pjztestgat-copy

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1 Hands on with Graph Neural Networks

1.1 Installing Pytorch Geometric and RDKit

- Pytorch Geometric => Build Graph Neural Network
- RDKit => Handle Molecule Data

Note: Currently I didn't find a way to set a specific python version in colab. However, when installing rdkit, only specific python versions are supported... Hence it cannot be ensured that the notebook runs properly. For example Python 3.7.10 is not supported.

```
[15]: # Enforce pytorch version 1.6.0
import torch
# if torch.__version__ != '1.6.0':
# !pip uninstall torch -y
# !pip uninstall torchvision -y
# !pip install torch=1.6.0
# !pip install torchvision==0.7.0

# # Check pytorch version and make sure you use a GPU Kernel
# !python -c "import torch; print(torch.__version__)"
# !python -c "import torch; print(torch.version.cuda)"
# !python --version
# !nvidia-smi
```

Make sure you clicked "RESTART RUNTIME" above (if torch version was different)!

```
[16]: #@title
    # # Install rdkit
    # import sys
# import os
# import requests
# import subprocess
# import shutil
# from logging import getLogger, StreamHandler, INFO
```

```
# logger = getLogger(__name__)
# logger.addHandler(StreamHandler())
# logger.setLevel(INFO)
# def install(
#
          chunk size=4096,
#
          file_name="Miniconda3-latest-Linux-x86_64.sh",
#
          url base="https://repo.continuum.io/miniconda/",
          conda\_path = os.path. \ expanduser(os.path.join("~", "miniconda")),
#
#
          rdkit version=None.
          add_python_path=True,
#
          force=False):
      """install rdkit from miniconda
#
#
#
      import\ rdkit\_installer
#
      rdkit_installer.install()
#
      11 11 11
#
#
      python_path = os.path.join(
#
          conda_path,
#
          "lib",
#
          "python{0}.{1}".format(*sys.version info),
#
          "site-packages",
#
      )
#
      if add_python_path and python_path not in sys.path:
#
          logger.info("add {} to PYTHONPATH".format(python_path))
#
          sys.path.append(python_path)
#
      if os.path.isdir(os.path.join(python_path, "rdkit")):
#
          logger.info("rdkit is already installed")
#
          if not force:
#
              return
          logger.info("force re-install")
#
#
      url = url base + file name
#
      python_version = "{0}.{1}.{2}".format(*sys.version_info)
      logger.info("python version: {}".format(python_version))
#
#
      if os.path.isdir(conda_path):
#
          logger.warning("remove current miniconda")
#
          shutil.rmtree(conda_path)
#
      elif os.path.isfile(conda_path):
```

```
logger.warning("remove {}".format(conda_path))
#
          os.remove(conda path)
      logger.info('fetching installer from {}'.format(url))
#
#
      res = requests.get(url, stream=True)
#
      res.raise_for_status()
#
      with open(file_name, 'wb') as f:
#
          for chunk in res.iter_content(chunk_size):
              f.write(chunk)
#
      logger.info('done')
      logger.info('installing miniconda to {}'.format(conda_path))
#
      subprocess.check_call(["bash", file_name, "-b", "-p", conda_path])
#
      logger.info('done')
      logger.info("installing rdkit")
#
#
      subprocess.check_call([
          os.path.join(conda_path, "bin", "conda"),
#
#
          "install",
#
          "--yes",
#
          "-c", "rdkit",
#
          "python == 3.7.3",
          "rdkit" if rdkit_version is None else "rdkit=={}".
 → format(rdkit version)])
      logger.info("done")
      import rdkit
      logger.info("rdkit-{} installation finished!".format(rdkit.__version__))
# if __name__ == "__main__":
      install()
⇔between the Python version, CUDA or torch
import torch
```

```
[17]: # If something breaks in the notebook it is probably related to a mismatch_
between the Python version, CUDA or torch
import torch
# pytorch_version = f"torch-{torch.__version__}.html"
# !pip install --no-index torch-scatter -f https://pytorch-geometric.com/whl/
$pytorch_version
# !pip install --no-index torch-sparse -f https://pytorch-geometric.com/whl/
$pytorch_version
# !pip install --no-index torch-cluster -f https://pytorch-geometric.com/whl/
$pytorch_version
# !pip install --no-index torch-spline-conv -f https://pytorch-geometric.com/
$whl/$pytorch_version
# !pip install torch-geometric
```

1.2 Background info on the Dataset

In the following we will use a dataset provided in the dataset collection of PyTorch Geometric (Here you find all datasets). The Dataset comes from the MoleculeNet collection, which can be found here.

"ESOL is a small dataset consisting of water solubility data for 1128 compounds. The dataset has been used to train models that estimate solubility directly from chemical structures (as encoded in SMILES strings). Note that these structures don't include 3D coordinates, since solubility is a property of a molecule and not of its particular conformers."

>>> Machine Learning task: How are different molecules dissolving in water?

Source: https://www.differencebetween.com/difference-between-solubility-and-solubility-product/

1.2.1 SMILES representation and important sidenodes

Source: https://medium.com/@sunitachoudhary 103/generating-molecules-using-a-char-rnn-in-pytorch-16885fd9394b

- Using the plain SMILES string as input is not suitable
- This will not consider the molecule structure but rather the grammar of the SMILES string
- The SMILES string can be different for a molecule, depending on the notation (a unique molecule can have multiple SMILES strings)
- Chemical graphs however, are invariant to permutations -> Graph Neural Networks

1.3 Looking into the Dataset

```
[18]: import rdkit
from torch_geometric.datasets import MoleculeNet

# Load the ESOL dataset
data = MoleculeNet(root=".", name="ESOL")
data
```

[18]: ESOL(1128)

Note: There seems to be a change in the Dataset class and somehow the target dim now equals 734 instead of one. You can simply ignore it at this point. :)

```
[19]: # Investigating the dataset
print("Dataset type: ", type(data))
print("Dataset features: ", data.num_features)
print("Dataset target: ", data.num_classes)
print("Dataset length: ", data.len)
```

```
print("Dataset sample: ", data[0])
      print("Sample nodes: ", data[0].num_nodes)
      print("Sample edges: ", data[0].num_edges)
      # edge_index = graph connections
      # smiles = molecule with its atoms
      \# x = node \ features \ (32 \ nodes \ have \ each \ 9 \ features)
      # y = labels (dimension)
     Dataset type: <class 'torch_geometric.datasets.molecule_net.MoleculeNet'>
     Dataset features: 9
     Dataset target: 734
     Dataset length: <bound method InMemoryDataset.len of ESOL(1128)>
     Dataset sample: Data(x=[32, 9], edge_index=[2, 68], edge_attr=[68, 3],
     smiles='OCC3OC(OCC2OC(OC(C#N)c1ccccc1)C(O)C(O)C2O)C(O)C(O)C3O', y=[1, 1])
     Sample nodes: 32
     Sample edges: 68
[20]: # Investigating the features
      # Shape: [num_nodes, num_node_features]
      data[0].x
[20]: tensor([[8, 0, 2, 5, 1, 0, 4, 0, 0],
              [6, 0, 4, 5, 2, 0, 4, 0, 0],
              [6, 0, 4, 5, 1, 0, 4, 0, 1],
              [8, 0, 2, 5, 0, 0, 4, 0, 1],
              [6, 0, 4, 5, 1, 0, 4, 0, 1],
              [8, 0, 2, 5, 0, 0, 4, 0, 0],
              [6, 0, 4, 5, 2, 0, 4, 0, 0],
              [6, 0, 4, 5, 1, 0, 4, 0, 1],
              [8, 0, 2, 5, 0, 0, 4, 0, 1],
              [6, 0, 4, 5, 1, 0, 4, 0, 1],
              [8, 0, 2, 5, 0, 0, 4, 0, 0],
              [6, 0, 4, 5, 1, 0, 4, 0, 0],
              [6, 0, 2, 5, 0, 0, 2, 0, 0],
              [7, 0, 1, 5, 0, 0, 2, 0, 0],
              [6, 0, 3, 5, 0, 0, 3, 1, 1],
              [6, 0, 3, 5, 1, 0, 3, 1, 1],
              [6, 0, 3, 5, 1, 0, 3, 1, 1],
              [6, 0, 3, 5, 1, 0, 3, 1, 1],
              [6, 0, 3, 5, 1, 0, 3, 1, 1],
              [6, 0, 3, 5, 1, 0, 3, 1, 1],
              [6, 0, 4, 5, 1, 0, 4, 0, 1],
              [8, 0, 2, 5, 1, 0, 4, 0, 0],
              [6, 0, 4, 5, 1, 0, 4, 0, 1],
              [8, 0, 2, 5, 1, 0, 4, 0, 0],
              [6, 0, 4, 5, 1, 0, 4, 0, 1],
```

```
[8, 0, 2, 5, 1, 0, 4, 0, 0], [6, 0, 4, 5, 1, 0, 4, 0, 1], [8, 0, 2, 5, 1, 0, 4, 0, 0], [6, 0, 4, 5, 1, 0, 4, 0, 1], [8, 0, 2, 5, 1, 0, 4, 0, 0], [6, 0, 4, 5, 1, 0, 4, 0, 1], [8, 0, 2, 5, 1, 0, 4, 0, 0]])
```

```
[21]: # Investigating the edges in sparse COO format
      # Shape [2, num_edges]
      data[0].edge_index.t()
[21]: tensor([[ 0,
                  1],
              [ 1,
                   0],
              [ 1,
                   2],
              [ 2,
                   1],
              [2, 3],
              [2,30],
              [ 3,
                   2],
              [3, 4],
              [4, 3],
              [4, 5],
              [4, 26],
              [5, 4],
              [5, 6],
              [6, 5],
              [6, 7],
              [7, 6],
              [7,
                   8],
              [7, 24],
              [8, 7],
              [8, 9],
              [9, 8],
              [9, 10],
              [9, 20],
              [10, 9],
              [10, 11],
              [11, 10],
              [11, 12],
              [11, 14],
              [12, 11],
              [12, 13],
```

[13, 12], [14, 11],

```
[14, 15],
[14, 19],
[15, 14],
[15, 16],
[16, 15],
[16, 17],
[17, 16],
[17, 18],
[18, 17],
[18, 19],
[19, 14],
[19, 18],
[20, 9],
[20, 21],
[20, 22],
[21, 20],
[22, 20],
[22, 23],
[22, 24],
[23, 22],
[24, 7],
[24, 22],
[24, 25],
[25, 24],
[26, 4],
[26, 27],
[26, 28],
[27, 26],
[28, 26],
[28, 29],
[28, 30],
[29, 28],
[30, 2],
[30, 28],
[30, 31],
[31, 30]])
```

[22]: data[0].y

[22]: tensor([[-0.7700]])

In the following we will perform predictions based on the graph level. This

- 1. List item
- 2. List item

means we have one y-label for the whole graph, as shown on the left image below. The right image would be node-level-predictions.

1.4 Converting SMILES to RDKit molecules - Visualizing molecules

Next we want to have our SMILES molecules as graphs...

```
[23]: data[0]["smiles"]
```

[23]: 'DCC3DC(DCC2DC(DC(C#N)c1ccccc1)C(D)C(D)C2D)C(D)C(D)C3D '

```
[24]: from rdkit import Chem
from rdkit.Chem.Draw import IPythonConsole
molecule = Chem.MolFromSmiles(data[0]["smiles"])
molecule
```

[24]:

```
[25]: type(molecule)
```

[25]: rdkit.Chem.rdchem.Mol

- We can also obtain the features from this RDKit representation
- It tells us everything we need to know e.g. atom features (type, ...), edges, ...
- In our case however, It's even easier as we have the information explicitly given already in the dataset
- Otherwise we would calculate the node features from those atom properties
- -> For datasets containing SMILES representations this would be the way to go

1.5 Implementing the Graph Neural Network

Building a Graph Neural Network works the same way as building a Convolutional

Neural Network, we simple add some layers.

The GCN simply extends torch.nn.Module. GCNConv expects: - in_channels = Size of each input sample. - out_channels = Size of each output sample.

We apply three convolutional layers, which means we learn the information about 3 neighbor hops. After that we apply a pooling layer to combine the information of the individual nodes, as we want to perform graph-level prediction.

Always keep in mind that different learning problems (node, edge or graph prediction) require different GNN architectures.

For example for node-level prediction you will often encounter masks. For graph-level predictions on the other hand you need to combine the node embeddings.

```
[26]: import torch
      import torch.nn as nn
      import torch.nn.functional as F
      from torch_geometric.nn import MessagePassing
      from torch_geometric.utils import add_self_loops
      class PJzGAT(MessagePassing):
          def __init__(self, in_channels, out_channels):
              super(PJzGAT, self).__init__(aggr='add')
              self.in_channels = in_channels
              self.out_channels = out_channels
              self.W = nn.Parameter(torch.zeros(size=(in_channels, out_channels)))
              nn.init.xavier uniform (self.W.data, gain=1.414)
              self.a = nn.Parameter(torch.zeros(size=(2*out_channels, 1)))
              nn.init.xavier_uniform_(self.a.data, gain=1.414)
              self.leakyrelu = nn.LeakyReLU(0.2)
              self.mlp = nn.Sequential(
                  nn.Linear(out_channels, out_channels), # Adjust input channels for_
       \hookrightarrowMLP
                  nn.ReLU(),
                  nn.Linear(out_channels, out_channels),
                  nn.ReLU()
              )
          def forward(self, x, edge index):
              h = torch.matmul(x, self.W) # Apply linear transformation
              edge_index, _ = add_self_loops(edge_index, num_nodes=x.size(0))
              # Compute attention coefficients
              a_input = torch.cat([h.repeat(1, x.size(0)).view(x.size(0) * x.size(0),__
       \rightarrow-1), h.repeat(x.size(0), 1)], dim=1).view(x.size(0), -1, 2 * self.
       →out_channels)
```

```
e = self.leakyrelu(torch.matmul(a_input, self.a).squeeze())
    # Apply mask and activation function
   row , col = edge_index
    zero_vec = -9e15 * torch.ones_like(e)
   mask = torch.zeros_like(e)
   mask[edge_index[0], edge_index[1]] = 1
   attention = mask * e
    attention = F.leaky_relu(attention, negative_slope=0.2)
    attention = custom_exp(edge_index, attention)
    # Perform message passing with attention
   out = self.propagate(edge_index, x=h, attention=attention)
    # Optionally apply MLP after aggregation
    out = self.mlp(out)
    return out
def message(self, x_j, attention):
    # Compute messages with attention coefficients
   attention = attention.view(-1)
    attention = attention[attention != 0].view(-1, 1)
   buffer = attention.view(-1, 1) * x_j
    return buffer
```

```
[27]: import torch
      from torch.nn import Linear
      import torch.nn.functional as F
      import torch.nn as nn
      from torch_geometric.nn import GCNConv, TopKPooling, global_mean_pool , GATConv
      from torch_geometric.nn import global_mean_pool as gap, global_max_pool as gmp
      embedding_size = 64
      import torch
      def custom_exp(edge_index, attention_scores):
          # Extract the number of nodes (N) from the shape of the attention score
       \hookrightarrow matrix
          N = attention_scores.size(0)
          # Initialize an empty tensor to store the attention coefficients
          attention_coefficients = torch.zeros_like(attention_scores)
          # Loop over each source node index (row index of edge_index)
          for i in range(edge_index.size(1)):
              src_idx, dst_idx = edge_index[:, i]
```

```
numerator = torch.exp(attention_scores[src_idx, dst_idx])
        # get an array with attention scores of neighbours of src_idx
        neighbours = edge_index[1, edge_index[0] == src_idx]
        denominator = torch.sum(torch.exp(attention_scores[src_idx,__
 →neighbours]))
        attention_coefficients[src_idx, dst_idx] = numerator / denominator
    return attention_coefficients
# Example usage:
# edge_index = torch.tensor([[0, 1, 1, 2], [1, 0, 2, 1]])
# attention scores = torch.randn(N, N) # Replace with your attention score_{\sqcup}
 \rightarrow matrix
# attention_coefficients = custom_exp(edge_index, attention_scores)
torch.manual_seed(42)
class GCN(torch.nn.Module):
    def __init__(self):
        # Init parent
        super(GCN, self).__init__()
        torch.manual_seed(42)
        # GCN layers
        self.initial_conv = GATConv(data.num_features, embedding_size)
        self.conv1 = GCNConv(embedding size, embedding size)
        self.conv2 = GCNConv(embedding_size, embedding_size)
        self.conv3 = GCNConv(embedding_size, embedding_size)
        # Output layer
        self.out = Linear(embedding_size*2, 1)
    def forward(self, x, edge_index, batch_index):
        # First Conv layer
        hidden = nn.Linear(data.num_features, data.num_features)(x)
        hidden = self.initial_conv(x, edge_index)
        hidden = F.tanh(hidden)
        # Other Conv layers
        # hidden = self.conv1(hidden, edge_index)
        # hidden = F.tanh(hidden)
        # hidden = self.conv2(hidden, edge_index)
        # hidden = F.tanh(hidden)
        # hidden = self.conv3(hidden, edge_index)
        # hidden = F.tanh(hidden)
```

```
# Global Pooling (stack different aggregations)
        hidden = torch.cat([gmp(hidden, batch_index),
                             gap(hidden, batch_index)], dim=1)
         # Apply a final (linear) classifier.
        out = self.out(hidden)
        return out, hidden
model = GCN()
print(model)
print("Number of parameters: ", sum(p.numel() for p in model.parameters()))
GCN(
  (initial_conv): GATConv(9, 64, heads=1)
  (conv1): GCNConv(64, 64)
  (conv2): GCNConv(64, 64)
  (conv3): GCNConv(64, 64)
  (out): Linear(in_features=128, out_features=1, bias=True)
Number of parameters:
```

- We could also reduce the embeddings, but as we have large molecules we use 64
- The more layers we add, the more information we get about the graph
- For the regression problem we use a Linear layer as final output layer
- We try to use not too many parameters, as we only have ~1k samples

1.6 Training the GNN

```
batch_size=NUM_GRAPHS_PER_BATCH, shuffle=True)
def train(data):
    # Enumerate over the data
    for batch in loader:
      # Use GPU
      batch.to(device)
      # Reset gradients
      optimizer.zero grad()
      # Passing the node features and the connection info
      pred, embedding = model(batch.x.float(), batch.edge_index, batch.batch)
      # Calculating the loss and gradients
      loss = loss_fn(pred, batch.y)
      loss.backward()
      # Update using the gradients
      optimizer.step()
    return loss, embedding
print("Starting training...")
losses = []
for epoch in range(2000):
    loss, h = train(data)
    losses.append(loss)
    if epoch % 100 == 0:
      print(f"Epoch {epoch} | Train Loss {loss}")
```

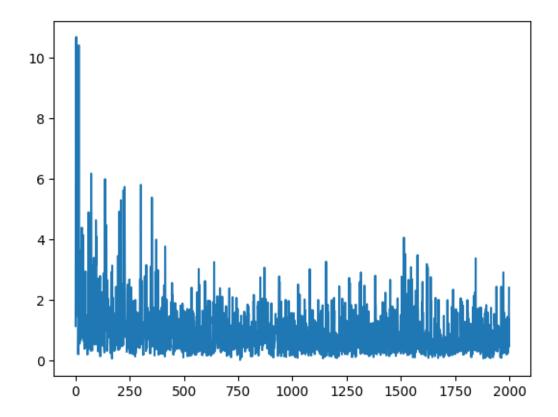
Starting training...

```
Epoch 0 | Train Loss 1.127703309059143
Epoch 100 | Train Loss 0.5818046927452087
Epoch 200 | Train Loss 1.2429836988449097
Epoch 300 | Train Loss 2.405256986618042
Epoch 400 | Train Loss 0.2629077732563019
Epoch 500 | Train Loss 0.7952205538749695
Epoch 600 | Train Loss 0.8865408301353455
Epoch 700 | Train Loss 0.8518815040588379
Epoch 800 | Train Loss 0.5335233807563782
Epoch 900 | Train Loss 0.1580432504415512
Epoch 1000 | Train Loss 0.5037855505943298
Epoch 1100 | Train Loss 0.6029763221740723
Epoch 1200 | Train Loss 0.9074375033378601
Epoch 1300 | Train Loss 1.4717516899108887
Epoch 1400 | Train Loss 0.5431175231933594
Epoch 1500 | Train Loss 0.22179903090000153
Epoch 1600 | Train Loss 2.588500738143921
Epoch 1700 | Train Loss 0.4345895051956177
Epoch 1800 | Train Loss 0.3560497760772705
Epoch 1900 | Train Loss 0.5723695755004883
```

1.6.1 Visualizing the Training loss

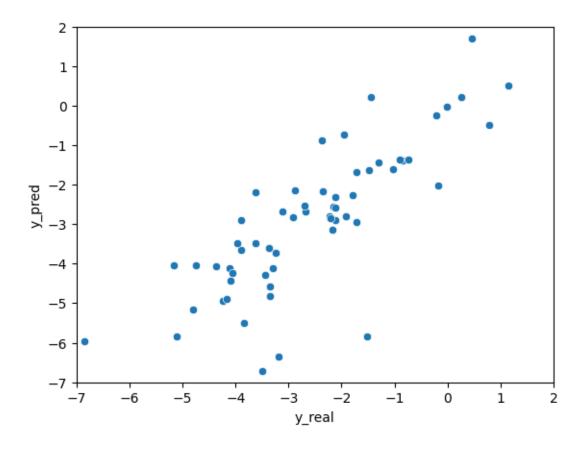
```
[29]: # Visualize learning (training loss)
import seaborn as sns
losses_float = [float(loss.cpu().detach().numpy()) for loss in losses]
loss_indices = [i for i,l in enumerate(losses_float)]
plt = sns.lineplot(losses_float)
plt
```

[29]: <Axes: >



1.6.2 Getting a test prediction

```
df["y_real"] = test_batch.y.tolist()
         df["y_pred"] = pred.tolist()
     df["y_real"] = df["y_real"].apply(lambda row: row[0])
     df["y_pred"] = df["y_pred"].apply(lambda row: row[0])
[30]:
         y_real
                  y_pred
     0
          -1.92 -2.806550
     1
          -1.72 -2.935613
     2
          -3.24 -3.730911
     3
          -6.86 -5.973111
          -0.85 -1.396756
     . .
           •••
     59
         -1.52 -5.830583
     60
         -2.38 -0.867452
     61
         -3.84 -5.504718
     62
         -2.18 -3.130320
          -0.22 -0.240277
     63
     [64 rows x 2 columns]
[31]: plt = sns.scatterplot(data=df, x="y_real", y="y_pred")
     plt.set(xlim=(-7, 2))
     plt.set(ylim=(-7, 2))
     plt
[31]: <Axes: xlabel='y_real', ylabel='y_pred'>
```



1.7 Improving the model / More to play around with

For example you can add: - Dropouts - Other (more intelligent) Pooling Layers (all layers here: https://pytorch-geometric.readthedocs.io/en/latest/modules/nn.html#) - Global Pooling Layers - Batch Normalization - More MP layers - Other hidden state sizes - Test metrics (test error) and Hyperparameter optimization - ...

https://github.com/rusty1s/pytorch_geometric/tree/master/examples

[]: