

Supplementary Material: Eighteen new aeruginosamide variants produced by the Baltic cyanobacterium *Limnraphis CCNP1324*

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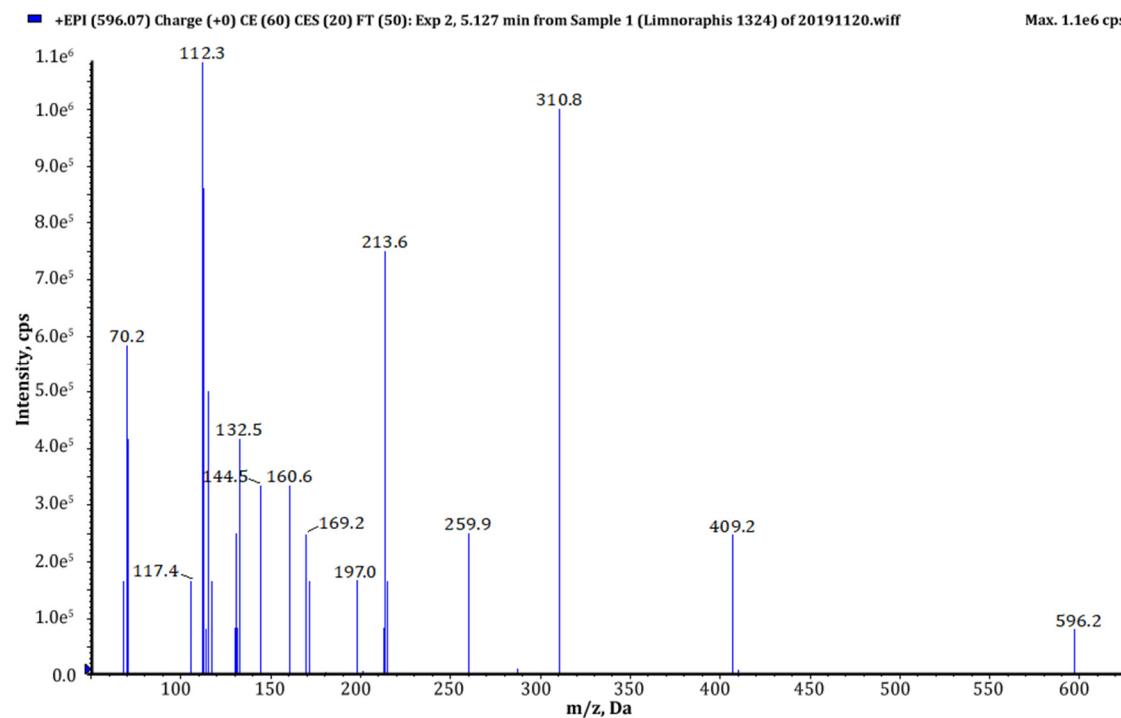


Figure S1. Enhanced product ion mass spectrum of aeruginosamide AER595 with general structure 187+Val+Pro+Pyr+TzlCOOMe identified based on the following fragment ions: 596 [M+H], 409 [Val+Pro+Pyr+TzlCOOMe], 310 [Pro+Pyr+TzlCOOMe+H], 259 [187+Val+H-CO], 213 [Pyr+TzlCOOMe+H], 197 [Val+Pro+H], 169 [Val+Pro+H-CO], 160 [187+H-CO], 144 [TzlCOOMe], 112 TzlCO, 70 Pro immonium ion.

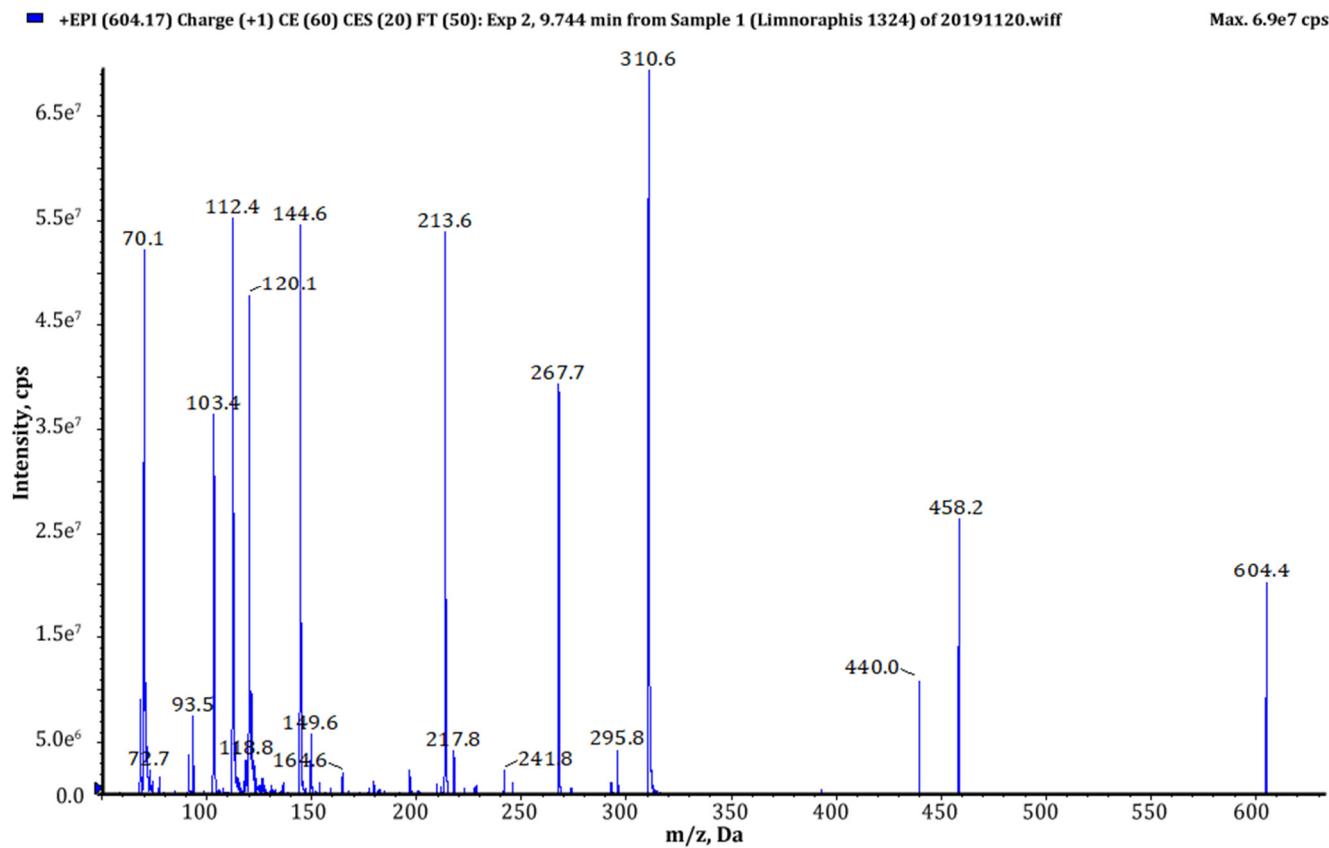


Figure S2. Enhanced product ion mass spectrum of aeruginosamide AEG603 with general structure Phe+Phe+Pro+Pyr+TzlCOOMe identified based on the following fragment ions: 604 [M+H], 458 [M+H-Phe], 310 [Pro+Pyr+TzlCOOMe+H], 295 [Phe+Phe+H], 267 [Phe+Phe+H-CO], 217 [Phe+Pro+H-CO], 213 [Pyr+TzlCOOMe+H], 144 [TzlCOOMe], 120 Phe immonium ion, 112 TzlCO, 70 Pro immonium ion.

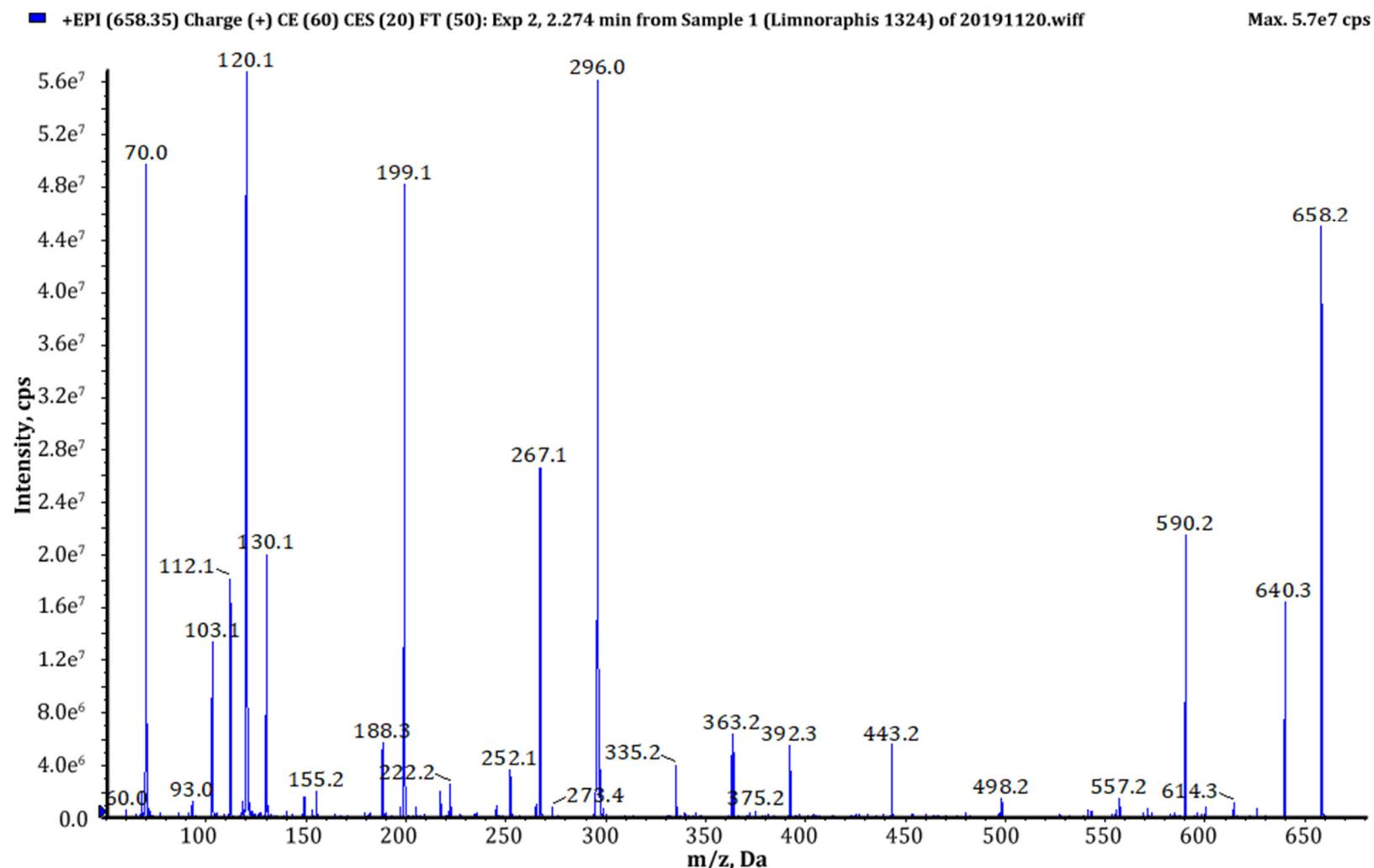


Figure S3. Enhanced product ion mass spectrum of aeruginosamide AEG657 with general structure Pre+Phe+Phe+Pro+Pyr+TzlCOOH identified based on the following fragment ions: 658 [M+H], 640 [M+H-H₂O], 590 [M+H-Pre], 443 [M+H-(Pre+Phe)], 392 [Phe+Phe+Pro+H], 363 [Pre+Phe+Phe+H], 335 [Pre+Phe+Phe+H-CO], 296 [Pro+Pyr+TzlCOOH], 267 [Phe+Phe+H-CO], 199 [Pyr+TzlCOOH+H], 188 [Pre+Phe+H-CO], 130 [TzlCOOH], 120 Phe immonium ion, 112 TzlCO, 70 Pro immonium ion.

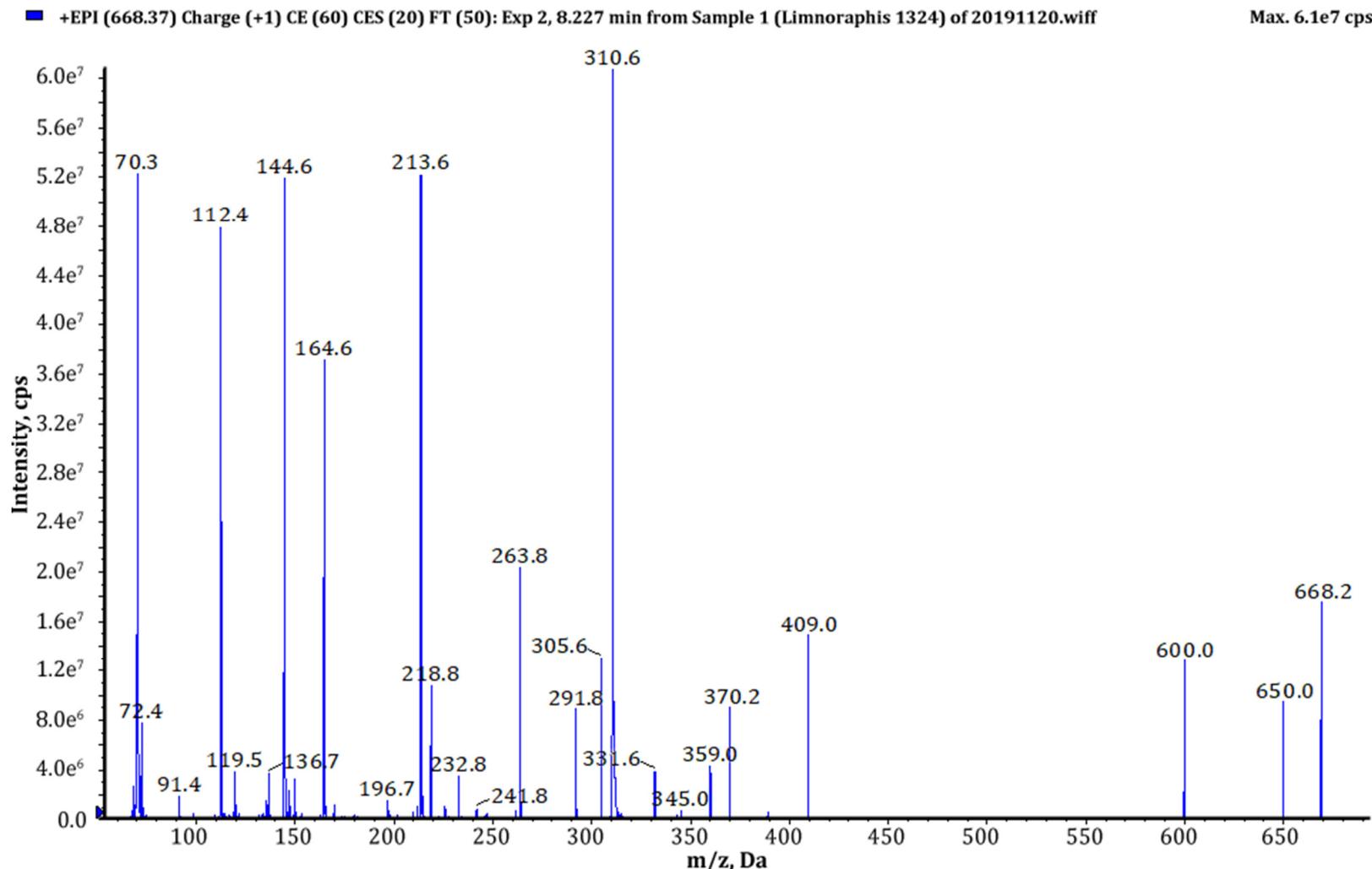


Figure S4. Enhanced product ion mass spectrum of aeruginosamide AER667 with general structure Pre+MeHty+Val+Pro+Pyr+TzlCOOMe identified based on the following fragment ions: 668 [M+H], 650 [M+H-H₂O], 600 [M+H-Pre], 409 [Val+Pro+Pyr+TzlCOOMe], 370 [MeHTyr+Val+Pro+H-H₂O], 359 [Pre+MeHTyr+Val+H], 331 [Pre+MeHTyr+Val+H-CO], 310 [Pro+Pyr+TzlCOOMe+H], 291 [MeHTyr+Val+H], 263 [MeHTyr+Val+H-CO], 213 [Pyr+TzlCOOMe+H], 164 MeHTyr immonium ion, 144 [TzlCOOMe], 112 TzlCO, 72 Val immonium ion, 70 Pro immonium ion.

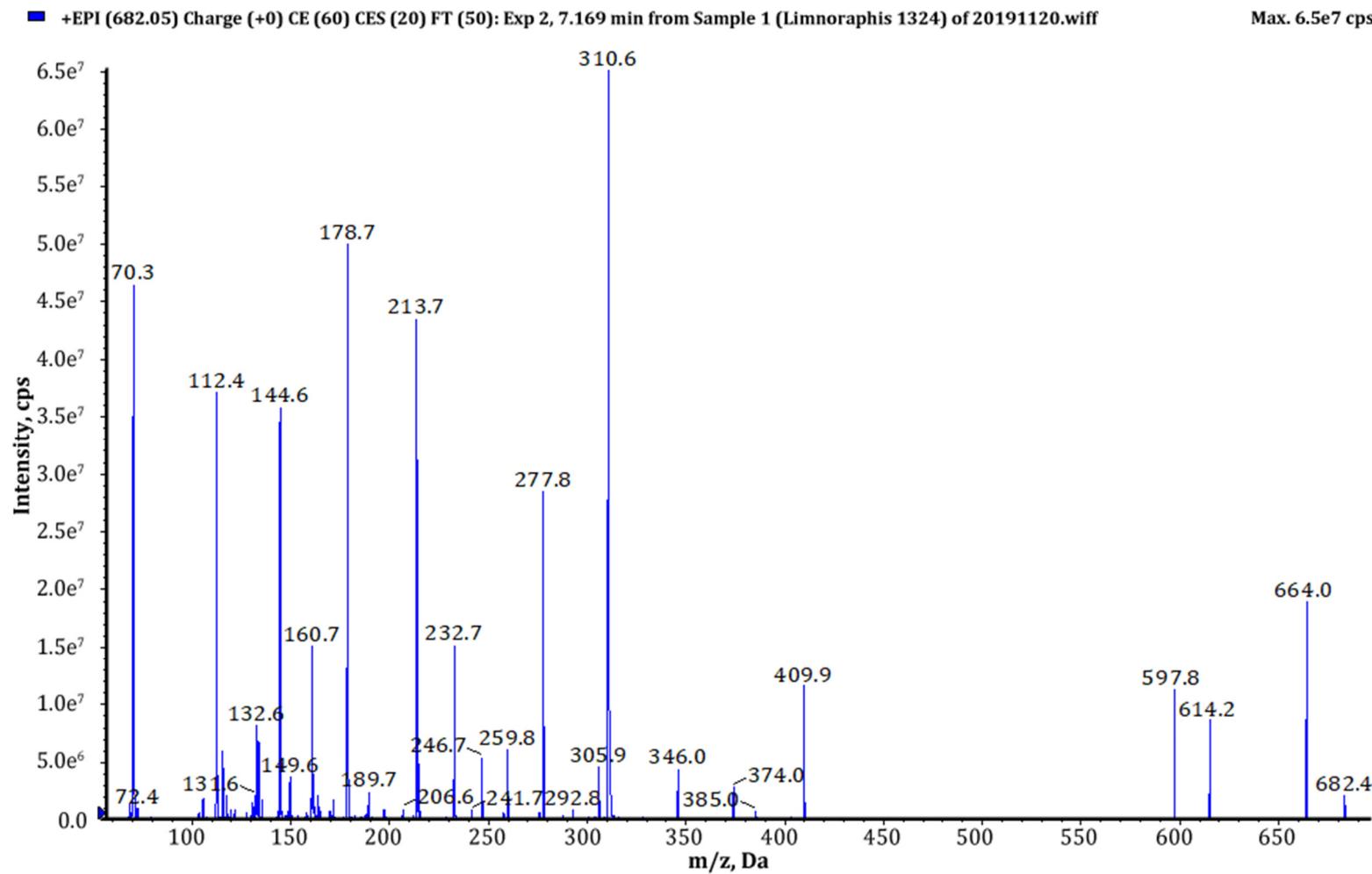


Figure S5. Enhanced product ion mass spectrum of aeruginosamide AER681a with general structure Pre+205+Val+Pro+Pyr+TzlCOOMe identified based on the following fragment ions: 682 [M+H], 664 [M+H-H₂O], 614 [M+H-Pre], 409 [M+H-(Pre+205)], 374 [205+Val+Pro+H-CO], 310 [Pro+Pyr+TzlOMe+H], 305 [205+Val+H], 277 [205+Val+H-CO], 259 [205+Val+H-CO-H₂O], 213 [Pyr+TzlCOOMe+H], 178 [205+H-CO], 144 [TzlCOOMe], 112 TzlCO, 72 Val immonium ion, 70 Pro immonium ion.

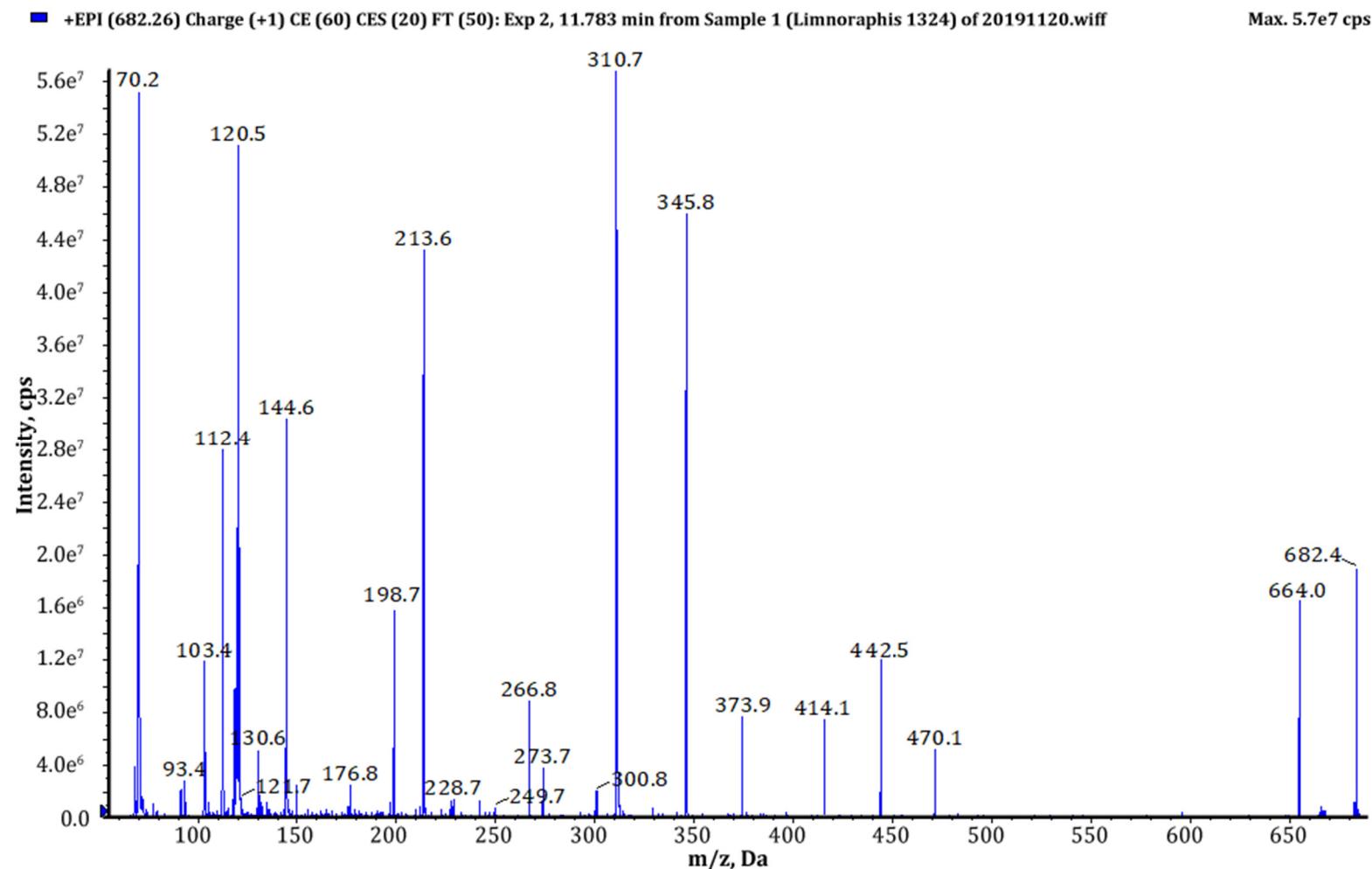


Figure S6. Enhanced product ion mass spectrum of aeruginosamide AER681b with general structure 225+Phe+Pro+Pyr+TzlCOOMe identified based on the following fragment ions: 682 [M+H], 664 [M+H-H₂O], 470 [225+Phe+Pro+H], 442 [225+Phe+Pro+H-CO], 373 [225+Phe+H], 345 [225+Phe+H-CO], 310 [Pro+Pyr+TzlCOOMe+H], 228 [Phe+187+H-CO], 213 [Pyr+TzlCOOMe+H], 198 [225+H-CO], 144 [TzlCOOMe], 120 Phe immonium ion, 112 TzlCO, 70 Pro immonium ion.

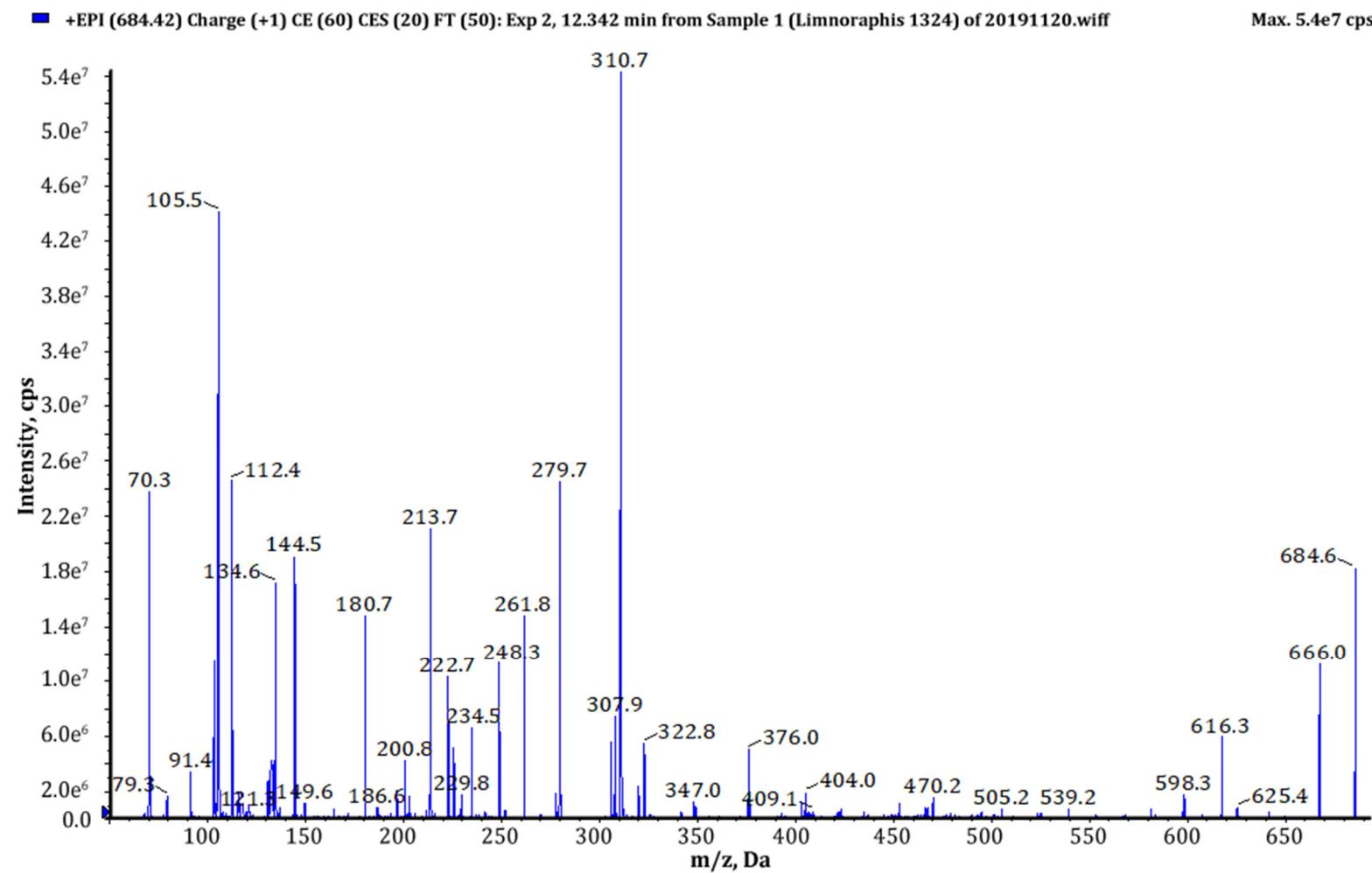


Figure S7. Enhanced product ion mass spectrum of aeruginosamide AER683 with general structure Pre+207+Val+Pro+Pyr+TzlCOOMe identified based on the following fragment ions: 684 [M+H], 666 [M+H-H₂O], 616 [M+H-Pre], 598 [M+H-Pre-H₂O], 409 [Val+Pro+Pyr+TzlCOOMe], 404 [207+Val+Pro+H], 376 [207+Val+Pro+H-CO], 310 [Pro+Pyr+TzlCOOMe+H], 307 [207+Val+H], 279 [207+Val+H-CO], 261 [207+Val+H-CO-H₂O], 213 [Pyr+TzlCOOMe+H], 180 [207+H-CO], 144 [TzlCOOMe], 112 TzlCO, 70 Pro immonium ion.

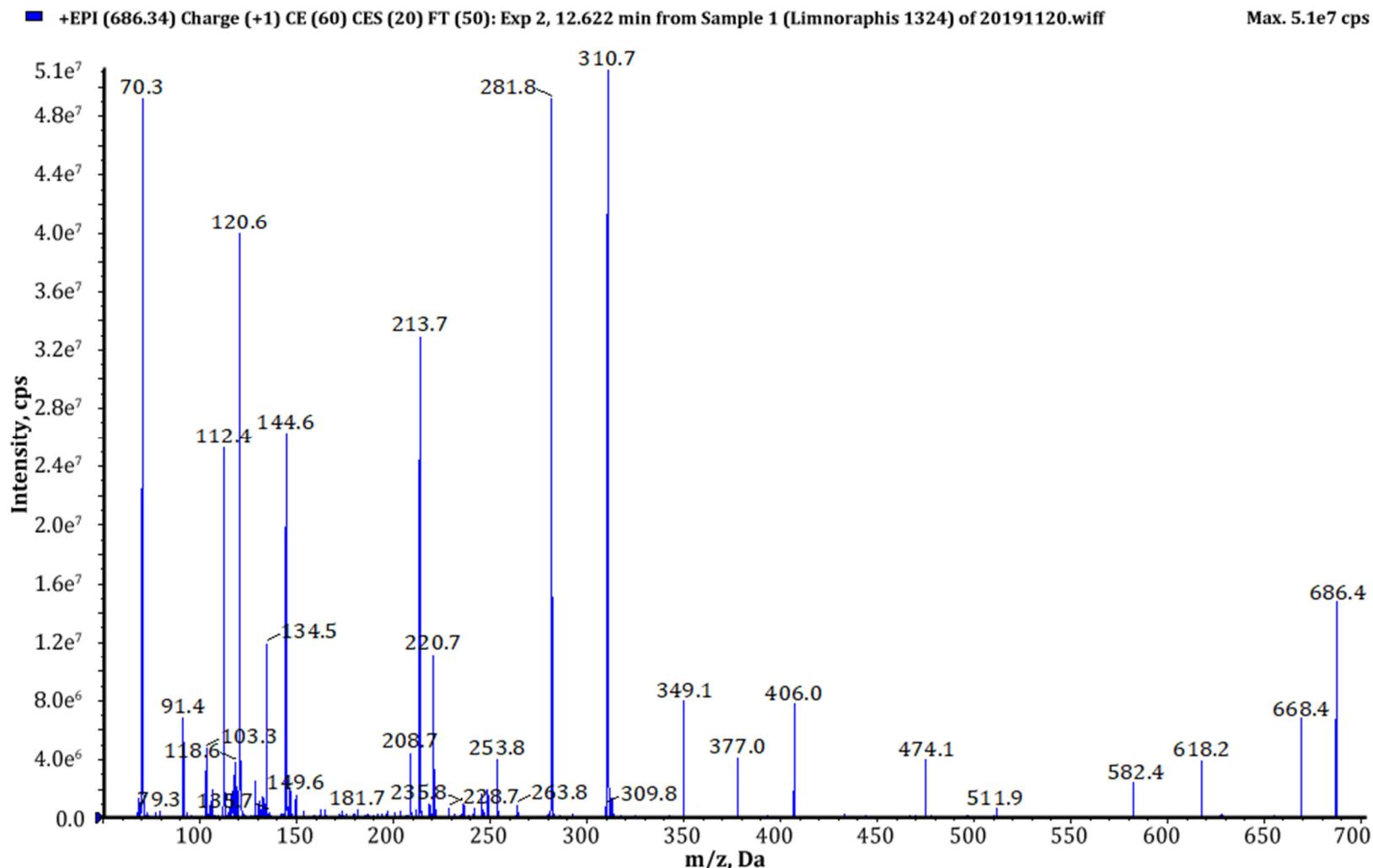


Figure S8. Enhanced product ion mass spectrum of aeruginosamide AER685 with general structure Pre+Phe+Hph/MeHph+Pro+Pyr+TzlCOOMe identified based on the following fragment ions: 686 [M+H], 668 [M+H-H₂O], 618 [M+H-Pre], 474 [Pre+Phe+Hph/MeHph+Pro+H], 406 [Phe+Hph/MeHph+Pro+H], 377 [Pre+Phe+Hph/MeHph+H], 349 [Pre+Phe+Hph/MeHph+H-CO], 310 [Pro+Pyr+TzlCOOMe+H], 309 [Phe+Hph/MeHph+H], 281 [Phe+Hph/MeHph+H-CO], 213 [Pyr+TzlCOOMe+H], 144 [TzlCOOMe], 134 Hph/MeHph immonium ion, 120 Phe immonium ion, 112 TzlCO, 70 Pro immonium ion.

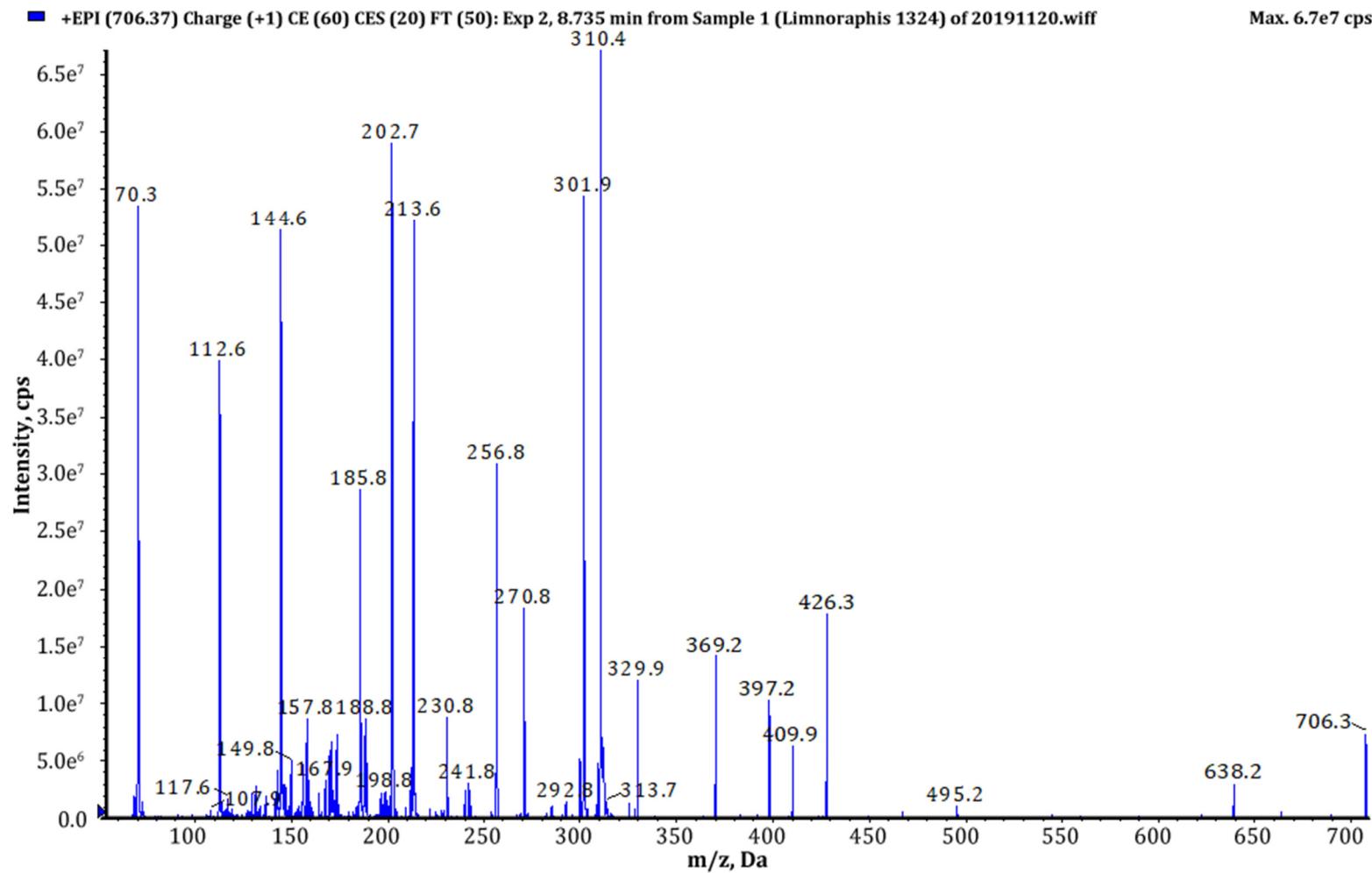


Figure S9. Enhanced product ion mass spectrum of aeruginosamide AEG705 with general structure (Pre)+Hph/MeHph+Val+Pro+Pyr+TzlCOOMe identified based on the following fragment ions: 706 [M+H], 638 [M+H–Pre], 495 [Pre+Hph/MeHph+Val+Pro+Pyr+H], 426 [Pre+Hph/MeHph+Val+Pro+H], 409 [Val+Pro+Pyr+TzlCOOMe], 397 [(Pre)₂+Hph/MeHph+Val+H], 369 [(Pre)₂+Hph/MeHph+Val+H–CO], 329 [Pre+Hph/MeHph+Val+H], 310 [Pro+Pyr+TzlCOOMe+H], 301 [Pre+Hph/MeHph+Val+H–CO], 270 [(Pre)₂+Hph/MeHph+H–CO], 230 [Pre+Hph/MeHph+H], 213 [Pyr+TzlCOOMe+H], 202 [Pre+Hph/MeHph+H–CO], 144 [TzlCOOMe], 112 TzlCO, 70 Pro immonium ion.

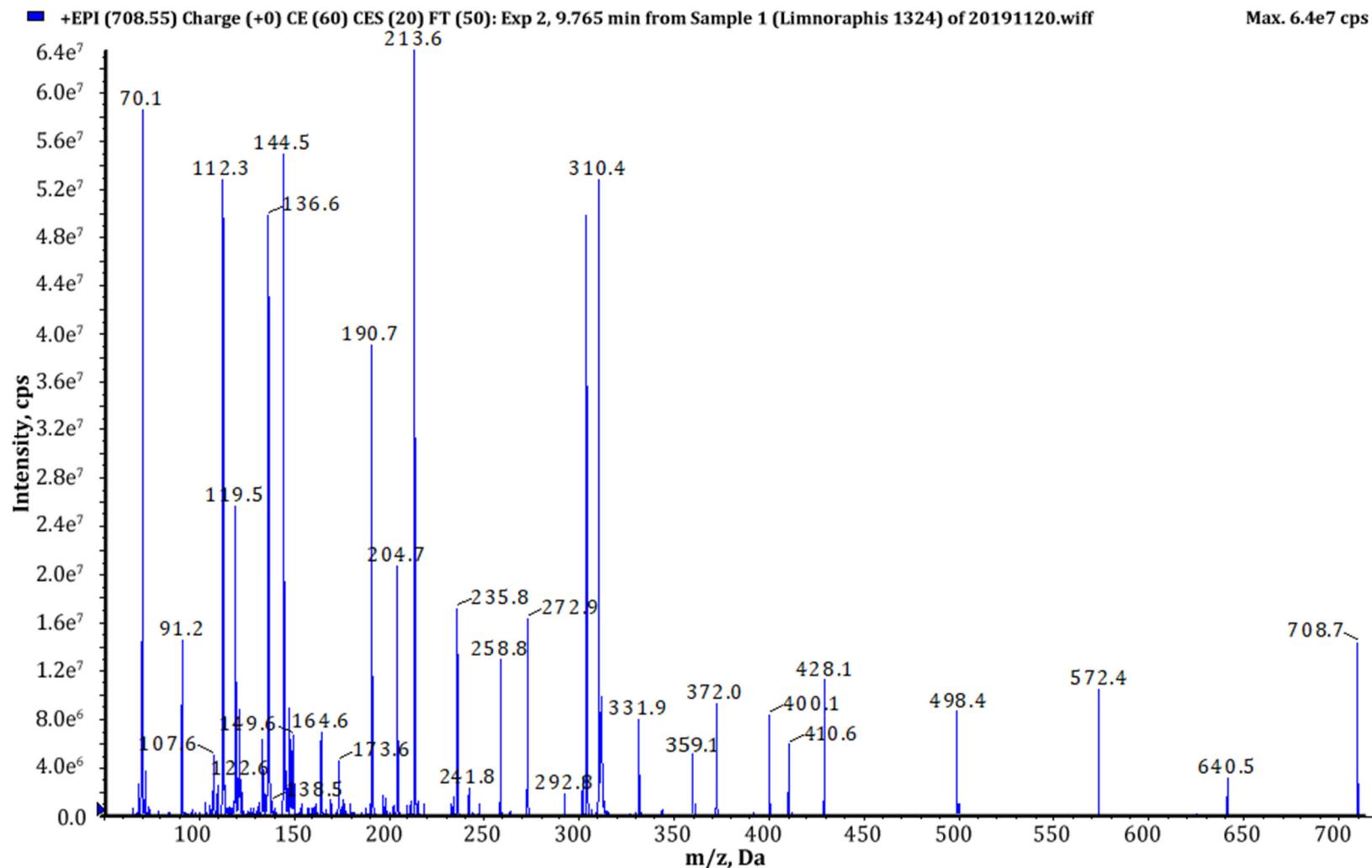


Figure S10. Enhanced product ion mass spectrum of aeruginosamide AEG707 with general structure (Pre)₂+Tyr+Val+Pro+Pyr+TzlCOOMe identified based on the following fragment ions: 708 [M+H], 640 [M+H-Pre], 572 [M+H-(Pre)₂], 498 [M+H-Pre-TzlCOOMe], 429 [Tyr+Val+Pro+Pyr+H], 409 [Val+Pro+Pyr+TzlCOOMe], 400 [Pre+Tyr+Val+Pro+H-CO], 370 [(Pre)₂+Tyr+Val+H-CO], 360 [Tyr+Val+Pro+H], 331 [Pre+Tyr+Val+H], 310 [Pro+Pyr+TzlCOOMe+H], 271 [(Pre)₂+Tyr+H-CO], 235 [Tyr+Val+H-CO], 213 [Pyr+TzlCOOMe+H], 204 [Pre+Tyr+H-CO], 144 [TzlCOOMe], 136 Tyr immonium ion, 112 TzlCO, 70 Pro immonium ion.

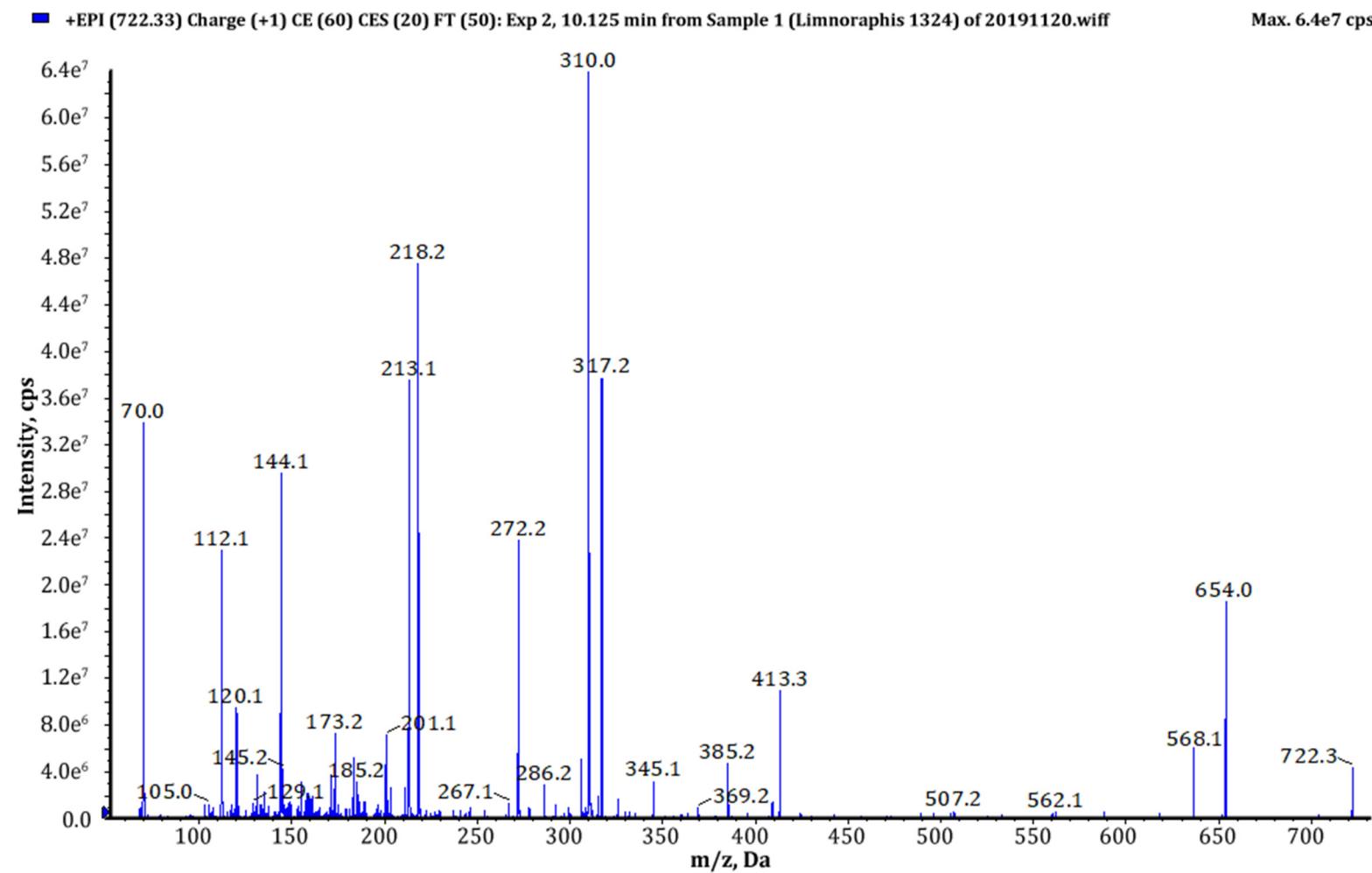


Figure S11. Enhanced product ion mass spectrum of aeruginosamide AEG721 with general structure (Pre)₂+Tyr+Ile/Leu+Pro+Pyr+TzlCOOMe identified based on the following fragment ions: 722 [M+H], 654 [M+H–Pre], 586 [M+H–(Pre)₂], 413 [(Pre)₂+Tyr+Ile/Leu+H], 385 [(Pre)₂+Tyr+Ile/Leu+H–CO], 345 [Pre+Tyr+Ile/Leu+H], 317 [Pre+Tyr+Ile/Leu+H–CO], 310 [Pro+Pyr+TzlOMe+H], 272 [(Pre)₂+Tyr+H–CO], 213 [Pyr+TzlCOOMe+H], 144 [TzlCOOMe], 112 TzlICO, 70 Pro immonium ion.

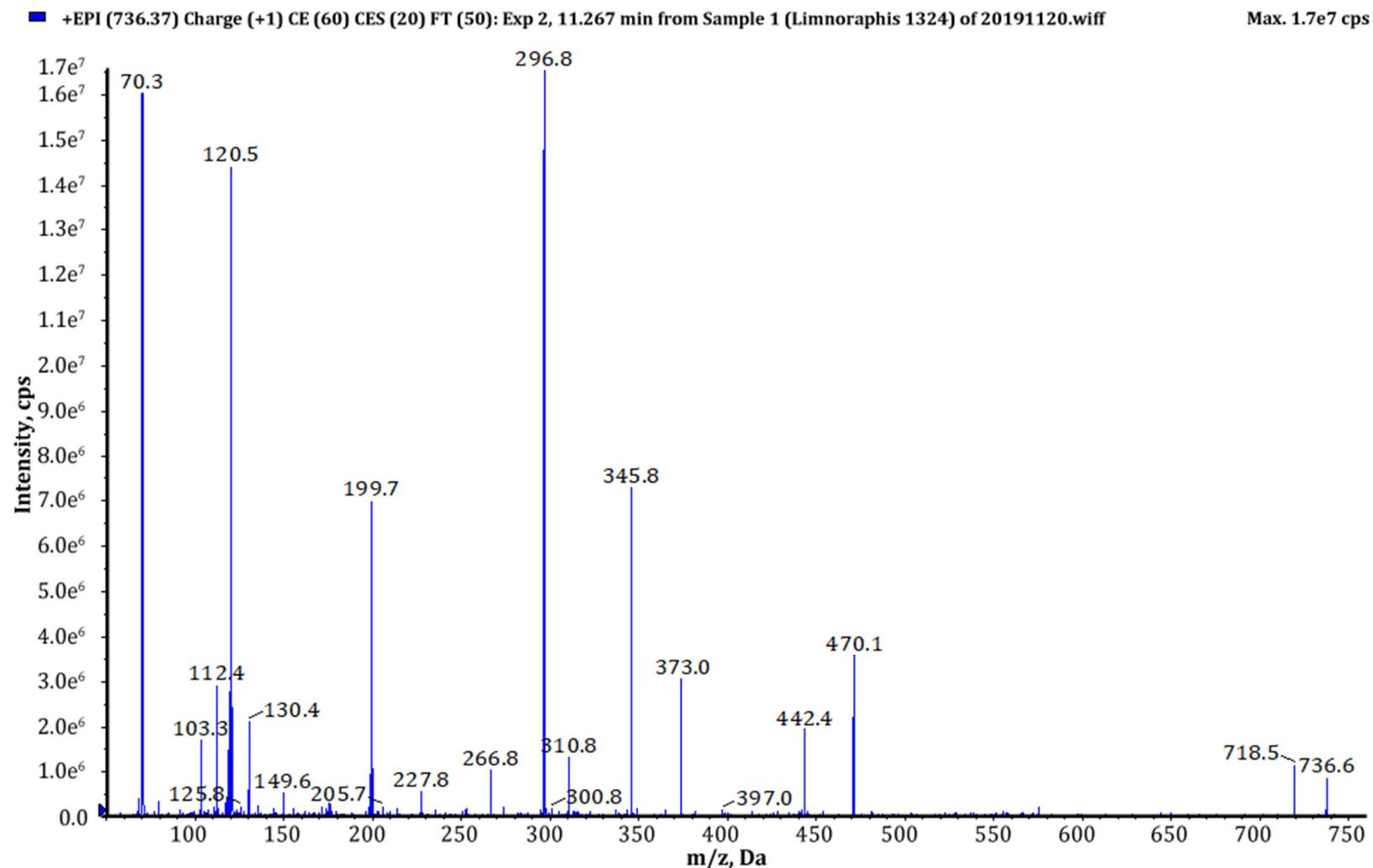


Figure S12. Enhanced product ion mass spectrum of aeruginosamide AER735 with general structure Pre+225+Phe+Pro+Pyr+TzlCOOH identified based on the following fragment ions: 736 [M+H], 718 [M+H-H₂O], 470 [225+Phe+Pro+H], 442 [225+Phe+Pro+H-CO], 373 [225+Phe+H], 345 [225+Phe+H-CO], 296 [Pro+Pyr+TzlCOOH+H], 266 [Pre+225+H-CO], 199 [Pyr+TzlCOOH+H], 130 [TzlCOOH], 120 Phe immonium ion, 112 TzlCO, 70 Pro immonium ion.

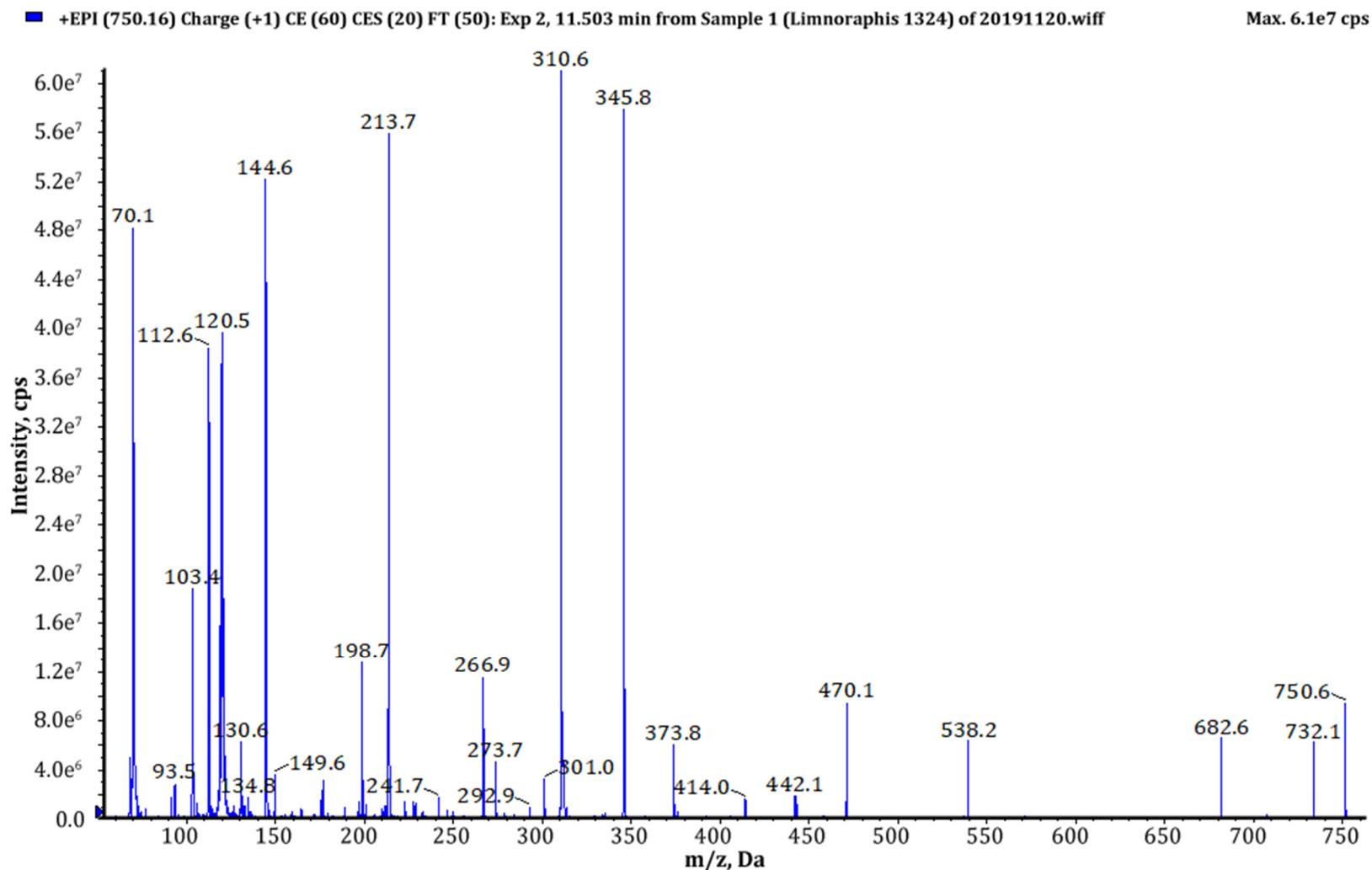


Figure S13. Enhanced product ion mass spectrum of aeruginosamide AER749 with general structure Pre+225+Phe+Pro+Pro+TzlCOOMe identified based on the following fragment ions: 750 [$M+H$], 732 [$M+H-H_2O$], 538 [$M+H-(Pyr+TzlCOOMe)$], 470 [225+Phe+Pro+H], 442 [225+Phe+Pro+H-CO], 373 [225+Phe+H], 345 [225+Phe+H-CO], 310 [Pro+Pyr+TzlCOOMe+H], 266 [Pre+225+H-CO], 213 [Pyr+TzlCOOMe+H], 198 [225+H-CO], 144 [TzlCOOMe], 120 Phe immonium ion, 112 TzlCO, 70 Pro immonium ion.

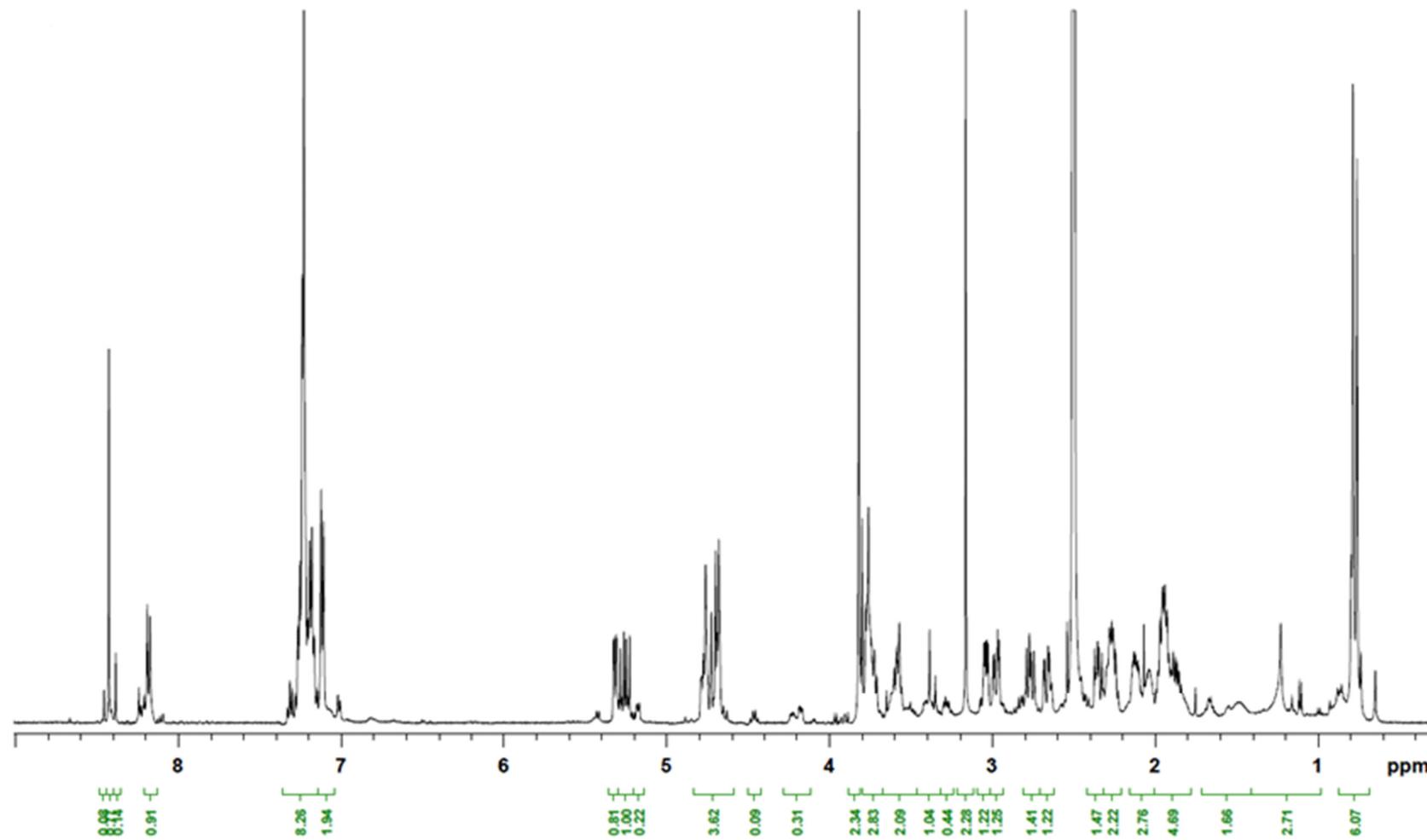


Figure 14. ¹H NMR Spectrum of aeruginosamide AEG671 in DMSO-d₆.

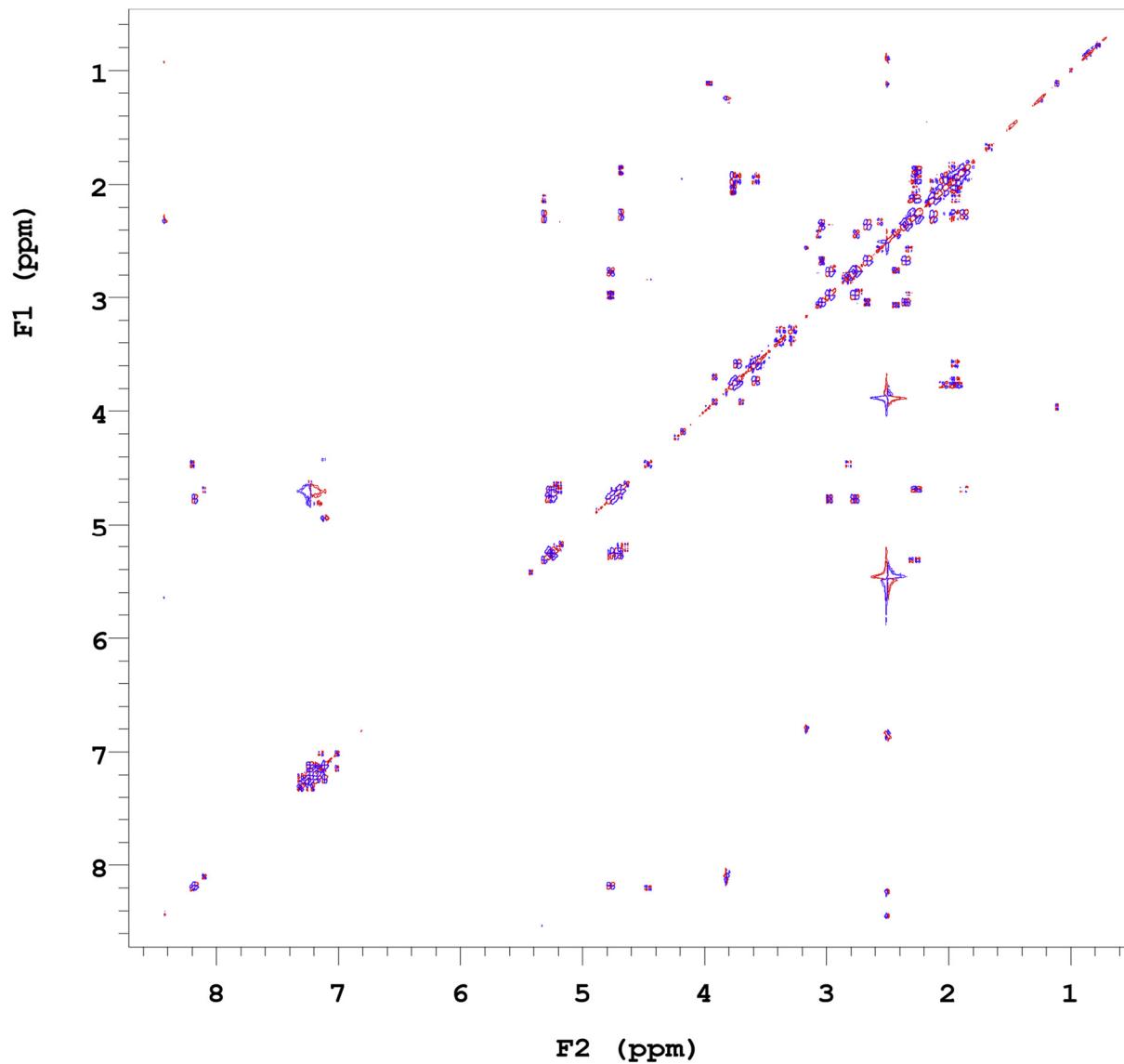


Figure 15. COSY Spectrum of aeruginosamide AEG671 in DMSO-d_6 .

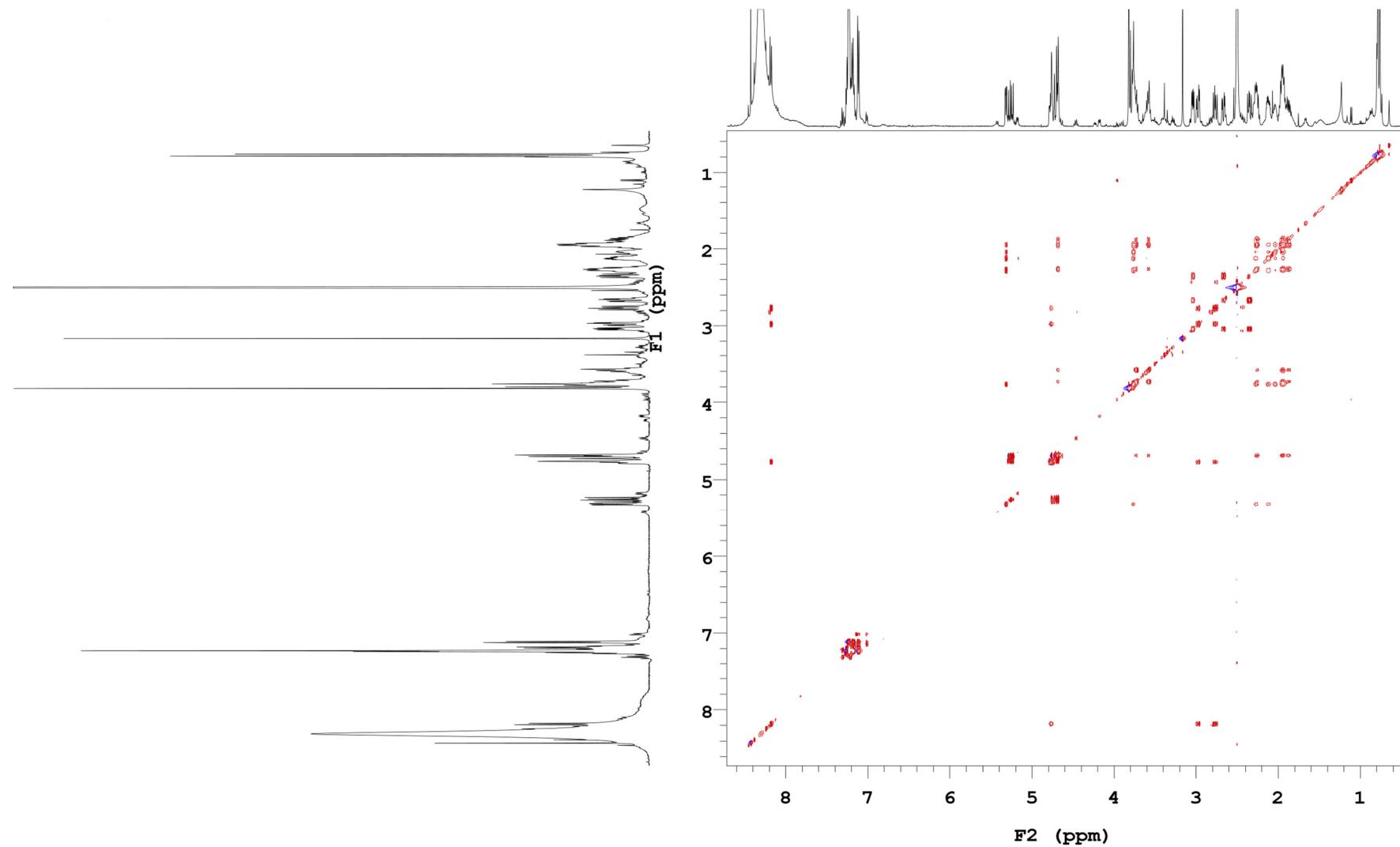


Figure 16. TOCSY Spectrum of aeruginosamide AEG671 in DMSO-d₆.

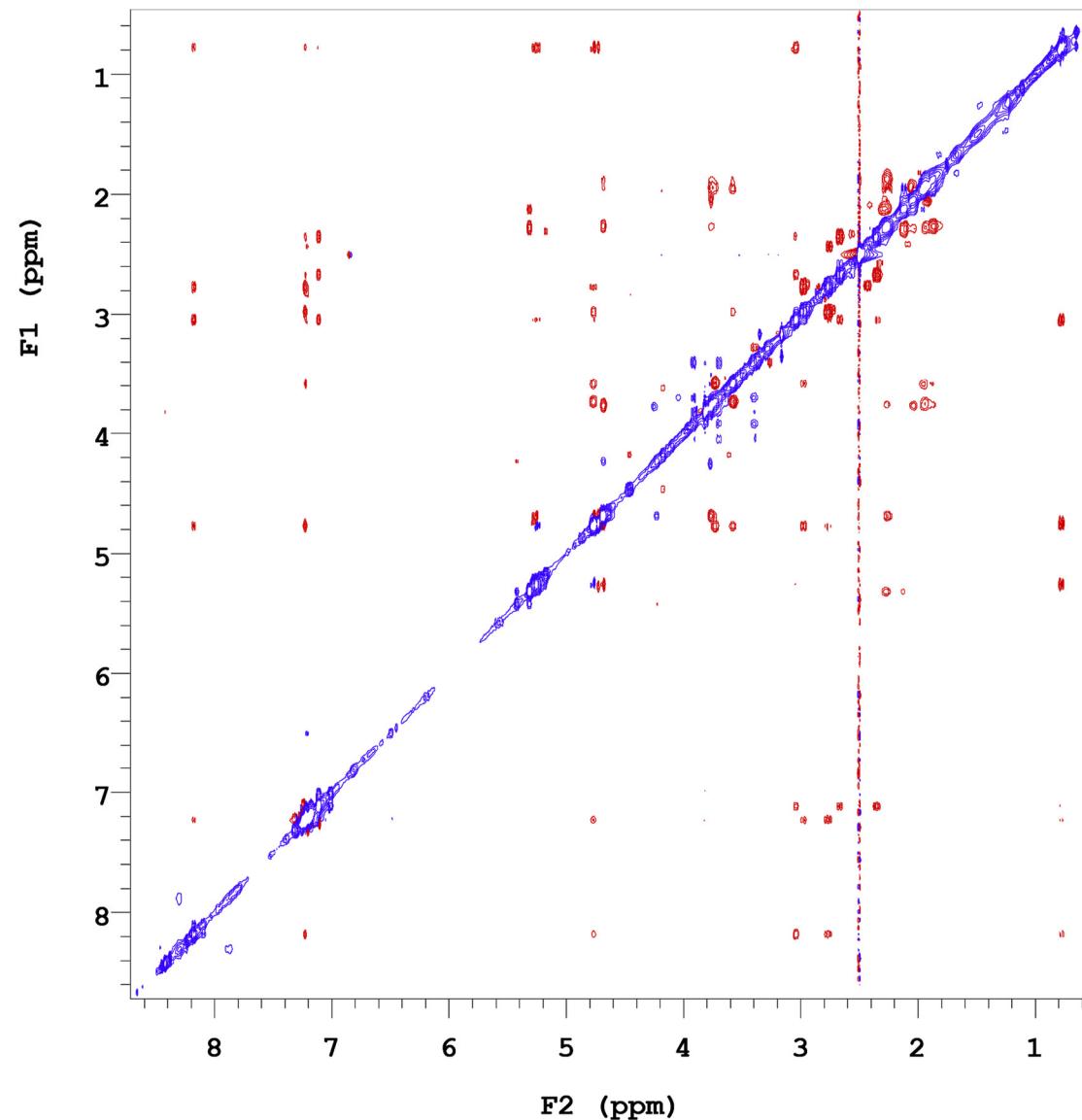


Figure 17. ROESY Spectrum of aeruginosamide AEG671 in DMSO-d₆.

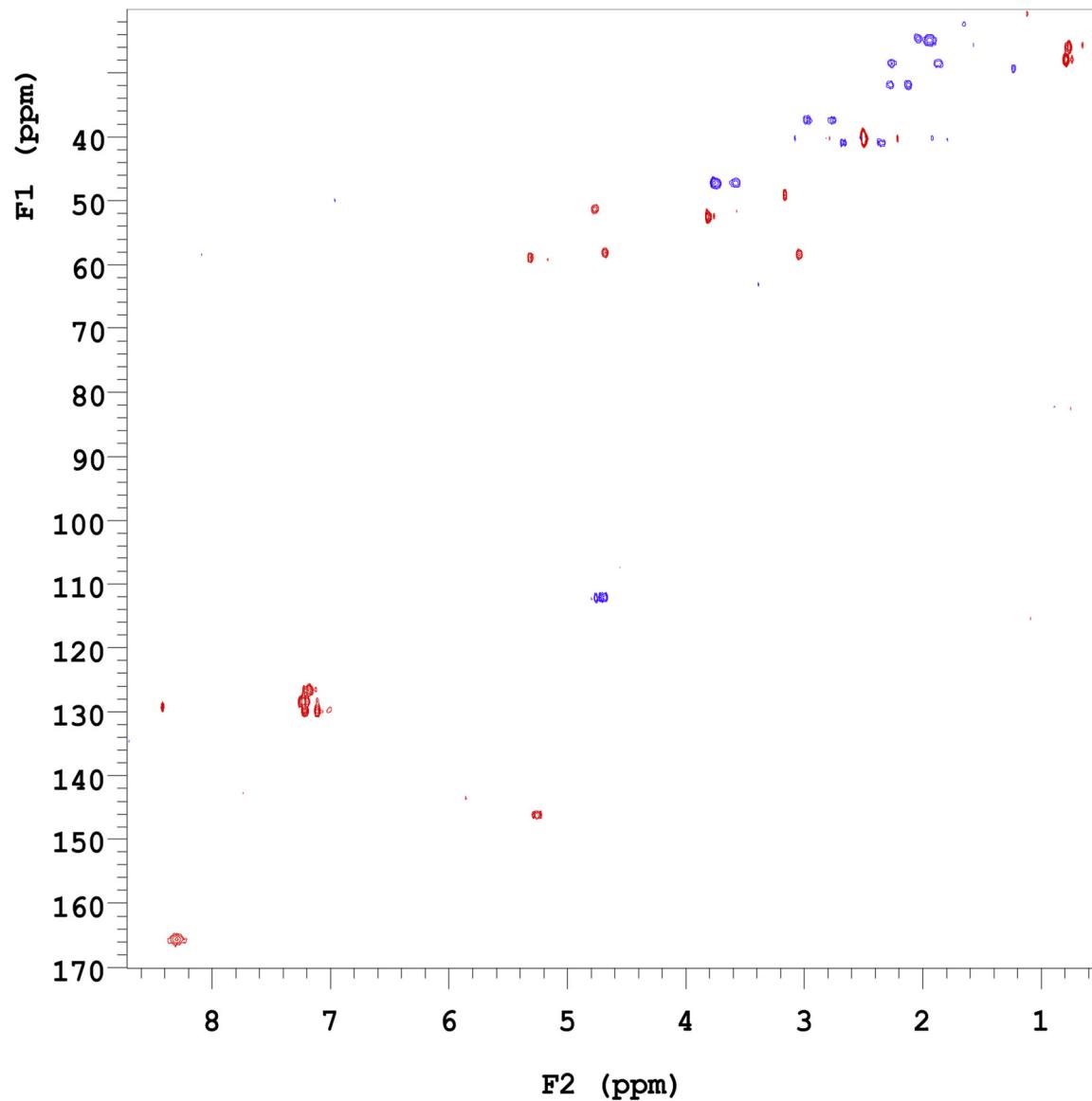


Figure 18. HSQC Spectrum of aeruginosamide AEG671 in DMSO-d_6 .

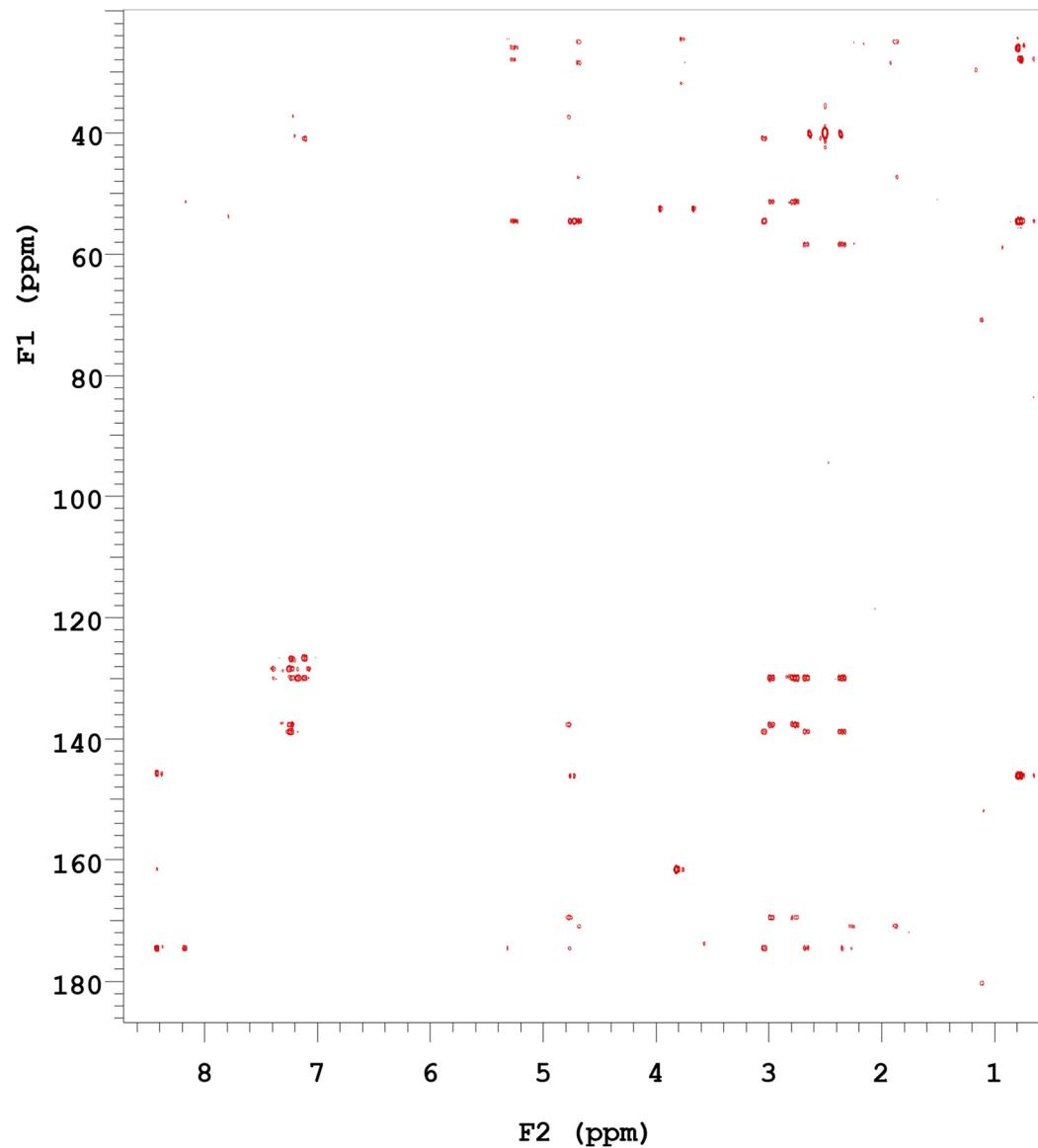


Figure 19. HMBC Spectrum of aeruginosamide AEG671 in DMSO-d_6 .



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