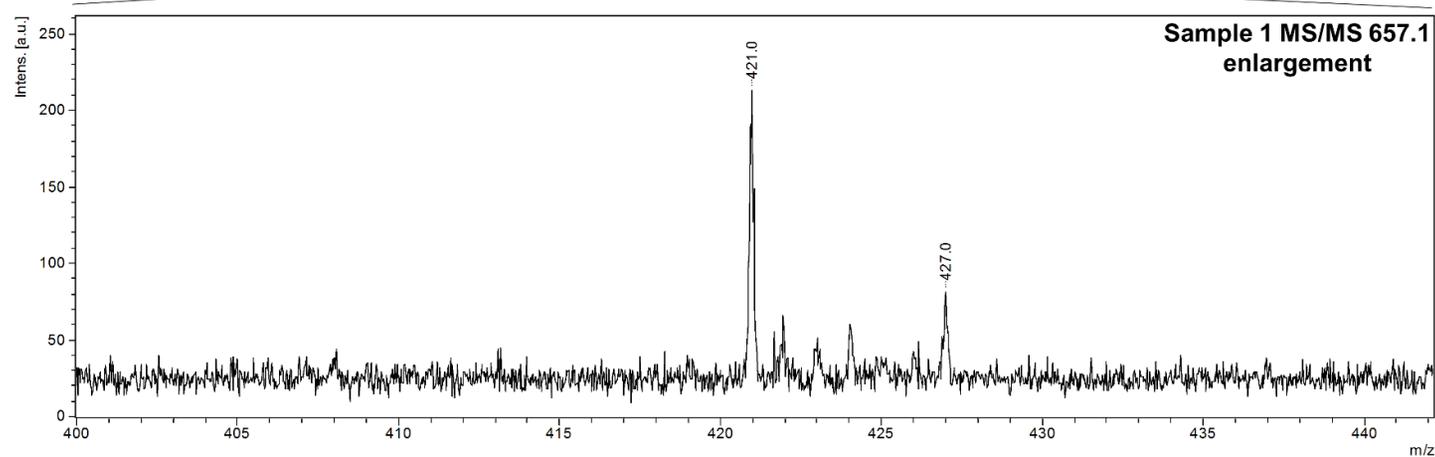
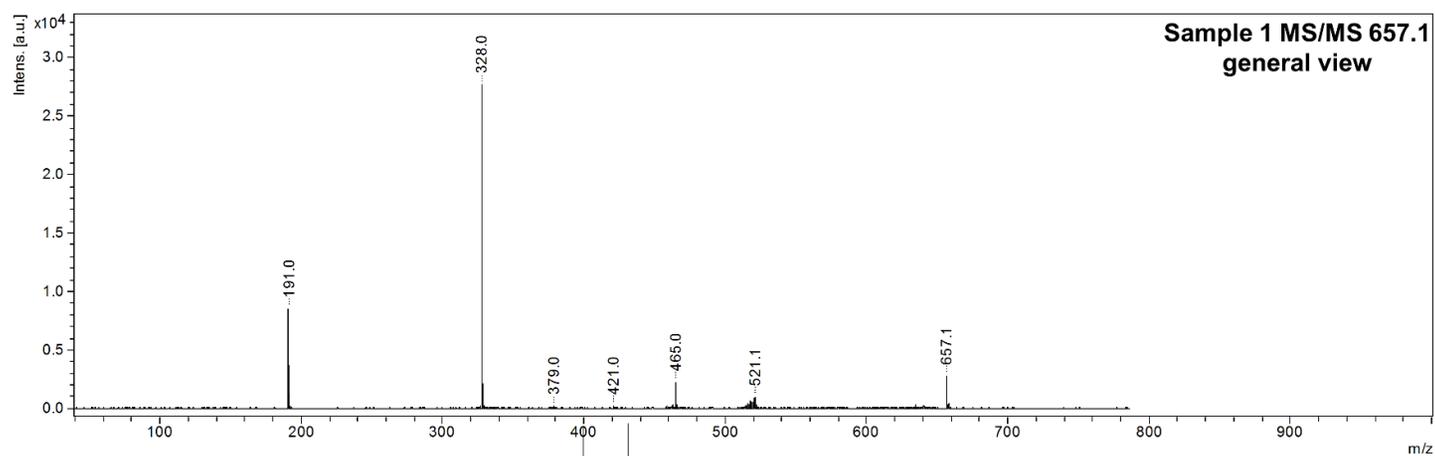


Fragmentation of the non-covalent cAMP dimer, MS/MS data, negative mode



The dominant fragment ($m/z = 328.0$) corresponds to free cAMP.
Signals indicating covalent binding ($m/z = 408.0, 426.0$) were not detected.