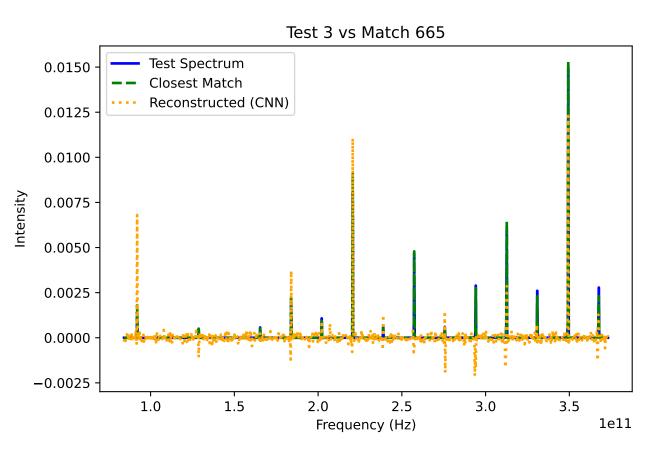


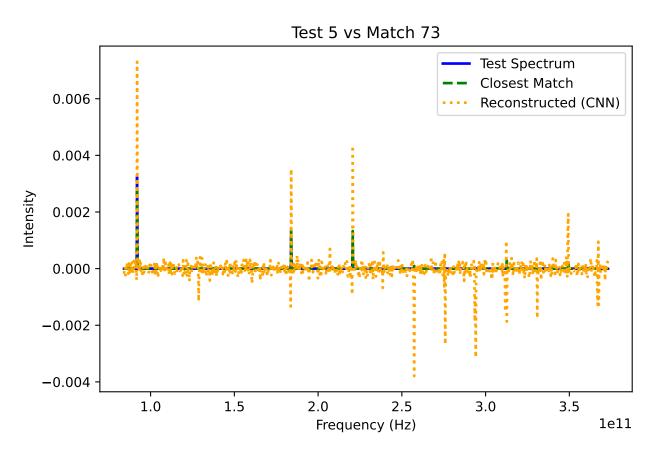
Test File: file\_JPL\_CH3CN\_T232.98085122481015\_N12.99999999999996\_simulate\_generate Header: //molecules='CH3CN|1' logn=13.0 tex=232.98085 velo=250.0 fwhm=50.0 sourcesize=10.0

Header: //molecules='CH3CN|1' logn=12.9 tex=179.21603 velo=250.0 fwhm=50.0 sourcesize=10.0



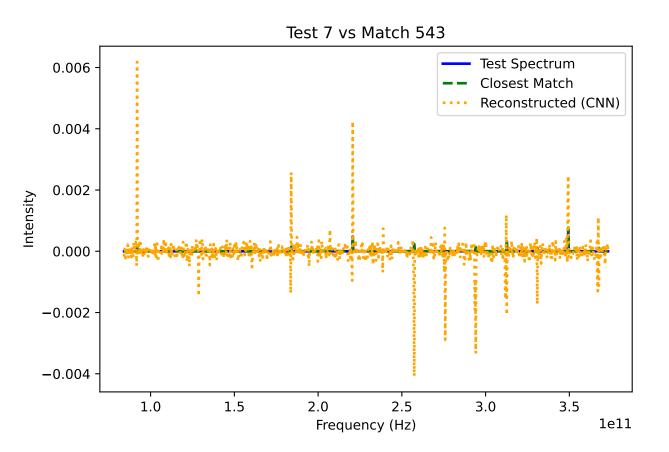
Test File: file\_JPL\_CH3CN\_T232.98085122481015\_N14.9999999999999\_simulate\_generate Header: //molecules='CH3CN|1' logn=15.0 tex=232.98085 velo=250.0 fwhm=50.0 sourcesize=10.0

Header: //molecules='CH3CN|1' logn=14.9 tex=179.21603 velo=250.0 fwhm=50.0 sourcesize=10.0



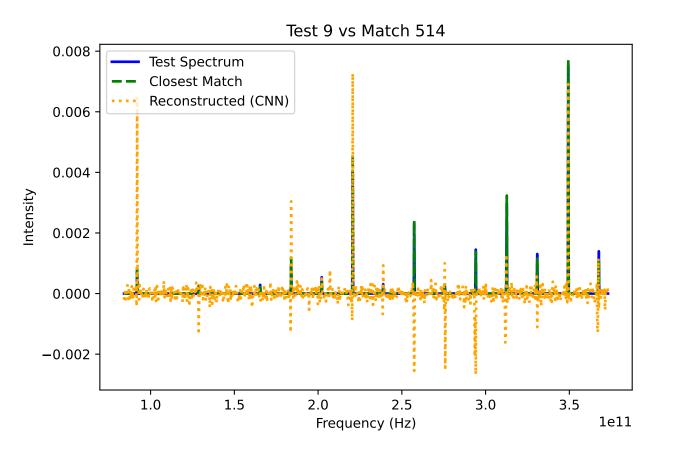
Test File: file\_JPL\_CH3CN\_T21.97000000000002\_N14.0999999999993\_simulate\_generate Header: //molecules='CH3CN|1' logn=14.1 tex=21.97 velo=250.0 fwhm=50.0 sourcesize=10.0

Match: file\_JPL\_CH3CN\_T28.56100000000003\_N14.0999999999993\_simulate\_generate Header: //molecules='CH3CN|1' logn=14.1 tex=28.561 velo=250.0 fwhm=50.0 sourcesize=10.0

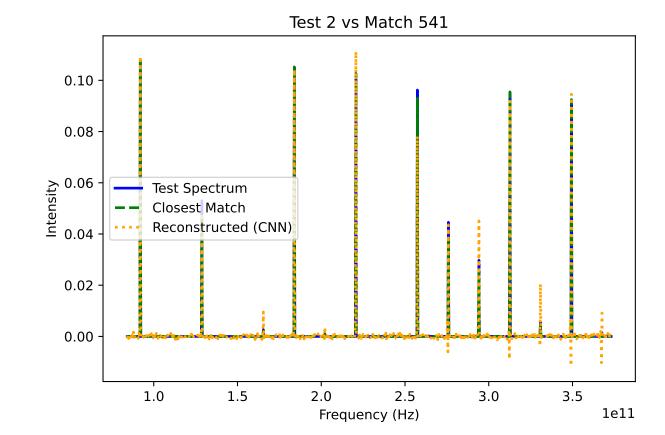


Test File: file\_JPL\_CH3CN\_T137.85849184900007\_N13.4999999999995\_simulate\_generate Header: //molecules='CH3CN|1' logn=13.5 tex=137.85849 velo=250.0 fwhm=50.0 sourcesize=10.0

Match: file JPL CH3CN T179.2160394037001 N13.599999999999999 simulate generate Header: //molecules='CH3CN|1' logn=13.6 tex=179.21603 velo=250.0 fwhm=50.0 sourcesize=10.0

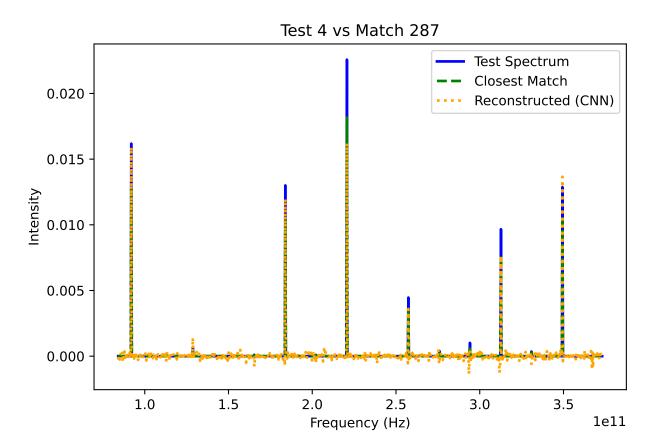


Test File: file JPL CH3CN T232.98085122481015 N14.6999999999999 simulate generate Header: //molecules='CH3CN|1' logn=14.7 tex=232.98085 velo=250.0 fwhm=50.0 sourcesize=10.0



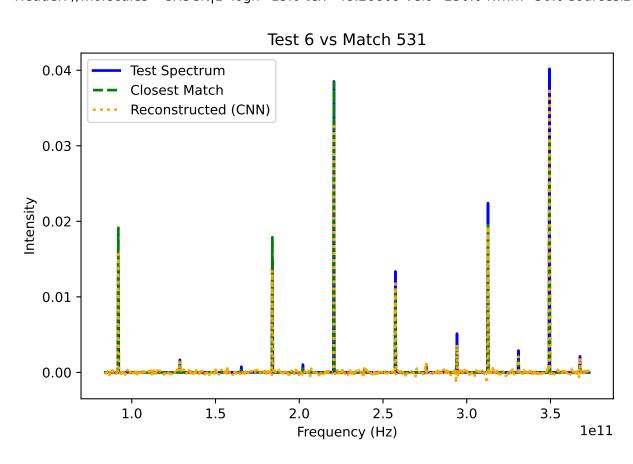
Test File: file\_JPL\_CH3CN\_T28.56100000000003\_N17.700000000001\_simulate\_generate Header: //molecules='CH3CN|1' logn=17.7 tex=28.561 velo=250.0 fwhm=50.0 sourcesize=10.0

Match: file JPL CH3CN T28.56100000000000 N17.600000000001 simulate generate Header: //molecules='CH3CN|1' logn=17.6 tex=28.561 velo=250.0 fwhm=50.0 sourcesize=10.0



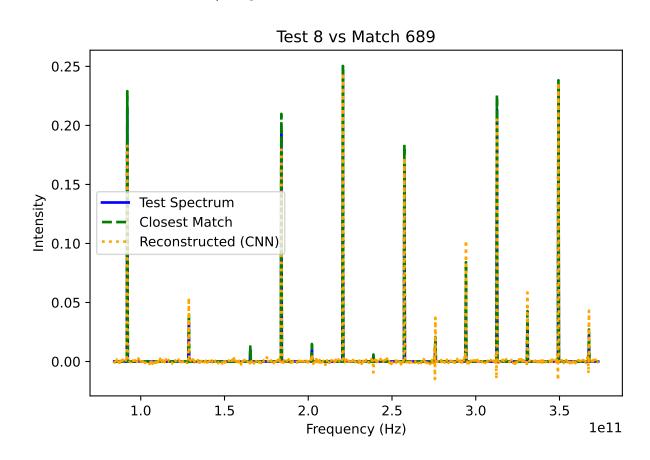
Test File: file\_JPL\_CH3CN\_T48.268090000000015\_N15.09999999999999 simulate\_generate Header: //molecules='CH3CN|1' logn=15.1 tex=48.26809 velo=250.0 fwhm=50.0 sourcesize=10.0

Match: file\_JPL\_CH3CN\_T48.26809000000015\_N14.99999999999999\_simulate\_generate Header: //molecules='CH3CN|1' logn=15.0 tex=48.26809 velo=250.0 fwhm=50.0 sourcesize=10.0



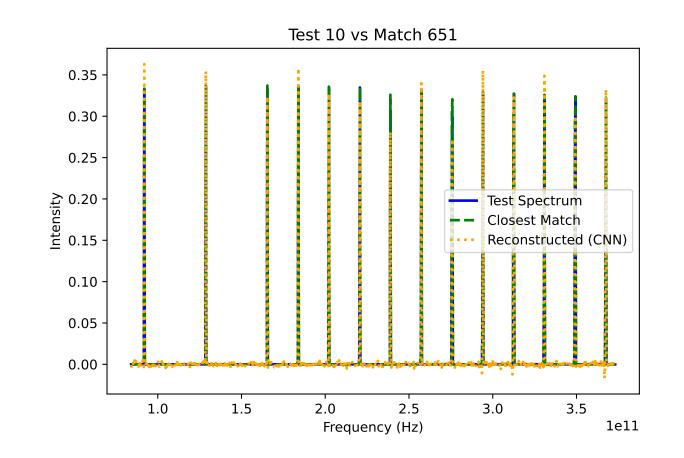
Test File: file\_JPL\_CH3CN\_T81.57307210000003\_N15.29999999999988\_simulate\_generate Header: //molecules='CH3CN|1' logn=15.3 tex=81.573074 velo=250.0 fwhm=50.0 sourcesize=10.0

Match: file\_JPL\_CH3CN\_T62.74851700000002\_N15.29999999999988\_simulate\_generate Header: //molecules='CH3CN|1' logn=15.3 tex=62.748516 velo=250.0 fwhm=50.0 sourcesize=10.0



Test File: file\_JPL\_CH3CN\_T62.74851700000002\_N16.69999999999996\_simulate\_generate Header: //molecules='CH3CN|1' logn=16.7 tex=62.748516 velo=250.0 fwhm=50.0 sourcesize=10.0

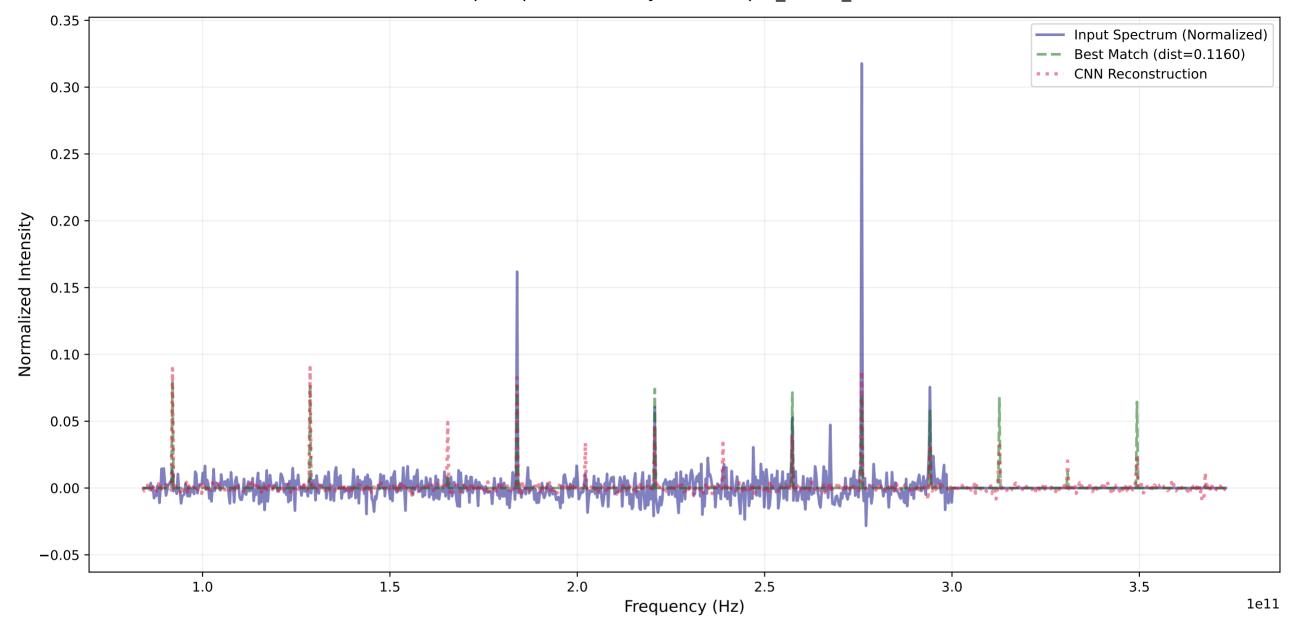
Match: file JPL CH3CN T62.74851700000002 N16.7999999999999 simulate generate Header: //molecules='CH3CN|1' logn=16.8 tex=62.748516 velo=250.0 fwhm=50.0 sourcesize=10.0



Test File: file JPL CH3CN T81.57307210000003 N18.600000000000023 simulate generate Header: //molecules='CH3CN|1' logn=18.6 tex=81.573074 velo=250.0 fwhm=50.0 sourcesize=10.0

Match: file JPL CH3CN T81.57307210000003 N18.800000000000026 simulate generate

## Input Spectrum Analysis: example\_4mols\_format



## INPUT FILE:

\_\_\_\_\_

Filename: example\_4mols\_format Header: //!xValues(GHz)□yValues(K)

## BEST MATCH:

\_\_\_\_\_

• File: file\_JPL\_CH3CN\_T21.970000000000002\_N19.1000000000003\_simulate\_generate

• Distance: 0.115987

• Parameters: //molecules='CH3CN|1' logn=19.1 tex=21.97 velo=250.0 fwhm=50.0 sourcesize=10.0