Aim/Hypothesis:

- Study about the NanoParticles and the important features which need to be encountered
- Trying to understand the Converter code and Implementing a converter method for the ISubstance method which is already implemented in the CDK.

Description:

 Starting of by writing the cdkSubstanceToCMLMolecule method under libio.cml.Convertor:

```
public CMLMolecule cdkSubstanceToCMLMolecule(ISubstance model) {
      CMLMolecule cmlMolecule = new CMLMolecule();
      cmlMolecule.setConvention("Substance");
        cmlMolecule.setDictRef("cml:Substance");
      return cmlMolecule;
    }
     Ended up with the code below
     * @Author : Sina M. Nick
     * TODO: Check if ontologies are right
    public CMLMolecule cdkSubstanceToCMLMolecule(ISubstance model) {
      CMLMolecule cmlMolecule = new CMLMolecule();
      cmlMolecule.setConvention("Substance");
      cmlMolecule.setDictRef("cml:Substance");
      if(model.getID() != null)
      cmlMolecule.setId(model.getID());
      else
            cmlMolecule.setId("AN ID");
      for (int j = 0; j < model.getAtomContainerCount(); j++){</pre>
            IAtomContainer atoms = model.getAtomContainer(j);
            for (int i = 0; i < atoms.getAtomCount(); i++) {</pre>
                  IAtom cdkAtom = atoms.getAtom(i);
                  CMLAtom cmlAtom = cdkAtomToCMLAtom(atoms, cdkAtom);
                  if (atoms.getConnectedSingleElectronsCount(cdkAtom) > 0) {
cmlAtom.setSpinMultiplicity(atoms.getConnectedSingleElectronsCount(cdkAtom) +
1);
                  cmlMolecule.addAtom(cmlAtom);
            for (int i = 0: i < atoms.getBondCount(): i++) {</pre>
                CMLBond cmlBond = cdkBondToCMLBond(atoms.getBond(i));
                cmlMolecule.addBond(cmlBond);
            }
      }
```

```
return cmlMolecule;
}
```

- Not sure if the conversion between a substance to a molecule is the right idea
- Since in this case for example if you have two molecules in your substance the bond between them is not encountered.
- Maybe the joint "moleculeRefs2" need to be added to the CDK "see http://pubs.acs.org/doi/pdf/10.1021/ci8002123"
- Any ways below you find the test method which was written for this.

```
@Test
    public void testCdkSubstanceToCMLMolecule() throws IOException {
        IChemObjectBuilder builder = DefaultChemObjectBuilder.getInstance();
        ISubstance substance = builder.newInstance(ISubstance.class);
      //S03
        IAtomContainer molecule = new AtomContainer();
        Atom S1 = new Atom("S");
        Atom 02 = \text{new Atom}("0");
        Atom 03 = \text{new Atom}("0");
        Atom 04 = \text{new Atom}("0");
        molecule.addAtom(S1);
        molecule.addAtom(02);
        molecule.addAtom(03);
        molecule.addAtom(04);
        Bond b1 = new Bond(S1, 02, IBond.Order.DOUBLE);
        Bond b2 = new Bond(S1, 03, IBond.Order.DOUBLE);
        Bond b3 = new Bond(S1, 04, IBond.Order.DOUBLE);
        molecule.addBond(b1);
        molecule.addBond(b2);
        molecule.addBond(b3);
        substance.addAtomContainer(molecule);
      //XeF4
        IAtomContainer moleculeTwo = new AtomContainer();
        Atom Xe1 = new Atom("Xe");
        Atom F2 = new Atom("F");
        Atom F3 = new Atom("F");
        Atom F4 = new Atom("F");
        Atom F5 = new Atom("F");
        moleculeTwo.addAtom(Xe1);
        moleculeTwo.addAtom(F2);
        moleculeTwo.addAtom(F3);
        moleculeTwo.addAtom(F4);
        moleculeTwo.addAtom(F5);
        Bond b_1 = new Bond(Xe1, F2, IBond.Order.SINGLE);
        Bond b 2 = new Bond(Xe1, F3, IBond.Order.SINGLE);
        Bond b 3 = new Bond(Xe1, F4, IBond.Order.SINGLE);
        Bond b 4 = new Bond(Xe1, F5, IBond.Order.SINGLE);
        moleculeTwo.addBond(b 1);
        moleculeTwo.addBond(b_2);
        moleculeTwo.addBond(b 3);
        moleculeTwo.addBond(b 4);
```

```
substance.addAtomContainer(moleculeTwo);
      //Molecule set as Molecule
        IAtomContainer moleculebounder = new AtomContainer();
       Bond MoleculeBound = new Bond(F5,S1);
       moleculebounder.addBond(MoleculeBound);
        substance.addAtomContainer(moleculebounder);
       Convertor convertor = new Convertor(true, null);
       CMLMolecule convertedSubstance =
convertor.cdkSubstanceToCMLMolecule(substance);
       ByteArrayOutputStream out = new ByteArrayOutputStream();
       Serializer serializer = new Serializer(out, "UTF-8");
        serializer.write(new Document(convertedSubstance));
       out.close();
       String expected = "Unkown";
       String actual = new String(out.toByteArray());
       System.out.println(out);
       Assert.assertTrue(actual.contains(expected));
   }
```

In order to create the bond between the molecules in this test method a molecule was added a representative of a bond to the substance, however in my opinion this is not the way this should be and the code needs to be extended.

Results:

The CMLfile created from the elaborated test is shown below:

```
<?xml version="1.0" encoding="UTF-8"?>
<molecule convention="Substance" dictRef="cml:Substance" id="AN ID"</pre>
 xmlns="http://www.xml-cml.org/schema">
  <atomArrav>
    <atom id="a932951280" elementType="S" formalCharge="0"/>
    <atom id="a1334056213" elementType="O" formalCharge="0"/>
    <atom id="a1753497211" elementType="O" formalCharge="0"/>
    <atom id="a1109537496" elementType="O" formalCharge="0"/>
    <atom id="a542149962" elementType="Xe" formalCharge="0"/>
    <atom id="a141297113" elementType="F" formalCharge="0"/>
    <atom id="a1811148256" elementType="F" formalCharge="0"/>
    <atom id="a1535526014" elementType="F" formalCharge="0"/>
```