

Supplementary Materials: Further Understanding of Degradation Pathways of Microcystin-LR by an Indigenous *Shingopyxis* sp. in Environmentally Relevant Pollution Concentrations

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Table S1. Specific primers sequences.

Primers	Sequence (5' to 3')
<i>mlrA</i> -F	GACCCGATGTTCAAGATACT
<i>mlrA</i> -R	CTCCTCCCACAAATCAGGAC
<i>mlrB</i> -F	CTCGATGCGGTATTTGCTG
<i>mlrB</i> -R	TCCAACGACCATCCCTTCTG
<i>mlrC</i> -F	CGAAGGCGAAAGGTGCAAC
<i>mlrC</i> -R	GAGCGCTTGTGATAGTGACG
<i>mlrD</i> -F	GTTCTCGGCCTAGCCT
<i>mlrD</i> -R	GCGACGAAGATCGTTGCT
16S-F-real	AGGATGAGCCCGCGTAAGAT
16S-R-real	TCAGTCCCAGTGTGGCTGATC
<i>mlrA</i> -F-real	GCACCATCATCGCTGTCTT
<i>mlrA</i> -R-real	ACCGTTGGAGCCCATTCCG
<i>mlrB</i> -F-real	CTATCAGGGCGGCTTTGGA
<i>mlrB</i> -R-real	GAGAGCGGCCGTGAACTG
<i>mlrC</i> -F-real	CAGCGTCGATCGCACAAAG
<i>mlrC</i> -R-real	CGCTCTGGGCCGTTACC
<i>mlrD</i> -F-real	ACGCCATCTTCTGCCTCC
<i>mlrD</i> -R-real	CGACCCGCAAGACGATTA

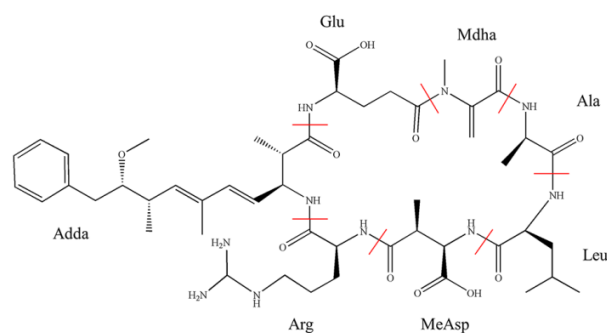


Figure S1. Chemical structure of MC-LR, cyclo-(Ala-Leu-MeAsp-Arg-Adda-Glu-Mdha).

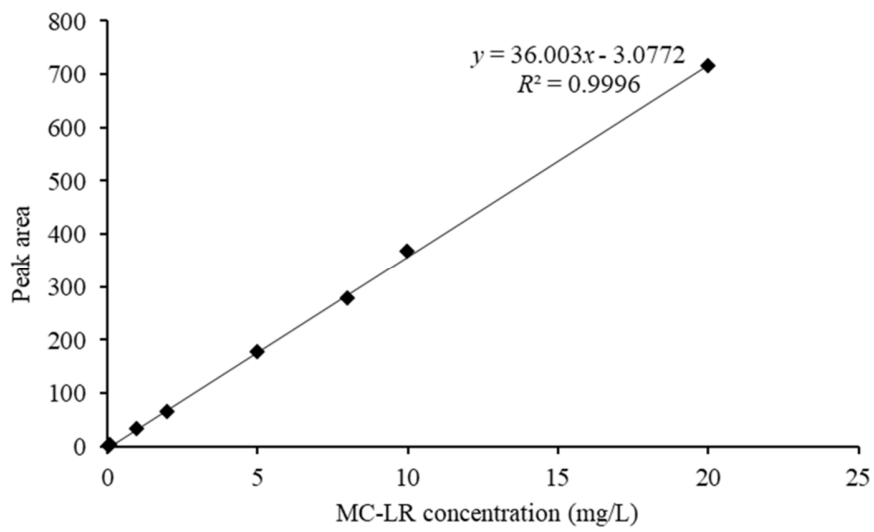


Figure S2. The standard curve of MC-LR quantitated by HPLC (50 µg/L–20 mg/L).

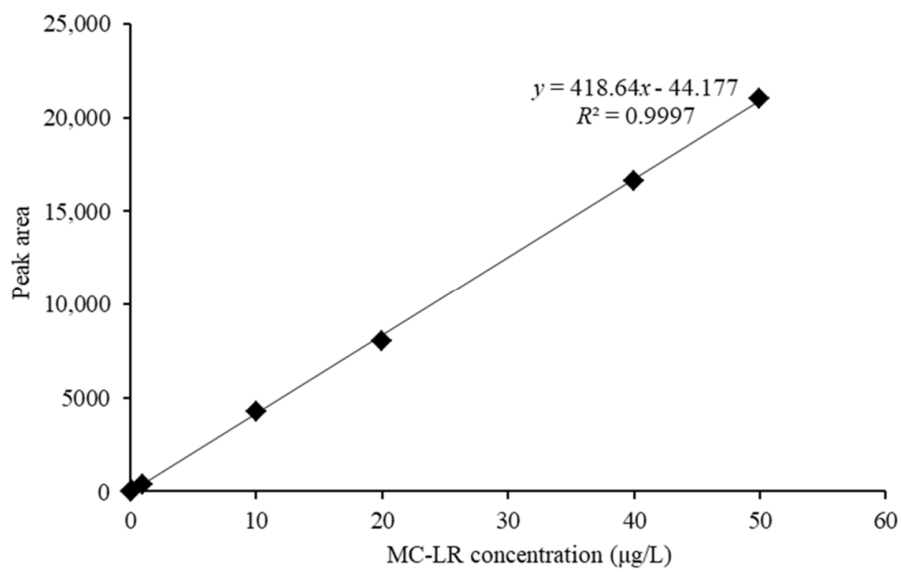


Figure S3. The standard curve of MC-LR quantitated by UPLC-MS/MS (0.01 µg/L–50µg/L).

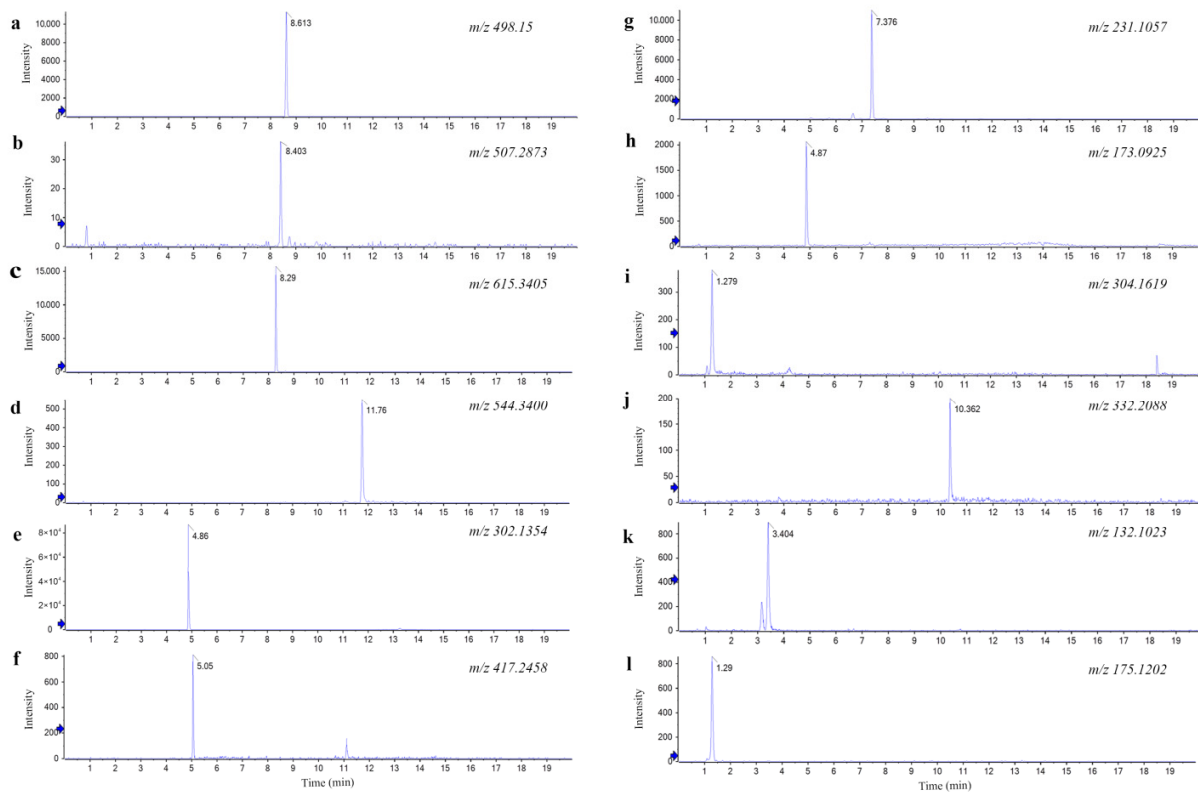


Figure S4. Chromatograms of MC-LR and its degradation products (a) MC-LR, (b) linearized MC-LR, (c) tetrapeptide, (d) Adda-Glu-Mdha, (e) Glu-Mdha-Ala, (f) Leu-MeAsp-Arg, (g) Glu-Mdha, (h) Mdha-Ala, (i) MeAsp-Arg, (j) Adda, (k) Leu, (l) Arg.

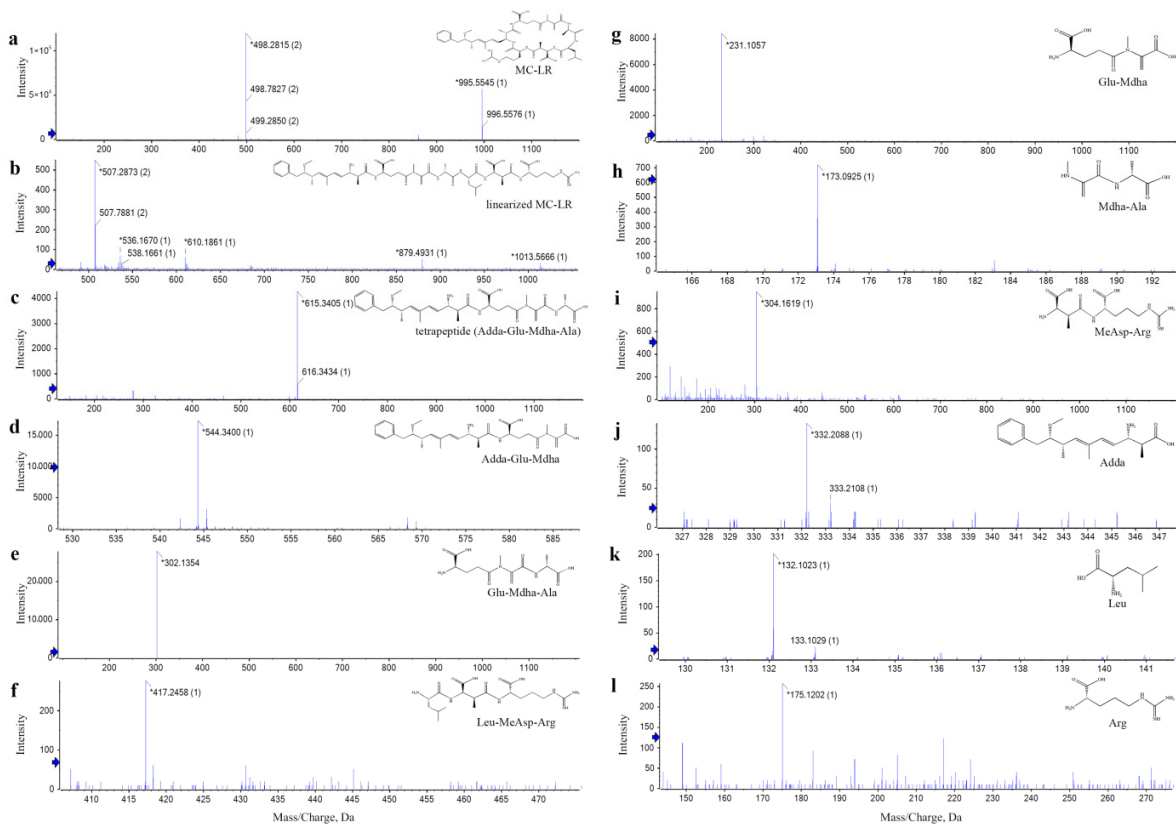


Figure S5. Mass spectrum of MC-LR and its degradation products (a) MC-LR, (b) linearized MC-LR, (c) tetrapeptide, (d) Adda-Glu-Mdha, (e) Glu-Mdha-Ala, (f) Leu-MeAsp-Arg, (g) Glu-Mdha, (h) Mdha-Ala, (i) MeAsp-Arg, (j) Adda, (k) Leu, (l) Arg.

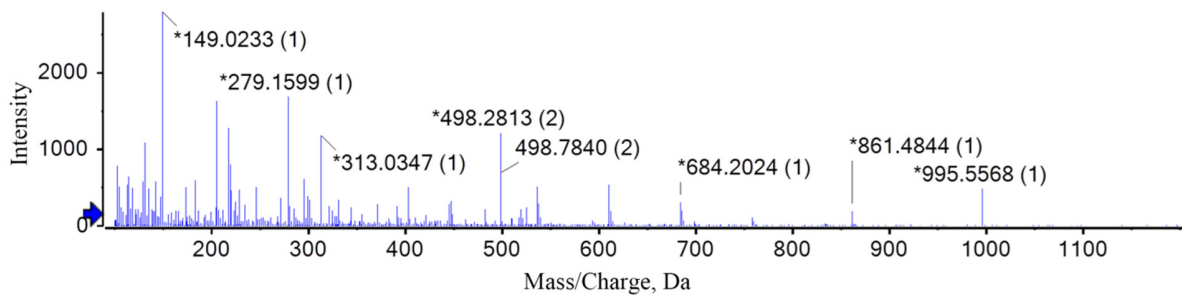


Figure S6. Fragment ions of standard MC-LR detected by UPLC-MS/MS.