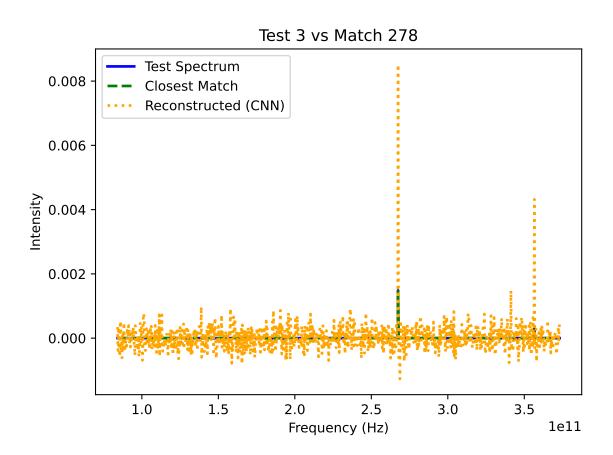
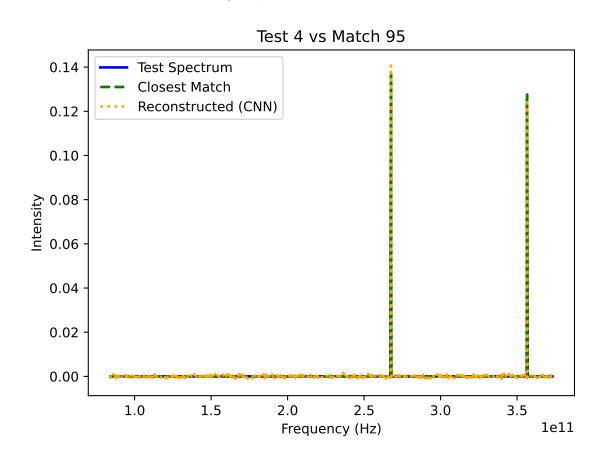


Test File: file_JPL_HCO+v=0,1,2_T232.98085122481015_N12.99999999999996_simulate_generate Test File: file_JPL_HCO+v=0,1,2_T81.57307210000003_N15.29999999999988_simulate_generate Header: //molecules='HCO+V=0,1,2|1' logn=13.0 tex=232.98085 velo=250.0 fwhm=50.0 sourcesize=10.@leader: //molecules='HCO+V=0,1,2|1' logn=15.3 tex=81.573074 velo=250.0 fwhm=50.0 sourcesize=10.0

Match: file JPL HCO+v=0,1,2 T179.2160394037001 N12.8999999999999 simulate generate Match: file JPL HCO+v=0,1,2 T28.56100000000000 N15.3999999999988 simulate generate Header: //molecules='HCO+V=0,1,2|1' logn=12.9 tex=179.21603 velo=250.0 fwhm=50.0 sourcesize=10.0 Header: //molecules='HCO+V=0,1,2|1' logn=15.4 tex=28.561 velo=250.0 fwhm=50.0 sourcesize=10.0

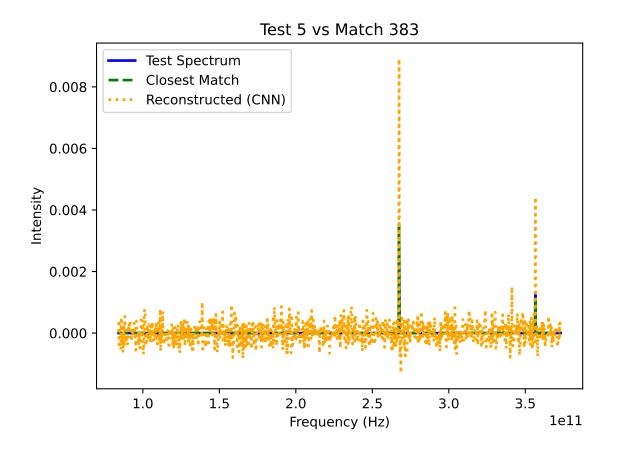


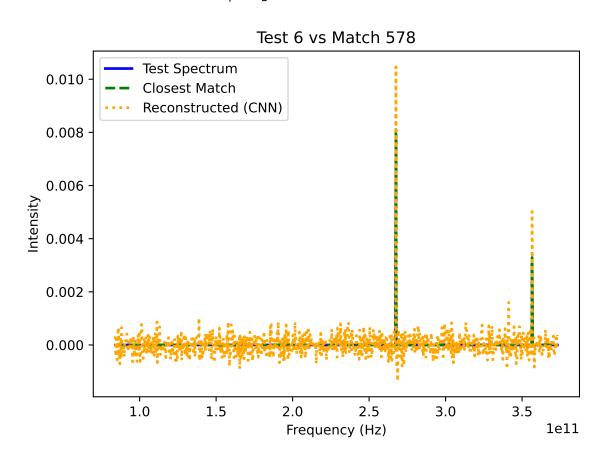


Test File: file JPL HCO+v=0,1,2 T28.56100000000000 N13.299999999999 simulate generate Header: $//molecules='HCO+V=0,\bar{1},2|1' logn=13.3 tex=28.\bar{5}61 velo=250.0 fwhm=50.0 sourcesize=10.0 Header: <math>//molecules='HCO+V=0,1,\bar{2}|1' logn=16.4 tex=37.\bar{1}293 velo=250.0 fwhm=50.0 fwhm=5$

Match: file JPL HCO+v=0,1,2 T21.9700000000000 N13.2999999999999 simulate generate Header: //molecules='HCO+V=0,1,2|1' logn=13.3 tex=21.97 velo=250.0 fwhm=50.0 sourcesize=10.0 Header: //molecules='HCO+V=0,1,2|1' logn=16.5 tex=37.1293 velo=250.0 fwhm=50.0 sourcesize=10.0

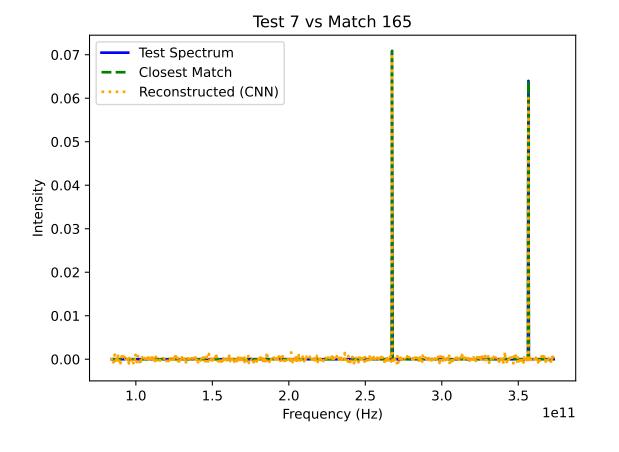
Match: file JPL HCO+v=0,1,2 T37.1293000000001 N16.4999999999999 simulate generate

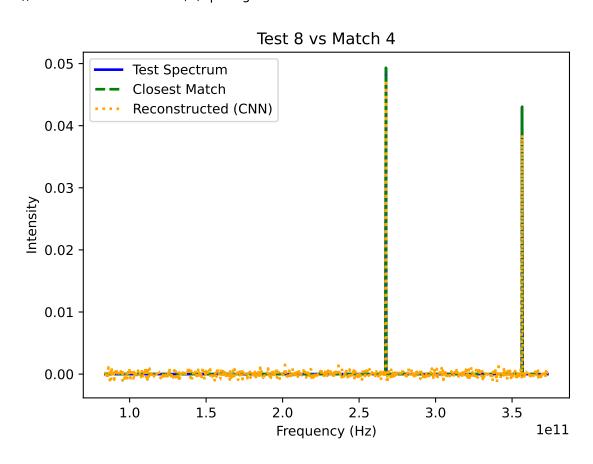




Test File: file_JPL_HCO+v=0,1,2_T62.74851700000002_N13.79999999999994_simulate_generate Test File: file_JPL_HCO+v=0,1,2_T106.04499373000004_N14.299999999999992_simulate_generate Header: //molecules='HCO+V=0,1,2|1' logn=13.8 tex=62.748516 velo=250.0 fwhm=50.0 sourcesize=10. Header: //molecules='HCO+V=0,1,2|1' logn=14.3 tex=106.04499 velo=250.0 fwhm=50.0 sourcesize=10.0 fwhm=50.0 sourcesize=10.0 fwhm=50.0 sourcesize=10.0 fwhm=50.0 sourcesize=10.0 fwhm=50.0 sourcesize=10.0 fwhm=50.0 fwhm=50.

Match: file_JPL_HCO+v=0,1,2_T37.1293000000001_N13.69999999999994_simulate_generate Header: $//molecules='HCO+V=0,\bar{1},2|1' logn=13.7 tex=3\bar{7}.1293 velo=250.0 fwhm=50.0 sourcesize=10.0 Header: <math>//molecules='HCO+V=0,\bar{1},2|1' logn=14.4 tex=13\bar{7}.85849 velo=250.0 fwhm=50.0 fwhm=$



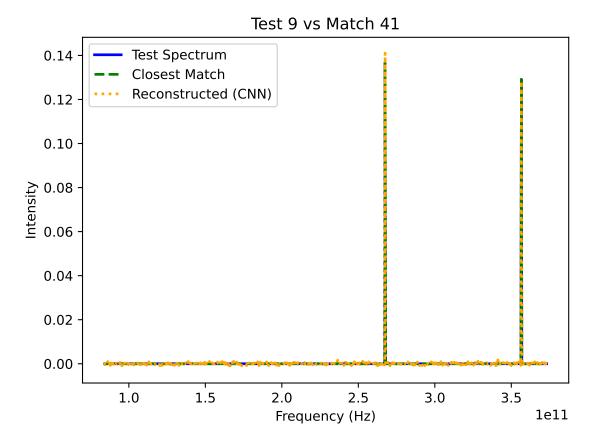


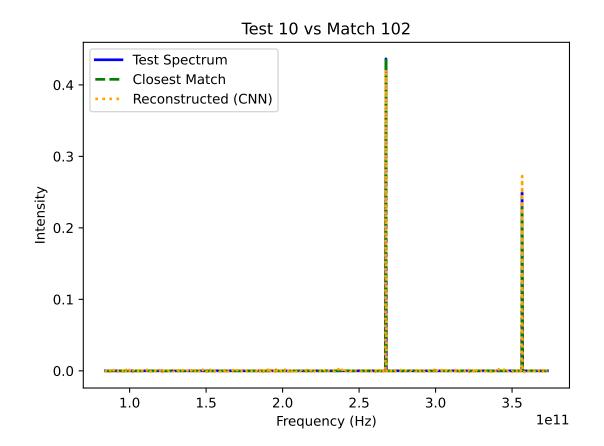
Header: $\frac{1}{100}$ Heade

Test File: file_JPL_HCO+v=0,1,2_T16.90000000000002_N19.100000000003_simulate_generate Header: //molecules='HCO+V=0,1,2|1' logn=19.1 tex=16.9 velo=250.0 fwhm=50.0 sourcesize=10.0

Match: file_JPL_HCO+v=0,1,2_T21.97000000000002_N16.6999999999996_simulate_generate Header: //molecules='HCO+V=0,1,2|1' logn=16.7 tex=21.97 velo=250.0 fwhm=50.0 sourcesize=10.0

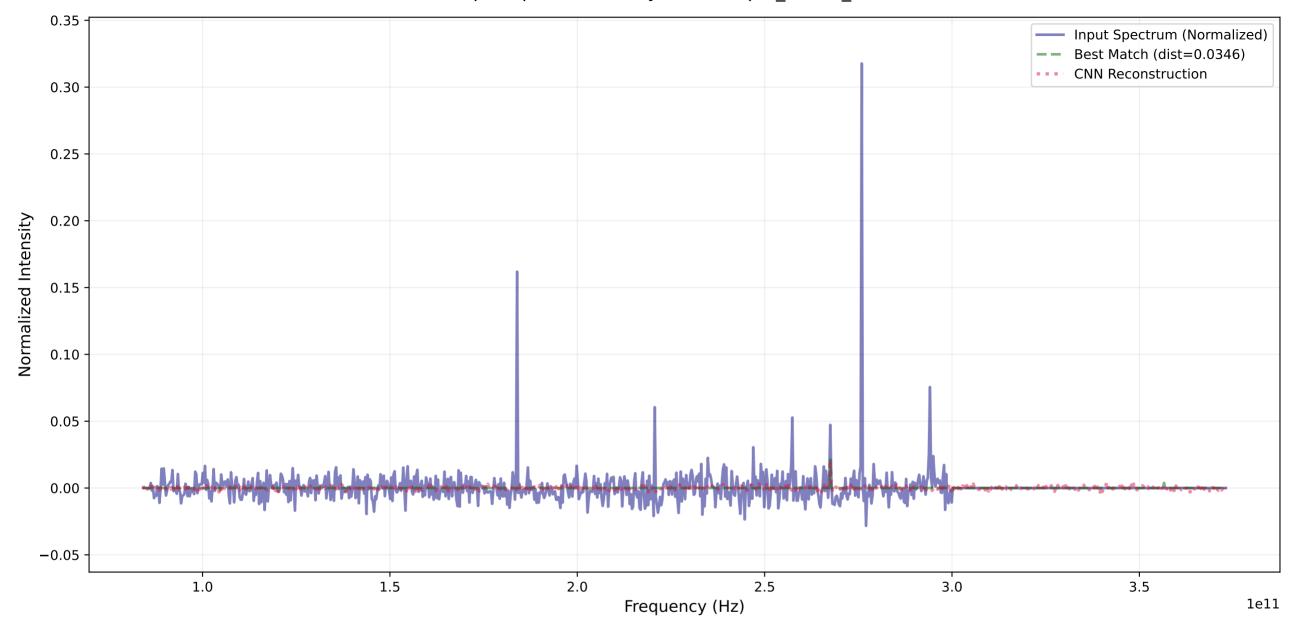
Match: file_JPL_HCO+v=0,1,2_T16.90000000000002_N17.600000000001_simulate_generate Header: $//molecules='HCO+V=\bar{0},1,2|1'|logn=17.6 tex=16.9 velo=250.0 fwhm=50.0 sourcesize=10.0$





Test File: file JPL HCO+v=0,1,2 T37.1293000000001 N18.7000000000024 simulate generate Header: //molecules='HCO+V=0,1,2|1' logn=18.7 tex=37.1293 velo=250.0 fwhm=50.0 sourcesize=10.0Header: //molecules='HCO+V=0,1,2|1' logn=16.4 tex=137.85849 velo=250.0 fwhm=50.0 sourcesize=10.0

Input Spectrum Analysis: example_4mols_format



INPUT FILE:

Filename: example_4mols_format Header: //!xValues(GHz)□yValues(K)

BEST MATCH:

• File: file_JPL_HCO+v=0,1,2_T13.0_N14.79999999999999_simulate_generate

• Distance: 0.034588

• Parameters: //molecules='HCO+V=0,1,2|1' logn=14.8 tex=13.0 velo=250.0 fwhm=50.0 sourcesize=10.0