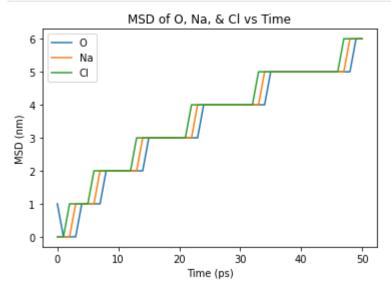
```
#Import modules
In [4]:
         import numpy as np
         import matplotlib.pyplot as plt
         import mdtraj as md
         #Load Files (xtc)
         traj = md.load_xtc('fit_unwrapped_water_nacl.xtc', top='step5_9.gro')
         #Input function to calculate MSD for a given atom
         def msd(traj, atom):
             # chlorine
             if atom == 0:
                 return md.rmsd(traj, traj, 0)
             # sodium
             elif atom == 1:
                 return md.rmsd(traj, traj, 1)
             # oxygen
             elif atom == 2:
                 return md.rmsd(traj, traj, 2)
         #Convert values to integers
         msd_cl = msd(traj, 0).astype(int)
         msd_na = msd(traj, 1).astype(int)
         msd_o = msd(traj, 2).astype(int)
         #Plot results vs time
         plt.plot(msd_o, label='0')
         plt.plot(msd_na, label='Na')
         plt.plot(msd cl, label='Cl')
         plt.xlabel('Time (ps)')
         plt.ylabel('MSD (nm)')
         plt.legend()
         plt.title('MSD of 0, Na, & Cl vs Time')
         plt.show()
```



```
In [ ]:
```