



Force Field Models in Polymer Informatics

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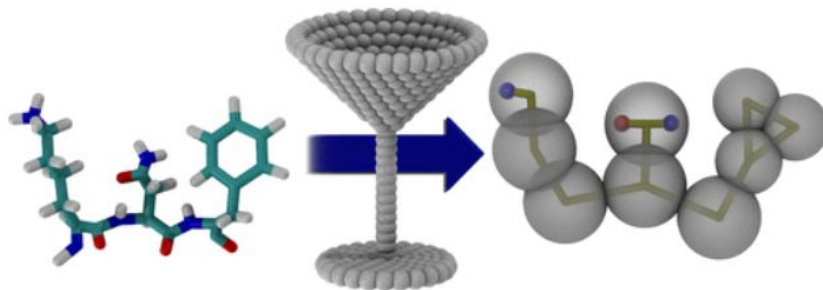
Motivation



- **Goal:** Predict the properties of a polymer given its conformation.
- Simulations are absolutely necessary.
 - Already existing options (e.g. all-atom simulations) are **slow**.
- Crystallization simulations are also slow, while involving mathematical assumptions.
- **We need a model for feature representation that allows for computational speed.**

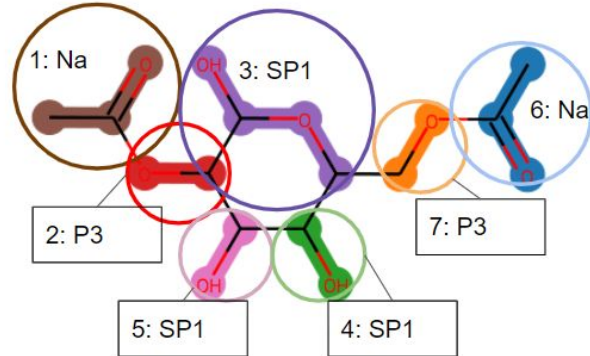
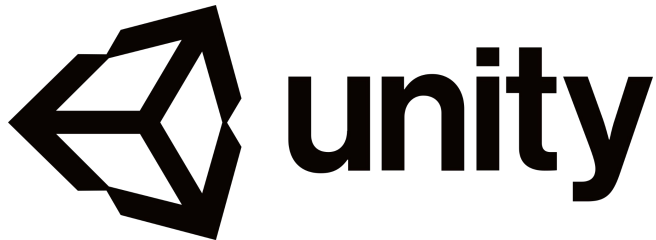
Objective: MARTINI Model

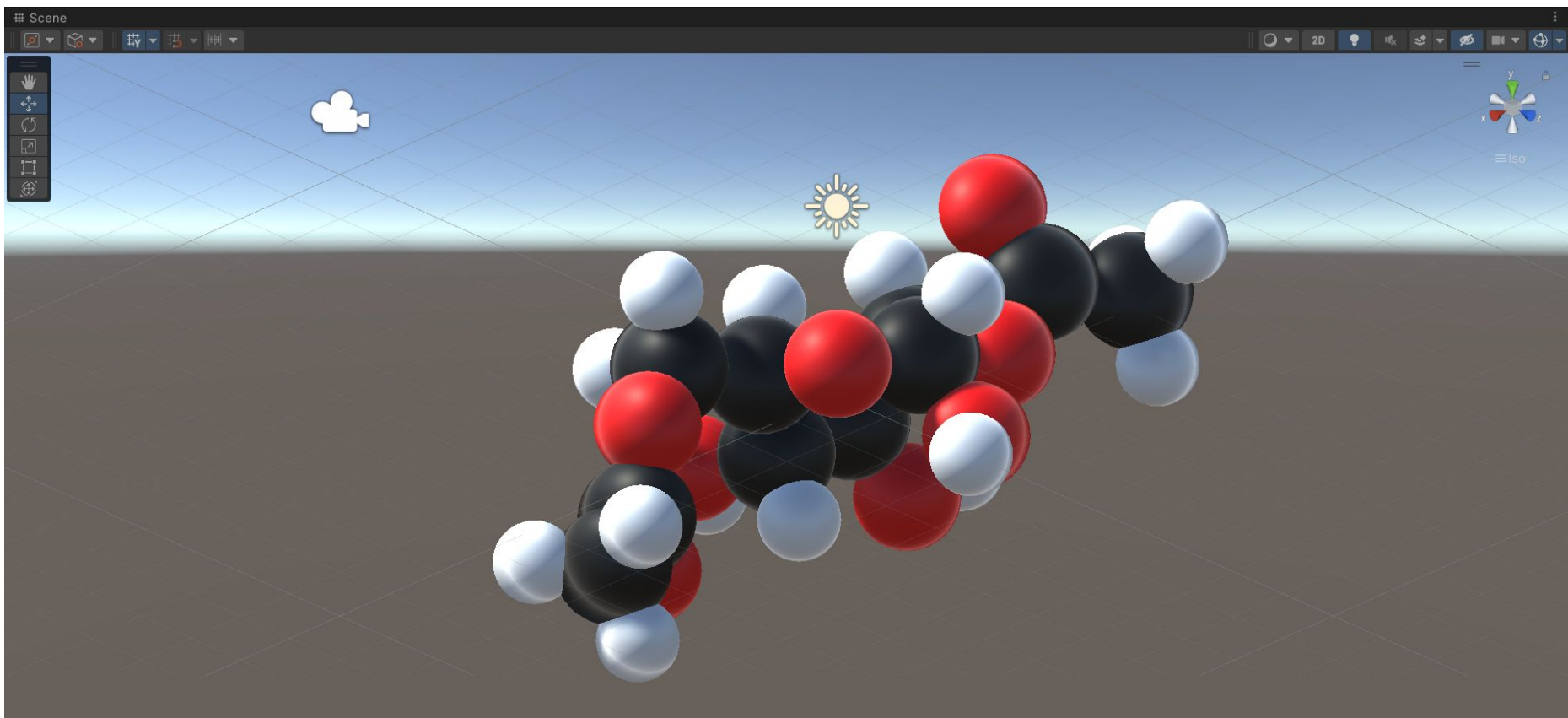
- **Main Objective:** Construct a generalize architecture to predict polymer predictions more efficiently.
- **Secondary Objective:** Decrease the resolution of our polymer chain and model molecular dynamics.
 - Use the MARTINI model; which categorizes groups of atoms (4-to-1) into certain types and subtypes. These categories are used to parametrize a set of bonded and non-bonded potential equations.
 - Model the molecular dynamics of the polymer through the use of MARTINI.



Approach

- **Construction of a pipeline:** GNN for the production of embeddings, and the use of a reinforcement learning architecture.
- **Tool:** Unity.
 - Representation of the whole molecule as a series of agents (beads) with scripts that describe the functionality of the MARTINI model.
 - Scripting code made use of C#.
 - Toy model: Cellulose Acetate





Approach

- **Embeddings:** Defined embeddings for each bead including the information necessary (i.e. type and subtype).
- **Non-bonded Interactions:**
 - Inclusion of interaction matrix the uses the embedded information to determine the parameters of Lennard-Jones equations for each pair of non-bonded beads.
- **Bonded Interactions:**
 - Inclusion of previous simulation data in order to parametrize the bonded interaction equations.
 - Use of graph-search algorithm in order to iterate through the bonded beads of the polymer chain to define both bond angle and dihedral angle conformations.

$$V_{LJ}(r_1 \dots r_n) + V_{Coul}(r_1 \dots r_n) = \sum_{i,j} 4\epsilon_{ij} \left[\left(\frac{\sigma_{ij}}{r_{ij}} \right)^{12} - \left(\frac{\sigma_{ij}}{r_{ij}} \right)^6 \right] + \sum_{i,j} \frac{q_i q_j}{4\pi\epsilon_0\epsilon_r r_{ij}}$$

$$V_b = 1/2 K_b (d_{ij} - d_b)^2,$$

$$V_a = 1/2 K_a [\cos(\varphi_{ijk}) - \cos(\varphi_a)]^2,$$


$$V_d = K_d [1 + \cos(\theta_{ijkl} - \theta_d)],$$

```

public static IReadOnlyDictionary<string, string>, int> typeMappings = new Dictionary<string, string>, int>() {
    {("Q", "da"), 0}, {("Q", "d"), 1}, {("Q", "a"), 2}, {("Q", "0"), 3},
    {("P", "5"), 4}, {("P", "4"), 5}, {("P", "3"), 6}, {("P", "2"), 7}, {("P", "1"), 8},
    {("N", "da"), 9}, {("N", "d"), 10}, {("N", "a"), 11}, {("N", "0"), 12},
    {("C", "5"), 13}, {("C", "4"), 14}, {("C", "3"), 15}, {("C", "2"), 16}, {("C", "1"), 17},
    {("Bp", "4"), 18}
};


public static readonly string[,] levels = new string[,]
{
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    {"O", "O", "O", "O", "O", "O", "O", "O", "O", "O", "O", "O", "O", "O", "O", "O", "O", "O"}
};

```


☒ Molecule
 ☐ Static

Tag Untagged
 Layer Default


Prefab Open Select Overrides


Transform

Position X 0 Y 0 Z 0

Rotation X 0 Y 0 Z 0


Scale X 1 Y 1 Z 1


☒ **New Molecule Creator (Script)**

Script NewMoleculeCreator

Creator Molecule (Martini Model)

Formula /StreamingAssets/SimInputs/am-8-04-22-A


Martini Model (Script)

Script MartiniModel

Num Monomer 75

```

public static void findAngles (Dictionary< GameObject, List<GameObject> > graph, GameObject bead, int length, List< List<GameObject> > paths, List<GameObject> path = null) {
    path = path ?? new List<GameObject>();
    if (path != null) {
        path.Add(bead);
        List<GameObject> pathr = new List<GameObject>(path);
        pathr.Reverse();
        if (path.Count == length) {
            if (!paths.Contains(path) && !paths.Contains(pathr)) {
                paths.Add(path);
            }
        }
        else {
            foreach (GameObject neighbour in graph[bead]) {
                findAngles(graph, neighbour, length, paths, path);
            }
        }
    }
}

```

Results



- Expanded my knowledge regarding molecular simulations.
 - In particular, obtained an improved understanding of the mathematical assumptions made in order to model simulations; both in atomistic and coarse-grained models.
- Improved my understanding of molecular property prediction architectures.
 - Reviewed the existing literature and gained an understanding of the advantages and disadvantages of existing approaches.
- Gained expertise in the use of Unity3D as a tool.
 - Learned how to develop C# scripts oriented towards the modelling of force fields and the implementation of reinforcement learning simulations.

Next Steps



- Finish the construction of the reinforcement learning implementation.
 - Construct a RL simulation through the use of the Unity ML-Agents toolkit by assigning C# scripts to our modeled polymer chain.
- Start the training stage of our architecture.
 - Introduce data to our model.
- Include the entanglement of polymer chains through reinforcement learning.
 - Model the dynamics involved between polymer chains in Unity.
- Compare results with the existing literature.