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Atomic partial charge prediction with graph neural networks

David Parker Dr. Cassiano Langini

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David

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Atomic partial charge

Coulomb's law:

$$|F| = k_{\rm e} \frac{|q_1||q_2|}{r^2} \tag{1}$$

Where

- Coulomb constant: $k_e = 9 \cdot 10^9 \frac{\text{N} \cdot \text{m}^2}{C^2}$
- q_1 and q_2 are the charges in Coulombs (C)
- \bullet r is the distance between the charges in meters (m)

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Atomic partial charge

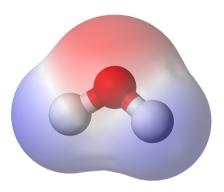


Figure: Representation of the electronic charge distribution of a water molecule. The oxygen atom has a partial negative charge, while the hydrogen atoms have partial positive charges[2].

Partial charge calculation

Experimentally (Spectroscopy):

• X-ray diffraction, Nuclear magnetic resonance, UV-Vis, etc

Computationally: Tradeoffs speed VS accuracy

• Quantum mechanics calculations: Evolution of a system through time

$$i\hbar \frac{\partial}{\partial t} |\psi(t)\rangle = \hat{H} |\psi(t)\rangle \qquad F = \frac{\partial}{\partial t} p$$

Quantum counterpart of Newton's second law of motion

• Force fields: Calculations based on a set of parameters

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Create a dataset of molecules and use the Charm General Force Field (CGenFF) to calculate partial charges for each atom

Train a graph neural network to predict atomic partial charges

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Can also be represented as a graph:

- Nodes with attributes
- Edges with attributes
- Global attributes

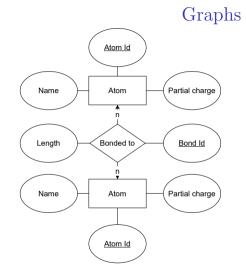


Figure: Data model for the mini-world

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Molecular Dataset: mol2 file

```
@cTRIPOS>MOLECULE
0002bf2c5ea40c3014dbcc0182ea4598
50.52
SMALL
CeenEE charge negative mean/may 2 42094 / 30 403
           0.1212 0.064 -0.3102 N.pl3 0.0
                                                  -0.163
           4.9006 -0.7556 0.5231 N.4
                                                  -0.426
            6.9083 -1.2723 0.7106 H
                                                   0.090
@<TRIPOS>BOND
1131
2 1 10 1
51 21 49 1
52 21 50 1
@/TRIPOS-MOLECULE
24d10877593ae175c68f8d409e6df5ae
26.25
```

Figure: Example of a mol2 file. Each molecule begins with a @<TRIPOS>MOLECULE tag, then the different fields are separated by their own tags.

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Molecular Dataset: Molecules

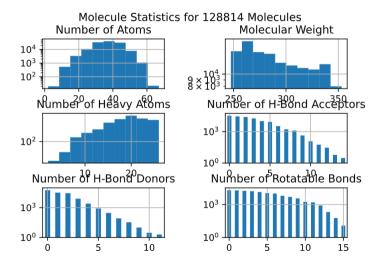


Figure: Distributions of different molecular properties in the dataset.

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Molecular Dataset: Atoms

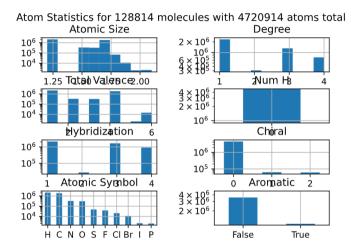


Figure: Distributions of different atomic properties in the dataset.

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Message-passing neural network

$$\mathbf{e}'_{k} = \phi^{e} \left(\mathbf{e}_{k}, \mathbf{v}_{r_{k}}, \mathbf{v}_{s_{k}}, \mathbf{u} \right) \quad \overline{\mathbf{e}}'_{i} = \rho^{e \to v} \left(E'_{i} \right)$$

$$\mathbf{v}'_{i} = \phi^{v} \left(\overline{\mathbf{e}}'_{i}, \mathbf{v}_{i}, \mathbf{u} \right) \qquad \overline{\mathbf{e}}' = \rho^{e \to u} \left(E' \right)$$

$$\mathbf{u}' = \phi^{u} \left(\overline{\mathbf{e}}', \overline{\mathbf{v}}', \mathbf{u} \right) \qquad \overline{\mathbf{v}}' = \rho^{v \to u} \left(V' \right)$$

$$m = 0 \qquad m = 1 \qquad m = 2 \qquad m = 3$$

Figure: Message-passing neural network. Three update functions ϕ and three aggregation functions ρ are used to update the edge, node and global attributes[1].

References

Message-passing neural network

Node features: Element type

- Preprocess layer: 2 linear layers with batch normalization and ReLU activation functions to embed the element type into an internal continuous representation.
- Edge function layer: several linear layers with batch normalization and ReLU activation functions to compute edge features based on the node features of the adjacent nodes.
- GNN layer: applies a convolution operation to update the node features using the edge features as weights.
- GRU layer: applies a gated recurrent unit to update the node features.
- Predict layer: 2 linear layers with batch normalization and ReLU activation functions, followed by a linear layer with a single output.

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Benchmarks

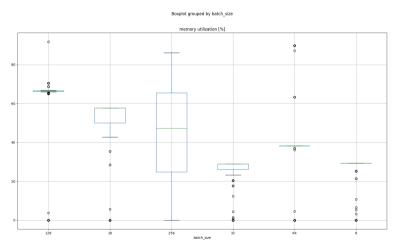


Figure: Benchmarks of GPU utilisation for different batch sizes.

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Absolute errors

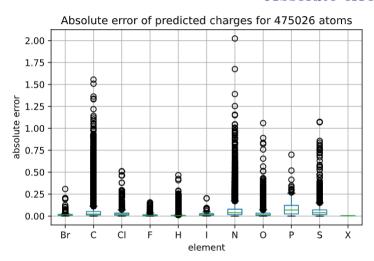


Figure: Distributions of the absolute errors of the predicted partial charges for each

type of atom.

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Root mean squared error

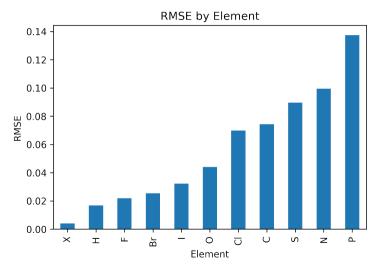


Figure: Root mean squared error for each type of atom.

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Mean absolute error

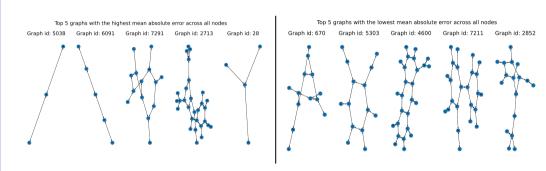


Figure: Visualizations of the graphs with the highest and lowest MAE.

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Our results compared to the literature

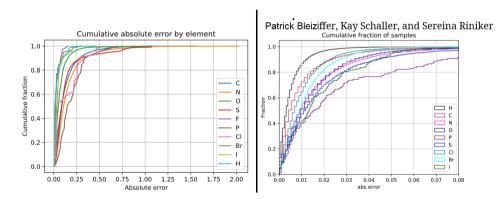


Figure: Cumulative fraction of each type of atom as a function of the absolute error. Our results on the left, the results of the literature on the right[3].

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Dataset generation

- The dataset is technically easy to generate:
 - SQL queries to extract the data from the database
 - Use a force field to compute the partial charges
- The hard part is to choose relevant molecules:
 - \bullet We chose drug-like molecules. Molecular weight around 300 g/mol, specific number of rotatable bonds, etc. (see Figure 4)

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- Choice of the model:
 - We went with a MPNN as introduced by Gilmer et al.[4]
- Achitechture of the model:
 - We would like to add physical constraints to the model, so that the predicted partial charges make more sense

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Conclusion

- We have generated a dataset of 8000 molecules with partial charges
- We have trained a MPNN to predict the partial charges
- The model has a RMSE of 0.054
 - Elementary charge of a proton: 1.6×10^{-19} C

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Atomic

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