



Discovering the intrinsic structure of ciliary motion



Joseph Monaco^{1*}, Maurice Marx², Chakra Chennubhotla^{2*}

¹TECBio REU @ Pitt, Dept. of Computational and Systems Biology, University of Pittsburgh, Pittsburgh, PA, 15260

²Dept. of Computational and Systems Biology, University of Pittsburgh, Pittsburgh, PA, 15260

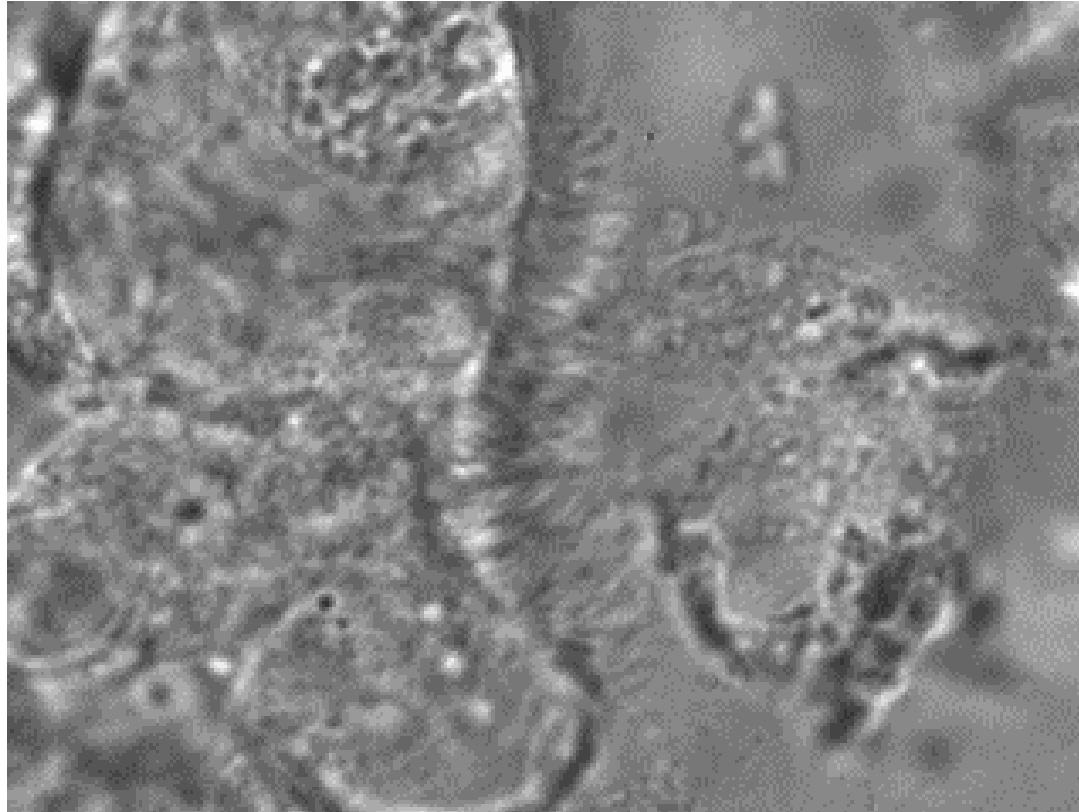
*Dept. of Computer Science, University of Pittsburgh, Pittsburgh, PA, 15260

Goal

Quantify ciliary motion in terms of motion primitives via clustering and
discover motion primitives indicative of abnormal ciliary motion

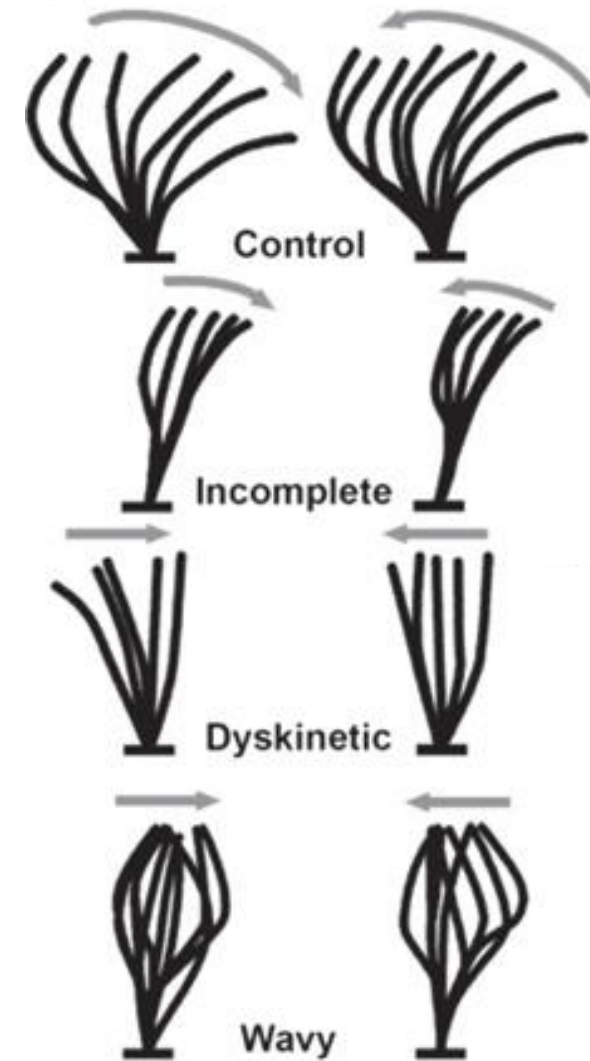
Types of Cilia

- Motile or immotile
- Motile cilia in respiratory tract beat in two steps
 - Power stroke
 - Recovery stroke



Ciliary Motion - Ciliopathies

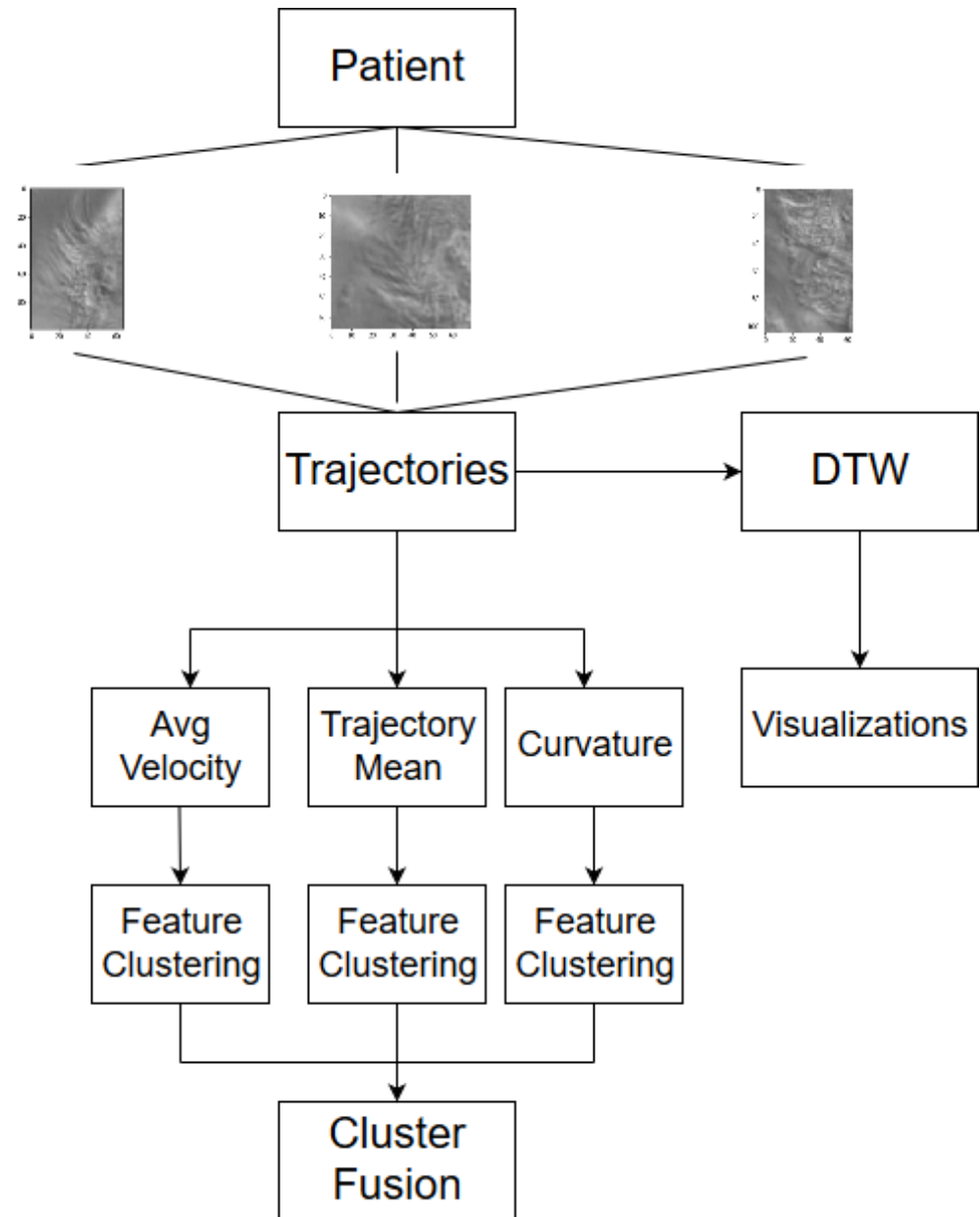
- Sinopulmonary disease
- Heterotaxy/Kartagener's
- Congenital heart disease



Manual detection of abnormal ciliary motion

- Visual examination by experts in detecting ciliary beat abnormalities
- Electron microscopy
- Ciliary beat frequency

Approach

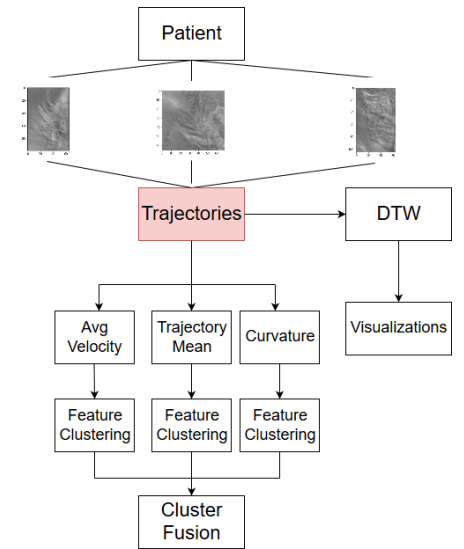
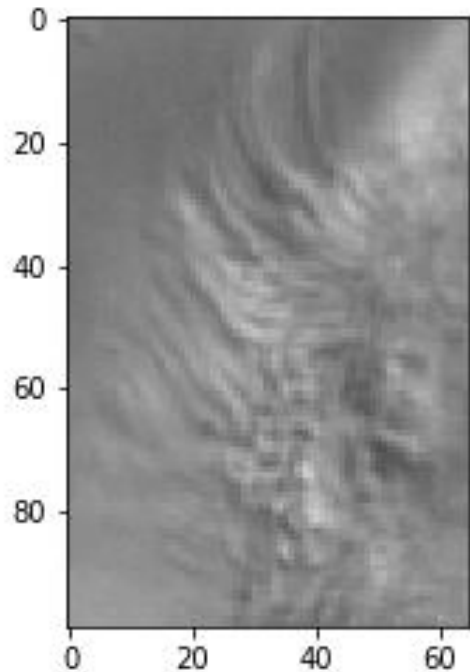


Dataset

- 78 patients
- Nasal brush biopsy
- Labeled by clinicians on a discrete scale from 1 – 4

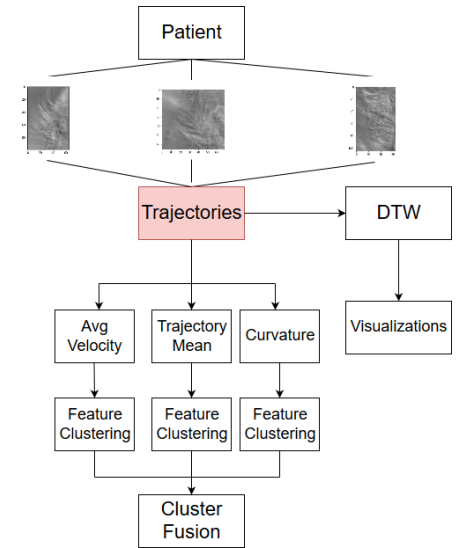
Optical Flow

- Apparent motion of objects between two frames
- Horizontal change and vertical change



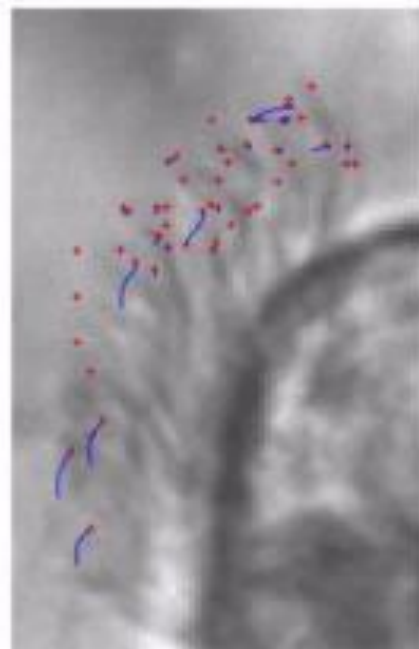
Trajectories

- Describe ciliary motion as trajectories
- Dense trajectories



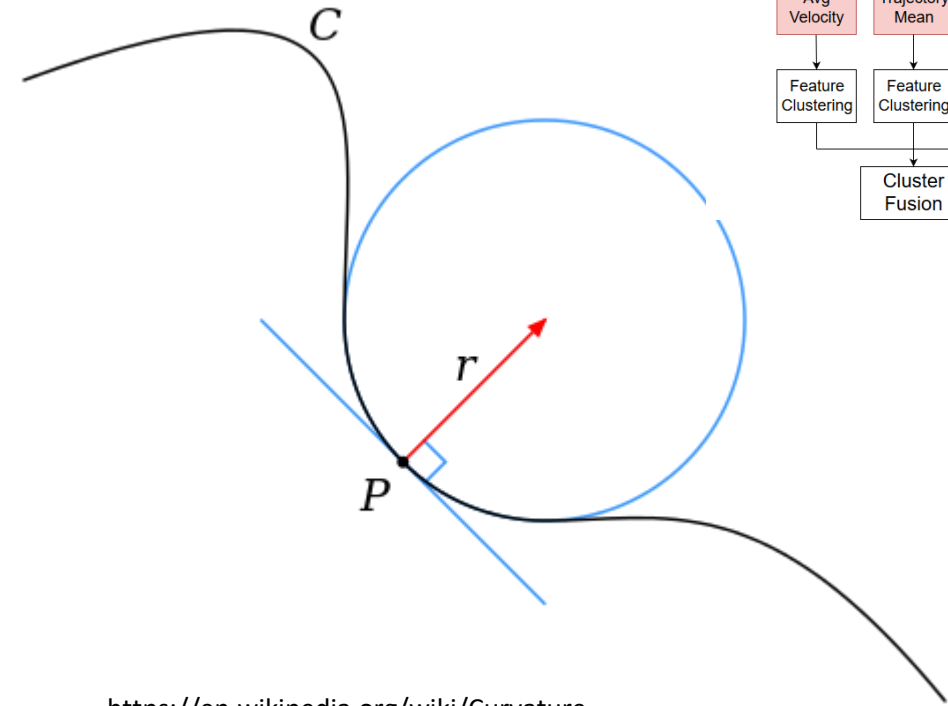
$$P_{t+1} = (x_{t+1}, y_{t+1}) = (x_t, y_t) + (M * \omega_t)|_{(x_t, y_t)}$$

$$P_1, P_2, P_3, \dots, P_M$$

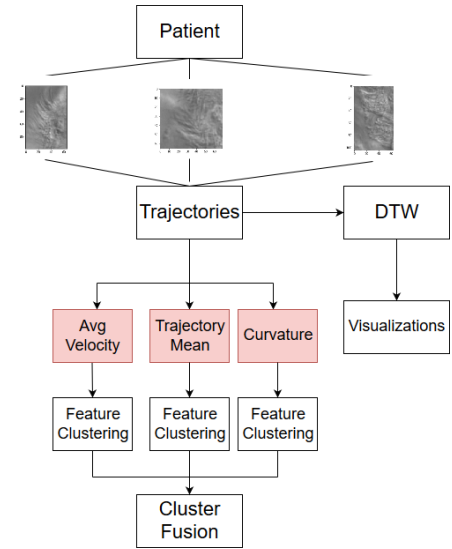


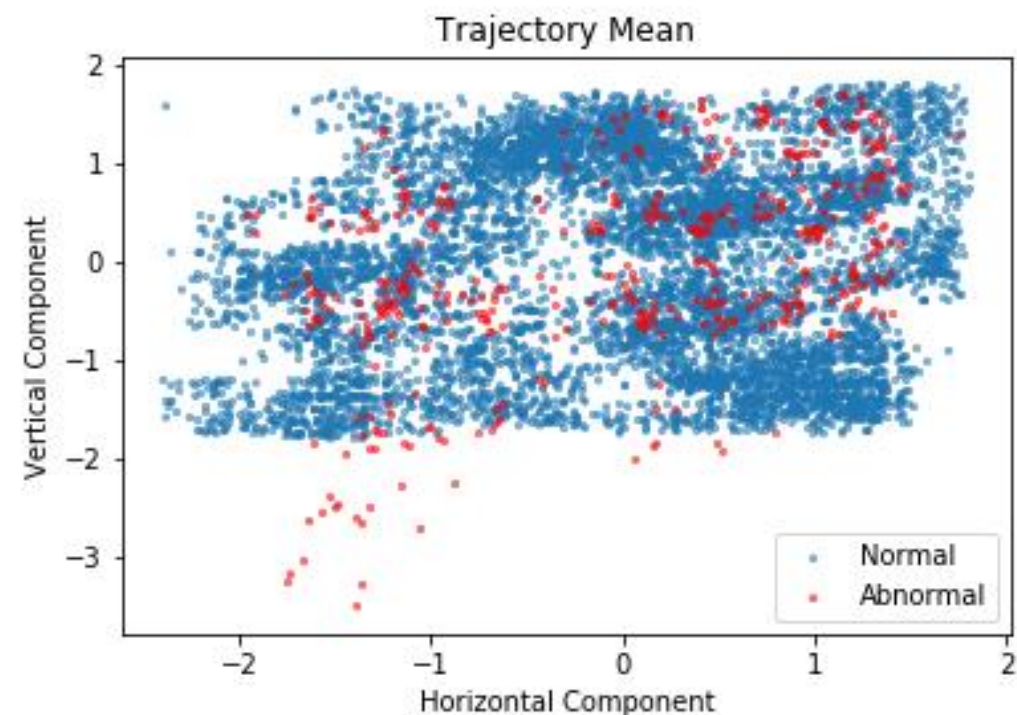
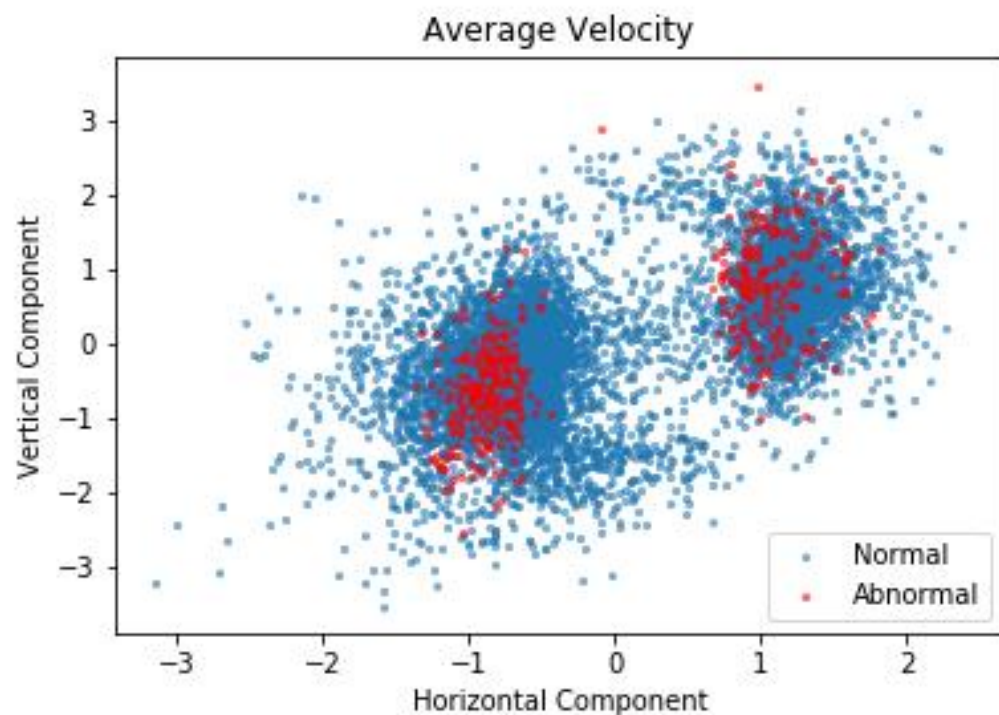
Feature Engineering

- Average velocity
- Trajectory mean
- Curvature



<https://en.wikipedia.org/wiki/Curvature>

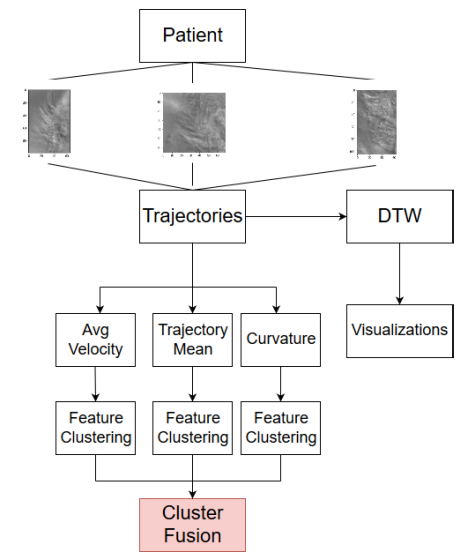




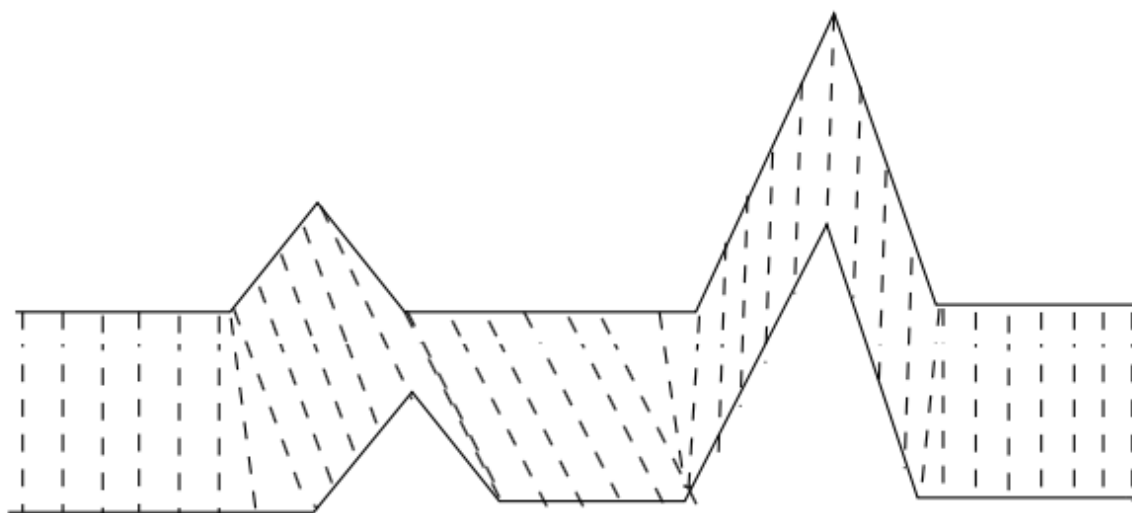
Features – average velocity/trajectory mean

Clustering

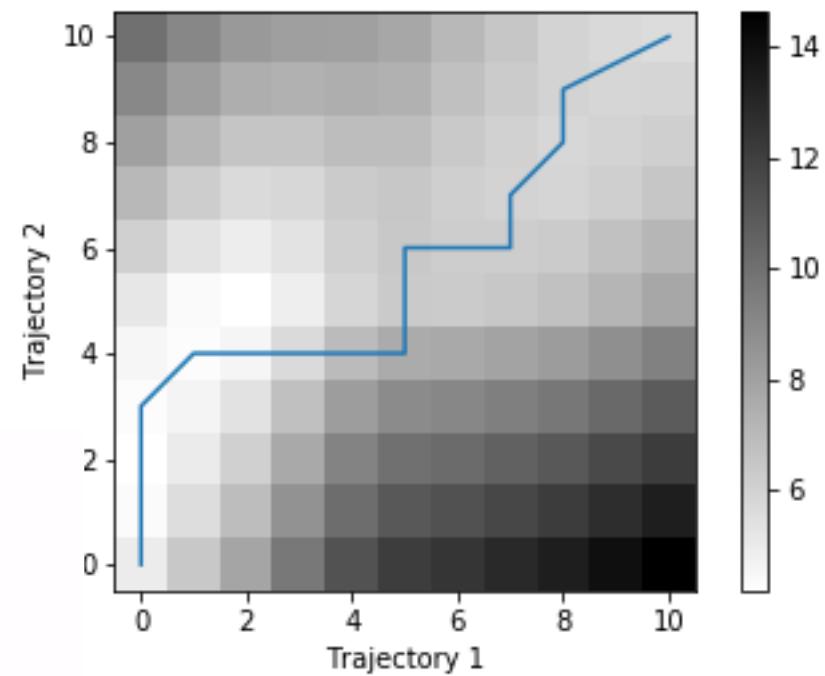
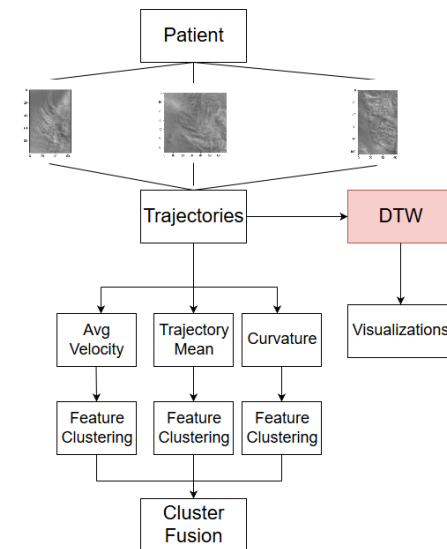
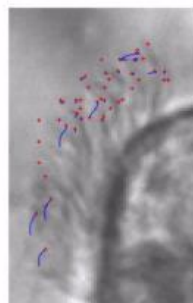
- dbscan for each feature space
- Cluster fusion
 - Pick median feature
 - Combine clusters by picking largest intersection with median feature



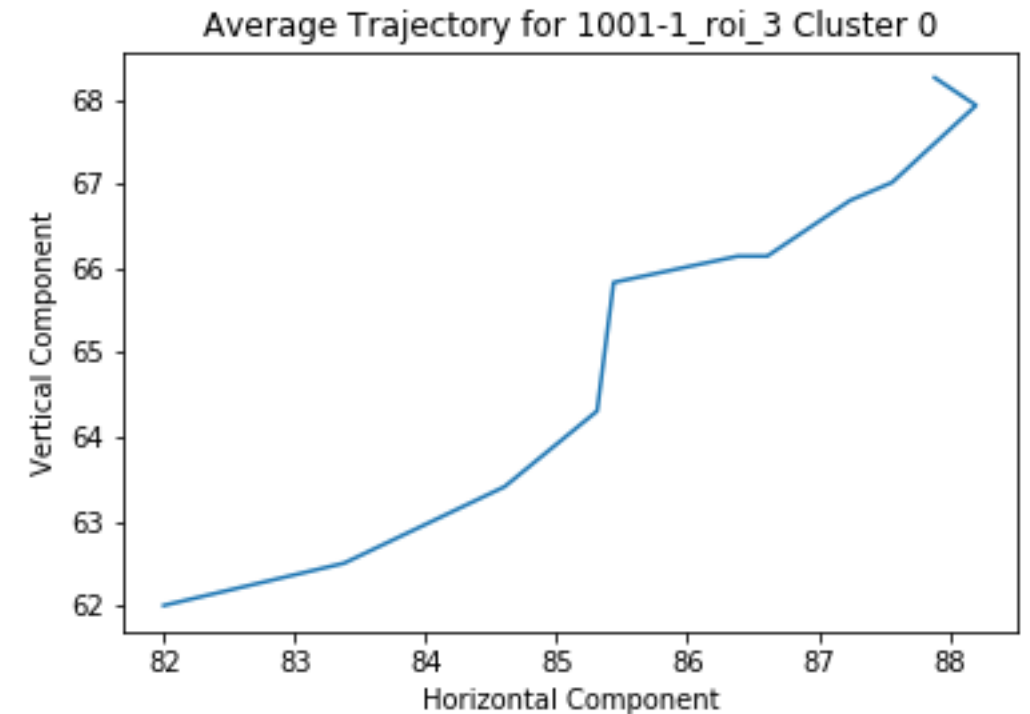
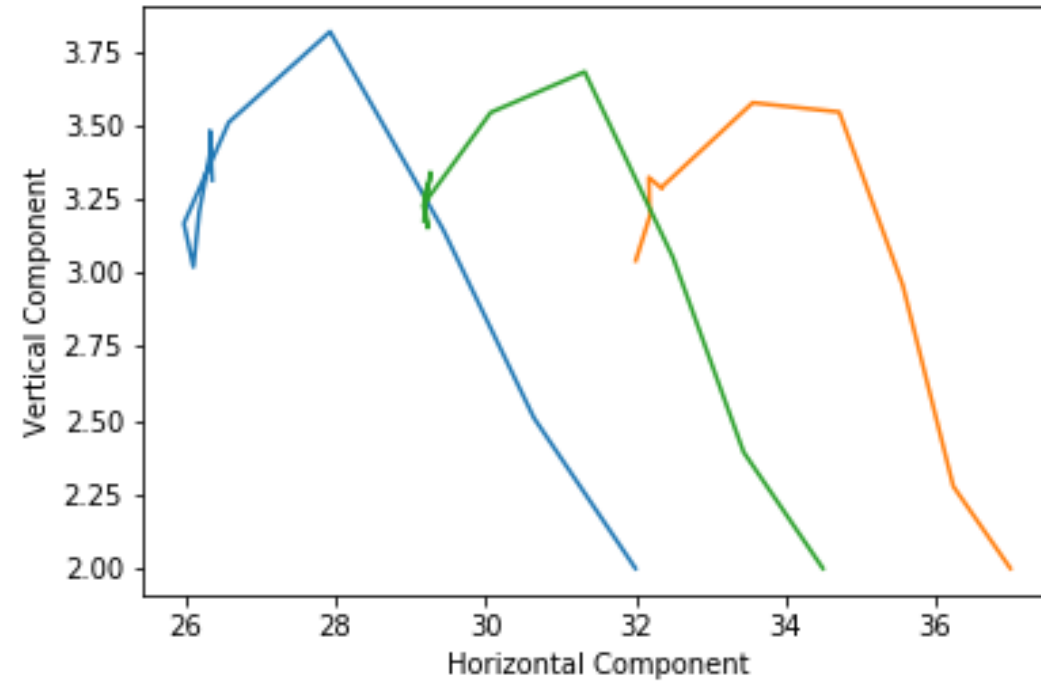
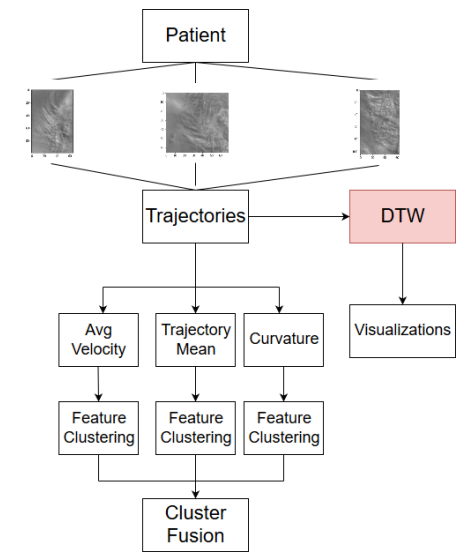
Dynamic time warping



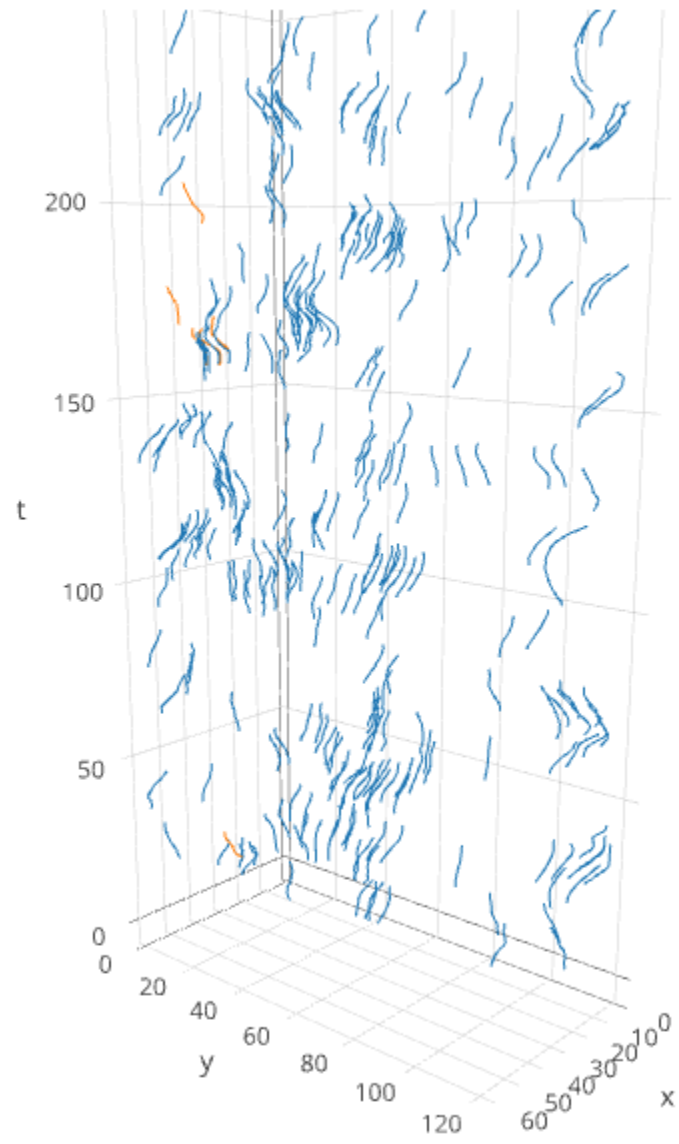
https://en.wikipedia.org/wiki/Dynamic_time_warping



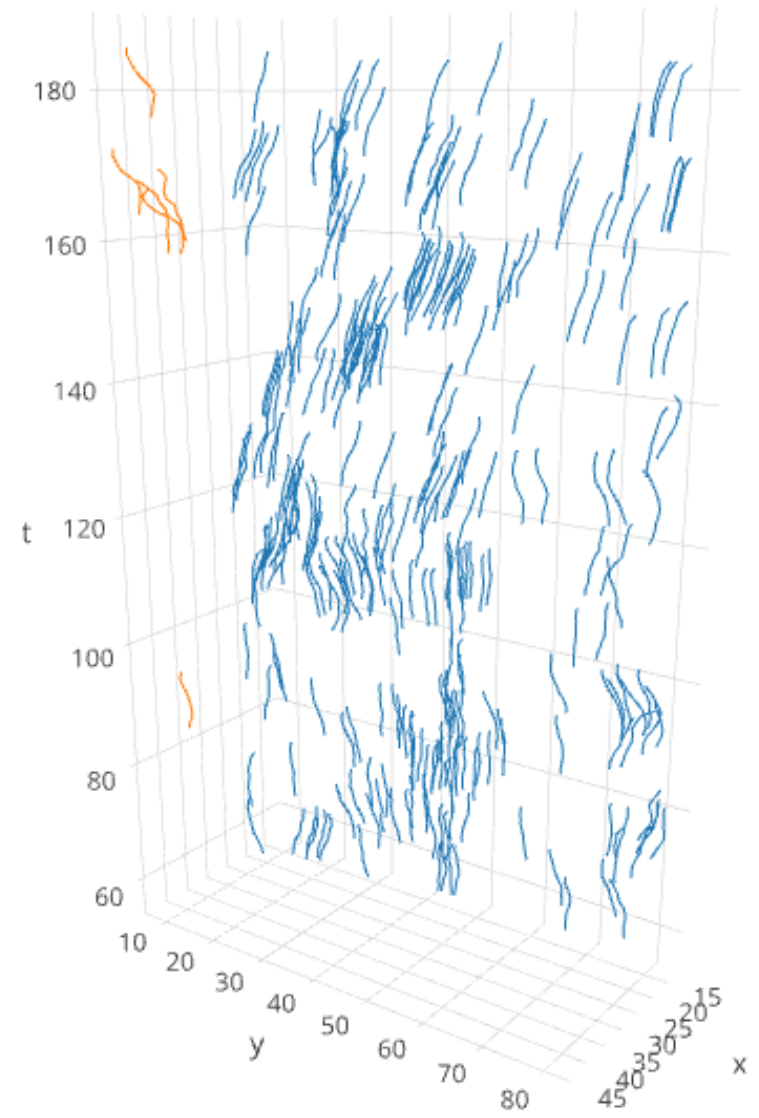
Dynamic time warping – avg trajectory



Trajectories before DTW

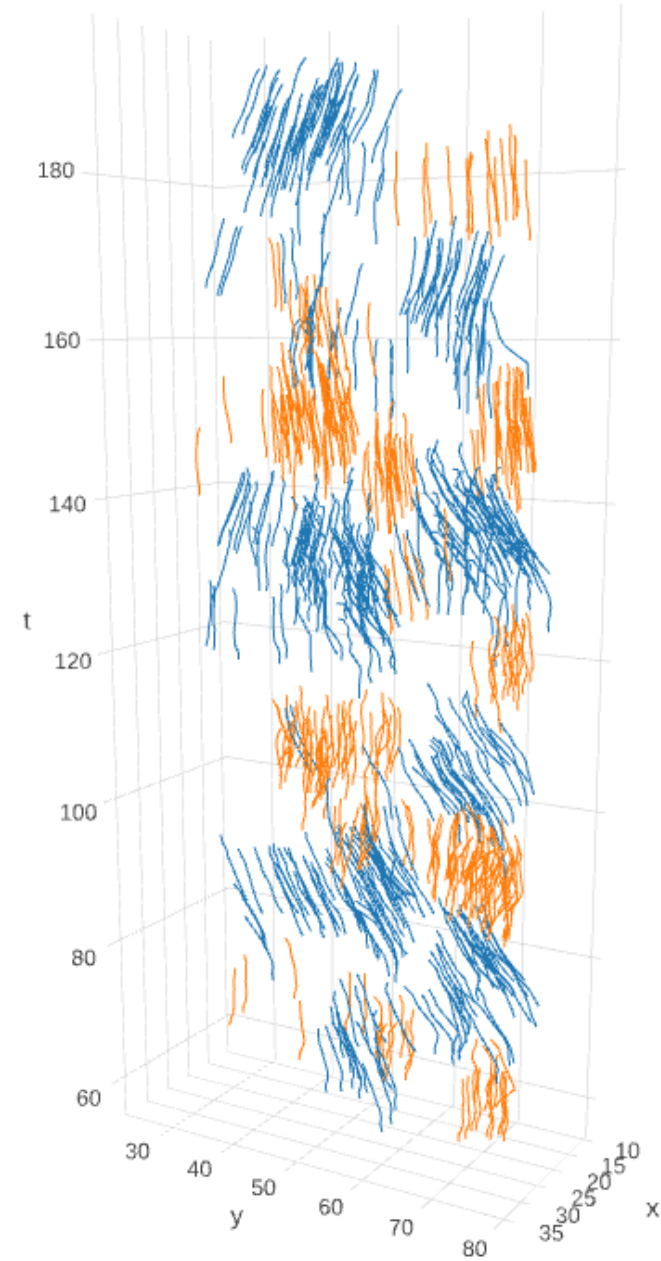


Trajectories after DTW



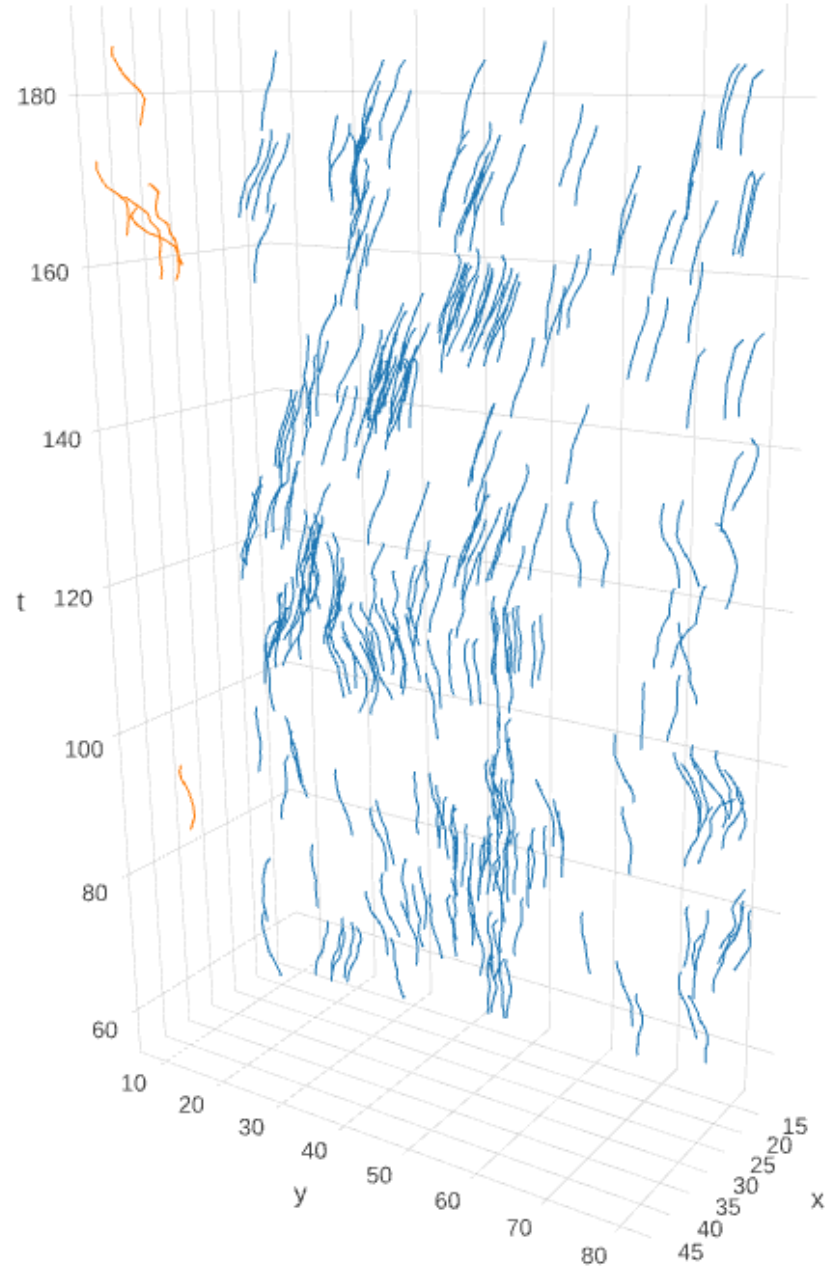
Clustering – trajectories average velocity

- Trajectory points (x, y, t)
- Labeled according to average velocity clustering
- Location and velocity related



Clustering – final trajectories

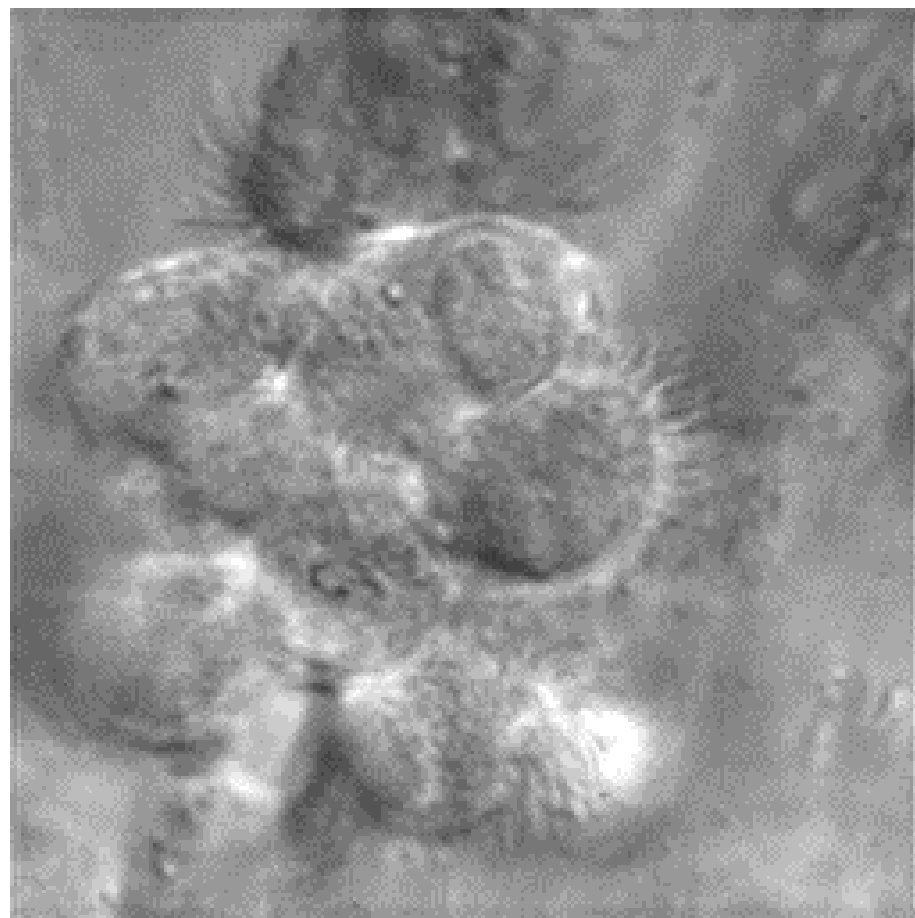
- Trajectory visualization
- Labeled according to their final cluster



Cilia ratio in each cluster

Label	Cluster 0	Cluster 1	Cluster 2	Cluster 3
1 – most normal	0.846	0.698	0.513	0.000
2	0.000	0.000	0.000	0.000
3	0.043	0.067	0.128	0.000
4 – most abnormal	0.110	0.235	0.358	1.000

- 4 clusters discovered – intrinsic motion patterns
- Little variability in abnormal motion
- Stronger variability in normal motion
 - Historically categorized as power/recovery stroke



Future Directions

- Show clusters to clinicians to refine clusters
- Use topic modeling for cluster ratio

Acknowledgements

- Mentor - Dr. Chakra Chennubhotla
 - Maurice Marx
 - Dr. Sherif Khattab
 - NSF
 - You!
-
- Questions?