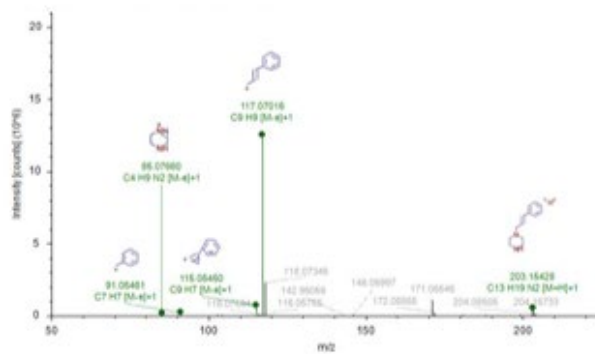
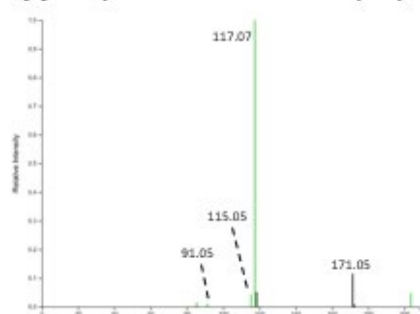
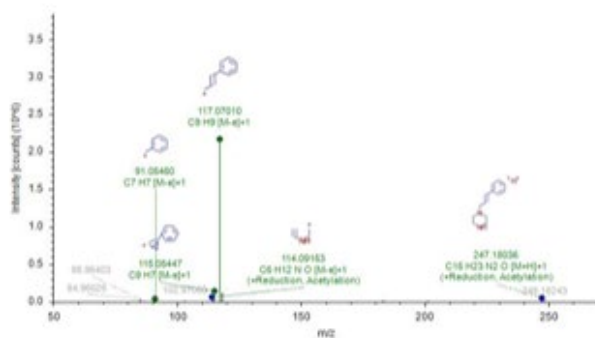
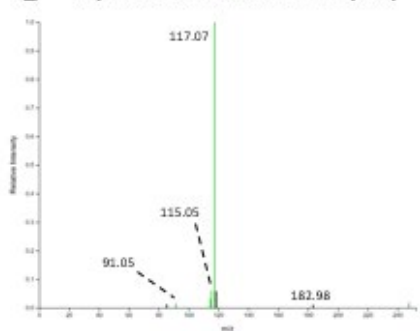
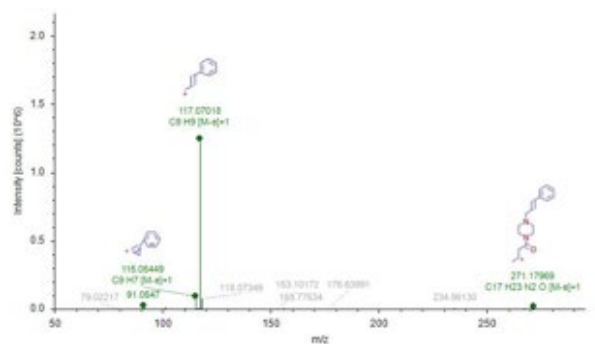
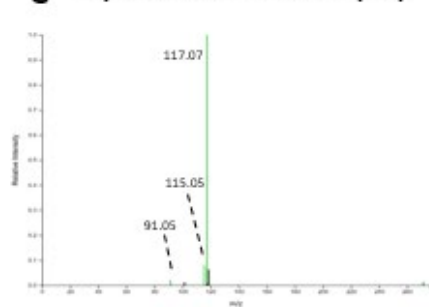
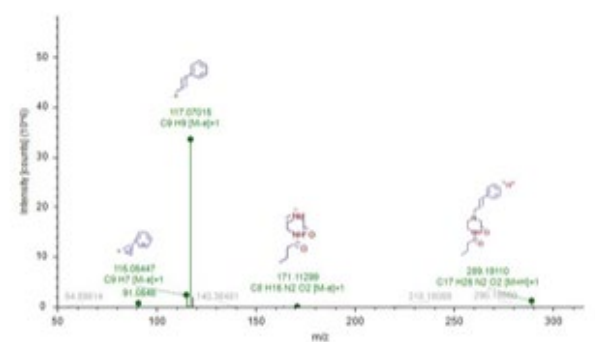
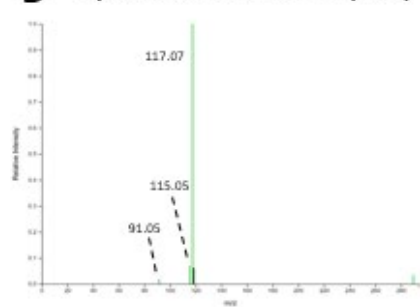


A m/z 203.1542 RT: 3.2 min (M1)**B** m/z 247.1804 RT: 5.6 min (M2)**C** m/z 271.1797 RT: 5.4 min (M3)**D** m/z 289.1911 RT: 4.8 min (M4a)

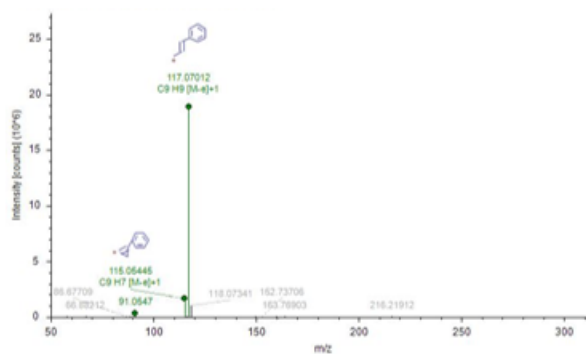
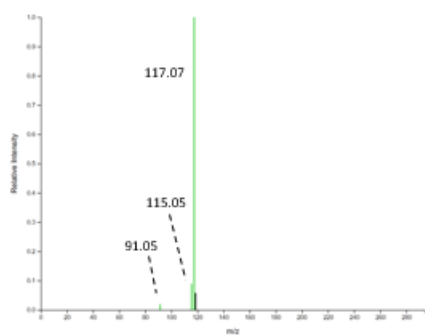
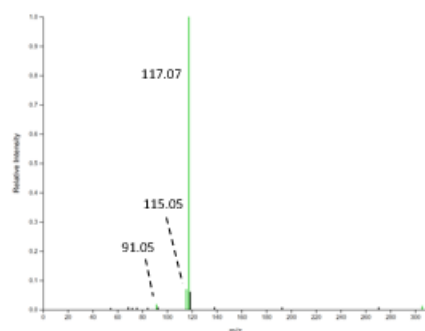
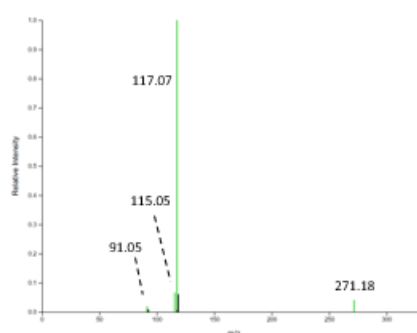
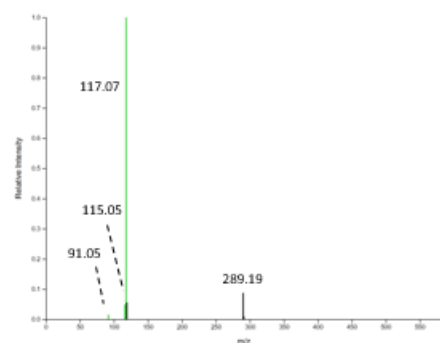
E m/z 289.1911 RT: 5.8 min (M4b)**F** m/z 305.1868 RT: 5.1 min**G** m/z 329.2225 RT: 6 min**H** m/z 577.3749 RT: 5.8 min

Figure S1: (A-E) Fragmentation spectrum of the most likely AP-237 metabolites. The spectra on the left were visualized using SIRIUS software, while the spectra on the right were visualized and annotated automatically using Compound Discoverer® software. (F-H) Fragmentation spectrum of compounds present in the molecular network but not proposed by Compound Discoverer®.