```
import torch
import torch.nn as nn
import tifffile
import torch.nn.parallel
import torch.utils.data
import os
import matplotlib.pyplot as plt
import numpy as np
import glob
#import torch.distributions as td
from reconstruction.dcgan test import Generator
import torch.backends.cudnn as cudnn
from skimage import measure
import h5py
from pylab import gca
from analyze_pore_samples.plotting utils import improve pairplot
def read file(fname):
   # fname = os.path.join(dir, filename)
    f = h5py.File(fname, 'r')
    return f['data']
def plt scatter matrices(prior realizations, post realizations, variable list,
plt prior=True, plt post=True, show = False):
   Plot the prior and posterior scatter matrices (on top of each other) for a
*variable list*.
    This is from a dictionary of *prior realizations* and *post realizations*.
    from pandas import DataFrame
    from pandas.plotting import scatter matrix
    import seaborn as sns
   def plt corr sns(corr):
        f, ax = plt.subplots(figsize=(10, 8))
        sns.heatmap(corr, mask=np.zeros like(corr, dtype=np.bool),
cmap=sns.diverging palette(220, 10, as cmap=True),
                    square=True, ax=ax, vmin = -1.0, vmax = 1.0)
    shape prior = np.array(prior realizations[variable list[0]]).shape
    shape post = np.array(post realizations[variable list[0]]).shape
   prior data = prior realizations.copy()
   prior data['Case'] = ['Ground Truth']*shape prior[0]#np.zeros(shape prior)
   post_data = post_realizations.copy()
   post data['Case'] = ['Reconstructed']*shape post[0]#np.ones(shape post)
   # breakpoint()
    if plt prior and plt post:
        for key in prior data:
            prior data[key] = np.append(prior data[key],post data[key])
        data = DataFrame(prior data)
   elif plt post:
        data = DataFrame(post_data)
   else:
        data = DataFrame(prior data)
```

```
52
53
       colors = ['blue', 'red']
54
      # breakpoint()
55
      plt.figure(figsize = [8,6], dpi = 150)
      pplot = sns.pairplot(data, hue = 'Case', kind = 'kde', dropna = True,
56
  diag kws=dict(common norm= 'False'))
57
      # breakpoint()
       replacements = {'Volume': r'$log {10} Volume$', 'Anisotropy':
58
                 'Orientation': r'Orientation, $\theta$ [rad]'}
59
       improve pairplot(pplot, replacements = replacements)
60
      plt.savefig('./reconstruction/gan/figures/pairplot gan.png')
61
62
           plt.show()
63
64 def save pore(pore identity, epoch = None, real = False, fname = '',
  pore matrix = None, value =0):
65
       if real:
66
           filename = './analyze pore samples/results/individual pore samples
  /partsample0/pore original {}.hdf5'.format(pore identity)
67
68
           filename = './reconstruction/gan/saved generated pores/generator '
  + str(pore identity) + '.hdf5'
69
70
          # breakpoint()
71
72
      data = np.array(read file(filename))
73
       labeled = measure.label(data)
74
       pores = measure.regionprops(labeled)
75
      pore idx = np.argmax([pore.area for pore in pores])
76
      data[labeled != pores[pore idx].label] = 0
77
       zmin = np.min(np.where(data)[2]) - 1
78
       zmax = np.max(np.where(data)[2]) + 1
79
      xmin = np.min(np.where(data)[0]) - 1
80
      xmax = np.max(np.where(data)[0]) + 1
81
      ymin = np.min(np.where(data)[1]) - 1
82
      ymax = np.max(np.where(data)[1]) + 1
83
      bounds min = np.min([xmin, ymin, zmin])
84
      bounds max = np.max([zmax, ymax, xmax])
85
86
      if not real:
87
           directory saved = './reconstruction/saved generated pores
  /pore_{}'.format(pore_identity) + fname
88
      else:
89
           directory_saved = './reconstruction/saved_example_pores
  /pore_{}'.format(pore_identity) + fname
90
91
      os.makedirs(directory saved, exist ok=True)
92
  plt.figure().add subplot(projection='3d').voxels(data[bounds min:bounds max,
  bounds min:bounds max, bounds min:bounds max], edgecolor = 'k')
93
       plt.title( '{0:.2f}'.format(value) + ' ' +
  str(pore matrix[pore identity]))
      plt.tight layout()
94
95
      plt.savefig(directory_saved+ '/pore.png')
96
      plt.clf()
```

```
97
98 def query_pore(volume, anis, angle, pore matrix):
        return np.argmin(np.abs(volume -
   pore matrix[:,0])/np.mean(pore matrix[:,0]) + np.abs(anis -
   pore matrix[:,2])/np.mean(pore matrix[:,2]) + np.abs(angle -
   pore matrix[:,3])/np.mean(pore matrix[:,3]))
100
101
102 params = {
103
        'imsize' : 64,# Spatial size of training images. All images will be
    resized to this size during preprocessing.
        'nc' : 2,# Number of channles in the training images. For coloured images
104
    this is 3.
105
        'nz' : 100,# Size of the Z latent vector
106
        'ngf' : 64,# Size of feature maps in the generator. The filtes will be
   multiples of this.
        'ndf' : 16, # Size of features maps in the discriminator. The filters will
107
   be multiples of this.
108
        'ngpu': 1, # Number of GPUs to be used
109
        'nepochs': 15,# Number of training epochs.
110
        'lr' : 0.0002,# Learning rate for optimizers
111
        'betal' : 0.5,# Betal hyperparam for Adam optimizer
112
        'alpha' : 1,# Size of z space
113
        'stride' : 16,# Stride on image to crop
        'num samples' : 179}# Save step.
114
115
116 def legend(location = 'best', fontsize = 8):
117
        plt.legend(loc = location, fontsize = fontsize, frameon = False)
118 def unit vector(vector):
       # """ Returns the unit vector of the vector. """
119
120
        return vector / np.linalg.norm(vector)
121 def angle between(v1, v2):
122
        """ Returns the angle in radians between vectors 'v1' and 'v2'::
123
124
                >>> angle_between((1, 0, 0), (0, 1, 0))
125
                1.5707963267948966
126
                >>> angle between((1, 0, 0), (1, 0, 0))
127
                0.0
128
                >>> angle between((1, 0, 0), (-1, 0, 0))
129
                3.141592653589793
        0.00
130
131
       v1 u = unit vector(v1)
132
       v2 u = unit vector(v2)
133
        return np.arccos(np.clip(np.dot(v1_u, v2_u), -1.0, 1.0))
134 def extract pore(im):
135
       size = 32
136
        im = np.array(im.cpu().detach().numpy(), dtype = 'int')
137
       props = measure.regionprops(measure.label(im))
138
       p idx = np.argmax([pore.area for pore in props])
139
       pore 3d = np.zeros((size*2, size*2, size*2))
140
       xdist = props[p idx].slice[0].stop - props[p idx].slice[0].start
141
       ydist = props[p idx].slice[1].stop - props[p idx].slice[1].start
        zdist = props[p idx].slice[2].stop - props[p idx].slice[2].start
142
143
144
       pore 3d[size - xdist//2: size+(xdist-xdist//2), size-ydist//2:size+(ydist-
```

```
ydist//2), size-zdist//2:size+(zdist-zdist//2)] =
   np.array(props[p_idx].image, dtype = 'float')
        # breakpoint()
145
146
        return pore 3d
147 def analyze pore(im):
148
       voxelsize = 3.49
149
150
        realvols = []
151
        realorientations = []
152
        realanisotropies = []
153
        realmin_axis_l = []
154
        realmaj axis l = []
155
        pores = measure.regionprops(measure.label(im))
156
        if len(pores) > 1:
157
            breakpoint()
       # breakpoint()
158
159
        if len(pores) == 0:
160
            # breakpoint()
161
            return [-1,0,0,0,0]
162
        # try1
163
        pore idx = np.argmax([pore.area for pore in pores])
164
        realvols.append(pores[pore idx].area)
165
        if pores[pore idx].area< 3:</pre>
166
            return [-1,0,0,0,0]
167
        realmaj axis l.append((pores[pore idx].major axis length)*voxelsize)
        realmin axis l.append((pores[pore idx].minor axis length)*voxelsize)
168
169
170
171
        pore = pores[pore idx]
172
173
        inertia_eigval = pore.inertia_tensor_eigvals
174
        inertia = pore.inertia tensor
175
        maxeig = np.argmax(inertia eigval)
176
        eigvec = np.linalg.eig(pore.inertia tensor)[1]
177
        eigvals = np.linalg.eig(pore.inertia_tensor)[0]
178
        if np.sum(eigvals) == 0:
179
            breakpoint()
180
        trv:
181
            anis = 1 - np.min(eigvals)/np.max(eigvals)
182
        except:
183
            anis = 0
184
185
        if np.max(eigvals) == 0:
186
            anis = 0
187
        realanisotropies.append(anis)
188
        maxvector = eigvec[:, maxeig]
189
        orientation = angle between(maxvector, np.array([0,0,1]))
190
        phi = angle between(maxvector, np.array([0,1,0]))
        realorientations.append(orientation)
191
192
193
        return [realvols[0], 1, anis, orientation, phi]
194 def save hdf5(ndarr, filename):
195
196
       # tensor = tensor.cpu()
197
       # ndarr = tensor.mul(255).byte().numpy()
```

```
198
       with h5py.File(filename, 'w') as f:
            f.create_dataset('data', data=ndarr, dtype="i8", compression="qzip")
199
200 def test generator(epoch = 38, folder index = 0, num samples = 100, show =
   False):
201
       pore list attr = []
202
       print("Epoch, ", epoch)
203
        cudnn.benchmark = True
204
205
       # Use GPU is available else use CPU.
       device = torch.device("cuda:0" if(torch.cuda.is available()) else "cpu")
206
       print(device, " will be used.\n")
207
208
       # out dir ='./reconstruction/generated pore samples' + str(epoch) +'/'
       out dir = './reconstruction/gan/saved_generated_pores/'
209
210
       os.makedirs(str(out dir), exist ok=True)
211
212
        checkpoint = torch.load('./reconstruction
   /gan/netG epoch 62.pth')##torch.load('./reconstruction/gan/saved model.pth')
213
       def frame_tick(frame_width = 2, tick_width = 1.5):
214
            ax = gca()
215
            for axis in ['top','bottom','left','right']:
216
                ax.spines[axis].set linewidth(frame width)
217
            plt.tick params(direction = 'in',
218
                            width = tick width)
219
220
221
       if('cuda' in str(device)):
222
            # Create the generator.
223
            netG = Generator(params['nz'], params['nc'], params['ngf'],
   params['ngpu'], size =64).to(device)
            netG.load state dict(checkpoint)
224
225
            netG = nn.DataParallel(netG)
226
227
       else:
228
            # Create the generator.
229
            netG = Generator(params['nz'], params['nc'], params['ngf'],
   params['ngpu']).to(device)
230
            netG.load state dict(checkpoint)
231
            netG = nn.DataParallel(netG)
232
233
            noise = torch.FloatTensor(1, params['nz'], params['alpha'],
   params['alpha'], params['alpha']).normal (0, 1)
234
            noise = noise.to(device)
235
236
       i = 0
237
       # Clean generated pores
238
       print("Removing previously generated pores")
239
       file list = glob.glob(os.path.join(out dir,'generator*hdf5'))
        for fname in file list:
240
            os.remove(fname)
241
242
243
       while i < num samples:#params['num samples']:</pre>
244
            noise = torch.FloatTensor(1, params['nz'], params['alpha'],
   params['alpha'], params['alpha']).normal (0, 1)
245
            noise = noise.to(device)
246
            fake data = netG(noise)
```

```
247
            fake argmax = fake data.argmax(dim=1)
248
            im = 1-fake argmax[0]
249
            if torch.sum(im) ==0:
250
                continue
251
           pore = extract pore(1-fake argmax[0])
252
            fake stats = analyze pore(pore)
253
            if fake stats[0] < 0:
254
               print('An unresolved pore filtered, continuing generation
   process')
255
           else:
256
257
                pore list attr.append(fake stats)
258
               if i%100 == 0:
259
                    print('======')
                   print(str(i) + ' pores processed')
260
                    print('======')
261
262
               save hdf5(pore, str(out dir)+'/generator {0}.hdf5'.format(i))
263
       print('{} pores generated, saved in '.format(i), out_dir)
264
265
       pore matrix = np.loadtxt('analyze pore samples/results
   /individual pore samples/partsample{}/pore matrix'.format(folder index))
266
       pore matrix generated = np.squeeze(np.array(pore list attr))
267
268
       np.savetxt('generated pore matrix.csv',pore matrix generated )
269
       pore identity = 100
270
       filename
                  = str(out dir)+'generator {0}.hdf5'.format(pore identity)
271
       data = read file(filename)
272
       pore_sizes = pore_matrix[:,0]
273
274
       vol target = np.sort(pore matrix[:,0])[int(9*len(pore matrix)//10)]
275
       anis =np.sort(pore matrix[:,2])[len(pore matrix)//2]
276
       angle = np.sort(pore matrix[:,3])[len(pore matrix)//2]
277
       pore identity real = query pore(vol target, anis, angle, pore matrix)
278
       pore_identity_generated = query_pore(vol_target, anis, angle,
   pore matrix generated)
279
280
       save pore(pore identity real, epoch = epoch, real = True, fname =
    'baseline', pore matrix=pore matrix, value = vol target)
        save pore(pore identity generated, epoch = epoch, fname = 'baseline',
281
   pore matrix = pore matrix generated, value = vol target)
282
283
284
285
       vol_target = np.sort(pore_matrix[:,0])[int(9.92*len(pore_matrix)//10)]
       anis =np.sort(pore matrix[:,2])[len(pore matrix)//2]
286
287
       angle = np.sort(pore matrix[:,3])[len(pore matrix)//2]
288
       pore identity real = query pore(vol target, anis, angle, pore matrix)
289
       pore identity generated = query pore(vol target, anis, angle,
   pore matrix generated)
290
291
        save_pore(pore_identity_real, epoch = epoch, real = True, fname =
    'bigvol', pore matrix=pore matrix, value = vol target)
       save pore(pore identity generated, epoch = epoch, fname = 'bigvol',
292
   pore matrix = pore matrix generated, value = vol target)
293
```

```
294
295
        vol target = np.sort(pore matrix[:,0])[int(9*len(pore matrix)//10)]
296
        anis =np.sort(pore matrix[:,2])[int(9.9*len(pore matrix)//10)]
297
        angle = np.sort(pore matrix[:,3])[len(pore matrix)//2]
298
        pore identity real = query pore(vol target, anis, angle, pore matrix)
299
        pore identity generated = query pore(vol target, anis, angle,
    pore matrix generated)
300
        save pore(pore identity real, epoch = epoch, real = True, fname =
301
    'biganis', pore matrix=pore matrix, value = anis)
302
        save pore(pore identity generated, epoch = epoch, fname = 'biganis',
    pore matrix = pore matrix generated, value = anis)
303
304
305
        vol target = np.sort(pore matrix[:,0])[int(9*len(pore matrix)//10)]
306
        anis =np.sort(pore matrix[:,2])[len(pore matrix)//2]
        angle = np.sort(pore matrix[:,3])[int(9.9*len(pore matrix)//10)]
307
308
        pore_identity_real = query_pore(vol_target, anis, angle, pore_matrix)
309
        pore identity generated = query pore(vol target, anis, angle,
    pore matrix generated)
310
        save_pore(pore_identity_real, epoch = epoch, real = True, fname =
311
    'bigangle', pore matrix=pore matrix, value = angle)
        save pore(pore identity generated, epoch = epoch, fname = 'bigangle',
312
    pore matrix = pore matrix generated, value = angle)
313
        os.makedirs('./reconstruction/gan/figures',exist ok = True)
314
        np.savetxt('./reconstruction/gan/figures
   /pore_matrix_updated.csv',pore_matrix_generated )
315
        plt.figure(figsize = [4,3], dpi = 150)
316
317
        plt.hist(pore matrix[:,0]*(3.49)**3, density = True, alpha =
    0.7, bins=np.logspace(np.log10(10e1),np.log10(10e5)), edgecolor = 'k', label =
    "Ground Truth")
        plt.hist(pore_matrix_generated[:,0]*(3.49)**3, density = True, alpha =
318
    0.7, bins=np.logspace(np.log10(10e1),np.log10(10e5)), edgecolor = \frac{k'}{k}, label =
    "Generated")
319
        plt.plot([8*3.49**3,8*3.49**3], [0, 0.004])
320
        # plt.ylim([0, 0.003])
321
        plt.xscale('log')
322
        plt.yscale('log')
323
324
        legend()
325
        plt.xlabel(r"Volume [$\mu m^3$]")
326
        plt.ylabel("Probability")
327
328
        frame tick()
329
        plt.tight layout()
330
       plt.savefig('./reconstruction/gan/figures/volume' +str(epoch))
331
        if show:
332
            plt.show()
333
        plt.clf()#savefig("vols new" + str(k+ 100) + ".png")
334
335
        plt.figure(figsize = [4,3], dpi = 150)
336
        frame tick()
337
        plt.hist((np.array(pore matrix[:,3])/np.pi)*180, density = True, bins=30,
```

```
edgecolor = 'k', alpha =0.7, label = "Ground Truth")
        plt.hist((np.array(pore_matrix_generated[:,3])/np.pi)*180,density = True,
338
   bins=30,alpha = 0.7, edgecolor = 'k', label = "Generated")
339
340
        legend()
341
       plt.xlabel(r"Angle [Degrees]")
342
       plt.ylabel("Probability")
343
       plt.tight layout()
       plt.savefig('./reconstruction/gan/figures/angle' +str(epoch))
344
345
       if show:
346
            plt.show()
       plt.clf()
347
348
349
       plt.figure(figsize = [4,3], dpi = 150)
350
        frame tick()
351
       plt.hist(pore matrix[:,2], density = True, bins=30, edgecolor = 'k', label
352
   = "Ground Truth", alpha = 0.7)
       plt.hist(pore matrix generated[:,2], density = True, bins=30, edgecolor =
353
    'k', label = "Generated", alpha = 0.7)
354
355
       legend()
        plt.xlabel(r"Anisotropy")
356
357
       plt.vlabel("Probability")
358
       plt.tight layout()
359
       plt.savefig('./reconstruction/gan/figures/anisotropy' +str(epoch))
360
       if show:
361
            plt.show()
       plt.clf()
362
363
364
       # plt.clf()
365
       dict one = {'Volume': np.log10(pore matrix[:,0]),
    'Anisotropy':pore matrix[:,2], 'Orientation':pore matrix[:,3]}#,'phis':
   gt total phi }
366
        dict two = {'Volume': np.log10(pore matrix generated[:,0]),
    'Anisotropy':pore matrix generated[:,2],
    'Orientation':pore matrix generated[:,3]}
367
        plt scatter matrices(dict one, dict two, list(dict one.keys()), show =
   show)
368
369
       plt.clf()
370 if name == " main ":
371
        test generator()
```