1	<i>S. dispersa</i>	KC596316
6	1	<i>S. abscondita</i>	MT441852
7	1	<i>S. transpacifica</i>	MT441857
8	1	<i>S. plantuloides</i>	MT441856
9	3	<i>S. longipedicellata</i>	KC511071, KC511072, KC511075
10	1	<i>S. murphyi</i>	MN974522
11	23	<i>S. dispersa</i>	KC596286, KC596287, KC596288, KC596289, KC596290, KC596291, KC596292, KC596293, KC596294, KC596297, KC596300, KC596303, KC596304, KC596305, KC596307, KC596308, KC596309, KC596310, KC596311, KC596312, KC596313, KC596314, KC596315

Data: Sheathia_COI.fasta

Histogram of distances [save]



Ranked distances [save]



Supplementary Fig. S2. Results from Automatic Barcode Gap Discovery (ABGD) using all 5' region of the cytochrome c oxidase subunit I (COI-5P) sequences for *Sheathia* species with homocortication in Supplementary Table S1.