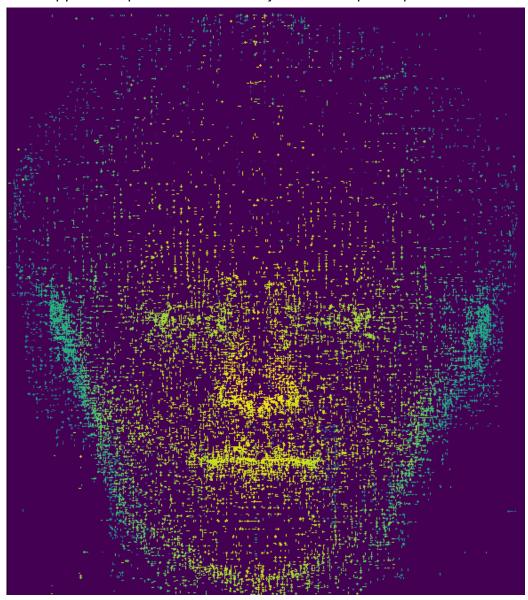
January 18th, 2022

- Met with Dr. Akbas and the team to discuss semester goals for the project.
- Decided to use machine learning to map the applicable significant landmarks on a patient's face.

January 23rd, 2022

- Researched possible solutions for automatically mapping facial landmarks.
- Found plenty of resources for mapping landmarks on 2D images, but very few resources for 3D objects.
- Wrote a python script to convert a 3D .obj file into a depth map of the face elements.



```
pixel_graph = np.full((2500, 2500), 0)
blur_scale = 5

for coordinate in test_array:
    pixel_graph[int(coordinate[0]), int(coordinate[1])] = coordinate[2]
    for i in range(blur_scale):
        pixel_graph[int(coordinate[0]) + i, int(coordinate[1])] = coordinate[2]
        pixel_graph[int(coordinate[0]) - i, int(coordinate[1])] = coordinate[2]
        pixel_graph[int(coordinate[0]), int(coordinate[1]) + i] = coordinate[2]
        pixel_graph[int(coordinate[0]), int(coordinate[1]) - i] = coordinate[2]

plt.figure(figsize=(12,12), dpi=80)
plt.imshow(pixel_graph, interpolation='nearest')
plt.show()
```

January 30th, 2022

- Wrote a python script to create clusters of similar looking faces.
- Created groups based on each patient's 10 most similar looking faces, allowing overlap.
- Created 10 groups of 8 faces without overlap

face_group1	12/6/2021 6:17 PM	File folder
face_group2	12/6/2021 6:17 PM	File folder
face_group3	12/6/2021 6:17 PM	File folder
邝 face_group4	12/6/2021 6:17 PM	File folder
face_group5	12/6/2021 6:17 PM	File folder
face_group6	12/6/2021 6:17 PM	File folder
face_group7	12/6/2021 6:17 PM	File folder
face_group8	12/6/2021 6:17 PM	File folder
face_group9	12/6/2021 6:17 PM	File folder
face_group10	12/6/2021 6:17 PM	File folder
face_group11	12/6/2021 6:17 PM	File folder
face_group12	12/6/2021 6:17 PM	File folder
face_group13	12/6/2021 6:17 PM	File folder
face_group14	12/6/2021 6:17 PM	File folder
face_group15	12/6/2021 6:17 PM	File folder
face_group16	12/6/2021 6:17 PM	File folder
face_group17	12/6/2021 6:17 PM	File folder
🖺 face_group18	12/6/2021 6:17 PM	File folder
face_group19	12/6/2021 6:17 PM	File folder
🖺 face_group20	12/6/2021 6:17 PM	File folder
face group21	12/6/2021 6:17 PM	File folder

邝 face_group1	1/30/2022 9:18 PM	File folder
face_group2	1/30/2022 9:18 PM	File folder
face_group3	1/30/2022 9:18 PM	File folder
邝 face_group4	1/30/2022 9:18 PM	File folder
face_group5	1/30/2022 9:18 PM	File folder
face_group6	1/30/2022 9:18 PM	File folder
face_group7	1/30/2022 9:18 PM	File folder
face_group8	1/30/2022 9:18 PM	File folder
face_group9	1/30/2022 9:18 PM	File folder
face_group10	1/30/2022 9:18 PM	File folder

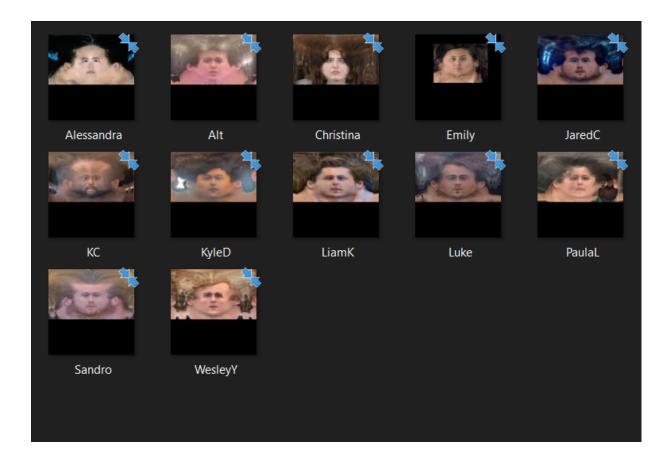
February 2nd, 2022

• Worked with Alessandra to complete sections 1 and 2 of the SRS and SDD documents

February 6th, 2022

- Wrote python script to create face clusters using a k-means algorithm
- Created 8 clusters containing varying amount of patients

face_group0	2/7/2022 6:05 PM	File folder
📑 face_group1	2/7/2022 6:05 PM	File folder
face_group2	2/7/2022 6:05 PM	File folder
face_group3	2/7/2022 6:05 PM	File folder
is face_group4	2/7/2022 6:05 PM	File folder
face_group5	2/7/2022 6:05 PM	File folder
📑 face_group6	2/7/2022 6:05 PM	File folder
face_group7	2/7/2022 6:05 PM	File folder



February 13th, 2022

 Reached out to GANFit database owners: the MICC, in order to download the dataset and generate a larger data pool. However, they were unable to help me and we had to scrap GANFit.

February 20th, 2022

- Researched different clustering algorithms to implement in order to optimize the face clusters.
- Decided upon 5 others in addition to Kmeans: affinity propagation, BIRCH, agglomerative, spectral, and Gaussian mixture clustering

March 2nd, 2022

• Wrote python script to generate clusters using multiple different clustering algorithms, including, affinity propagation, BIRCH, agglomerative, spectral, and Gaussian mixture.

```
def gaussian_mixture():
              df = pd.read_csv('database.csv')
              ratios = df.drop(['patient', 'class'], axis=1)
              gaussian = GaussianMixture(n_components=7).fit(ratios).predict(ratios)
              df['gaussian'] = gaussian
              \textbf{newpath} = r'C: \begin{tabular}{l} \textbf{Newpath
              if os.path.exists(newpath):
                             shutil.rmtree(newpath)
              os.makedirs(newpath)
               for i in range(len(gaussian)):
                               folder_location = (newpath + '\\face_group' + str(df.loc[i, 'gaussian']))
                             file_location = "C:\\Users\\jared\\Documents\\DigitizedRhinoplasty\\DigitizedRhinoplasty-main\\ScanII
                             if not (os.path.exists(folder_location)):
                                             os.makedirs(folder_location)
                                              shutil.copy(file_location, folder_location)
                             except (FileNotFoundError):
gaussian_mixture()
spectral()
agglomerative()
birch()
affinity_propagation()
kmeans()
```

face_group_affinity	3/2/2022 6:43 PM	File folder
face_group_agglomerative	3/2/2022 6:43 PM	File folder
face_group_birch	3/2/2022 6:43 PM	File folder
🌅 face_group_gaussian	3/2/2022 6:43 PM	File folder
face_group_kmeans	3/2/2022 6:43 PM	File folder
🖺 face_group_spectral	3/2/2022 6:43 PM	File folder

• Each folder contains a varying number of clusters depending on the settings of the algorithm.

March 6th

Updated the SRS and SDD documents to align with new goals for the 3rd Sprint.

March 16th

- Created sprint 3 demo presentation.
- Identified all the nose points to plot and manipulate

Nose Points

- Alar base junction ac_l ac_r
- Alar rim's highest point armax_l armax_r
- Alar flare al I al r
- Anterior point of nostril stn_l stn_r
- · Columellar break point cb
- Columellar rim cmin_l cmin_r
- Maxilloanteriorale ma_l ma_r
- Maxillofrontale mf I mf r
- Nasal parenthesis np_l np_r
- Posterior point of nostril itn_l itn_r
- Pronasale prn
- Rhinion r
- · Sellion se
- Sill-base junction sbj_l sbj_r
- Subalare sbal_l sbal_r
- Subnasale sn_l sn_r
- Supratip break point s
- Tip defining point td_l td_r