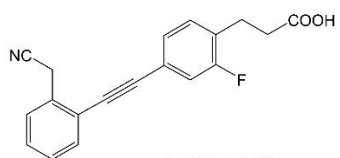


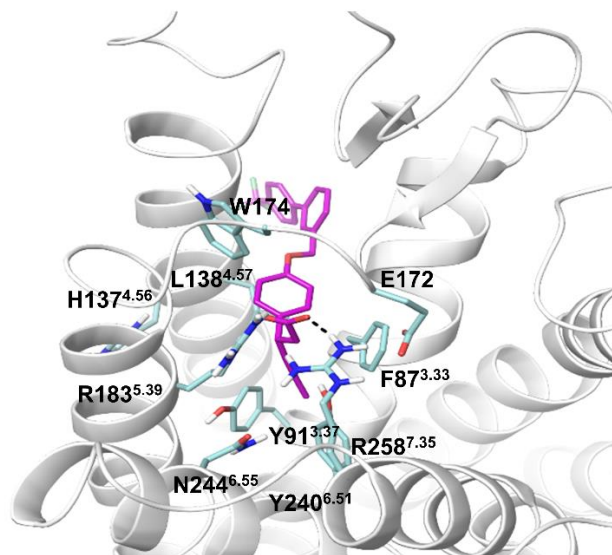
Mode 2



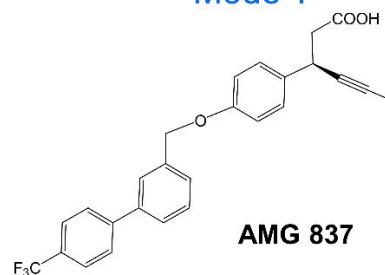
TUG-770

pEC₅₀ = 8.2

agonist



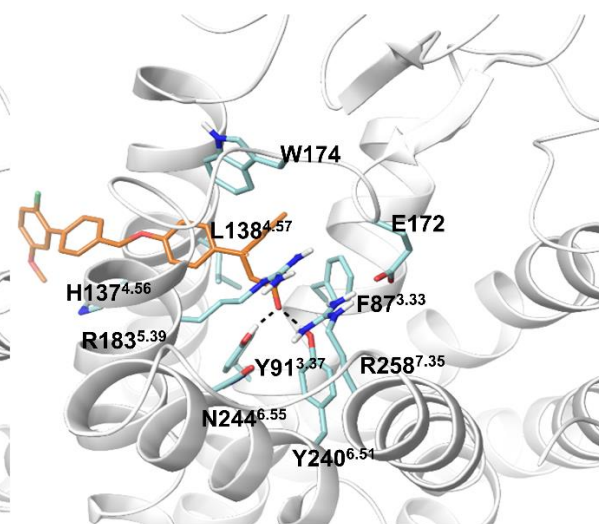
Mode 1



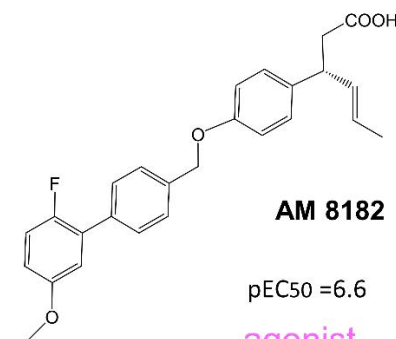
AMG 837

pEC₅₀ = 8.5

partial agonist



Mode 2



AM 8182

pEC₅₀ = 6.6

agonist

Additional Figure 5S. Agonist binding at the FFA1 crystal structure.