

```
import numpy as np
from matplotlib import pyplot as plt
import pickle
import pandas
import numpy as np
import matplotlib.pyplot as plt
from skimage import measure
import imageio
import skimage
from scipy import spatial
import time
import h5py
import os
import numpy as np
import sys
from mpl_toolkits.mplot3d import Axes3D
from skimage import measure, transform, segmentation, morphology
from pylab import gca
from scipy.ndimage.interpolation import geometric_transform
import numpy as np
import torch
import os
from reconstruction.polar_cartesian_convert import linear_polar,
polar_linear, map_pixel, calc_theta, unmap_pixel
from skimage.transform import resize
from mpl_toolkits.axes_grid1 import make_axes_locatable
from scipy.signal import savgol_filter
import numpy as np
import scipy.ndimage as ndimage

import pandas as pd

import numpy as np
import matplotlib.pyplot as plt
import os

import numpy as np
def unit_vector(vector):
    """ Returns the unit vector of the vector. """
    return vector / np.linalg.norm(vector)

def read_file(fname):
    f = h5py.File(fname, 'r')
    return f['data']
def angle_between(v1, v2):
    """ Returns the angle in radians between vectors 'v1' and 'v2': """

    >>> angle_between((1, 0, 0), (0, 1, 0))
    1.5707963267948966
    >>> angle_between((1, 0, 0), (1, 0, 0))
    0.0
    >>> angle_between((1, 0, 0), (-1, 0, 0))
```

```
54         3.141592653589793
55         """
56         v1_u = unit_vector(v1)
57         v2_u = unit_vector(v2)
58         return np.arccos(np.clip(np.dot(v1_u, v2_u), -1.0, 1.0))
59
60
61 def set_axes_equal(ax):
62     '''Make axes of 3D plot have equal scale so that spheres appear as
63     spheres,
64     cubes as cubes, etc.. This is one possible solution to Matplotlib's
65     ax.set_aspect('equal') and ax.axis('equal') not working for 3D.
66
67     Input
68     ax: a matplotlib axis, e.g., as output from plt.gca().
69
70     x_limits = ax.get_xlim3d()
71     y_limits = ax.get_ylim3d()
72     z_limits = ax.get_zlim3d()
73
74     x_range = abs(x_limits[1] - x_limits[0])
75     x_middle = np.mean(x_limits)
76     y_range = abs(y_limits[1] - y_limits[0])
77     y_middle = np.mean(y_limits)
78     z_range = abs(z_limits[1] - z_limits[0])
79     z_middle = np.mean(z_limits)
80
81     plot_radius = 0.5*max([x_range, y_range, z_range])
82
83     ax.set_xlim3d([x_middle - plot_radius, x_middle + plot_radius])
84     ax.set_ylim3d([y_middle - plot_radius, y_middle + plot_radius])
85     ax.set_zlim3d([z_middle - plot_radius, z_middle + plot_radius])
86
87
88
89
90 '''
91 Make frame thicker, make tick pointing inside, make tick thicker
92 default frame width is 2, default tick width is 1.5
93 '''
94 def frame_tick(frame_width = 2, tick_width = 1.5):
95     ax = gca()
96     for axis in ['top', 'bottom', 'left', 'right']:
97         ax.spines[axis].set_linewidth(frame_width)
98     plt.tick_params(direction = 'in',
99                     width = tick_width)
100
101 '''
102 legend:
103 default location : upper left
104 default fontsize: 8
105 Frame is always off
106 '''
107 def legend(location = 'upper left', fontsize = 8):
```

```
108     plt.legend(loc = location, fontsize = fontsize, frameon = False)
109
110     '''
111     savefig:
112     bbox_inches is always tight
113     '''
114     def savefig(filename):
115         plt.savefig(filename, bbox_inches = 'tight')
116
117
118     def topolar(img, order=1):
119         """
120         Transform img to its polar coordinate representation.
121
122         order: int, default 1
123             Specify the spline interpolation order.
124             High orders may be slow for large images.
125         """
126         # max_radius is the length of the diagonal
127         # from a corner to the mid-point of img.
128         max_radius = 0.5*np.linalg.norm( img.shape )
129
130         def transform(coords):
131             theta = 2*np.pi*coords[1] / (img.shape[1] - 1.)
132
133             radius = max_radius * coords[0] / img.shape[0]
134
135             i = 0.5*img.shape[0] - radius*np.sin(theta)
136             j = radius*np.cos(theta) + 0.5*img.shape[1]
137             return i,j
138
139         polar = geometric_transform(img, transform, order=order)
140
141         rads = max_radius * np.linspace(0,1,img.shape[0])
142         ang = np.linspace(0, 2*np.pi, img.shape[1])
143
144         return polar, (rads, ang)
145     def improve_pairplot(g, replacements):
146         g.fig.set_dpi(300)
147         for idx,i in enumerate(g.axes[0]):
148             for idx_j,j in enumerate(g.axes):
149                 g.axes[idx_j][idx].spines['left'].set_linewidth(2)
150                 g.axes[idx_j][idx].spines['bottom'].set_linewidth(2)
151                 g.axes[idx_j][idx].tick_params(direction = 'in', width = 1.5)
152                 xlabel = g.axes[idx_j][idx].get_xlabel()
153                 ylabel = g.axes[idx_j][idx].get_ylabel()
154                 if xlabel in replacements.keys():
155                     g.axes[idx_j][idx].set_xlabel(replacements[xlabel], fontsize
156 = 18)
157                     if ylabel in replacements.keys():
158                         g.axes[idx_j][idx].set_ylabel(replacements[ylabel], fontsize
159 = 18)
160                 return g
161     def analyze_pore(im):
162         voxelsize = 1
```

```
161
162     realvols = []
163     realorientations = []
164     realanisotropies = []
165     realmin_axis_l = []
166     realmaj_axis_l = []
167     im = np.array(im, dtype = 'int')
168     pores = measure.regionprops(im)
169     if len(pores) == 0:
170         return np.array([0,0,0])
171     pore_idx = np.argmax([pore.area for pore in pores])
172     if pores[pore_idx].area > 1:
173         realvols.append(pores[pore_idx].area)
174         realmaj_axis_l.append((pores[pore_idx].major_axis_length)*voxelsize)
175         realmin_axis_l.append((pores[pore_idx].minor_axis_length)*voxelsize)
176
177
178     pore = pores[pore_idx]
179
180     inertia_eigval = pore.inertia_tensor_eigvals
181     inertia = pore.inertia_tensor
182     maxeig = np.argmax(inertia_eigval)
183     eigvec = np.linalg.eig(pore.inertia_tensor)[1]
184     eigvals = np.linalg.eig(pore.inertia_tensor)[0]
185     anis = 0
186     realanisotropies.append(anis)
187     maxvector = eigvec[:, maxeig]
188     orientation = angle_between(maxvector, np.array([0,0,1]))
189     realorientations.append(orientation)
190     return np.array([anis, realvols[0]**(1/3), orientation/np.pi])
191
192 def load_image(fname, dim = 4):
193     im = np.loadtxt(fname).reshape(566,571,dim)
194     img = np.copy(im)
195     boundary = np.zeros((im.shape))
196     boundary[img == [1, 1, 1]] = 1
197     boundary = boundary[:, :, 0]
198     img[img == [1, 1, 1]] = 0
199     return boundary, img
200
201 def load_shell(fname,num_frames = 200, dim = 4, start = 0):
202     total = num_frames
203     im_stack = np.zeros((566,571, total, dim))
204     shells = np.zeros((566,571, total))
205     for i in range(start, start+total):
206         im =
np.loadtxt(fname+'tiff_threephase_id_'+str(i)).reshape(566,571,dim)
207         print(np.unique(im[:, :, 2]))
208         print(fname+'tiff_threephase_id_'+str(i))
209
210         img = np.copy(im)
211         boundary = np.zeros((im.shape))
212         boundary[img == [1, 1, 1]] = 1
213
214         img[img == [1, 1, 1]] = 0
```

```
215         boundary = boundary[:, :, 0]
216         shells[:, :, i-start] = boundary
217         im_stack[:, :, i-start] = img
218
219         print(i)
220
221     return shells ,im_stack
222
223
224
225 def compute_statistics(pore_dataset, voxelsize, imstack = None):
226     anisotropies = []
227     orientations = []
228     vols = []
229     sphericity = []
230     x_locs = []
231     y_locs = []
232     z_locs = []
233     maj_axis_l = []
234     min_axis_l = []
235     phis = []
236     for i in range(len(pore_dataset)):
237         pore = pore_dataset[i]
238
239         if pore_dataset[i]['area'] == 1:
240             continue
241         vols.append(pore_dataset[i]['area']*voxelsize*voxelsize*voxelsize)
242         y_locs.append(pore_dataset[i]['centroid'][1]*voxelsize)
243         x_locs.append(pore_dataset[i]['centroid'][0]*voxelsize)
244         z_locs.append(pore_dataset[i]['centroid'][2]*voxelsize)
245         maj_axis_l.append((pore_dataset[i].major_axis_length)*voxelsize)
246         thresh = 0.3
247
248         inertia_eigval = pore.inertia_tensor_eigvals
249         inertia = pore.inertia_tensor
250         maxeig = np.argmax(inertia_eigval)
251         eigvec = np.linalg.eig(pore.inertia_tensor)[1]
252         eigvals = np.linalg.eig(pore.inertia_tensor)[0]
253         anis = 1 - np.min(eigvals)/np.max(eigvals)
254         anisotropies.append(anis)
255         maxvector = eigvec[:, maxeig]
256         orientation = angle_between(maxvector, np.array([0,0,1]))
257         orientations.append(orientation)
258
259         phi = angle_between(maxvector, np.array([0,1,0]))
260         phis.append(phi)
261     stats = {
262         'anisotropies': anisotropies,
263         'orientations': orientations,
264         'vols': vols,
265         'x_locs': x_locs,
266         'y_locs': y_locs,
267         'maj_axis_l': maj_axis_l,
268         'z_locs': z_locs,
269         'phis': phis
```

```

270     }
271     if not len(np.unique([len(stat) for stat in stats.values()])) == 1:
272         print([len(stat) for stat in stats.values()])
273         breakpoint()
274
275     return stats#, examples
276 def extract_pores(imstack):
277     boundary_imstack = np.copy(imstack)
278     boundary_imstack[imstack != 255] = 0
279     boundary_imstack[imstack == 255] = 1
280
281     imstack[imstack == 255] = 0
282     imstack[imstack == 159] = 1
283     imstack = np.array(imstack, dtype = 'uint8')
284     im = measure.label(imstack[:,:,:])
285     props = skimage.measure.regionprops(im[:,:,:])
286     return props, im, boundary_imstack
287
288
289 def draw_centroid(label_img, frame_id = 0):
290     plt.imshow(label_img, cmap = 'binary')
291
292     props = measure.regionprops(label_img)
293     prop_labels = [prop.label for prop in props]
294     for idx in np.unique(label_img):
295         if idx != 0:
296
297             prop_idx = np.where(prop_labels == idx)[0][0]
298             centroid = props[prop_idx].centroid
299             plt.text(centroid[1], centroid[0], idx)
300     plt.title('Frame ' + str(frame_id))
301     plt.savefig('centroidgen'+ str(frame_id) + '.png')
302     plt.clf()
303
304 def plot_image(images,shell, title, global_idx = 0):
305     os.makedirs(title, exist_ok = True)
306     print("SAVING TO", title)
307     for p in range(images.shape[2]):
308         plt.imsave(title + str(global_idx+p) + '.png', np.array((images[:,:,:
309 p]*(255//2) + shell[:,:,:p]*255), dtype = int), cmap = 'gist_gray')
310     plt.clf()
311
312 def load_existing(start = 0, num_frames = 200):
313     total = num_frames
314     seconds = time.time()
315     shell, im_stack = load_shell('/media/cmu/DATA/francis/pore_gan
316 /8approxstyleganthreechannelboundaryfulltest_threephase_2D/', dim = 4, start
317 = start, num_frames = num_frames)
318     pore_part = np.zeros((566,571, total)) #shell.shape
319     sum_tot = np.sum(im_stack[:,:,:], axis = 3)
320     sum_tot[sum_tot > 0] = 1
321     return pore_part, shell, im_stack
322
323 def read_from_file(filename, iteration = None, restart = None):

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322     tensors = torch.load(filename)
323     im_opt = np.squeeze(tensors['tensor_opt'])
324     plt.imshow(im_opt, cmap = 'binary')
325     plt.colorbar()
326     plt.title('MST generated surface roughness profiles, iteration: ' +
str(iteration) + ' instance: ' + str(restart))
327     plt.savefig('modelCfigs' + str(restart) + '_' + str(iteration)
+'profiletest' + '.png')
328
329     plt.clf()
330
331 def load_boundary(num_frames = 500, start = 0, pore_part_shape = (566, 571),
return_profile = False, im_opt = None, resultsdir = None, folder_index = 0):
332     i = folder_index
333
334     if resultsdir is None:
335         resultsdir = 'make_surface/results
/sample_number_0original_folder_{}'.format(i)
336     # i = 0
337     xmean = np.loadtxt(resultsdir+ '/xmean{}'.format(i))
338     ymean = np.loadtxt(resultsdir+ '/ymean{}'.format(i))
339     minmax = pd.read_csv(resultsdir+ '/minmax_values{}.csv'.format(i,i))
340     print(minmax['max'])
341     print(minmax['min'])
342     maxim = np.array(minmax['max'])
343     minim = np.array(minmax['min'])
344     ratio = minim/maxim
345
346
347     if im_opt is None:
348         fname = resultsdir + '/modelC_krec' + str(0) + '_start' + str(1) +
'.pt'
349         tensors = torch.load(fname)
350         im_opt = np.array(np.squeeze(tensors['tensor_opt']) + xmean+ymean)
351     print(i)
352
353     polar_index_r = linear_polar(np.zeros(pore_part_shape)).shape[0]
354     polar_index_theta = linear_polar(np.zeros(pore_part_shape)).shape[1]
355     new_im = resize(im_opt, (2000, polar_index_theta), order = 3)
356     line = new_im[0]
357
358     line_int = np.array(new_im[0], dtype = 'int')
359     polar_image = np.zeros((polar_index_r, polar_index_theta))
360     idxs = (line_int, np.arange(polar_index_theta, dtype = int))
361     polar_image[idxs] = 1
362     shells = np.zeros((pore_part_shape[0], pore_part_shape[1], num_frames),
dtype = 'uint8')
363
364
365
366
367     xs = np.linspace(0, pore_part_shape[0], pore_part_shape[0])
368     ys = np.linspace(0, pore_part_shape[1], pore_part_shape[1])
369
370     # full coordinate arrays

```



```

371
372     xx, yy = np.meshgrid(xs, ys)
373     zz = np.sqrt((xx - pore_part_shape[0]//2)**2 + (yy -
pore_part_shape[1]//2)**2)
374     toprow = zz[0]
375     bottomrow = zz[-1]
376     left = zz[:,0]
377
378
379     right = zz[:, -1]
380     continuous = np.hstack((toprow, bottomrow, left, right))
381     # Calculate the radius at which it would
382     edge_radii = np.array(continuous)
383     max_valid_radius = np.min(edge_radii)
384     min_valid_radius = ratio*max_valid_radius
385
386     for k in range(start, num_frames+start):
387         line = new_im[k]*(max_valid_radius - min_valid_radius) +
min_valid_radius
388         line[-1] = line[0]
389         line = savgol_filter(line, 27, 3)
390         line_int = np.array(line, dtype = 'int')
391         polar_image = np.zeros((polar_index_r, polar_index_theta))
392         idxs = (line_int, np.arange(polar_index_theta, dtype = int))
393         polar_image[idxs] = 1
394         shell_img = polar_linear(polar_image, output = (pore_part_shape[0],
pore_part_shape[1]))
395         shells[:, :, k-start] = np.array(shell_img > 0, dtype='uint8')
396
397     if return_profile:
398         return shells, new_im*(max_valid_radius - min_valid_radius) +
min_valid_radius
399     return shells
400
401 def analyze_results(original, pore_reconstruct, fname = None,
lists_all_original = None, lists_all_new = None, voxelsize = 3.49):
402     print('FINISHED RECONSTRUCTION')
403     title = fname
404     frame_tick()
405     os.makedirs(title, exist_ok = True)
406     if lists_all_original == None:
407         props, _, _ = extract_pores(pore_reconstruct)
408         props_orig, _, _ = extract_pores(original)
409         stats = compute_statistics(props, voxelsize=voxelsize)
410         # stats_orig = compute_statistics(props_orig, voxelsize=3.49)
411
412
413         total_anisotropies = []
414         total_x = []
415         total_y = []
416         total_orientations = []
417         total_z = []
418         total_maj = []
419         total_vols = []
420

```



```
421     total_phis = []
422     gt_total_phis = []
423     total_x.extend(stats['x_locs'])
424     total_y.extend(stats['y_locs'])
425     total_maj.extend(stats['maj_axis_l'])
426     total_vols.extend(stats['vols'])
427     total_anisotropies.extend(stats['anisotropies'])
428     total_orientations.extend(stats['orientations'])
429     total_phis.extend(stats['phis'])
430     total_z.extend(stats['z_locs'])
431     stats = compute_statistics(props_orig, voxelsize=voxelsize)
432
433
434     gt_total_anisotropies = []
435     gt_total_x = []
436     gt_total_y = []
437     gt_total_orientations = []
438     gt_total_z = []
439     gt_total_maj = []
440
441     gt_total_vols = []
442
443     gt_total_x.extend(stats['x_locs'])
444     gt_total_y.extend(stats['y_locs'])
445     gt_total_maj.extend(stats['maj_axis_l'])
446     gt_total_vols.extend(stats['vols'])
447     gt_total_anisotropies.extend(stats['anisotropies'])
448     gt_total_orientations.extend(stats['orientations'])
449     gt_total_phis.extend(stats['phis'])
450     gt_total_z.extend(stats['z_locs'])
451
452
453     lists_all_original = {'x_locs': gt_total_x, 'y_locs': gt_total_y,
454 'maj_axis_l': gt_total_maj, 'vols': gt_total_vols,
455 'anisotropies': gt_total_anisotropies, 'orientations': gt_total_orientations,
456 'z_locs': gt_total_z, 'phis': gt_total_phis }
457     lists_all_new = {'x_locs': total_x, 'y_locs': total_y, 'maj_axis_l':
458 total_maj, 'vols': total_vols, 'anisotropies': total_anisotropies,
459 'orientations': total_orientations, 'z_locs': total_z, 'phis': total_phis }
460
461     density = True
462     fig = plt.figure(figsize=[4,3], dpi = 300)
463     histogram = plt.hist((np.array(lists_all_new['orientations'])/np.pi)*180,
464 density = density, bins=30, edgecolor = 'k', label = 'reconstructed', alpha
465 = 0.7)
466     histogram2 =
467     plt.hist((np.array(lists_all_original['orientations'])/np.pi)*180, density =
468 density, bins=30, edgecolor = 'k', label = 'original', alpha = 0.7)
469     plt.title("Orientation")
470
471     np.savetxt(title+ 'num_pores',
472 np.array([len(lists_all_original['orientations']),
473 len(lists_all_new['orientations'])]))
474     plt.xlabel(r"Angle [Degrees]")
```

```
465     plt.ylabel("Probability")
466     frame_tick()
467     legend()
468     plt.tight_layout()
469     plt.savefig(title + "orientation" + ".png")
470     plt.show()
471     plt.clf()
472
473
474
475
476     fig = plt.figure(figsize=[4,3], dpi = 300)
477     histogram = plt.hist((np.array(lists_all_new['phis'])/np.pi)*180, density
= density, bins=30, edgecolor = 'k', label = 'reconstructed', alpha = 0.7)
478     histogram2 = plt.hist((np.array(lists_all_original['phis'])/np.pi)*180,
density = density, bins=30, edgecolor = 'k', label = 'original', alpha = 0.7)
479     plt.title("Phi")
480     np.savetxt(title+ 'num_pores', np.array([len(lists_all_original['phis']),
len(lists_all_new['phis'])]))
481     plt.xlabel(r"Angle [Degrees]")
482     plt.ylabel("Probability")
483     frame_tick()
484     legend()
485     plt.tight_layout()
486     plt.savefig(title + "phi" + ".png")
487     plt.show()
488     plt.clf()
489
490
491     fig = plt.figure(figsize=[4,3], dpi = 300)
492     histogram = plt.hist((np.array(lists_all_new['anisotropies'])), density =
density, bins=30, edgecolor = 'k', label = 'reconstructed', alpha = 0.7)
493     histogram2 = plt.hist(np.array(lists_all_original['anisotropies']),
density = density, bins=30, edgecolor = 'k', label = 'original', alpha = 0.7)
494     plt.title("Anisotropy")
495     plt.xlabel(r"Anisotropy")
496     plt.ylabel("Probability")
497     frame_tick()
498     legend()
499     plt.tight_layout()
500     plt.savefig(title+"anisotropy" + ".png")
501     plt.show()
502     plt.clf()
503     fig = plt.figure(figsize=[4,3], dpi = 300)
504     histogram = plt.hist((np.array(lists_all_new['y_locs'])), density =
density, bins=30, edgecolor = 'k', label = 'reconstructed', alpha = 0.7)
505     histogram2 = plt.hist((np.array(lists_all_original['y_locs'])), density =
density, bins=30, edgecolor = 'k', label = 'original', alpha = 0.7)
506     plt.title("Y location")
507     plt.xlabel(r"Y location [micrometers]")
508     plt.ylabel("Probability")
509     frame_tick()
510     legend()
511     plt.tight_layout()
512     plt.savefig(title+"yloc" + ".png")
```

```
513     plt.show()
514
515     plt.clf()
516     fig = plt.figure(figsize=[4,3], dpi = 300)
517
518     histogram = plt.hist((np.array(lists_all_new['x_locs'])), density =
density, bins=30, edgecolor = 'k', label = 'reconstructed', alpha = 0.7)
519     histogram2 = plt.hist((np.array(lists_all_original['x_locs'])), density =
density, bins=30, edgecolor = 'k', label = 'original', alpha = 0.7)
520     plt.title("X location")
521     plt.xlabel(r"X location [micrometers]")
522     plt.ylabel("Probability")
523     frame_tick()
524     legend()
525     plt.tight_layout()
526     plt.savefig(title+"xloc" + ".png")
527     plt.show()
528
529     plt.clf()
530     fig = plt.figure(figsize=[4,3], dpi = 300)
531
532     histogram = plt.hist(lists_all_new['vols'], density = density, alpha =
0.7, bins=np.logspace(np.log10(10e1), np.log10(10e5)), edgecolor = 'k', label =
'reconstructed')
533     histogram2 = plt.hist(lists_all_original['vols'], density = density,
alpha = 0.7, bins=np.logspace(np.log10(10e1), np.log10(10e5)), edgecolor = 'k',
label = 'original')
534
535     plt.title("Volume")
536
537     plt.xscale('log')
538     plt.xlabel(r"Volume [ $\mu\text{ m}^3$ ]")
539     plt.ylabel("Probability")
540     frame_tick()
541     legend()
542     plt.tight_layout()
543     plt.savefig(title + "vols" + ".png")
544     plt.show()
545     plt.clf()
546
547     print("Pores in the original sample: " +
str(len(lists_all_original['vols'])))
548     print("Pores in the new sample: " + str(len(lists_all_new['vols'])))
549 def replace_sampling(pore_part, generated_boundary, n_bins = 30, window_size
= 100, properties_folder = './analyze_pore_samples/results/pore_properties
/probability_matrices/', use_generated = True, use_gt = True):
550     import scipy
551
552     prob_matrix_volume=np.load(properties_folder + str(n_bins) +
'_{0}allprob_matrix_volume.npy'.format(0))
553     prob_matrix_num = np.load(properties_folder + str(n_bins) +
'_{0}allprob_matrix_num.npy'.format(0))*(window_size/100)#/2
554     bin_edges_vols = np.load(properties_folder + str(n_bins) +
'_{0}allbin_edges_vols.npy'.format(0))
555     bin_edges_anis = np.load(properties_folder + str(n_bins) +
```

```

'_{allbin_edges_anis.npy}'.format(0))
556     bin_edges_phis = np.load(properties_folder + str(n_bins) +
'_{allbin_edges_phis.npy}'.format(0))
557     prob_matrix_phis = np.load(properties_folder + str(n_bins) +
'_{allprob_matrix_phis.npy}'.format(0))
558
559     prob_matrix_anis = np.load(properties_folder + str(n_bins) +
'_{allprob_matrix_anis.npy}'.format(0))
560     bin_edges_orientations = np.load(properties_folder + str(n_bins) +
'_{allbin_edges_orientations.npy}'.format(0))
561     prob_matrix_orientation = np.load(properties_folder + str(n_bins) +
'_{allprob_matrix_orientations.npy}'.format(0))
562
563     target_list_size = []
564     target_anis = []
565     target_vols = []
566     actual_vols = []
567     generated_pore_matrix = np.loadtxt('./reconstruction/gan/figures
/pore_matrix_updated.csv')
568     gt_pore_matrix = np.loadtxt('./analyze_pore_samples/results
/individual_pore_samples/partsample0/pore_matrix')
569     x_extent = np.linspace(0, pore_part.shape[0], n_bins)
570     y_extent = np.linspace(0, pore_part.shape[1], n_bins)
571     z_extent = np.arange(0, pore_part.shape[2], window_size)
572     gt_pores_used = []
573     gen_pores_used = []
574     gen_losses = []
575     gt_losses = []
576     for idx_x, x_sample in enumerate(x_extent):
577         for idx_y, y_sample in enumerate(y_extent):
578             for idx_z, z_sample in enumerate(z_extent):
579
580                 vol_hist = prob_matrix_volume[idx_x, idx_y, :]
581
582                 vol_hist_dist = scipy.stats.rv_histogram((vol_hist,
bin_edges_vols))
583                 num = prob_matrix_num[idx_x, idx_y]
584                 if num == 0:
585                     continue
586                 if num < 1:
587                     unif_sample = np.random.uniform()
588                     if num > unif_sample:
589                         num = 1
590                 else:
591                     num = 0
592                     continue
593                 elif num > 1:
594                     num = int(np.around(num))
595
596                 volumes = vol_hist_dist.rvs(size=int(num))
597                 ani_hist = prob_matrix_anis[idx_x, idx_y]
598                 ani_hist_dist = scipy.stats.rv_histogram((ani_hist,
bin_edges_anis))
599                 anisotropies = ani_hist_dist.rvs(size=int(num))
600

```

```

601
602         angle_hist = prob_matrix_orientation[idx_x, idx_y]
603         angle_hist_dist = scipy.stats.rv_histogram((angle_hist,
bin_edges_orientations))
604         angles = angle_hist_dist.rvs(size=int(num))
605         phi_hist = prob_matrix_phis[idx_x, idx_y]
606         phi_hist_dist = scipy.stats.rv_histogram((phi_hist,
bin_edges_phis))
607         phis = phi_hist_dist.rvs(size = int(num))
608         target_anis.extend(anisotropies)
609         target_vols.extend(np.array(volumes)**3)
610         for idx_pore, gen_pore in enumerate(volumes):
611
612             gen_pore = np.max([gen_pore, 2])
613
614             tmp_x =
np.random.randint(0, int(pore_part.shape[0]/n_bins))
615             tmp_y =
np.random.randint(0, int(pore_part.shape[1]/n_bins))
616
617             curr_x = int(x_sample-tmp_x)
618             curr_y = int(y_sample - tmp_y)
619
620             curr_z = z_sample+np.random.randint(0, window_size)
621
622             target_pore= gen_pore**3#/25
623             target_list_size.append(target_pore)
624             anis = anisotropies[idx_pore]
625             angle = angles[idx_pore]
626             phi = phis[idx_pore]
627
628             pore_matrix = generated_pore_matrix
629             pore_matrix = generated_pore_matrix
630             pore_sizes = (pore_matrix[:,0])
631
632             gen_pore_identity = np.argmin(np.abs(target_pore -
pore_matrix[:,0])/np.mean(pore_matrix[:,0]) + np.abs(anis -
generated_pore_matrix[:,2])/np.mean(generated_pore_matrix[:,2]) +
np.abs(angle -
generated_pore_matrix[:,3])/np.mean(generated_pore_matrix[:,3])+ np.abs(phi
- generated_pore_matrix[:,4])/np.mean(generated_pore_matrix[:,4]))# +
np.abs(phi - pore_matrix[:,4])/np.mean(pore_matrix[:,4]))
633             gen_loss = np.min(np.abs(target_pore -
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])+ np.abs(phi -
generated_pore_matrix[:,4])/np.mean(generated_pore_matrix[:,4]))# +
np.abs(phi - pore_matrix[:,4])/np.mean(pore_matrix[:,4]))
634
635             pore_matrix = gt_pore_matrix
636             gt_pore_identity = np.argmin(np.abs(target_pore -
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])+ np.abs(phi -
pore_matrix[:,4])/np.mean(pore_matrix[:,4]))# + np.abs(phi -

```

```

    pore_matrix[:,4])/np.mean(pore_matrix[:,4]))
637         gt_loss = np.min(np.abs(target_pore -
    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]) + np.abs(phi -
    pore_matrix[:,4])/np.mean(pore_matrix[:,4]))# + np.abs(phi -
    pore_matrix[:,4])/np.mean(pore_matrix[:,4]))
638
639         gen_losses.append(gen_loss)
640         gt_losses.append(gt_loss)
641
642         if use_generated and use_gt: # Use both generated and
ground truth pores
643             if gen_loss > gt_loss:
644                 generated = False
645                 pore_identity = gt_pore_identity
646                 gt_pores_used.append(pore_identity)
647             else:
648                 generated = True
649                 pore_identity = gen_pore_identity
650                 gen_pores_used.append(pore_identity)
651         elif use_generated: # only use generated
652             generated = True
653             pore_identity = gen_pore_identity
654             gen_pores_used.append(pore_identity)
655         else: # only use ground truth
656
657             generated = False
658             pore_identity = gt_pore_identity
659             gt_pores_used.append(pore_identity)
660
661             if generated:
662                 filename = './reconstruction
/gan/saved_generated_pores/generator_' + str(pore_identity) + '.hdf5'
663             else:
664                 filename = './analyze_pore_samples/results
/individual_pore_samples/partsample0/pore_original_' + str(pore_identity) +
'.hdf5'
665
666                 data = read_file(filename)
667
668                 if len(np.where(data)[0]) == 0:
669                     print('continue 1 activated')
670                     continue
671                 center = 32
672                 size = 32
673                 if generated:
674                     data = np.array(data)/255
675                 else:
676                     data = np.array(data)
677                 xmin = np.min(np.where(data)[0])
678                 ymin = np.min(np.where(data)[1])
679                 zmin = np.min(np.where(data)[2])
680
681                 xmax = np.max(np.where(data)[0])+1

```

```

682         ymax = np.max(np.where(data)[1])+1
683         zmax = np.max(np.where(data)[2])+1
684
685
686         xmin_slice = np.min(np.where(data[:, :, zmin])[0])
687         ymin_slice = np.min(np.where(data[:, :, zmin])[1])
688         lowerlim_x = 0
689         lowerlim_y = 0
690         data_ylower = 0
691         data_xlower = 0
692
693
694         data_yupper = 64-np.abs(np.min((0, pore_part.shape[1] -
(lowerlim_y + 64)))) # in case of negative indices
695         data_xupper = 64-np.abs(np.min((0, pore_part.shape[0] -
(lowerlim_x + 64))))
696         target_z = curr_z
697         if int(target_z) + (zmax-zmin) > pore_part.shape[2]:
698             print('continue 2 activated: pore on back surface')
699             continue
700         if int(target_z) < 0:
701             print('continue 3 activated: pore on front surface')
702             continue
703         try:
704             test_window = pore_part[curr_x: curr_x+ int(xmax-
xmin), curr_y:curr_y + int(ymax-ymin), int(target_z):int(target_z)+(zmax-
zmin)] + data[xmin:xmax , ymin:ymax, zmin:zmax]
705         except Exception as e:
706             print(e)
707             print('continue 4 activated: indexing exception')
708             continue
709         if 0 in test_window.shape:
710             print('continue 5 activated: unresolved pore')
711             continue
712
713         elif np.max(test_window) > 1:
714
715             print('continue 6 activated: Collision')
716             collision_z = np.where(test_window > 1)[2][0]
717             continue
718
719         try:
720             xdatastart = np.min(np.where(data)[0])
721             labelpore = measure.label(data[xmin:xmax , ymin:ymax,
zmin:zmax])
722
723             oldpore = pore_part[curr_x: curr_x+ int(xmax-xmin),
curr_y:curr_y + int(ymax-ymin), int(target_z):int(target_z)+(zmax-zmin)]
724
725             newpore = data[xmin:xmax , ymin:ymax, zmin:zmax]
+pore_part[curr_x: curr_x+ int(xmax-xmin), curr_y:curr_y + int(ymax-ymin),
int(target_z):int(target_z)+(zmax-zmin)]
726             if len(measure.regionprops(measure.label(newpore)))
!= len(measure.regionprops(measure.label(oldpore))) + 1:
727

```



```

print(len(measure.regionprops(measure.label(newpore))), len(measure.regionprop
s(measure.label(oldpore))), 'merged')
728         print('continue 7 activated: Collision')
729         continue
730
731         if generated:
732             pore_part[curr_x: curr_x+ int(xmax-xmin),
curr_y:curr_y + int(ymax-ymin), int(target_z):int(target_z)+(zmax-zmin)] =
2*data[xmin:xmax , ymin:ymax, zmin:zmax]+ pore_part[curr_x: curr_x+ int(xmax-
xmin), curr_y:curr_y + int(ymax-ymin), int(target_z):int(target_z)+(zmax-
zmin)]
733         else:
734             pore_part[curr_x: curr_x+ int(xmax-xmin),
curr_y:curr_y + int(ymax-ymin), int(target_z):int(target_z)+(zmax-zmin)] =
data[xmin:xmax , ymin:ymax, zmin:zmax]+ pore_part[curr_x: curr_x+ int(xmax-
xmin), curr_y:curr_y + int(ymax-ymin), int(target_z):int(target_z)+(zmax-
zmin)]
735
736         except Exception as f:
737             print(f)
738             print('continue 8 activated: Insertion process
failed')
739
740             continue
741
742     actual_vols.append(measure.regionprops(measure.label(np.array(data)))
[0].area)
743     # breakpoint()
744     return pore_part, target_list_size
745
746
747 def replace_sampling_polar(pore_part, generated_boundary, profile_2d, n_bins
= 30, window_size = 100, properties_folder = './analyze_pore_samples/results
/pore_properties/probability_matrices/', folder_index = 0):
748     import scipy#.stats
749     prob_matrix_dir = properties_folder
750     np.load(properties_folder + str(n_bins) +
'_{0}allprob_matrix_volume.npy'.format(0))
751     prob_matrix_volume=np.load(prob_matrix_dir + str(n_bins) +
'_{0}polarprob_matrix_volume.npy'.format(folder_index))
752     prob_matrix_num = np.load(prob_matrix_dir + str(n_bins) +
'_{0}polarprob_matrix_num.npy'.format(folder_index))*(window_size/100)#/2
753     bin_edges_vols = np.load(prob_matrix_dir + str(n_bins) +
'_{0}polarbin_edges_vols.npy'.format(folder_index))
754     bin_edges_anis = np.load(prob_matrix_dir + str(n_bins) +
'_{0}polarbin_edges_anis.npy'.format(folder_index))
755     bin_edges_phis = np.load(prob_matrix_dir + str(n_bins) +
'_{0}polarbin_edges_phis.npy'.format(folder_index))
756     prob_matrix_phis = np.load(prob_matrix_dir + str(n_bins) +
'_{0}polarprob_matrix_phis.npy'.format(folder_index))
757     prob_matrix_anis = np.load(prob_matrix_dir + str(n_bins) +
'_{0}polarprob_matrix_anis.npy'.format(folder_index))
758     bin_edges_orientations = np.load(prob_matrix_dir + str(n_bins) +
'_{0}polarbin_edges_orientations.npy'.format(folder_index))

```

```

759     prob_matrix_orientation= np.load(prob_matrix_dir + str(n_bins) +
    '_{}_polarprob_matrix_orientations.npy'.format(folder_index))
760     target_list_size = []
761     target_anis = []
762     target_vols = []
763     actual_vols = []
764     generated_pore_matrix = np.loadtxt('./reconstruction/gan/figures
    /pore_matrix_updated.csv')
765     gt_pore_matrix = np.loadtxt('./analyze_pore_samples/results
    /individual_pore_samples/partsample0/pore_matrix')
766     r_extent = np.linspace(0,1.1, n_bins)
767     theta_extent = np.linspace(0, np.pi*2, n_bins)
768     z_extent = np.arange(0, pore_part.shape[2],window_size)
769     gt_pores_used = []
770     gen_pores_used = []
771     gen_losses = []
772     gt_losses = []
773     angles = np.linspace(0, np.pi*2, profile_2d.shape[1])
774     radii_totals = []
775     for idx_x,x_sample in enumerate(r_extent):
776         for idx_y,y_sample in enumerate(theta_extent):
777             for idx_z, z_sample in enumerate(z_extent):
778
779                 vol_hist = prob_matrix_volume[idx_x, idx_y, :]
780
781                 vol_hist_dist = scipy.stats.rv_histogram((vol_hist,
    bin_edges_vols))
782                 num = prob_matrix_num[idx_x, idx_y]
783                 if num == 0:
784                     continue
785                 if num < 1:
786                     unif_sample = np.random.uniform()
787                     if num > unif_sample:
788                         num = 1
789                 else:
790                     num = 0
791                     continue
792                 elif num > 1:
793                     num = int(np.around(num))
794
795                 volumes = vol_hist_dist.rvs(size=int(num))
796                 ani_hist = prob_matrix_anis[idx_x, idx_y]
797                 ani_hist_dist = scipy.stats.rv_histogram((ani_hist,
    bin_edges_anis))
798                 anisotropies = ani_hist_dist.rvs(size=int(num))
799
800
801                 angle_hist = prob_matrix_orientation[idx_x, idx_y]
802                 angle_hist_dist = scipy.stats.rv_histogram((angle_hist,
    bin_edges_orientations))
803                 angles = angle_hist_dist.rvs(size=int(num))
804                 phi_hist = prob_matrix_phis[idx_x, idx_y]
805                 phi_hist_dist = scipy.stats.rv_histogram((phi_hist,
    bin_edges_phis))
806                 phis = phi_hist_dist.rvs(size = int(num))

```

```

807
808     target_anis.extend(anisotropies)
809     target_vols.extend(np.array(volumes)**3)
810     print(x_sample, "RADIUS SAMPLE")
811     for idx_pore, gen_pore in enumerate(volumes):
812
813         gen_pore = np.max([gen_pore, 2])
814         tmp_x = np.random.uniform(0, r_extent[1])
815         tmp_y = np.random.uniform(0, theta_extent[1])
816         curr_radius = (x_sample - tmp_x)
817         curr_angle = (y_sample - tmp_y)
818         idx_angle = np.argmin(np.abs(curr_angle -
np.linspace(0, 2*np.pi, profile_2d.shape[1])))
819
820         curr_z = z_sample + np.random.randint(0, window_size)
821         radius_total = profile_2d[curr_z, idx_angle]
822         full_radii = curr_radius * radius_total
823         if x_sample == r_extent[-1]:
824             print(curr_radius, "Current Radius, final evolution")
825         curr_x, curr_y = unmap_pixel(full_radii, theta_idx =
idx_angle, theta = curr_angle, output= pore_part[:, :, 0].shape)
826         radii_totals.append(curr_radius)
827
828         target_pore = gen_pore**3
829         target_list_size.append(target_pore)
830         anis = anisotropies[idx_pore]
831         angle = angles[idx_pore]
832         phi = phis[idx_pore]
833
834         pore_matrix = generated_pore_matrix
835         pore_matrix = generated_pore_matrix
836         pore_sizes = (pore_matrix[:, 0])
837
838         gen_pore_identity = np.argmin(np.abs(target_pore -
pore_matrix[:, 0])/np.mean(pore_matrix[:, 0]) + np.abs(anis -
generated_pore_matrix[:, 2])/np.mean(generated_pore_matrix[:, 2]) +
np.abs(angle -
generated_pore_matrix[:, 3])/np.mean(generated_pore_matrix[:, 3]) + np.abs(phi
- generated_pore_matrix[:, 4])/np.mean(generated_pore_matrix[:, 4]))# +
np.abs(phi - pore_matrix[:, 4])/np.mean(pore_matrix[:, 4]))
839         gen_loss = np.min(np.abs(target_pore -
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]) + np.abs(phi -
generated_pore_matrix[:, 4])/np.mean(generated_pore_matrix[:, 4]))# +
np.abs(phi - pore_matrix[:, 4])/np.mean(pore_matrix[:, 4]))
840
841         pore_matrix = gt_pore_matrix
842         gt_pore_identity = np.argmin(np.abs(target_pore -
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]) + np.abs(phi -
pore_matrix[:, 4])/np.mean(pore_matrix[:, 4]))# + np.abs(phi -
pore_matrix[:, 4])/np.mean(pore_matrix[:, 4]))
843         gt_loss = np.min(np.abs(target_pore -

```

```

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]) + np.abs(phi -
pore_matrix[:,4])/np.mean(pore_matrix[:,4]))# + np.abs(phi -
pore_matrix[:,4])/np.mean(pore_matrix[:,4]))
844         gen_losses.append(gen_loss)
845         gt_losses.append(gt_loss)
846
847         if gen_loss > gt_loss:
848             generated = False
849             pore_identity = gt_pore_identity
850             gt_pores_used.append(pore_identity)
851         else:
852
853             generated = True
854             pore_identity = gen_pore_identity
855             gen_pores_used.append(pore_identity)
856
857         if generated:
858             filename = './reconstruction
/gan/saved_generated_pores/generator_' + str(pore_identity) + '.hdf5'
859         else:
860             filename = './analyze_pore_samples/results
/individual_pore_samples/partsample0/pore_original_' + str(pore_identity) +
'.hdf5'
861
862         data = read_file(filename)
863         if len(np.where(data)[0]) == 0:
864             print('Empty pore, skipped')
865             continue
866
867         center = 32
868         size = 32
869         if generated:
870             data = np.array(data)#/255
871         else:
872             data = np.array(data)
873
874         xmin = np.min(np.where(data)[0])
875         ymin = np.min(np.where(data)[1])
876         zmin = np.min(np.where(data)[2])
877
878         xmax = np.max(np.where(data)[0])+1
879         ymax = np.max(np.where(data)[1])+1
880         zmax = np.max(np.where(data)[2])+1
881
882         xmin_slice = np.min(np.where(data[:, :, zmin])[0])
883         ymin_slice = np.min(np.where(data[:, :, zmin])[1])
884
885
886         lowerlim_x = 0
887         lowerlim_y = 0
888         data_ylower = 0
889         data_xlower = 0
890

```

```

891
892         data_yupper = 64-np.abs(np.min((0, pore_part.shape[1] -
(lowerlim_y + 64)))) # in case of negative indices
893         data_xupper = 64-np.abs(np.min((0, pore_part.shape[0] -
(lowerlim_x + 64))))
894         target_z = curr_z
895         if int(target_z) + (zmax-zmin) > pore_part.shape[2]:
896             print('continue 2 activated')
897             continue
898         if int(target_z) < 0:
899             print('continue 3 activated')
900             continue
901         try:
902             test_window = pore_part[curr_x: curr_x+ int(xmax-
xmin), curr_y:curr_y + int(ymax-ymin), int(target_z):int(target_z)+(zmax-
zmin)] + data[xmin:xmax , ymin:ymax, zmin:zmax]
903             except Exception as e:
904                 print(e)
905                 print('continue 4 activated')
906                 continue
907             if 0 in test_window.shape:
908                 print('continue 5 activated')
909                 continue
910
911             elif np.max(test_window) > 1:
912                 print('continue 6 activated')
913                 collision_z = np.where(test_window >1)[2][0]
914                 continue
915             try:
916                 xdatastart = np.min(np.where(data)[0])
917                 labelpore = measure.label(data[xmin:xmax , ymin:ymax,
zmin:zmax])
918
919                 oldpore = pore_part[curr_x: curr_x+ int(xmax-xmin),
curr_y:curr_y + int(ymax-ymin), int(target_z):int(target_z)+(zmax-zmin)]
920                 newpore = data[xmin:xmax , ymin:ymax, zmin:zmax]
+ pore_part[curr_x: curr_x+ int(xmax-xmin), curr_y:curr_y + int(ymax-ymin),
int(target_z):int(target_z)+(zmax-zmin)]
921                 if len(measure.regionprops(measure.label(newpore)))
!= len(measure.regionprops(measure.label(oldpore))) + 1:
922                     print(len(measure.regionprops(measure.label(newpore))),len(measure.regionprop
s(measure.label(oldpore))), 'cmerged')
923                     print('continue 7 activated')
924                     continue
925                 if generated:
926                     pore_part[curr_x: curr_x+ int(xmax-xmin),
curr_y:curr_y + int(ymax-ymin), int(target_z):int(target_z)+(zmax-zmin)] =
data[xmin:xmax , ymin:ymax, zmin:zmax]+ pore_part[curr_x: curr_x+ int(xmax-
xmin), curr_y:curr_y + int(ymax-ymin), int(target_z):int(target_z)+(zmax-
zmin)]
927                 else:
928                     pore_part[curr_x: curr_x+ int(xmax-xmin),
curr_y:curr_y + int(ymax-ymin), int(target_z):int(target_z)+(zmax-zmin)] =
data[xmin:xmax , ymin:ymax, zmin:zmax]+ pore_part[curr_x: curr_x+ int(xmax-

```

```

xmin), curr_y:curr_y + int(ymax-ymin), int(target_z):int(target_z)+(zmax-
zmin)]
929         except Exception as f:
930             print(f)
931             print('continue 8 activated')
932             continue
933
934     actual_vols.append(measure.regionprops(measure.label(np.array(data)))
[0].area)
934     breakpoint()
935     return pore_part, target_list_size
936 def nan_helper(y):
937     """Helper to handle indices and logical indices of NaNs.
938
939     Input:
940         - y, 1d numpy array with possible NaNs
941     Output:
942         - nans, logical indices of NaNs
943         - index, a function, with signature indices= index(logical_indices),
          to convert logical indices of NaNs to 'equivalent' indices
944     Example:
945         >>> # linear interpolation of NaNs
946         >>> nans, x= nan_helper(y)
947         >>> y[nans]= np.interp(x(nans), x(~nans), y[~nans])
948     """
949
950
951     return np.isnan(y), lambda z: z.nonzero()[0]
952 def trim_boundary(shell, pore_part):
953     for sample in range(pore_part.shape[2]):
954
955         oldtime = time.time()
956         coords = np.array(np.where(shell[:, :, sample]))
957         index_anglesort = np.argsort(calc_theta(coords, shell[:, :, sample]))
958         coords_pores = np.array(np.where(pore_part[:, :, sample]))
959         outside_bounds = measure.points_in_poly(coords_pores.T,
np.array(coords)[:, index_anglesort].T)
960         test = np.copy(pore_part[:, :, sample])
961
962         test[coords_pores[0], coords_pores[1]] = np.array(outside_bounds,
dtype = 'int')
963         pore_part[:, :, sample] = test
964
965     return pore_part
966 def threshold_pore_size(profile_2d, pore_part, boundary = None, name = ''):
967     im_label = measure.label(pore_part)
968     props = measure.regionprops(im_label)
969     small_props= [prop for prop in props if prop.area < 8]
970     centroids = [prop.centroid for prop in props]
971     polar, rs, ts, o, r, out_h, out_w = linear_polar(pore_part[:, :, 0],
verbose = 1)
972     radii = []
973     angles = []
974     for centroid in centroids:
975
976         x = centroid[0]

```

```

977         y = centroid[1]
978         z = int(centroid[2])
979         r_index, theta_index, theta =
map_pixel(int(x),int(y),pore_part[:,:,:z], o = o, r=r, out_h = out_h, out_w =
out_w, debug= False )
980         radius = profile_2d[z, theta_index] - r_index
981         oldtime = time.time()
982         r_index, theta_index, theta =
map_pixel(int(x),int(y),pore_part[:,:,:z], o = o, r=r, out_h = out_h, out_w =
out_w, debug= False )
983         newtime = time.time()
984         print(newtime-oldtime)
985         radii.append(radius)
986         angles.append(theta)
987
988
989
990         plt.imshow(pore_part[:,:,:z]+boundary[:,:,:z])
991         plt.scatter(int(y), int(x))
992         plt.title(str(radius))
993         plt.savefig('./failures/polar' + name + 'frame'+str(z))
994         plt.clf()
995         plt.imshow(linear_polar(boundary[:,:,:z]))
996         plt.scatter(theta_index, r_index)
997         plt.title(str(profile_2d[z, theta_index]) + ' radius: ' +
str(r_index) + ' calc diff: ' + str(radius))
998         plt.savefig('./failures/polar' + name + 'polarframe'+str(z))
999         plt.clf()
1000
1001 def plt_scatter_matrices(prior_realizations, post_realizations,
variable_list, plt_prior=True, plt_post=True, plt_corr=True,
print_corr=True):
1002     """
1003     Plot the prior and posterior scatter matrices (on top of each other) for
a *variable_list*.
1004     This is from a dictionary of *prior_realizations* and
*prior_realizations*.
1005     """
1006     from pandas import DataFrame
1007     from pandas.plotting import scatter_matrix
1008     import seaborn as sns
1009
1010     def plt_corr_sns(corr):
1011
1012         f, ax = plt.subplots(figsize=(10, 8))
1013         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)
1014
1015         shape_prior = np.array(prior_realizations[variable_list[0]]).shape
1016         shape_post = np.array(post_realizations[variable_list[0]]).shape
1017         prior_data = prior_realizations.copy()
1018         prior_data['Case'] = ['Ground
Truth']*shape_prior[0]#np.zeros(shape_prior)
1019         post_data = post_realizations.copy()

```



```

1021     post_data['Case'] = ['Reconstructed']*shape_post[0]*np.ones(shape_post)
1022     plt.figure(figsize = [8,6], dpi = 150)
1023
1024     if plt_prior and plt_post:
1025         for key in prior_data:
1026             prior_data[key] = np.append(prior_data[key],post_data[key])
1027             data = DataFrame(prior_data)
1028     elif plt_post:
1029         data = DataFrame(post_data)
1030     else:
1031         data = DataFrame(prior_data)
1032     data.to_csv('Generated_data.csv')
1033     return
1034 def fill_boundary(shell, pore_part):
1035     oldtime = time.time()
1036     test = np.zeros(np.shape(pore_part[:,:,:]))
1037     for sample in range(pore_part.shape[2]):
1038         center = (pore_part.shape[0]//2, pore_part.shape[1]//2)
1039         filled = np.array(segmentation.flood(shell[:,:,:sample], (310,310),
connectivity = 0), dtype = int)
1040         pores = np.array(pore_part[:,:,:sample]>0, dtype = int)
1041         test[:,:,:sample] = np.array((filled - pores) > 0, dtype= int)
1042         quadrant_check = len(np.where(test[0,0,:])[0]) +
len(np.where(test[-1,0,:])[0])+ len(np.where(test[0,-1,:])[0])+
len(np.where(test[-1,-1,:])[0])
1043
1044         if quadrant_check > 0:
1045             print("Segmentation didn't work, trying dilation ", sample)
1046
1047             dilated = morphology.dilation(shell[:,:,:sample])
1048             skeleton = morphology.skeletonize(dilated)
1049             filled = morphology.flood(np.array(skeleton,dtype = 'uint8'),
(310,310), connectivity = 0)
1050             pores = np.array(pore_part[:,:,:sample]>0, dtype = int)
1051             test[:,:,:sample] = np.array((filled - pores) > 0, dtype= int)
1052             quadrant_check = len(np.where(test[0,0,:])[0]) +
len(np.where(test[-1,0,:])[0])+ len(np.where(test[0,-1,:])[0])+
len(np.where(test[-1,-1,:])[0])
1053
1054             if quadrant_check >0:
1055                 print("Segmentation didn't work, trying angle based method ",
sample)
1056                 coords = np.array(np.where(shell[:,:,:sample]))
1057                 index_anglesort = np.argsort(calc_theta(coords,
shell[:,:,:sample]))
1058                 coords_pores = np.array(np.where(pore_part[:,:,:sample]))
1059
1060                 solid =
measure.grid_points_in_poly(pore_part[:,:,:sample].shape, np.array(coords)[: ,
index_anglesort].T)
1061                 test[:,:,:sample] = np.array((np.array(solid, dtype = int) -
np.array(pore_part[:,:,:sample]>0, dtype = int)) > 0, dtype = int)
1062
1063     return test
1064

```

```

1065 def n_bins_study(profile_2d, pore_part, generated_boundary, gt_pores):
1066     threshold_pore_size(profile_2d, pore_part, generated_boundary)
1067     list_means = []
1068     for n_bins in [5, 10, 20, 30, 50, 100]:
1069         pore_part_test, _ = replace_sampling(np.zeros((pore_part.shape[0],
1070 pore_part.shape[1], 200)), n_bins = n_bins)
1071         list_means.append(np.mean(pore_part_test, axis = 2))
1072         plt.clf()
1073         plt.figure(figsize = [4,3], dpi = 150)
1074         pore_dataset = measure.regionprops(measure.label(pore_part_test))
1075         pore_gt_dataset, _, _ =
1076 extract_pores(gt_pores[:, :, :200]) #measure.regionprops(measure.label())
1077         locations = [pore.centroid for pore in pore_dataset]
1078         loc_pores = np.array(locations)
1079         kdtree = spatial.KDTree(locations)
1080         dd, ii = kdtree.query(locations, len(locations))
1081         voxelsize = 3.49
1082         nearest_neighbor = dd[:, 1:] * voxelsize
1083         neighbors = dd[:, 1] * voxelsize
1084         edges, hist = np.histogram(dd[:, 1:] * voxelsize, bins =
1085 np.arange(0, 800, 25) * voxelsize)
1086         plt.plot(hist[:-1], edges, linewidth = 2.0, label= "Original
1087 Distribution")
1088         gtlocations = [pore.centroid for pore in pore_gt_dataset]
1089         gtloc_pores = np.array(gtlocations)
1090         gtkdtree = spatial.KDTree(gtlocations)
1091         gtdd, gtii = kdtree.query(gtlocations, len(gtlocations))
1092         gtnearest_neighbor = gtdd[:, 1:] * voxelsize
1093         gtneighbors = gtdd[:, 1] * voxelsize
1094         gtedges, gthist = np.histogram(gtdd[:, 1:] * voxelsize, bins =
1095 np.arange(0, 800, 25) * voxelsize)
1096         plt.plot(gthist[:-1], gtedges, linewidth = 2.0, label = 'Generated
1097 Distribution')
1098         plt.xlabel(r"Distance [ $\mu$  m]")
1099         plt.ylabel("Number of Pores")
1100         plt.tight_layout()
1101         legend(location='best')
1102         frame_tick()
1103         plt.savefig('matriximproved_boundaryaddednbins' + "/rdf" + str(n_bins)
1104 + ".png")
1105         plt.clf()
1106         plt.imshow('matriximproved_boundaryaddednbins'
1107 + '/truedensity'+str(n_bins)+ 'example.png',
1108 np.sum(pore_part_test+generated_boundary, axis = 2), cmap= 'binary')
1109         plt.imshow('matriximproved_boundaryaddednbins'
1110 + '/truedensity'+str(n_bins)+ 'gt.png', np.sum(gt_pores[:, :, :500], axis = 2),
1111 cmap= 'binary')
1112         plt.colorbar()

```

```

1109         plt.savefig()
1110         plt.clf()
1111
1112
1113 def save_binary_segment(segment, fname, voxelsize, index, vname = None):
1114     voxel_conversions=[vname]
1115     if segment.shape[0] > segment.shape[1]:
1116         l_pad = (segment.shape[0] - segment.shape[1])//2
1117         r_pad = (segment.shape[0] - segment.shape[1]) - l_pad
1118         padstack = np.pad(segment, ((0,0),(l_pad,r_pad), (0,0)))
1119
1120     elif segment.shape[0] < segment.shape[1]:
1121         l_pad = (segment.shape[1] - segment.shape[0])//2
1122         r_pad = (segment.shape[1] - segment.shape[0]) - l_pad
1123         padstack = np.pad(segment, ((l_pad,r_pad),(0,0), (0,0)))
1124     else:
1125         padstack = segment
1126     for size in [64, 128, 256, 512, padstack.shape[0]]:
1127
1128         ratio = size/padstack.shape[0]
1129         z_size = np.max([int(padstack[:, :, :100].shape[2]*ratio), 1])
1130         resized_imstack = resize(np.array(padstack[:, :, :100]),
1131 (size,size,z_size), anti_aliasing=False, order =0 )
1132         print(ratio,padstack.shape[2],resized_imstack.shape,
1133 int(padstack.shape[0]*ratio), (padstack.shape[0]*ratio),
1134 size*voxelsize/ratio, "diameter of bounding box")
1135         resized_arr = np.array(resized_imstack>0,dtype='uint8')
1136         if size == padstack.shape[0]:
1137             np.save(fname + str(index) + '_' + "fullres" , resized_arr*255)
1138         else:
1139             np.save(fname + str(index) + '_' +str(size) , resized_arr*255)
1140         voxel_conversions.append(voxelsize/ratio)
1141     return padstack.shape, voxel_conversions
1142
1143
1144 def make_caps(segments, capsize = 100):
1145     props = measure.regionprops(measure.label(segments[:, :, 0]))
1146     prop_max = props[np.argmax([prop.area for prop in props])]
1147     centroid = prop_max.centroid
1148     plt.plot(centroid[1], centroid[0], 'r.')
1149     plt.plot(centroid[1]+ prop_max.major_axis_length//2, centroid[0] , 'r.')
1150     from skimage.draw import circle
1151
1152     rr, cc= circle(centroid[0], centroid[1],
1153 (prop_max.major_axis_length*1.03)//2, shape =segments[:, :, 0].shape )
1154     circ_image = np.zeros(segments[:, :, 0].shape)
1155     circ_image[rr,cc] = 1
1156     start = np.repeat(circ_image[:, :, np.newaxis], capsize, axis = 2)
1157
1158     props = measure.regionprops(measure.label(segments[:, :, -1]))
1159     prop_max = props[np.argmax([prop.area for prop in props])]
1160     centroid = prop_max.centroid
1161     plt.plot(centroid[1], centroid[0], 'r.')

```

```

1160     plt.plot(centroid[1]+ prop_max.major_axis_length//2, centroid[0] , 'r.')
1161     from skimage.draw import circle
1162     rr, cc= circle(centroid[0], centroid[1],
    (prop_max.major_axis_length*1.03)//2,shape =segments[:, :,0].shape)
1163     circ_image = np.zeros(segments[:, :, -1].shape)
1164     circ_image[rr,cc] = 1
1165     end = np.repeat(circ_image[:, :, np.newaxis], capsize, axis = 2)
1166
1167     return np.array(start, dtype = 'uint8'), np.array(end, dtype = 'uint8')
1168
1169
1170 def combine_segments(fname, num, actual_size, caps = True, folder_index =
    0):
1171     capsize = 100
1172     for resolution in [64, 128, 256, 512, "fullres"]:
1173
1174         pname_test = fname+'{}_'.format(0) + str(resolution) + '.numpy'
1175         segment_part_test = np.array(np.load(pname_test, allow_pickle =
    True), dtype = 'uint8')
1176         segments = np.zeros((segment_part_test.shape[0],
    segment_part_test.shape[1], 0), dtype = 'uint8')
1177         for part in range(0, num, 100):
1178             pname = fname+'{}_'.format(part) + str(resolution) + '.numpy'
1179             segment_part = np.array(np.load(pname, allow_pickle = True),
    dtype = 'uint8')
1180             segments = np.dstack((segments, segment_part))
1181             if caps:
1182                 start,end = make_caps(segments, capsize = capsize)
1183                 full_segments = np.dstack((start, segments, end))
1184             if caps:
1185                 save_name = fname.split('partial')[0] + "padded"
1186                 if resolution == "fullres":
1187                     assert full_segments.shape == (actual_size[0],
    actual_size[1], actual_size[2] + capsize*2)
1188                 else:
1189                     save_name = fname.split('partial')[0]
1190                     full_segments = segments
1191                 if resolution == 'fullres':
1192                     assert full_segments.shape == actual_size
1193                 np.save(save_name+'{}.numpy'.format(resolution), np.array(full_segments,
    dtype = 'uint8'))
1194                 if resolution == 64:
1195                     os.makedirs(save_name + '/64datasamples', exist_ok=True)
1196                     np.save(save_name+'/64datasamples
    /{}.numpy'.format(folder_index), np.array(full_segments, dtype = 'uint8'))
1197
1198
1199 def clean_segments(fname):
1200     command = 'rm ' + fname + '*.numpy'
1201     print(command)
1202     os.system(command)
1203
1204 def save_sample(folder_index, generated_dir, imstack_all = None, voxelsize =
    None, generated = True, boundary_stack_all = None):
1205     frame_window = 100

```

```
1206     shift = 100
1207     num = None
1208     num = imstack_all.shape[2]
1209     shell_fragment = 'segment_{}_plane_removed_partial_'
1210     fragment = 'segment_{}_partial__'
1211     index = 0
1212     folder_name = generated_dir + '/Part{}/'.format(folder_index)
1213
1214     os.makedirs(folder_name, exist_ok=True)
1215     while index < num:
1216         pores_total = []
1217         print("Processing pores, index = " + str(index) + " out of " +
1218 str(num) + " ...")
1218         im = np.copy(imstack_all[:, :, index:index + frame_window])
1219         boundary_stack = np.copy(boundary_stack_all[:, :,
1220 index:index+frame_window])
1220         fractured = np.array(fill_boundary(boundary_stack, im), 'uint8')
1221         pores_removed = np.array(fill_boundary(boundary_stack,
1222 np.zeros(im.shape)), 'uint8')
1222
1223         square_shape, voxel_conversions =
1224 save_binary_segment(fractured, folder_name + fragment.format(folder_index),
1225 voxelsize = voxelsize, index = index)
1224         square_shape, _ = save_binary_segment(pores_removed, folder_name +
1225 shell_fragment.format(folder_index), voxelsize = voxelsize, index = index)
1225         index = index + shift
1226
1227         combine_segments(os.path.join(folder_name,
1228 fragment.format(folder_index)), num = num, actual_size = (square_shape[0],
1229 square_shape[1], imstack_all.shape[2]), caps = True, folder_index =
1230 folder_index)
1231
1232         combine_segments(os.path.join(folder_name,
1233 fragment.format(folder_index)), num = num, actual_size = (square_shape[0],
1234 square_shape[1], imstack_all.shape[2]), caps = False, folder_index =
1235 folder_index)
1236
1237         clean_segments(os.path.join(folder_name, fragment.format(folder_index)))
1238
1239         combine_segments(os.path.join(folder_name,
1240 shell_fragment.format(folder_index)), num = num, actual_size =
1241 (square_shape[0], square_shape[1], imstack_all.shape[2]), caps = True,
1242 folder_index = folder_index)
1243
1244         combine_segments(os.path.join(folder_name,
1245 shell_fragment.format(folder_index)), num = num, actual_size =
1246 (square_shape[0], square_shape[1], imstack_all.shape[2]), caps = False,
1247 folder_index = folder_index)
1248
1249         clean_segments(os.path.join(folder_name,
1250 shell_fragment.format(folder_index)))
1251
1252         return voxel_conversions
1253
1254
1255 def test_sample(sample, generated_dir):
1256     pdir = generated_dir + '/Part{}/'.format(sample)
1257     npyfiles = [os.path.join(pdir, file).split('.')[0] for file in
1258 os.listdir(pdir) if file.endswith('.npy')]
```

```

1241     for npyfile in npyfiles:
1242         sample_dir = './reconstruction/full/testing_binarization/part{}
./format(sample) + npyfile.split('/')[ -1]
1243         os.makedirs(sample_dir, exist_ok = True)
1244         npy = np.load(npyfile+'.npy', allow_pickle = "True")
1245         plt.clf()
1246         plt.title(npyfile.split('/')[ -1] + "beginning")
1247         plt.imshow(npy[:, :, 0])
1248         plt.savefig(sample_dir + '/' + npyfile.split('/')[ -1] + 'begin.png')
1249         plt.clf()
1250         plt.title(npyfile.split('/')[ -1] + "middle")
1251         plt.imshow(npy[:, :, npy.shape[2]//2])
1252         plt.savefig(sample_dir + '/' + npyfile.split('/')[ -1] + 'middle.png')
1253         plt.clf()
1254         print(npyfile.split('/')[ -1])
1255         print("done case " + str(sample))
1256     os.makedirs('./reconstruction/full/testing_binarization/part{}
/tiff_stack'.format(sample), exist_ok= True)
1257     npy = np.load(generated_dir + '/Part{}
/segment_{}_fullres.npy'.format(sample, sample), allow_pickle = True)
1258     for i in range(npy.shape[2]):
1259         plt.imsave('./reconstruction/full/testing_binarization/part{}
/tiff_stack'.format(sample) + "/frame{:05}.png".format(i), npy[:, :, i], cmap
= 'gist_gray')
1260         plt.clf()
1261     os.system('./Fiji.app/ImageJ-linux64 -macro ./reconstruction
/full/testing_binarization/Video.ijm ' + str(sample))
1262     os.system('rm ' + './reconstruction/full/testing_binarization/part{}
/tiff_stack'.format(sample) + "/frame*png")
1263
1264 def main():
1265     tot_frames= 2000
1266     num_frames = 2000
1267     interval = 1
1268     generated_dir  = './reconstruction/results/GeneratedPartSamples'
1269
1270     groundtruth_stack_shape = (566, 571) # example shape from dataset
1271
1272     n_bins = 30
1273     import time
1274     oldtime = time.time()
1275     start = 0#int(sys.argv[1])
1276     for iter in range(start, start+ 100):
1277
1278
1279         for total_frame in range(0, tot_frames, num_frames):
1280             pore_part =
np.zeros((groundtruth_stack_shape[0],groundtruth_stack_shape[1], num_frames),
dtype = 'uint8')
1281
1282             generated_boundary, profile_2d = load_boundary(num_frames =
num_frames, start = total_frame, pore_part_shape = (pore_part.shape[0],
pore_part.shape[1]), return_profile=True)
1283
1284

```



```
1285         before_polar_time = time.time()
1286         pore_part,_ = replace_sampling(pore_part, generated_boundary =
generated_boundary,n_bins = 30)
1287         after_polar_time = time.time()
1288         print(after_polar_time - before_polar_time, "Time taken polar")
1289
1290
1291
1292         before_cartesian_time = time.time()
1293         pore_part,_ = replace_sampling(pore_part, generated_boundary =
generated_boundary,n_bins = 30)
1294         after_cartesian_time = time.time()
1295         print(after_cartesian_time - before_cartesian_time, "Time taken
cartesian")
1296         trim_boundary(generated_boundary, pore_part)
1297
1298         plt.close('all')
1299         voxel_to_micron = {}
1300         voxel_to_micron['Resolution'] = ["Name", 64, 128, 256, 512,
"Full"]
1301         num_samples = 12
1302
1303         conversion = save_sample(iter, voxelsize = 4.87,
boundary_stack_all=generated_boundary, imstack_all = pore_part, generated_dir
= generated_dir)
1304
1305         os.system("7z a "+ generated_dir + " /Part{}.zip ".format(iter) +
generated_dir + " /Part{}/".format(iter, iter))
1306
1307         if os.path.exists(generated_dir + '/Part{}.zip'.format(iter)):
1308             os.system("rm " + generated_dir + " /Part{}/
*.npy".format(iter))
1309             print(time.time() - oldtime, "TIME ELAPSED PER ITERATION")
1310             oldtime = time.time()
1311
1312
1313 if __name__ == "__main__":
1314     main()
1315
```