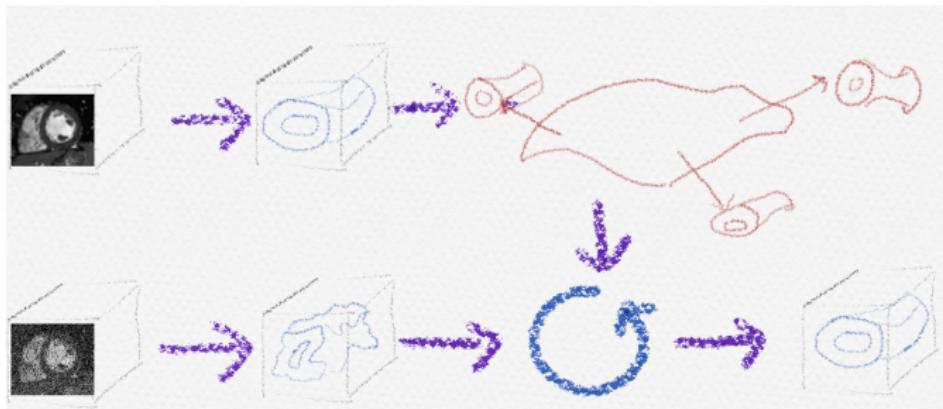


# P1 Cardiac motion estimation

January 24, 2024

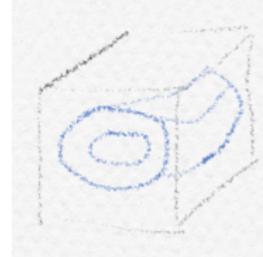
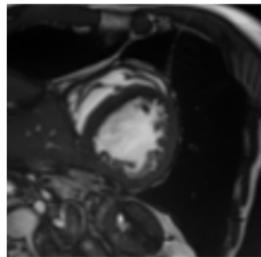


# Setup

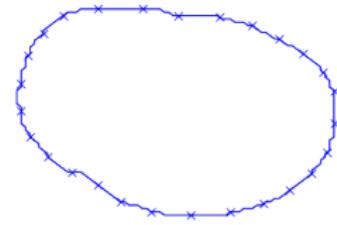
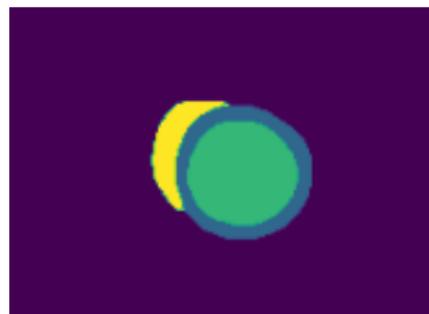
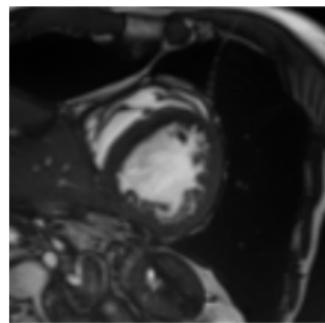
- Needed data: batch of images of the cardiac cycle sliced in  $z$ - and  $t$ -dimension. Typical batch will have a shape like
$$(30, 8, 256, 256),$$

with

- 30  $t$  slices.
- 8  $z$  slices → take middle one → can be omitted.
- 256 × 256 pixel images.



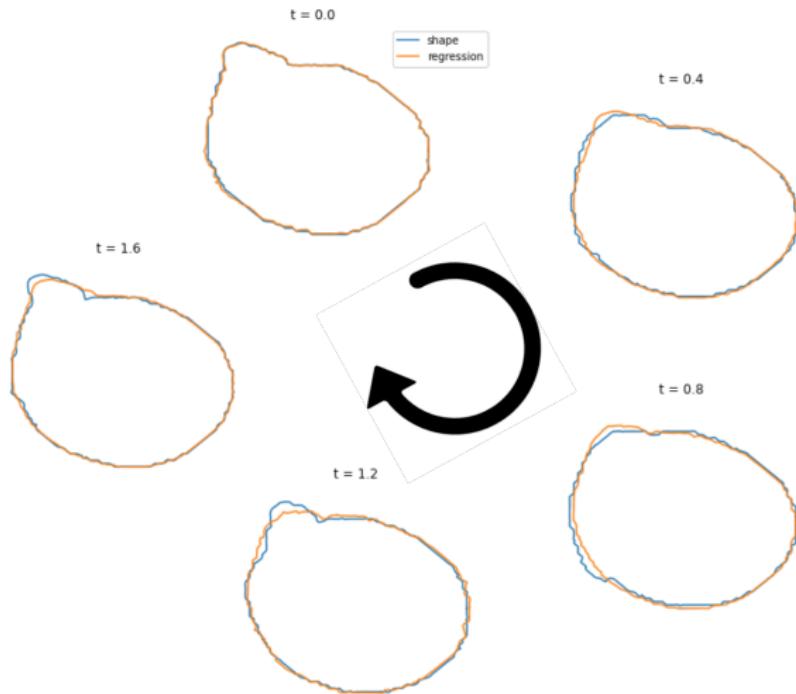
## Register data



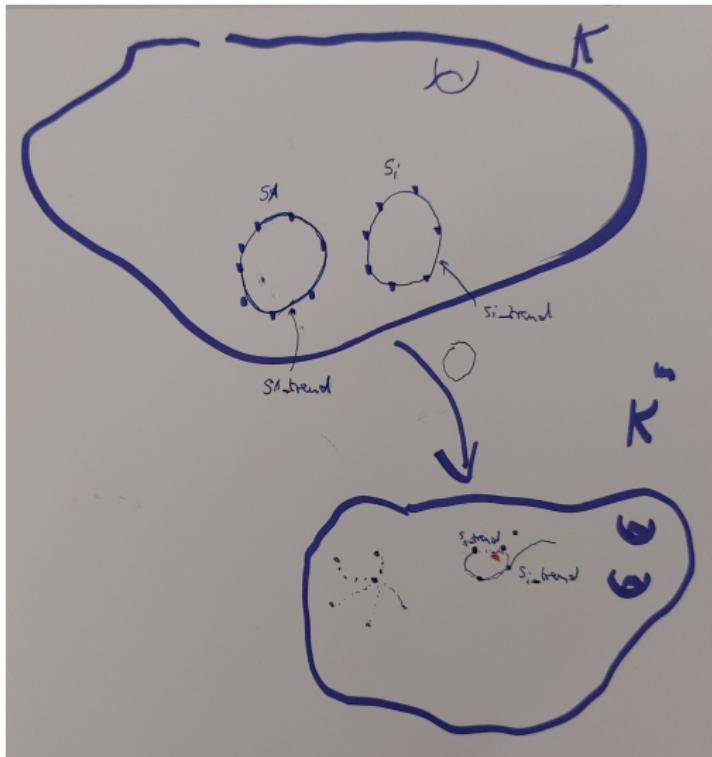
(with skimage.measure.find\_contours)

Blue: Cardiac contour evolution over time

Orange: Regressed shape interpolation

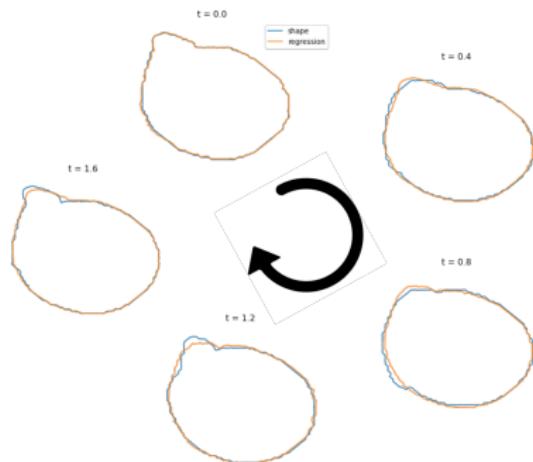


# The idea



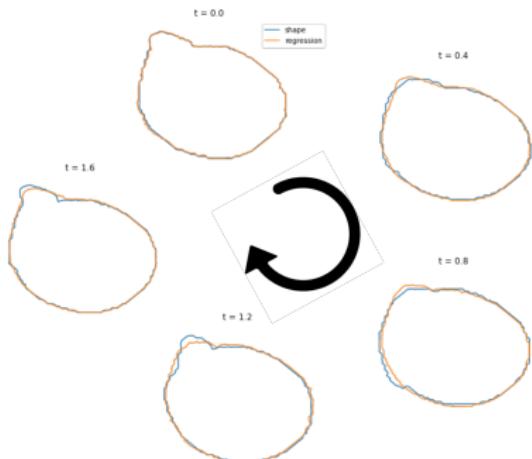
# The idea

regressed shape interpolations are elements on the manifold

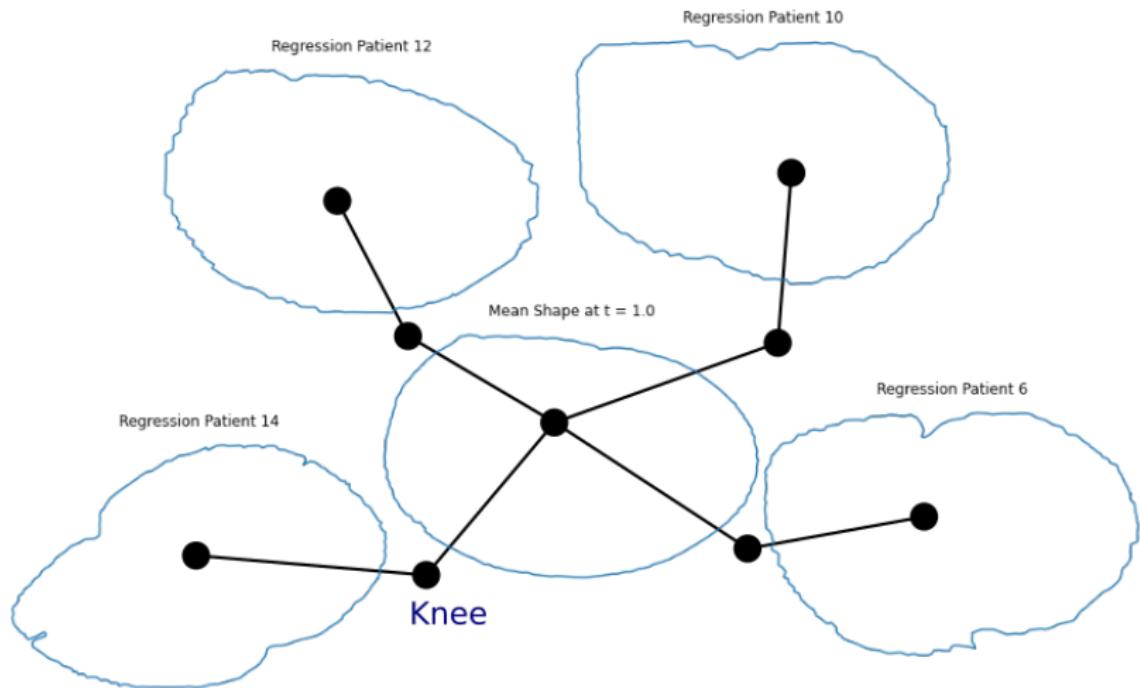


# The idea

the Bezier splines are again points on a manifold



# Approximating Mean Shape on a Manifold



# Classify diseases using mean shape trajectories

Class DCM

Class HCM

Class MINF

Class NOR

Test data: which class?

# Results

- compute distance (on manifold) between Bezier spline of test data and of corresponding means

$$\text{dist}^2(x, y) = \int d^2(x(t), y(t)) dt$$

$$d(a, b) = \min_{R \in SO_3} \cos^{-1} \langle a, Rb \rangle$$

- classify test data to class with smallest distance to corresponding mean
- accuracy of 45%
- potential problems:
  - used only 2D data at specific slide → maybe slide differs not too much between the classes
  - small data set