NetSci Documentation

1 Installation

```
wget https://repo.anaconda.com/miniconda/Miniconda3-latest-Linux-x86_64.sh

chmod +x Miniconda3-latest-Linux-x86_64.sh

source ~/.bashrc

conda install -c conda-forge git

git clone https://github.com/amstokely/netsci.git

cd netsci

NETSCI_ROOT=$(pwd)

conda env create -f netsci.yml

source activate netsci

mkdir ${NETSCI_ROOT}/build

cd ${NETSCI_ROOT}/build

nvcc ${NETSCI_ROOT}/build_scripts/cuda_architecture.cu -o cuda_architecture

CUDA_ARCHITECTURE=$(./cuda_architecture)
```

```
cmake .. -DCONDA_DIR=$CONDA_PREFIX -DCUDA_ARCHITECTURE=${CUDA_ARCHITECTURE}}

cmake --build . -j

make python

ctest

cd ${NETSCI_ROOT}/tests/cuarray/python

pytest

cd ${NETSCI_ROOT}/tests/netcalc/python
```

2 CuArray Tutorial

```
for i in range(len(cu_a)):
    for j in range(len(cu_a[0])):
       print(f'{cu_a[i][j]:.4f}', end=' ')
   print()
cu_a_row0_sorted = cu_a.sort(0)
for i in range(len(cu_a_row0_sorted[0])):
   print(f'{cu_a_row0_sorted[0][i]:.4f}', end=' ')
cu_a_row0_sorted_indices = cu_a.argsort(0)
for i in range(len(cu_a_row0_sorted_indices[0])):
    sorted_index = cu_a_row0_sorted_indices[0][i]
   print(f'{sorted_index} {cu_a[0][sorted_index]:.4f}', end='\n')
cu_a_sorted = cuarray.FloatCuArray()
cu_a_sorted.init(cu_a.m(), cu_a.n())
for i in range(len(cu_a)):
    cu_a_sorted[i] = cu_a.sort(i)[0]
for i in range(len(cu_a)):
   for j in range(len(cu_a[0])):
       print(f'{cu_a_sorted[i][j]:.4f}', end=' ')
   print()
cu_a_sorted_indices = cuarray.IntCuArray()
cu_a_sorted_indices.init(cu_a.m(), cu_a.n())
for i in range(len(cu_a)):
    cu_a_sorted_indices[i] = cu_a.argsort(i)[0]
for i in range(len(cu_a)):
   for j in range(len(cu_a[0])):
        sorted_index = cu_a_sorted_indices[i][j]
        print(f'\{cu_a\_sorted[i][j]:.4f\}', end=' ')
   print()
cu_a_fname = f'{os.getcwd()}/cu_a.npy'
cu_a.save(cu_a_fname)
```

```
cu_a_from_load = cuarray.FloatCuArray()
cu_a_from_load.load(cu_a_fname)
np_a = cu_a.toNumpy2D()
for i in range(m):
   for j in range(n):
      print(f'{cu_a[i][j]:.4f}', end=' ')
   print(end=' ')
   for j in range(n):
      print(f'{np_a[i][j]:.4f}', end=' ')
   print()
np_a_linear = cu_a.toNumpy1D()
for i in range(m):
   for j in range(n):
      print(f'{cu_a[i][j]:.4f}', end=' ')
   print(end=' ')
   for j in range(n):
       linear_index = i*n + j
       print(f'{np_a_linear[linear_index]:.4f}', end=' ')
   print()
cu_a_np = cuarray.FloatCuArray()
cu_a_np.fromNumpy2D(np_a)
for i in range(m):
   for j in range(n):
   print(f'{cu_a[i][j]:.4f}', end=' ')
print(end=' ')
   for j in range(n):
       print(f'{cu_a_np[i][j]:.4f}', end=' ')
   print()
cu_a_np_linear = cuarray.FloatCuArray()
cu_a_np_linear.fromNumpy1D(np_a_linear)
for i in range(m):
   for j in range(n):
       linear_index = i*n + j
       print(f'{cu_a_np_linear[0][linear_index]:.4f}', end=' ')
   print(end=' ')
   for j in range(n):
       linear_index = i*n + j
       print(f'{np_a_linear[linear_index]:.4f}', end=' ')
   print()
```

```
cu_a_copy = cuarray.FloatCuArray()
cu_a_copy.fromCuArray(
   cuArray=cu_a,
   start=0,
   end=len(cu_a)-1,
   m=cu_a.m(),
   n=cu_a.n(),
for i in range(len(cu_a)):
   for j in range(len(cu_a[0])):
       print(f'{cu_a_copy[i][j]:.4f}', end=' ')
   print(end=' ')
   for j in range(len(cu_a[0])):
       print(f'{cu_a[i][j]:.4f}', end=' ')
   print()
cu_a_from_linear = cuarray.FloatCuArray()
cu_a_from_linear.fromCuArray(
    cuArray=cu_a_np_linear,
   start=0,
   end=0,
   m=cu_a.m(),
   n=cu_a.n(),
for i in range(len(cu_a)):
   for j in range(len(cu_a[0])):
       print(f'{cu_a_from_linear[i][j]:.4f}', end=' ')
   print(end=' ')
   for j in range(len(cu_a[0])):
       print(f'{cu_a[i][j]:.4f}', end=' ')
   print()
```

3 NetChem Tutorial

```
import os
import tarfile
import netchem

# NETSCI_ROOT=<path to netsci root directory>
tutorial_directory = f'{NETSCI_ROOT}/tutorial'
os.chdir(tutorial_directory)

tutorial_tarball = tarfile.open(f'{os.getcwd()}/tutorial.tar.gz')
tutorial_tarball.extractall()
tutorial_tarball.close()
```

```
first_frame = 0
last_frame = 999
trajectory_file = f'{tutorial_directory}/pyro.dcd'
topology_file = f'{tutorial_directory}/pyro.pdb'
graph = netchem.Graph()
graph.init(
   trajectoryFile=trajectory_file,
   topologyFile=topology_file,
    firstFrame=first_frame,
   lastFrame=last_frame,
num_nodes = graph.numNodes()
print(num_nodes)
num_frames = graph.numFrames()
print(num_frames)
num_atoms = atoms.numAtoms()
print(num_atoms)
for atom in atoms:
   print(atom.index())
atom0 = atoms.at(0)
atom0_index = atom0.index()
atomO_serial = atomO.serial()
atom0_name = atom0.name()
atom0_element = atom0.element()
atom0_residue_id = atom0.residueId()
atomO_residue_name = atomO.residueName()
atom0_chain_id = atom0.chainId()
atom0_segment_id = atom0.segmentId()
atom0_mass = atom0.mass()
atom0_properties_str = f'Index: {atom0_index}\n'
atomO_properties_str += f'Serial: {atomO_serial}\n'
atomO_properties_str += f'Name: {atomO_name}\n'
atom0_properties_str += f'Element: {atom0_element}\n'
atom0_properties_str += f'Residue ID: {atom0_residue_id}\n'
atomO_properties_str += f'Residue Name: {atomO_residue_name}\n'
atomO_properties_str += f'Chain ID: {atomO_chain_id}\n'
atomO_properties_str += f'Segment ID: {atomO_segment_id}\n'
atomO_properties_str += f'Mass: {atomO_mass:.4f}'
```

```
print(atom0_properties_str)
nodes = graph.nodes()
for node in nodes:
   print(node.index())
for node in graph:
   print(node.index())
node0_tag = node0.tag()
node0_num_atoms = node0.numAtoms()
node0_index = node0.index()
node0_total_mass = node0.totalMass()
node0_atoms = node0.atoms()
node0_properties_str = f'Index: {node0_index}\n'
node0_properties_str = f'Number of Atoms: {node0_num_atoms}\n'
node0_properties_str += f'Total Mass: {node0_total_mass:.4f}\n'
node0_properties_str += f'Tag: {node0_tag}\n'
nodeO_properties_str += f'Node O Atoms:\n'
for atom in node0_atoms:
   node0_properties_str += f'
                                  Index: {atom.index()}\n'
   node0\_properties\_str \; += \; f' \qquad Residue \; Name: \; \{atom.residueName()\} \backslash n' \\
   node0_properties_str += f'
                                  Residue ID: {atom.residueId()}\n'
   node0_properties_str += f'
                                   Chain ID: {atom.chainId()}\n
   node0_properties_str += f'
                                   Segment ID: {atom.segmentId()}\n'
   node0_properties_str += f'
                                  Mass: \{atom.mass():.4f}\n\n'
print(node0_properties_str)
node_coordinates = graph.nodeCoordinates()
node0_coordinates = cuarray.FloatCuArray()
node0_coordinates.fromCuArray(
    cuArray=node_coordinates,
    start=0,
   end=0,
   m=3,
   n=num_frames,
```