

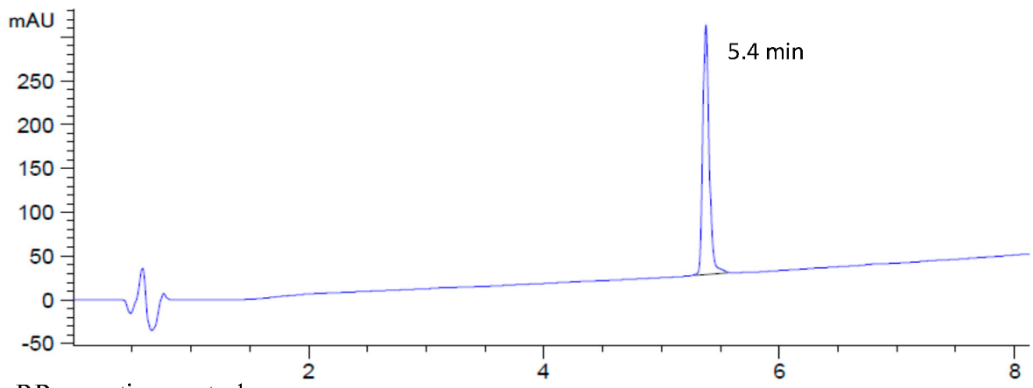
# Supplementary Materials: Characterization of Enzymatic Activity of MlrB and MlrC Proteins Involved in Bacterial Degradation of Cyanotoxins Microcystins

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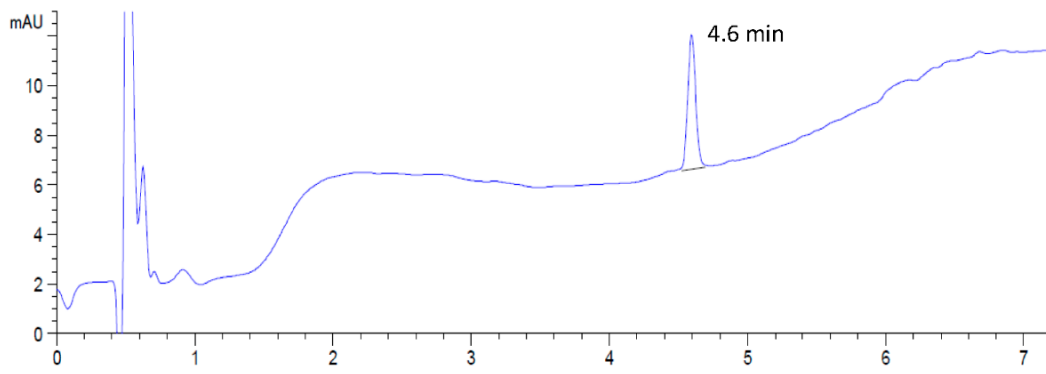


**Figure S1.** Alignment of MlrB protein amino acid sequences from bacterial strains of *Sphingomonaceae* family (AAL10287—*Sphingomonas* sp. ACM-3962 (Bourne, 2001), BAI47771—*Sphingopyxis* sp. C-1, AGG86526—*Sphingomonas* sp. USTB-05, KR150744—*Sphingomonas* sp. ACM-3962) verified in the present study.

acLR\_negative control



acRR\_negative control



acRR\_MlrC

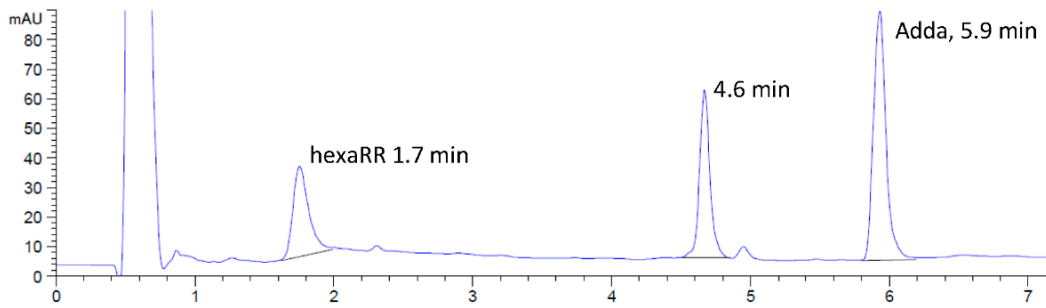
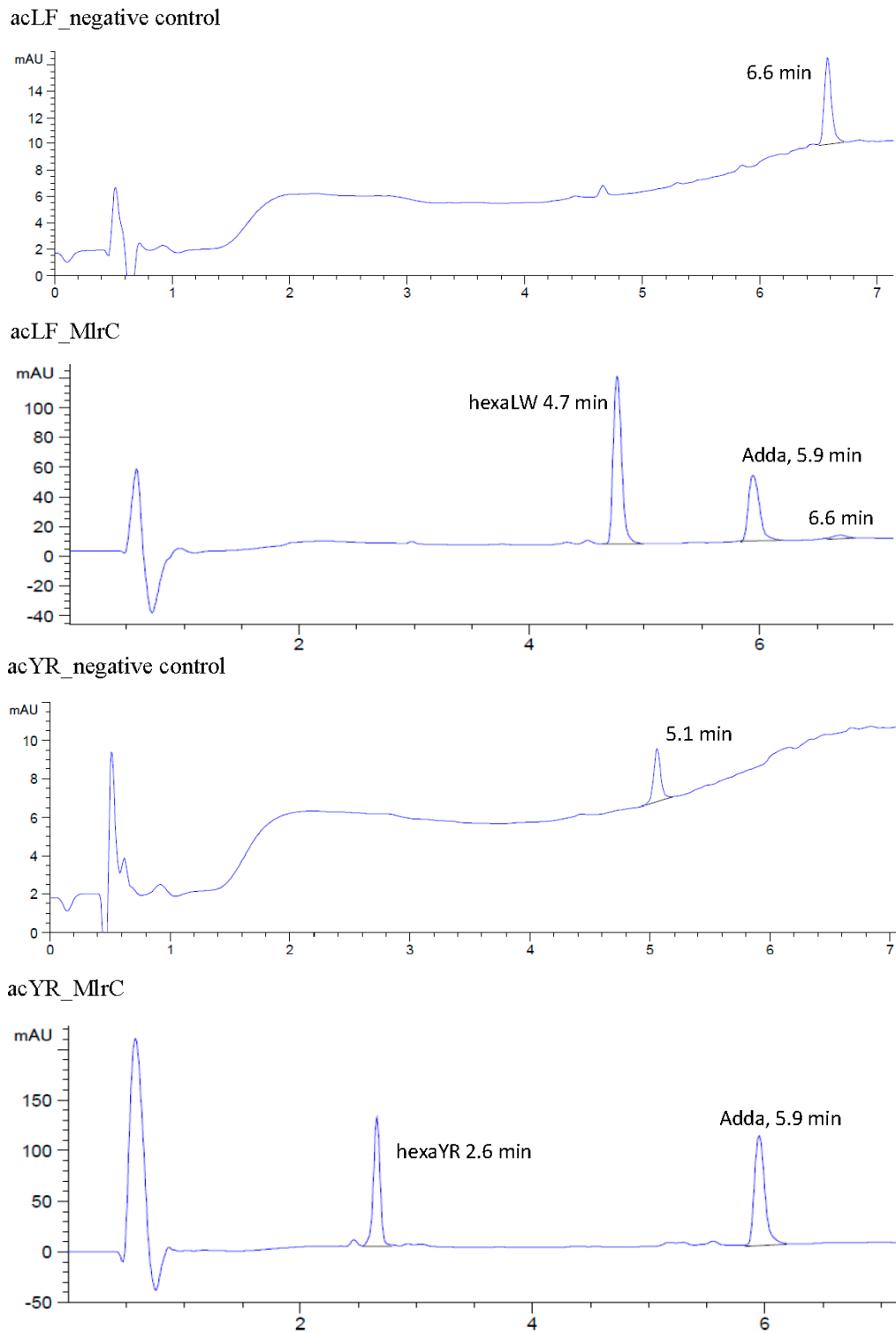


Figure S2. Cont.



**Figure S2.** The examples of chromatograms indicating the formation of hexapeptides from different linearized MC variants by MlrC and the lack of degradation of linearized variants in the presence of extract of *E. coli* with empty plasmid pET21a (negative controls).

hexaMC-LF:

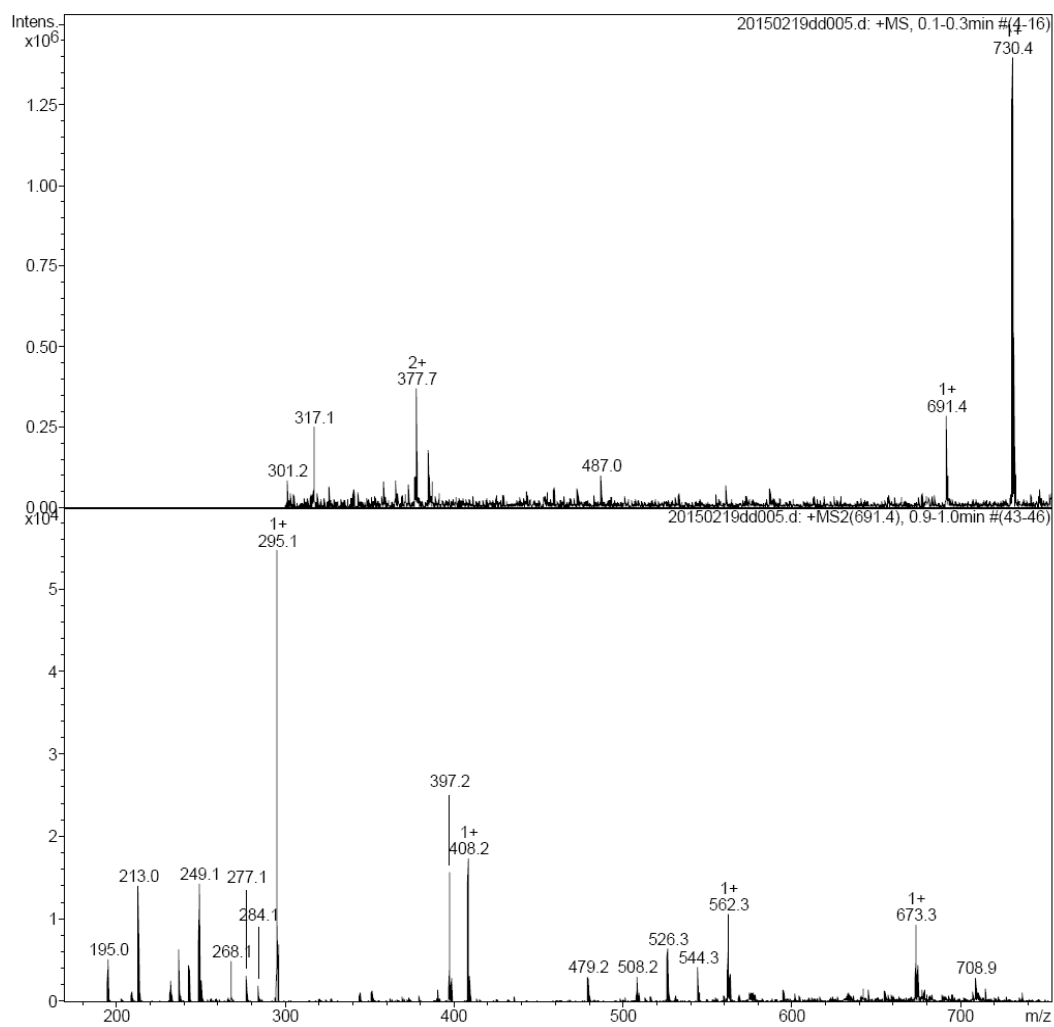


Figure S3. Cont.

hexaMC-LW:

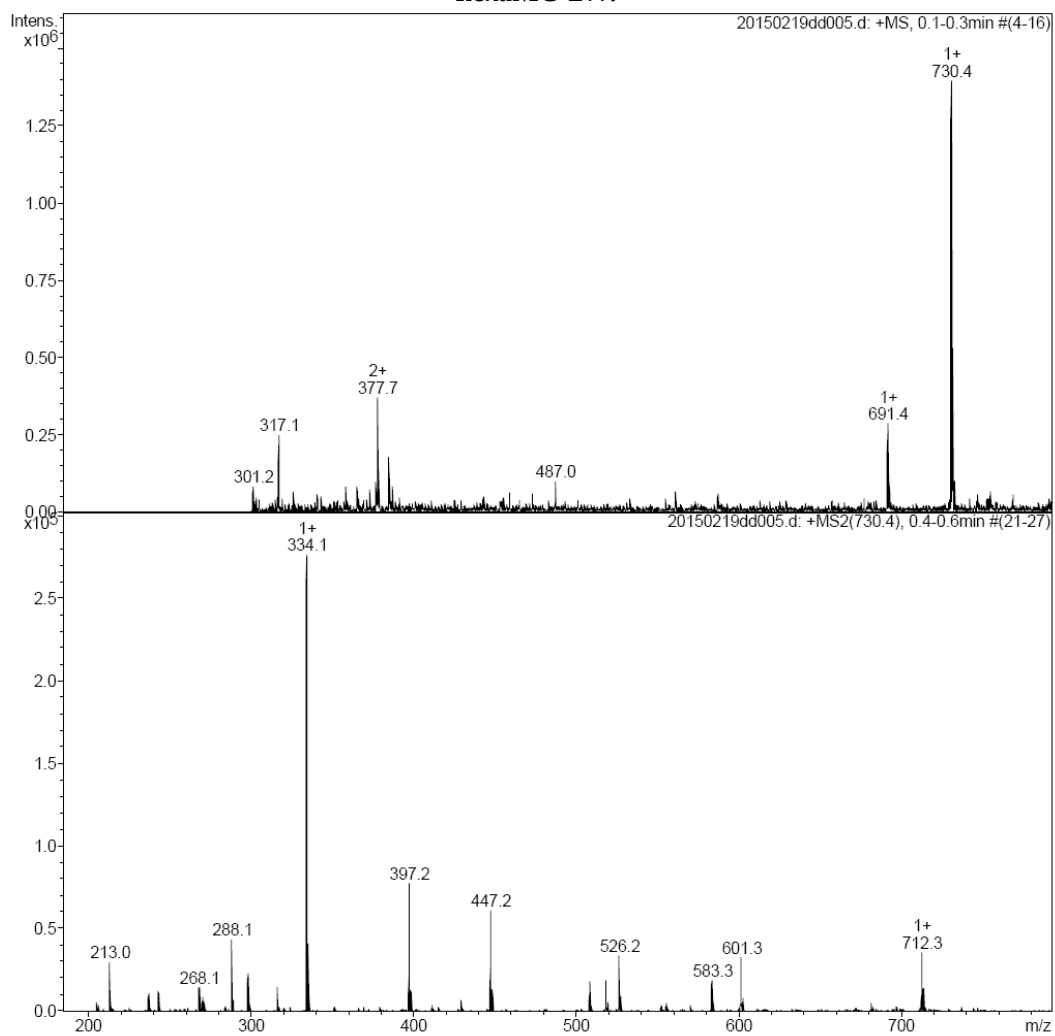


Figure S3. Cont.

hexaMC-LY:

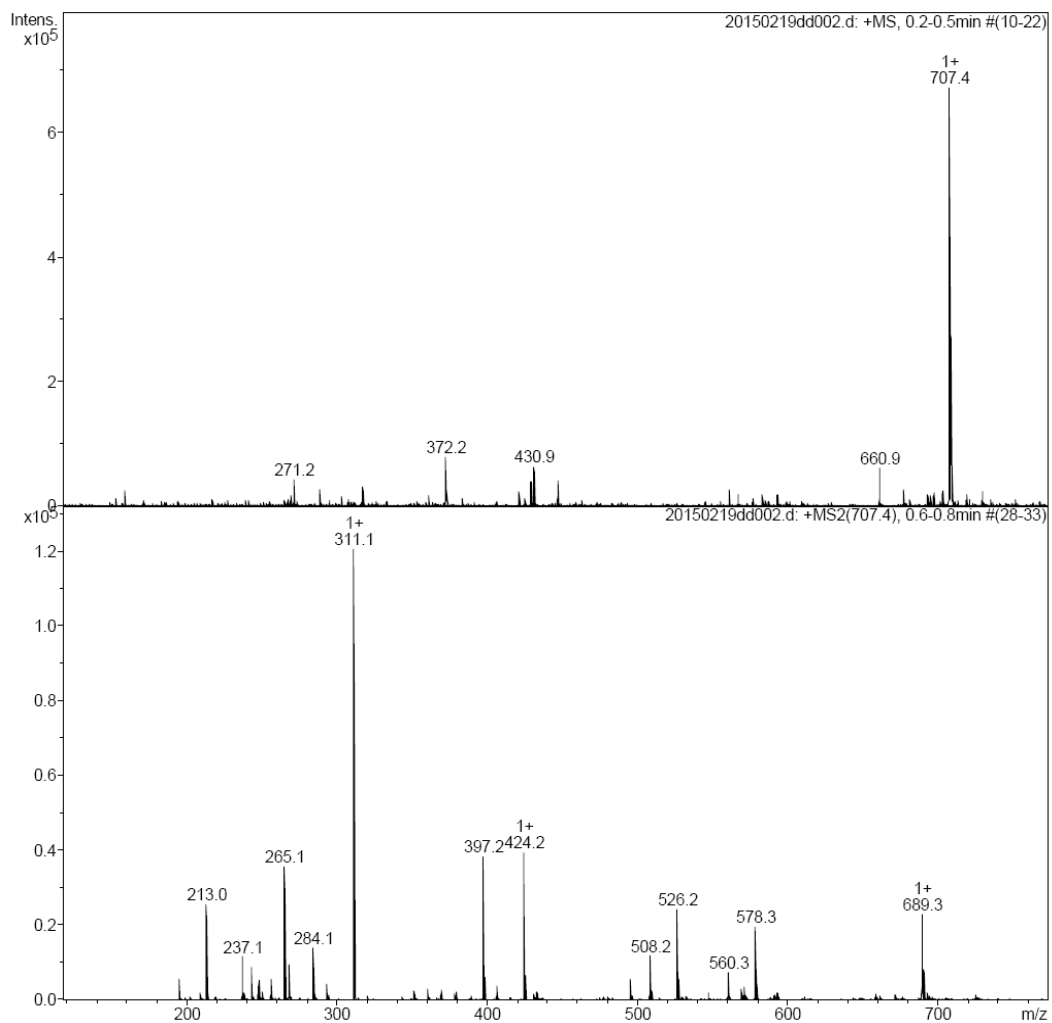


Figure S3. Cont.

hexaMC-RR:

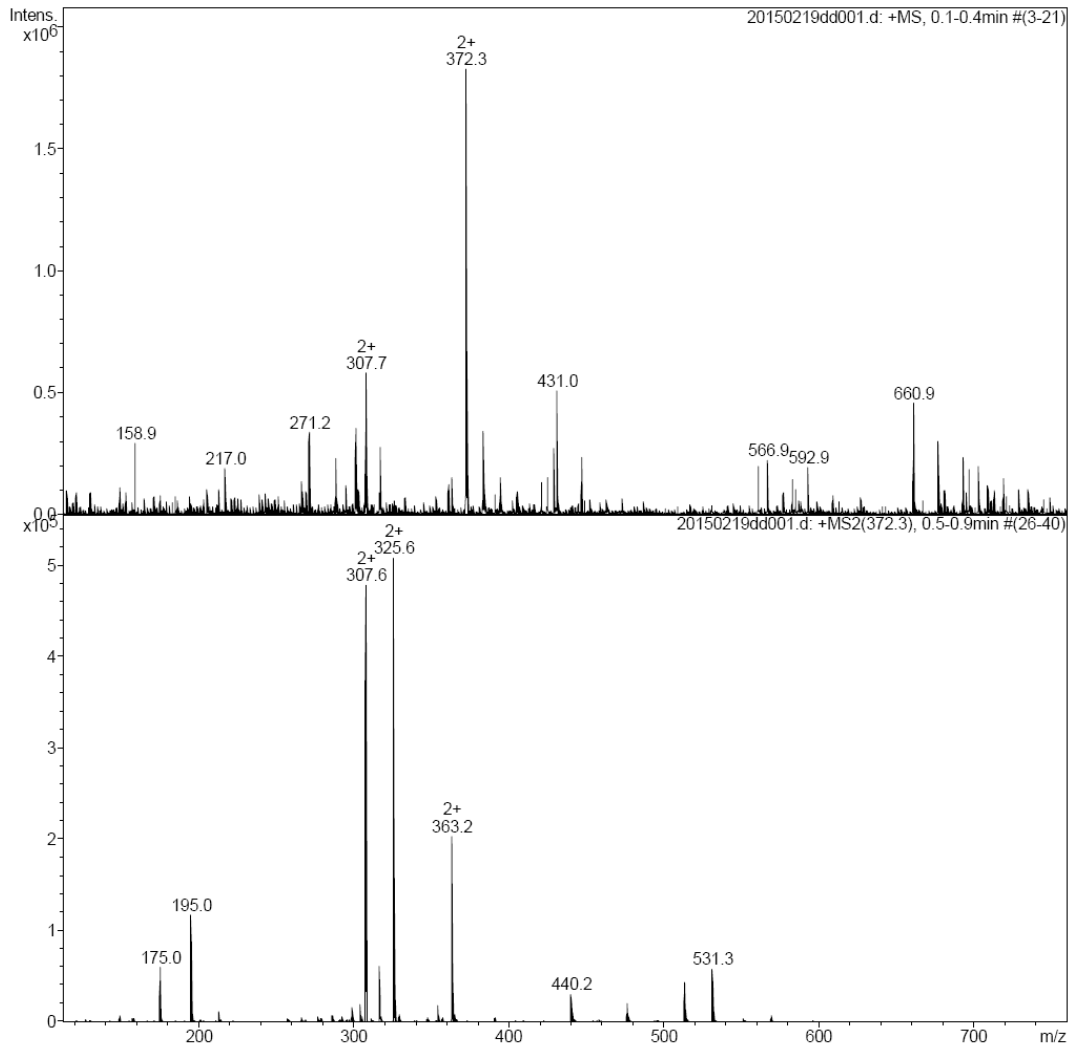
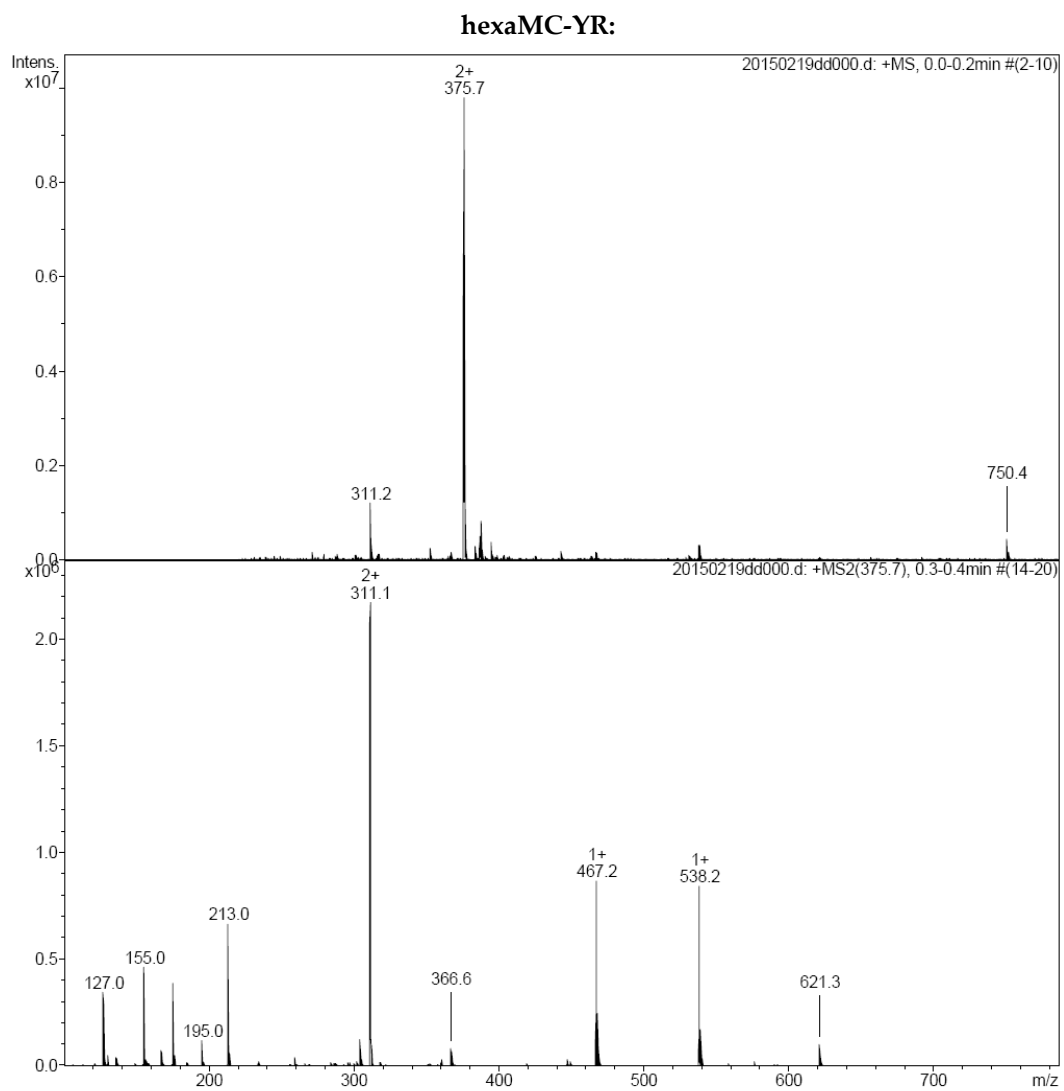


Figure S3. Cont.



**Figure S3.** MS<sup>2</sup> of the hexapeptides originated from other MC variants.