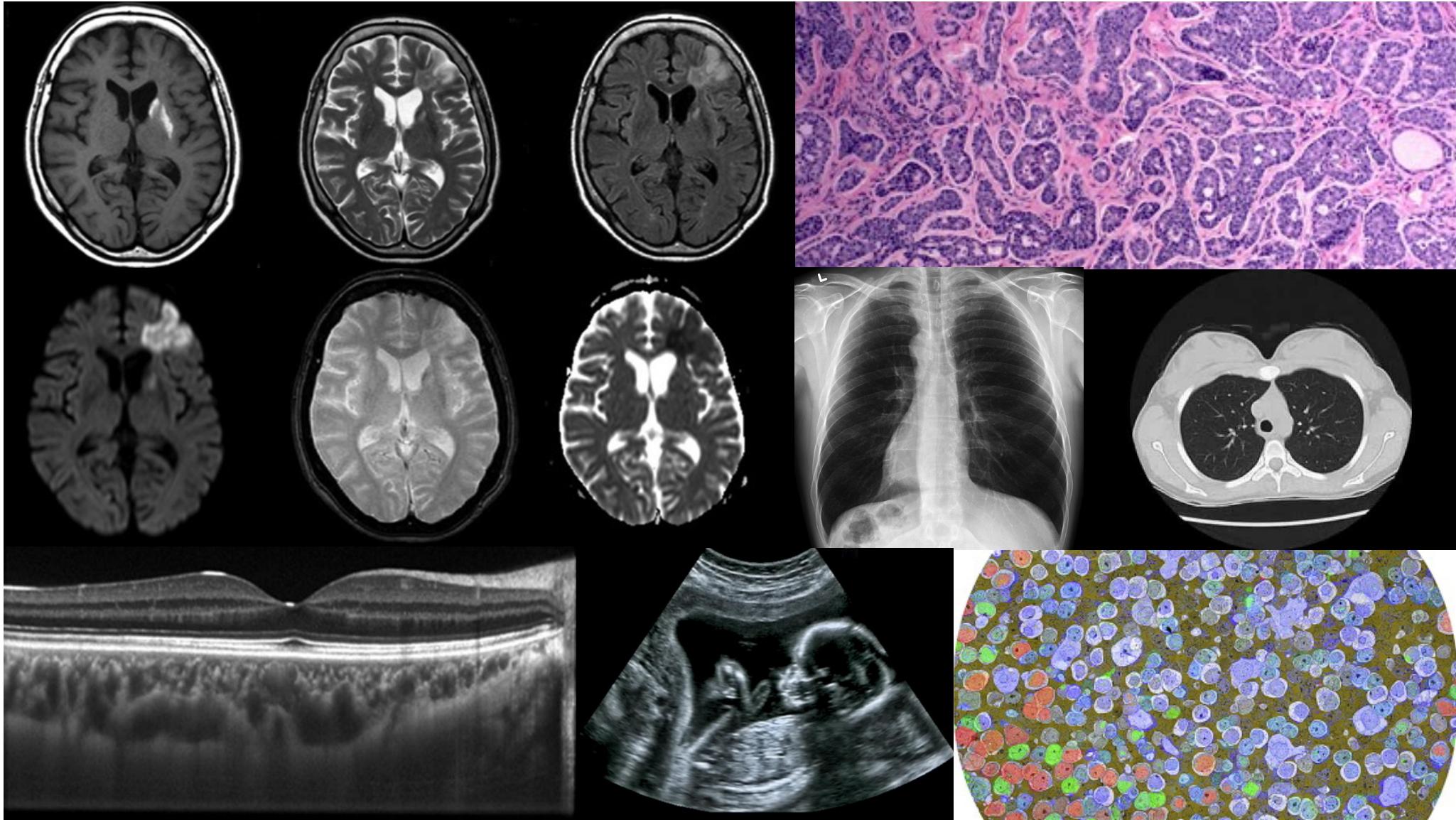


# Machine Learning for Medical Image Analysis

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Adrian V. Dalca

MIT CSAIL and  
Massachusetts General Hospital, Harvard Medical School



# Outline

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- Overview of Medical Imaging
  - Utility and properties
- Example: Segmentation
  - *Classical* and deep learning approaches
- Example: Registration (alignment):
  - Optimization and learning approaches
- Example: Imaging genetics
- Takeaways

# Takeaway Goals

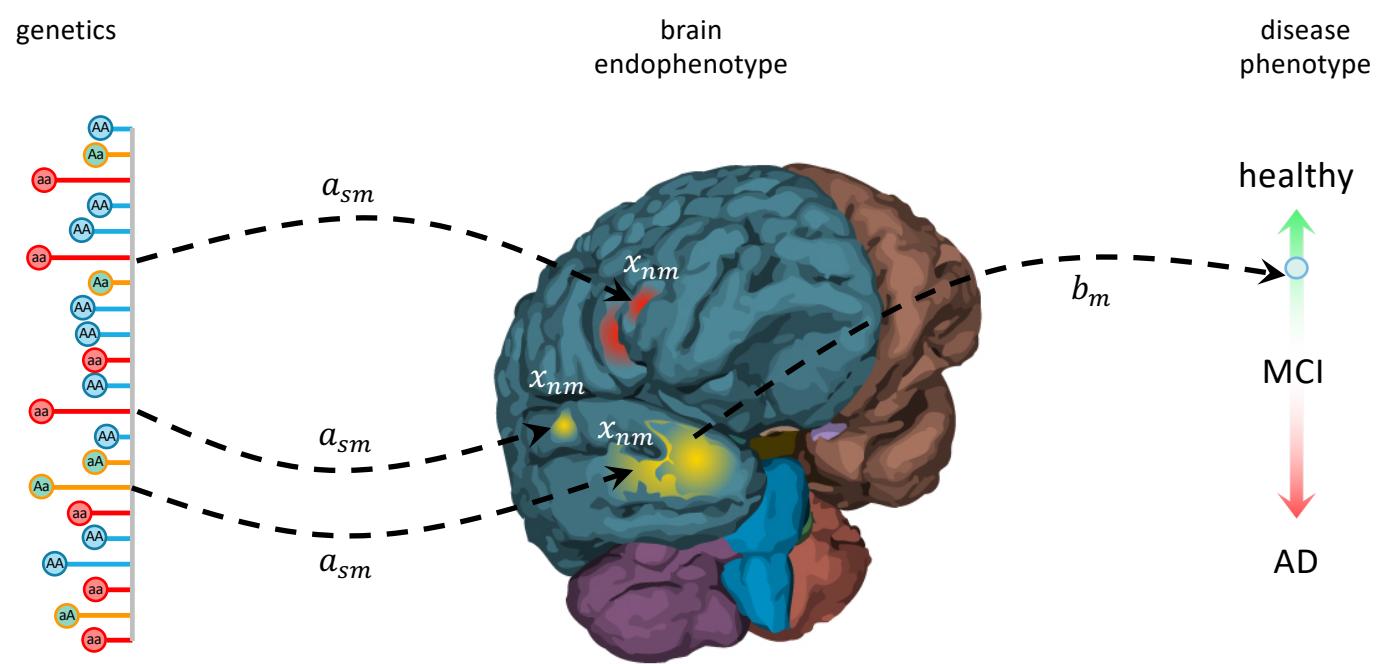
---

- Problems
  - Help the clinicians or scientists (don't replace them)
- Tools and approaches
  - Probabilities, convolutions, and anatomical models
  - Clinical interpretation
- Challenges
  - The systems don't really work (yet)
- Opportunity
  - Impact healthcare (and research)!

# Medical Imaging

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- Crucial tool in clinical practice
  - Diagnostic (and incidental findings)
  - Planning treatment
  - Guide small and large interventions
  - ...



# Medical Imaging

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- Crucial tool in clinical practice
  - Diagnostic (and incidental findings)
  - Planning treatment
  - Guide small and large interventions
  - ...
- Research
  - Clinical studies
  - Scientific studies

# Medical Image Analysis (or: how can we help?)

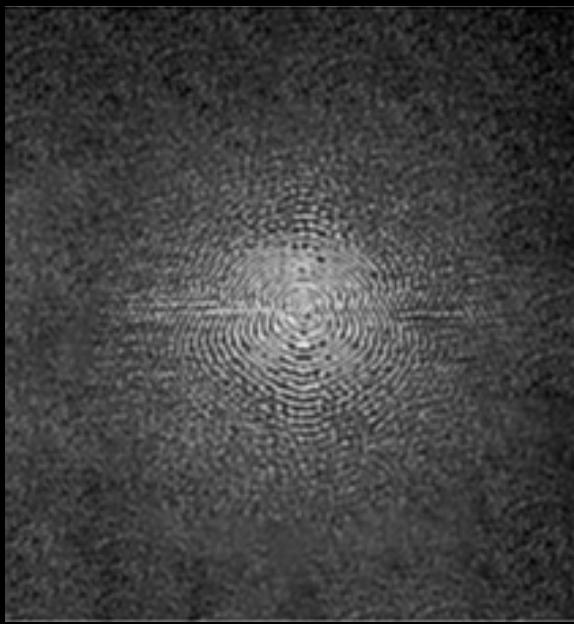
---

- Diagnosis algorithms - require large datasets
- Visualization - learn what to show, widely overlooked?
- **Segmentation** - outline, measure anatomy and pathology
- **Registration** - alignment for treatment planning, population analysis
- Acquisition - faster, better
- Abnormality detection - pathology
- Shape modelling
- Joint inference with other clinical data
- ...

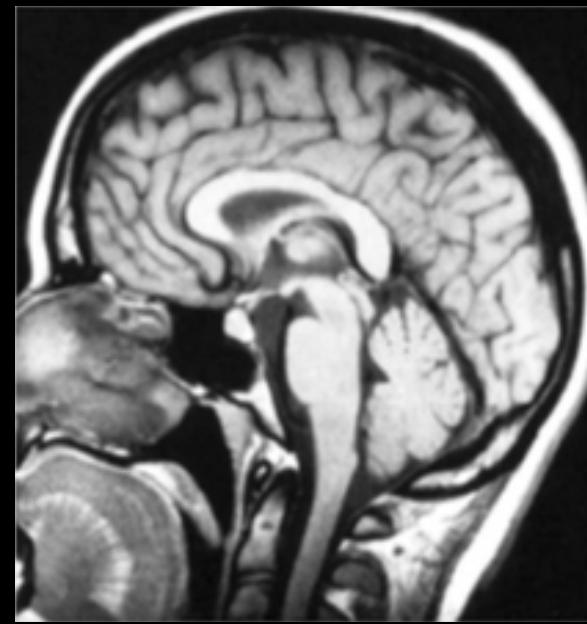
# Properties of Medical Images

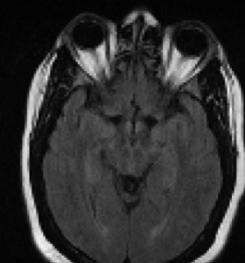
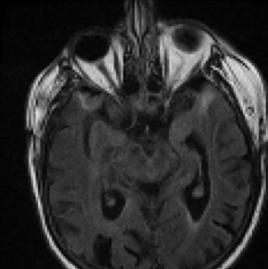
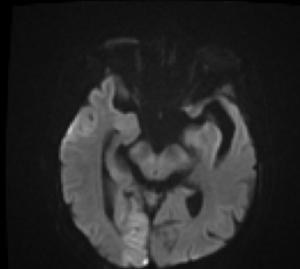
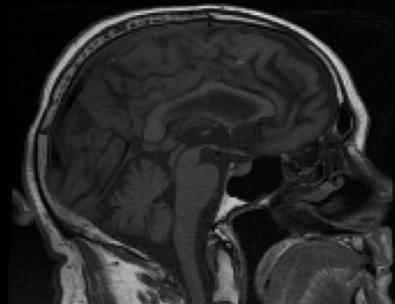
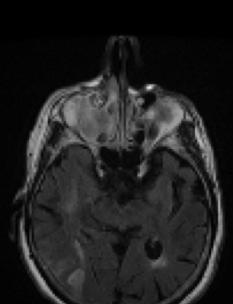
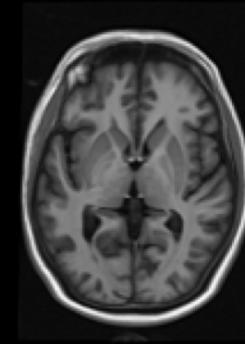
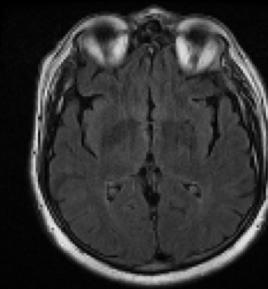
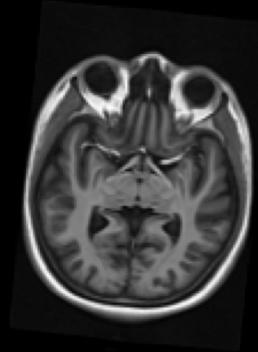
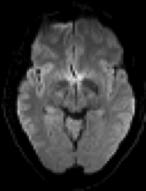
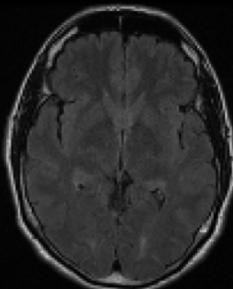
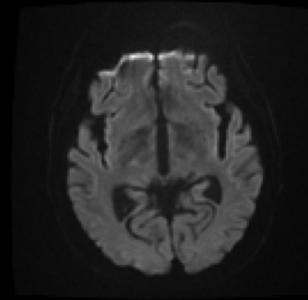
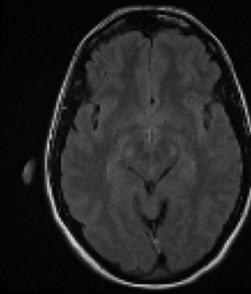
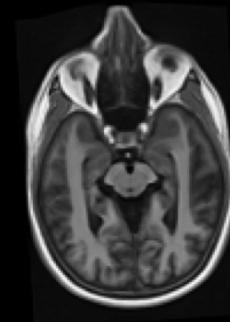
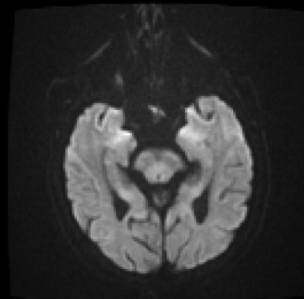
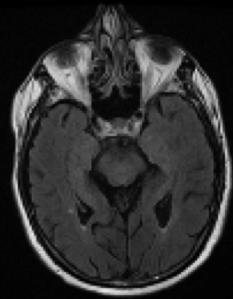
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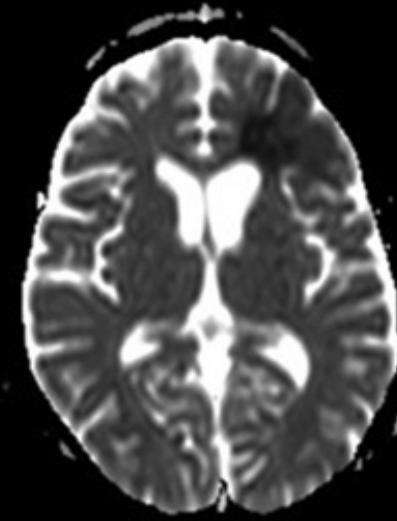
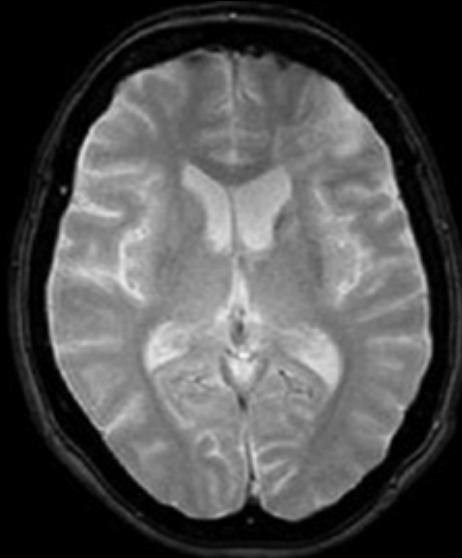
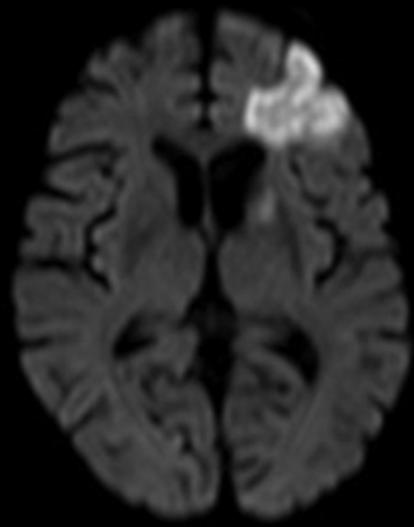
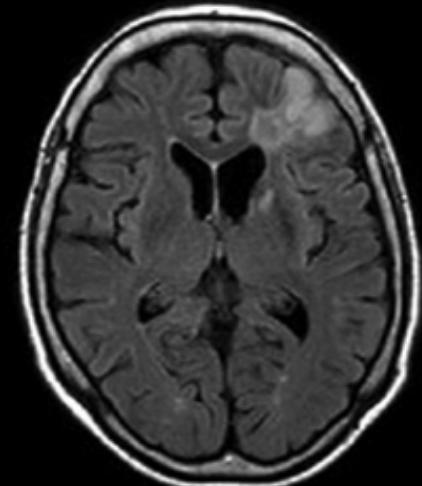
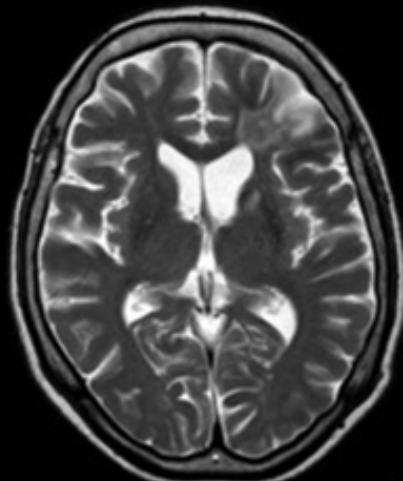
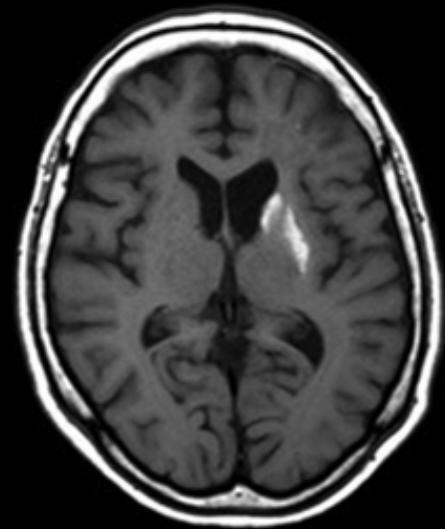
- Varies dramatically by image type

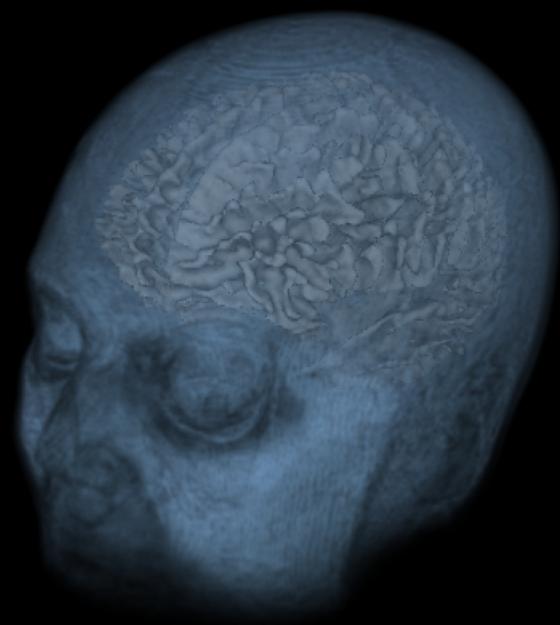


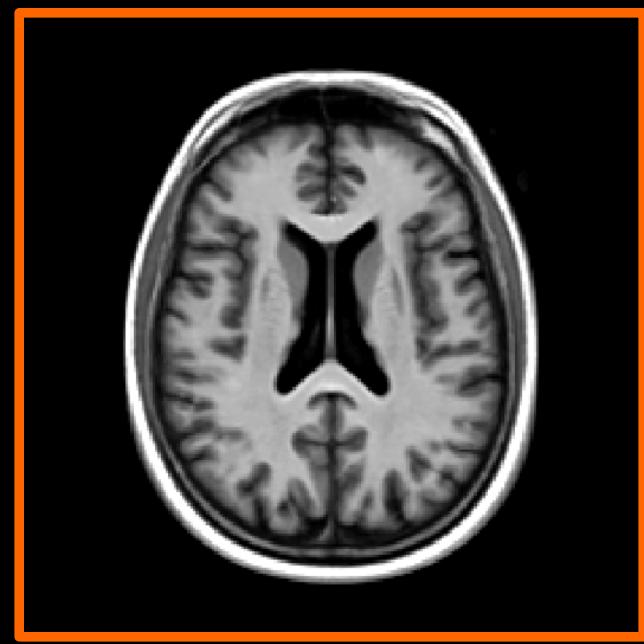
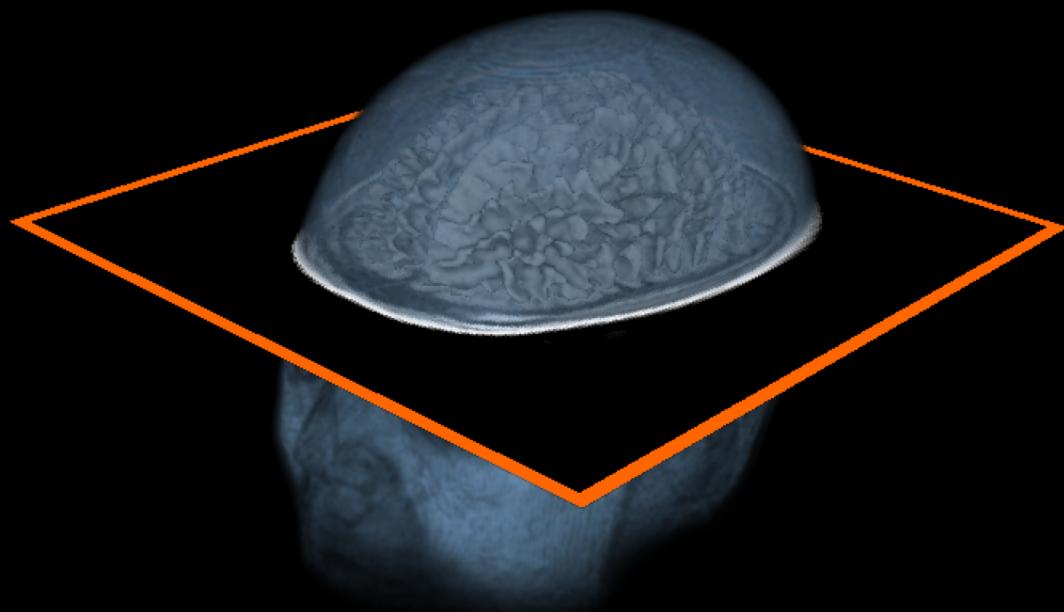
Fourier  
Transform



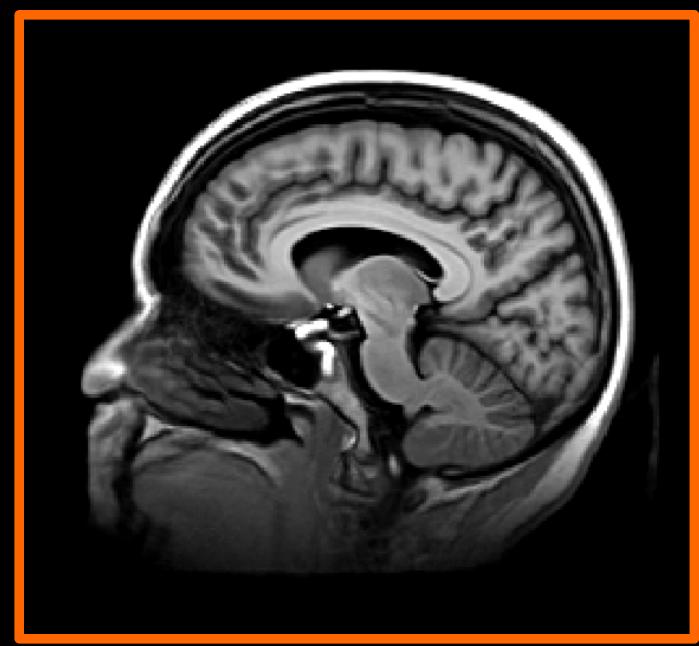
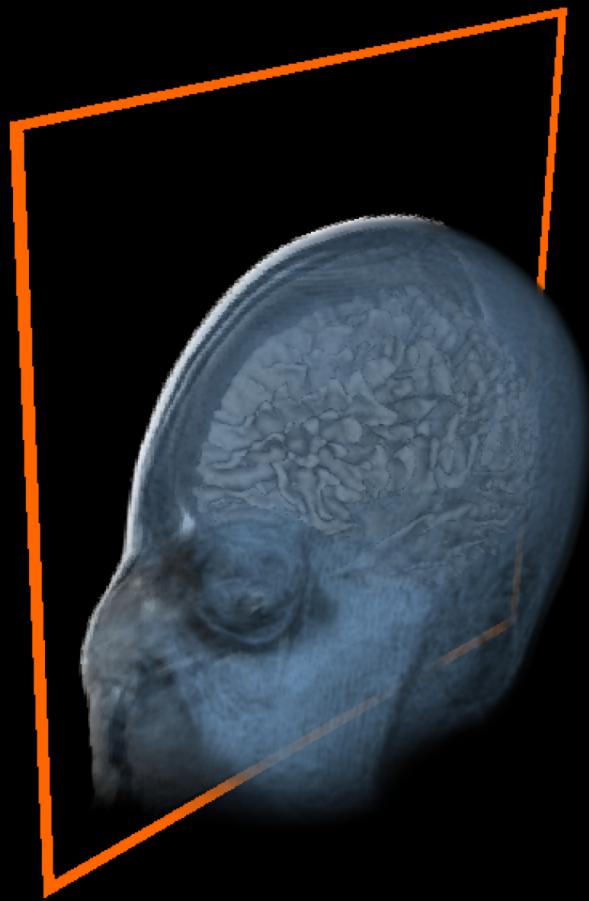




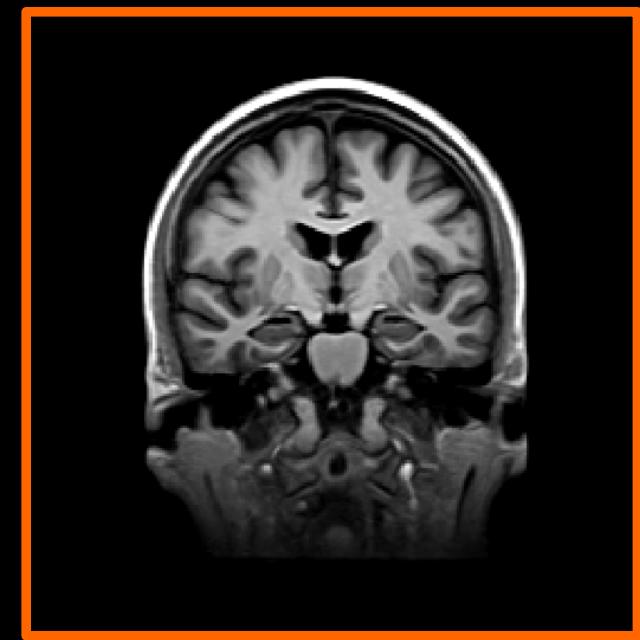




axial



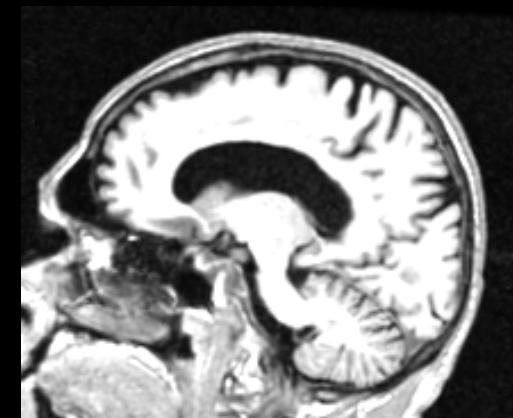
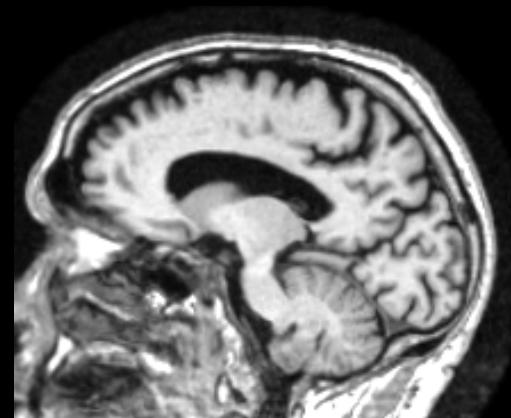
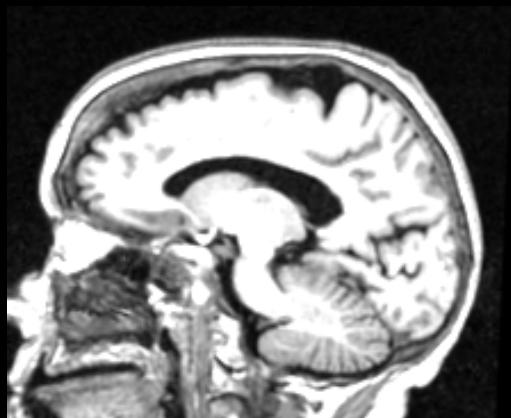
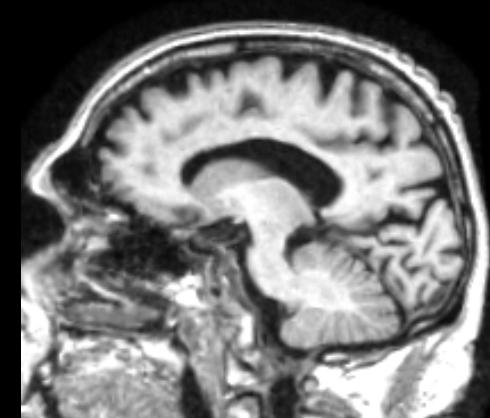
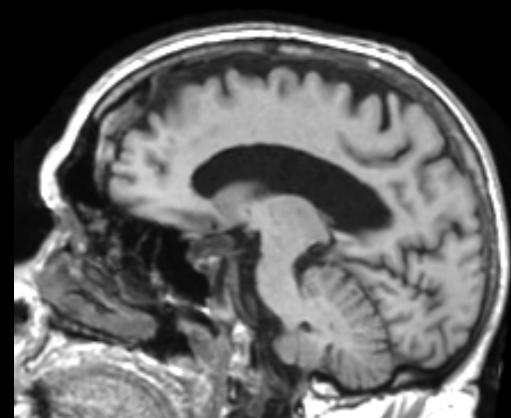
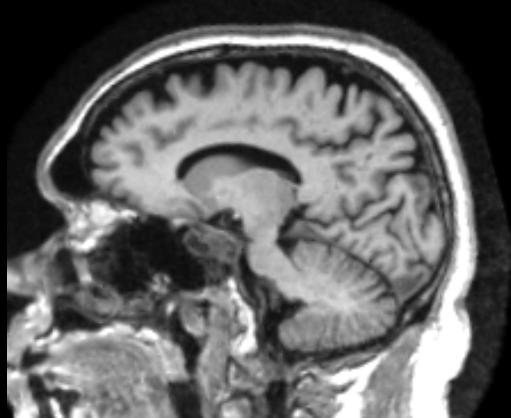
sagittal



coronal

# Variability and similarity

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# Properties

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- Vary dramatically by image type
- MR Image quality:
  - Different noise patterns, patient motion, disease, many modalities
- Commonality of anatomy
- Pathology
  - can be big and obvious (e.g. tumor)...
  - ... or very small and subtle (e.g. neurodegeneration)
- A lot of 3+ dimensions
  - So '**voxel**' (**volume element**) instead of 'pixel' (picture element)

# Questions?

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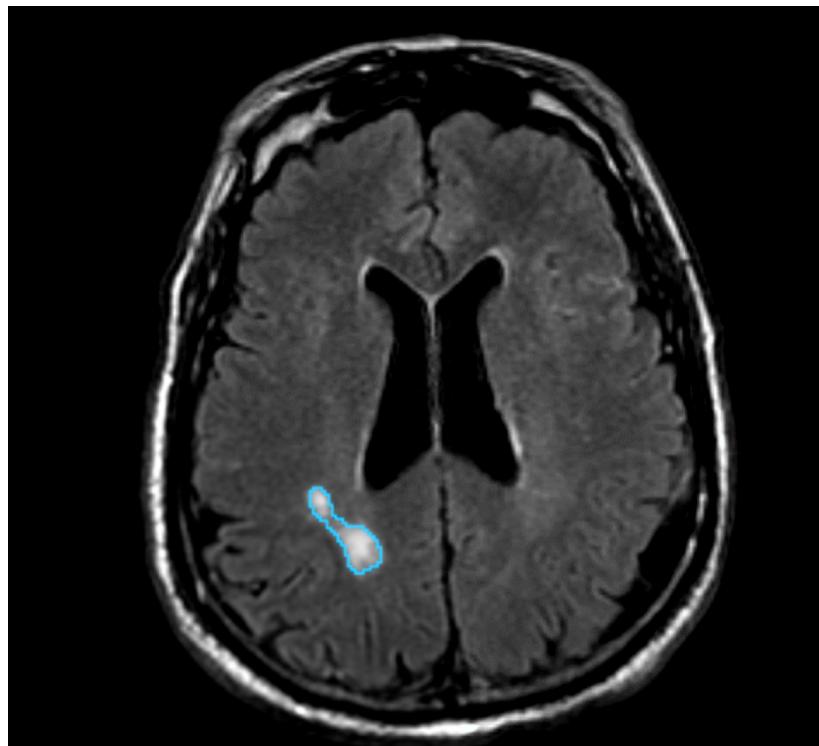
# Outline

---

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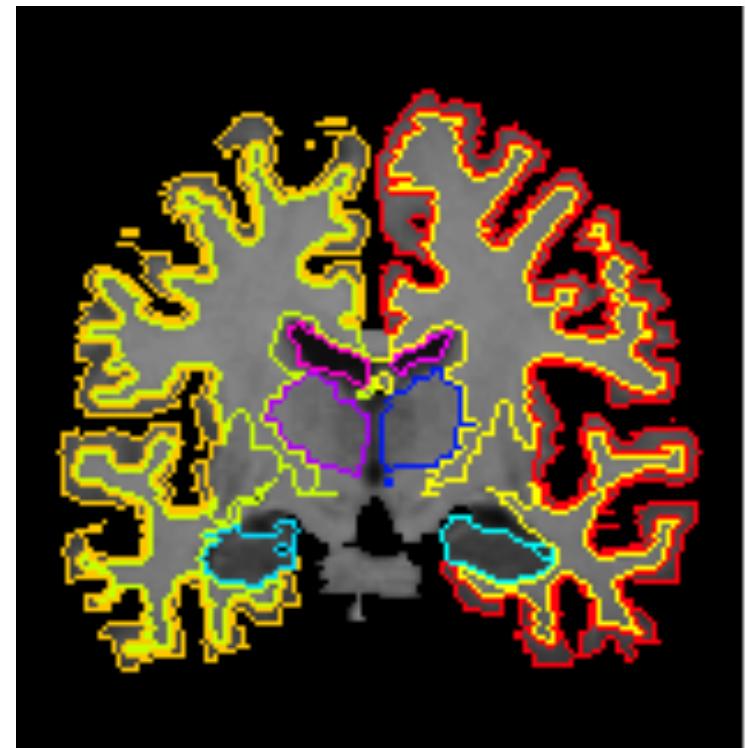
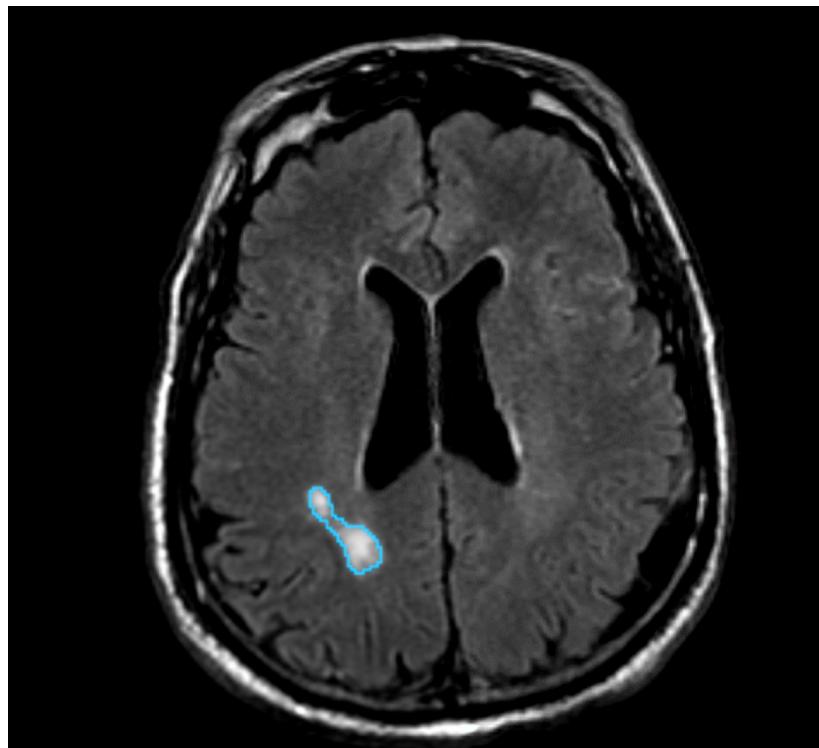
# Image Segmentation

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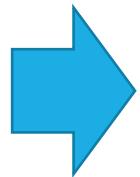
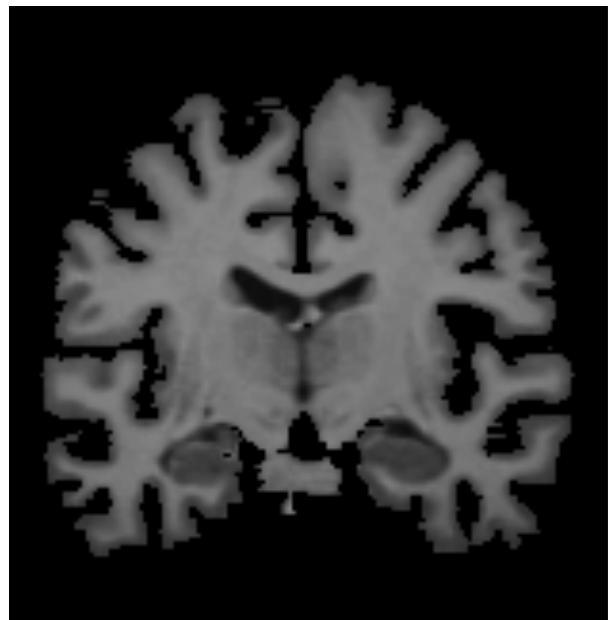
# Image Segmentation

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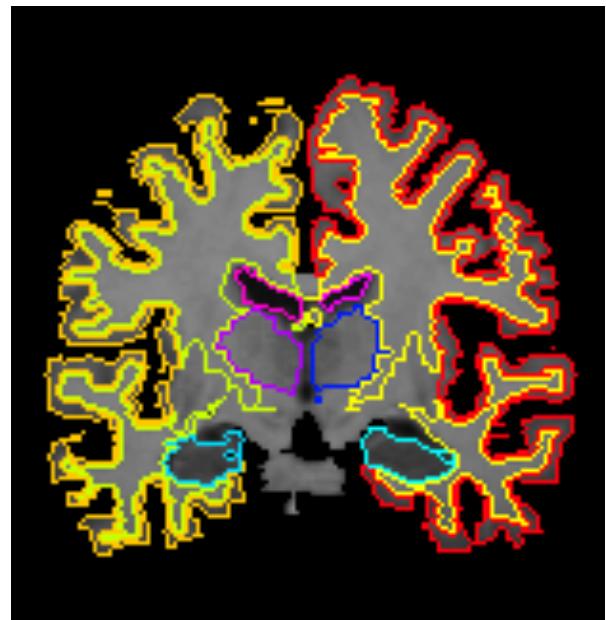


# Supervised segmentation

---



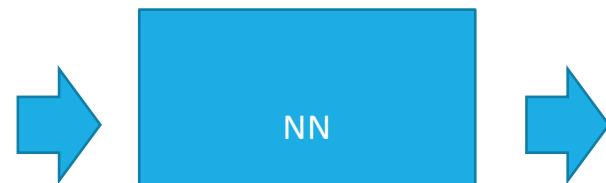
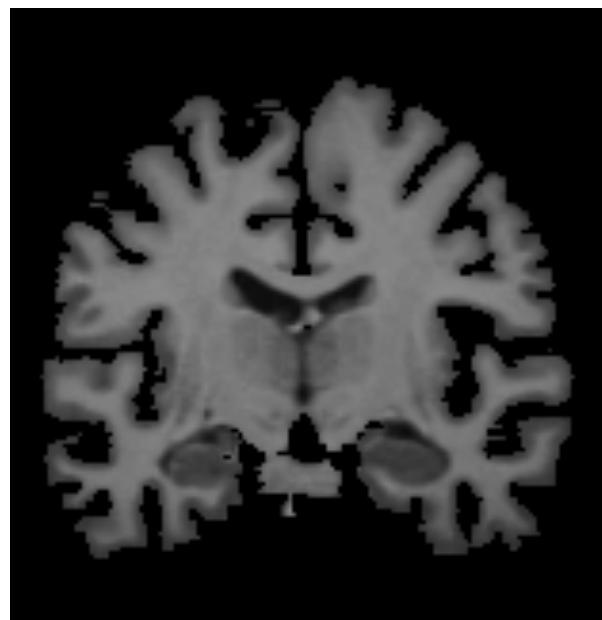
$$seg = f_{\phi}(image)$$



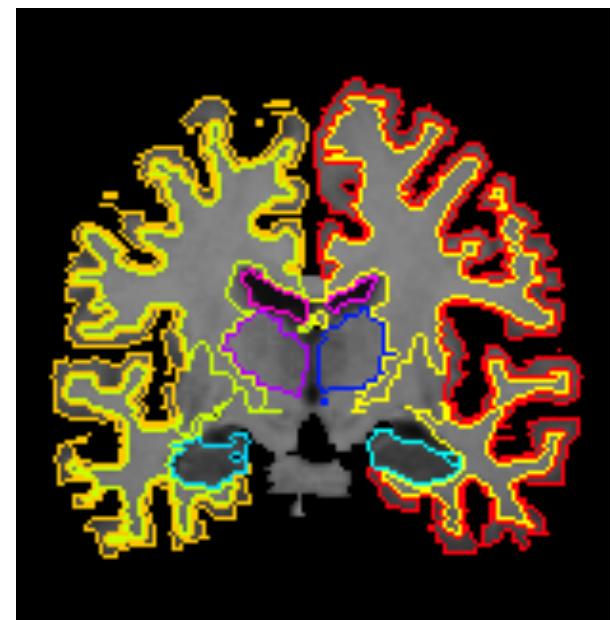
# Supervised segmentation

---

Large example dataset: solved problem by DL?



$$seg = f_{\phi}(image)$$



# What kind of NN?

---

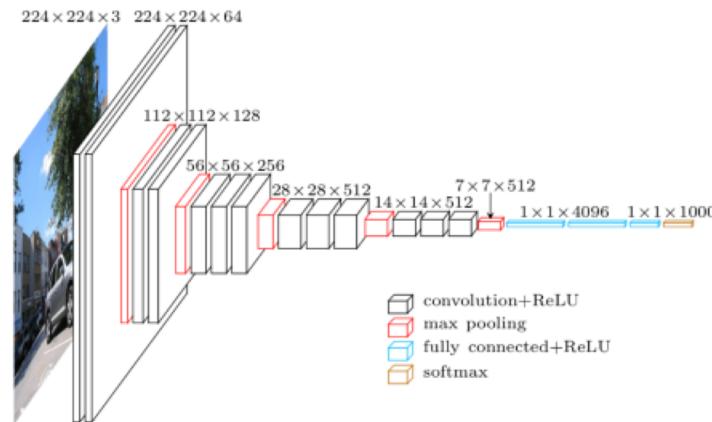
# VGG, etc?

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Use existing multi-label networks

Architecture: convolutions, max-pools, fully connected, etc.

But need to output 8 million voxels! – hours!



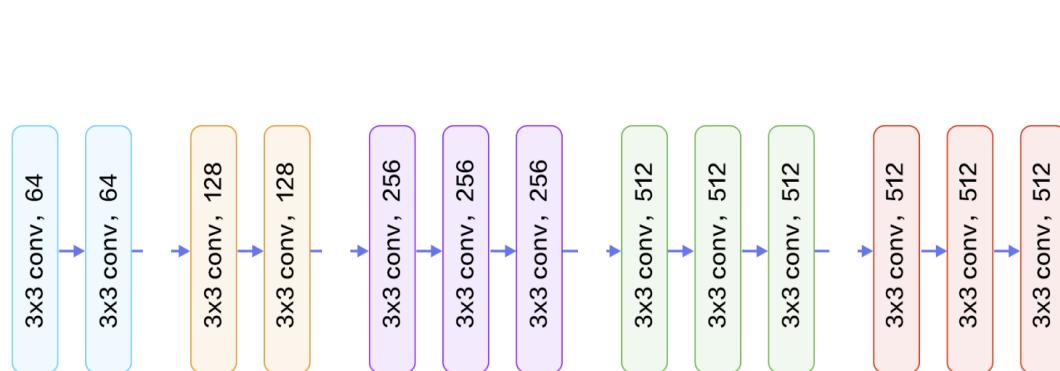
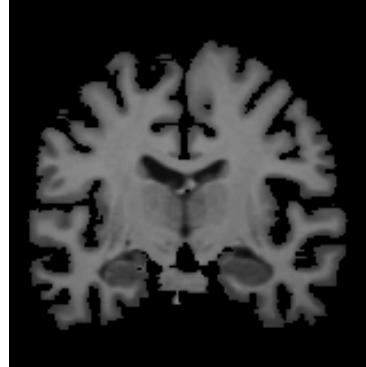
anatomical  
label  
(one-hot encoding)

# Fully convolutional?

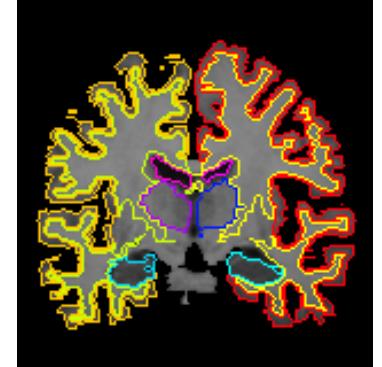
Input-output both high dimensional, no max-pooling (make 3D)

10-layer network: don't have enough context to predict anatomy

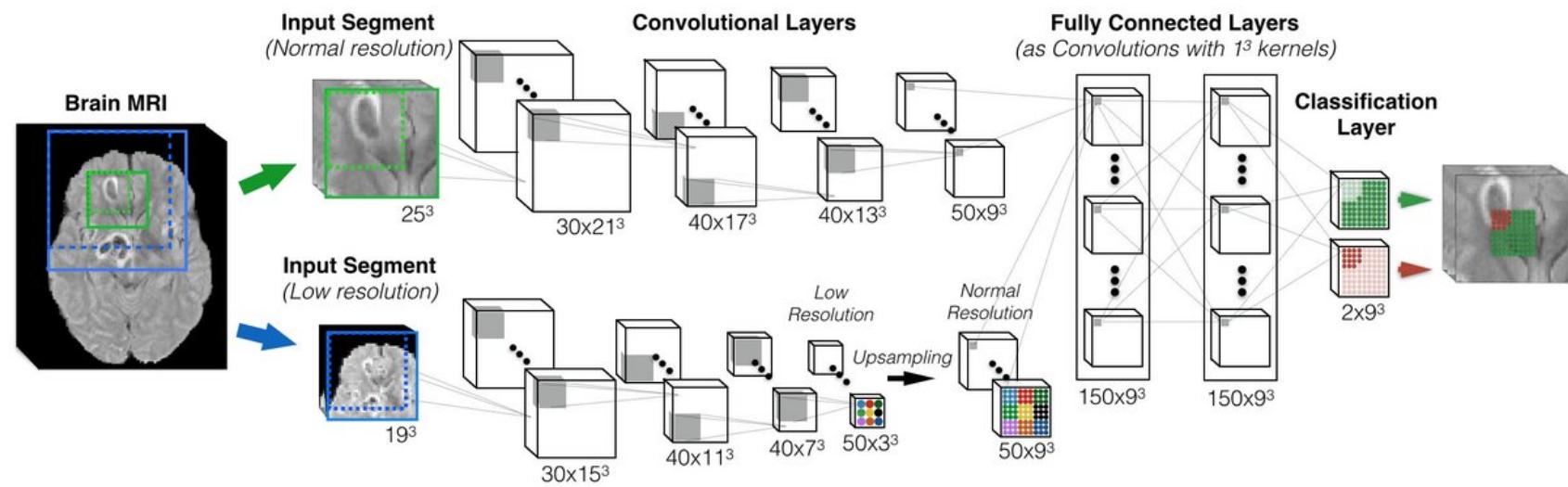
Deep networks (100 layers) – require too many parameters



# channels = # labels

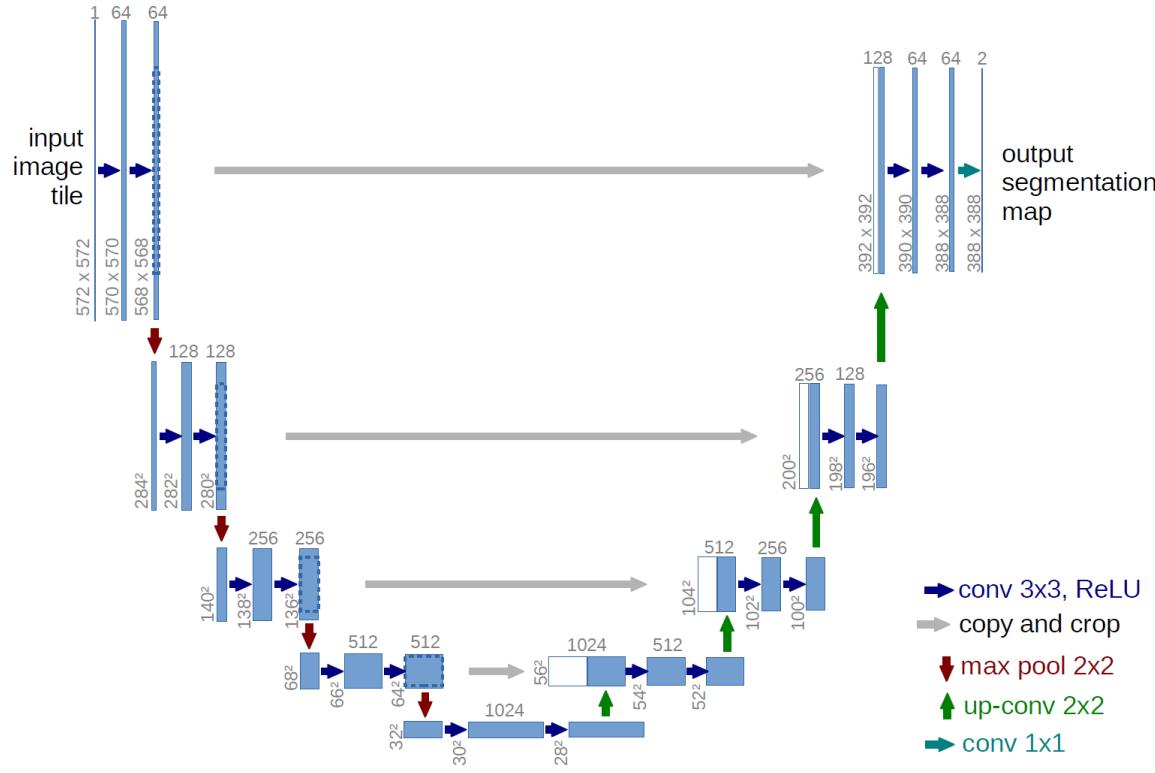


# Multi-scale inputs



Kamnitsas, et al., Media 2016

# U-Net



Ronneberger et al, 2015

# What kind of CNN?

---

## Network architecture

- Predict each voxel (e.g. 3D **VGG**)? too slow, cumbersome
- Fully Convolutional? Large memory, parameter space, not enough **field of view**
- Multiscale input?
- UNet!

# Results

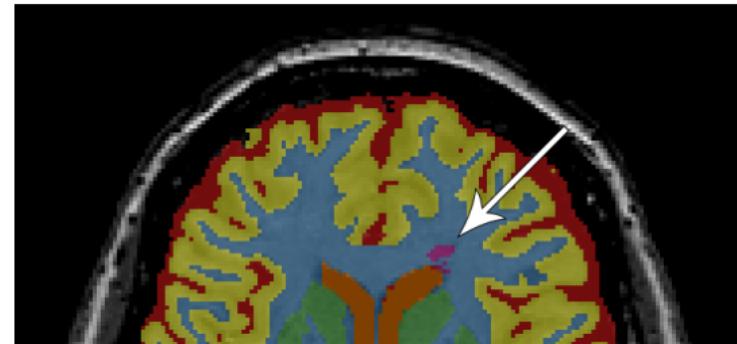
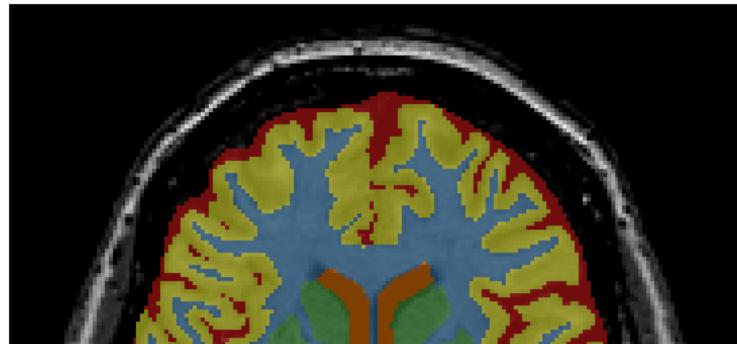
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Dice (Volume Overlap)	Dice (Volume Overlap)	Runtime
FreeSurfer (classical state of the art)	~80	~6-24 hours
Deep Methods	~85-91	~1 second-1 hour

# Problems

---

- Often don't actually have these **segmented** data
  - Long time to segment for experts!
  - Too many modalities
  - Too much variation (especially pathologies)
- Our metrics
  - Easy to compute, differentiate
  - Often not **anatomically** meaningful

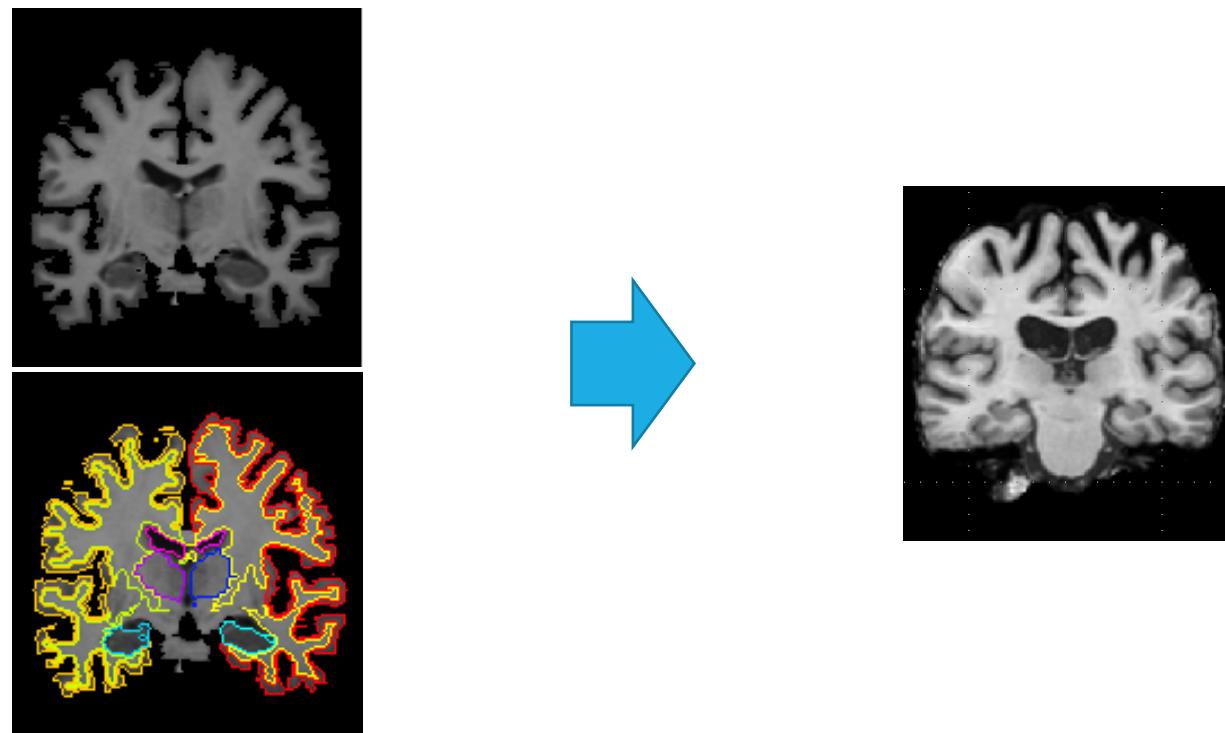


P. Moeskops et al, DLMIA, 2017

# Segmentation in a more realistic setting

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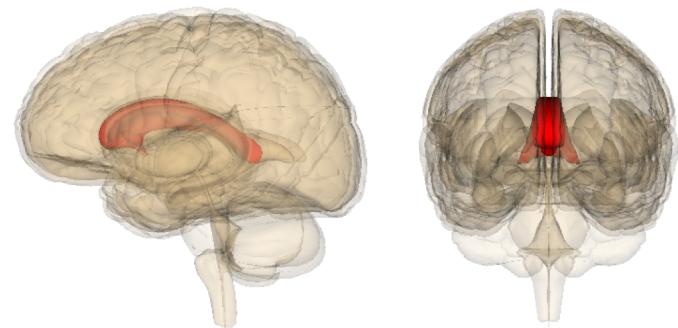
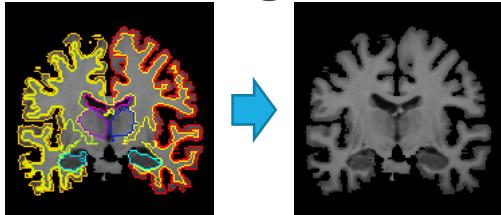
Few (one) segmented example



# Probabilistic (Generative) Model

---

- Define segmentation  $\rightarrow$  image model  $p(I|S) * P(S)$

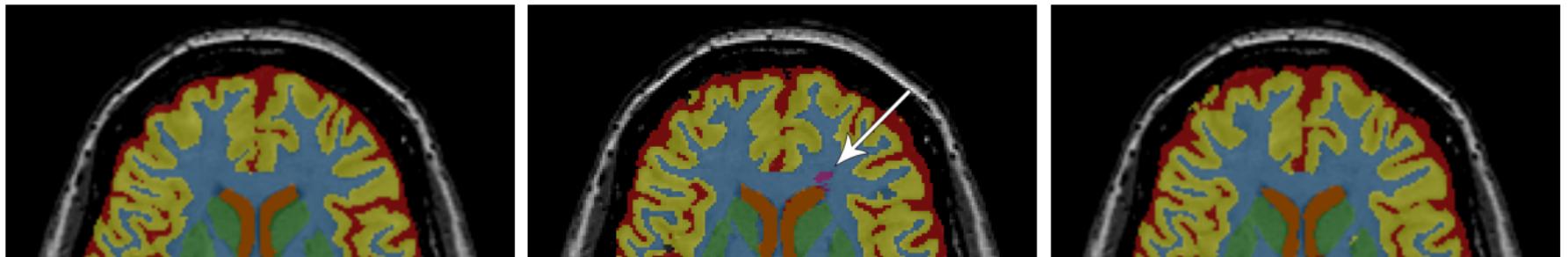


- Enables knowledge (priors) into segmentation model  $p(S)$ 
  - $p(S)$  defined based on likely *\*shapes\** of each label
  - $P(I|S)$  is the intensity (distribution) for each label
  - **Inference:**  $p(S|I)$  at each voxel: **label** matches the **intensity** such that **shapes** make sense.

# Probabilistic (Generative) Model

---

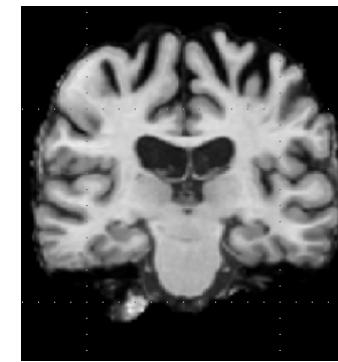
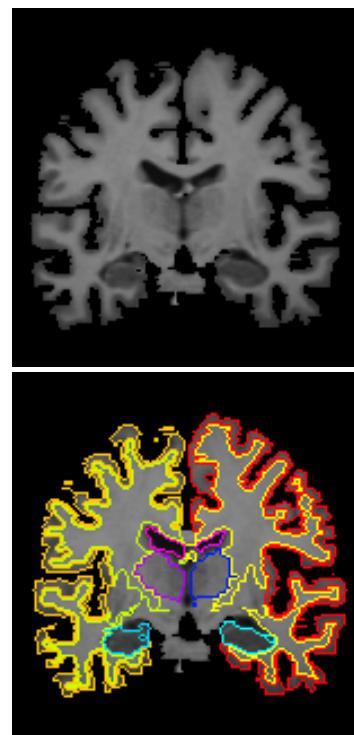
- Combine with deep learning predictions:  
 $p(S)$  can be anatomically specified  
or learned from another distribution
- Attach prior to network, or modelling through VAEs, etc...



# Brains are similar!

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- Can similarity of brains help?



# Questions?

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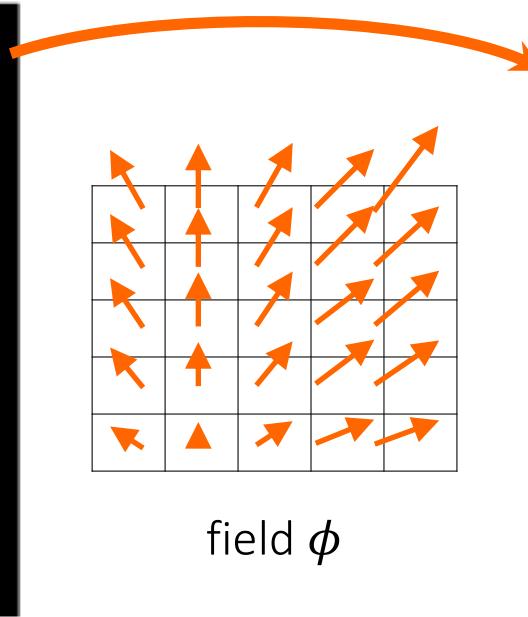
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# Image Registration

---



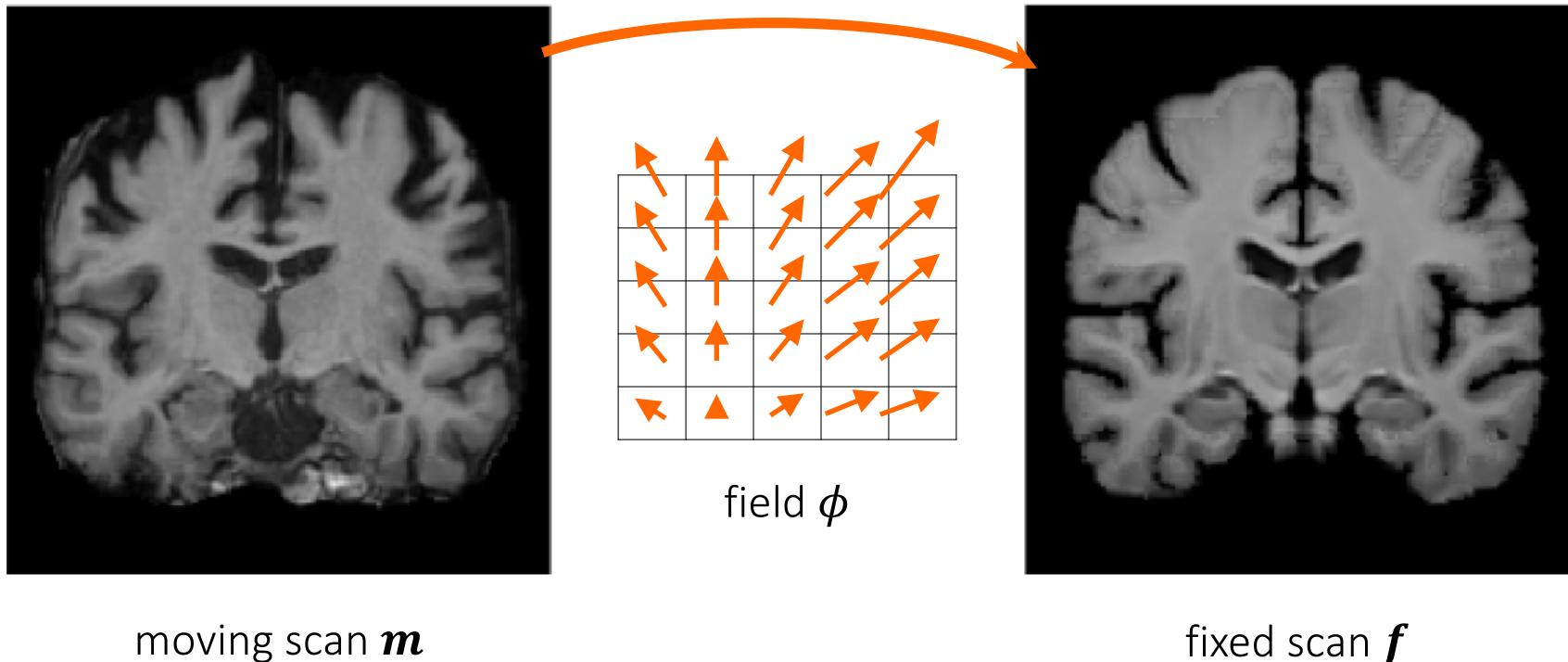
moving scan  $m$



fixed scan  $f$

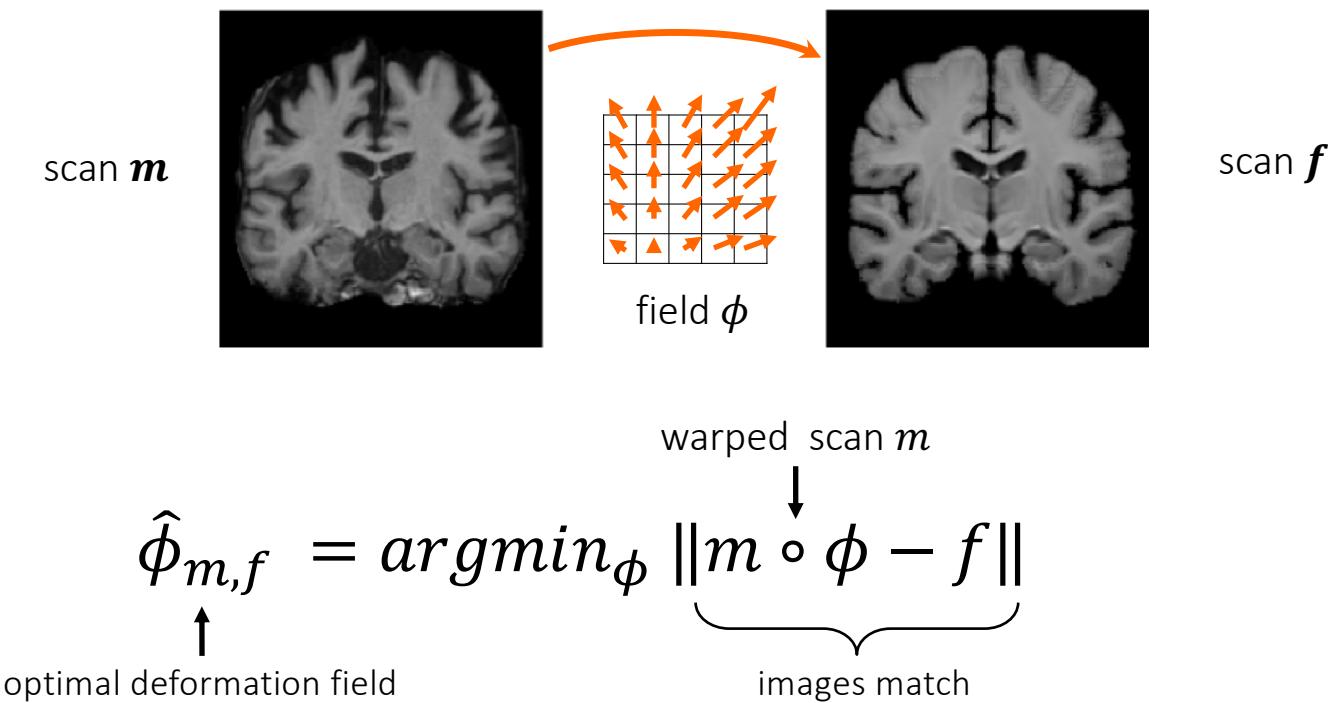
# Image Registration

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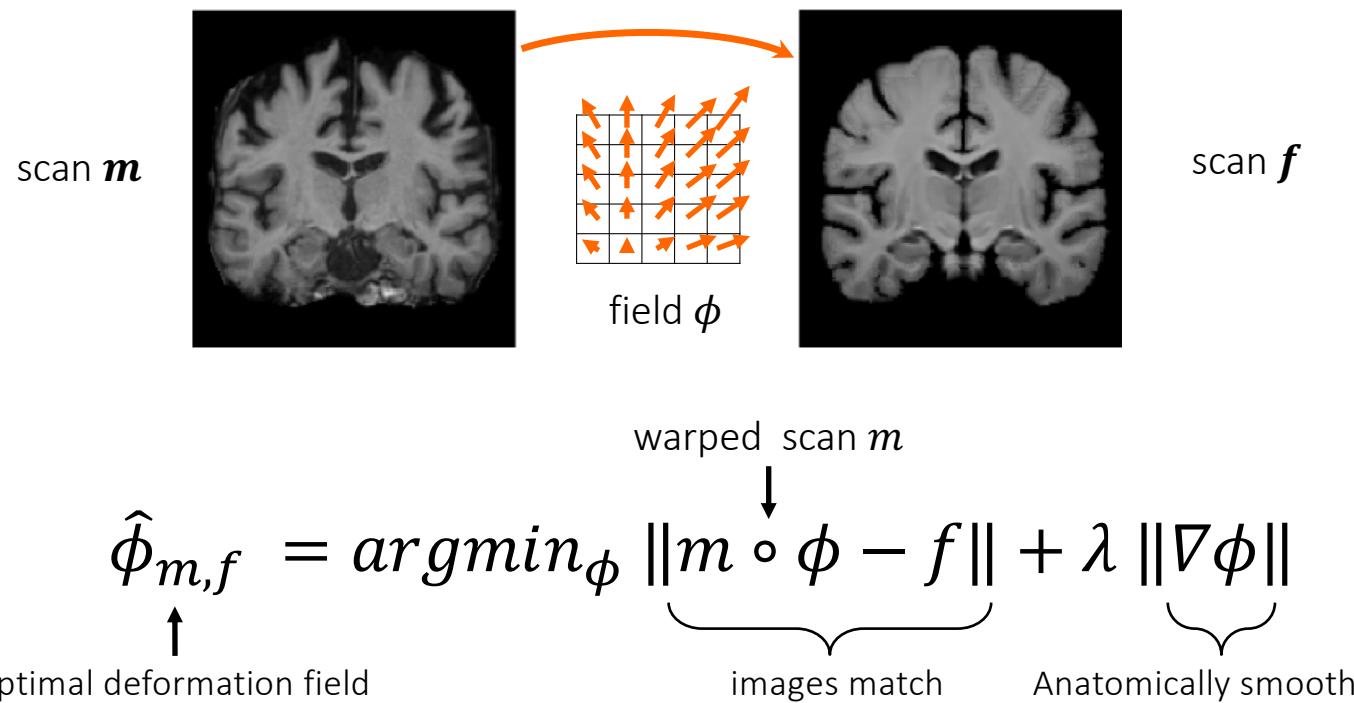


# Traditional approach

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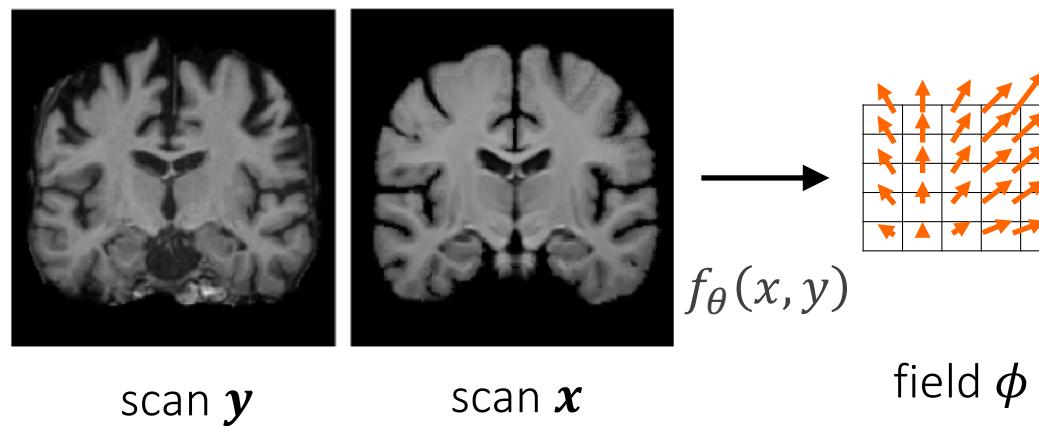
# Traditional approach



Pairwise optimization: slow (hours per image on CPU)

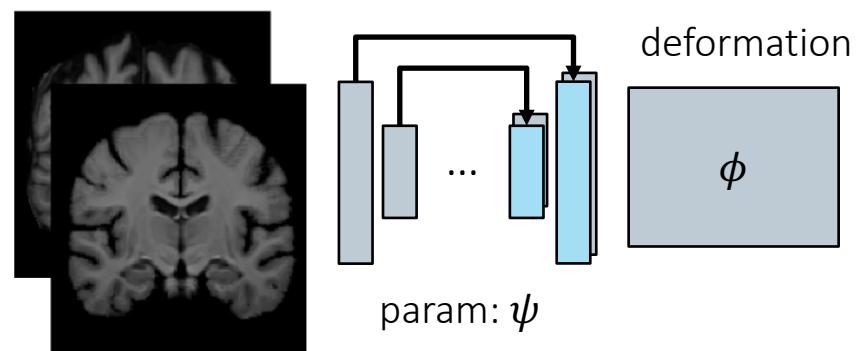
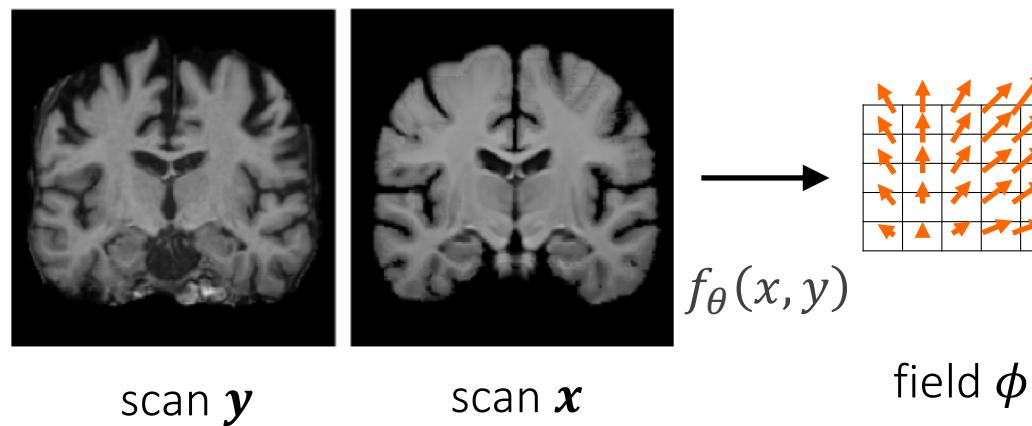
# How can machine learning help?

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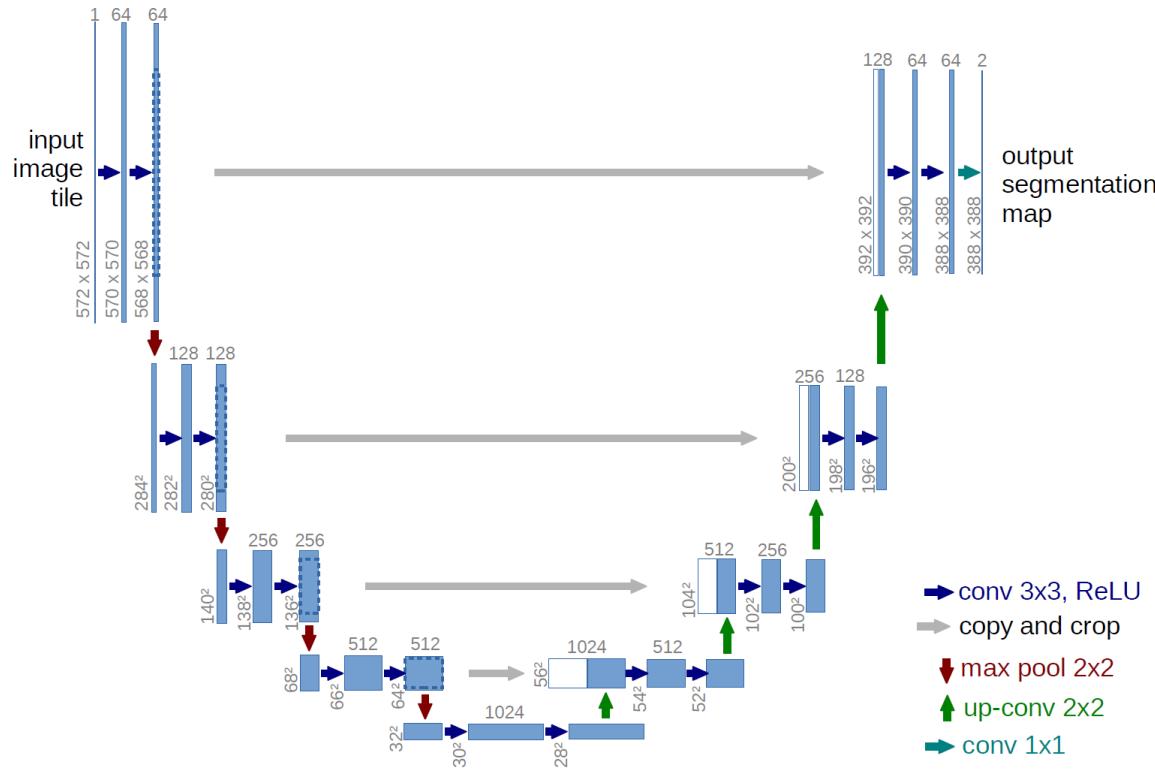


# Supervised Learning

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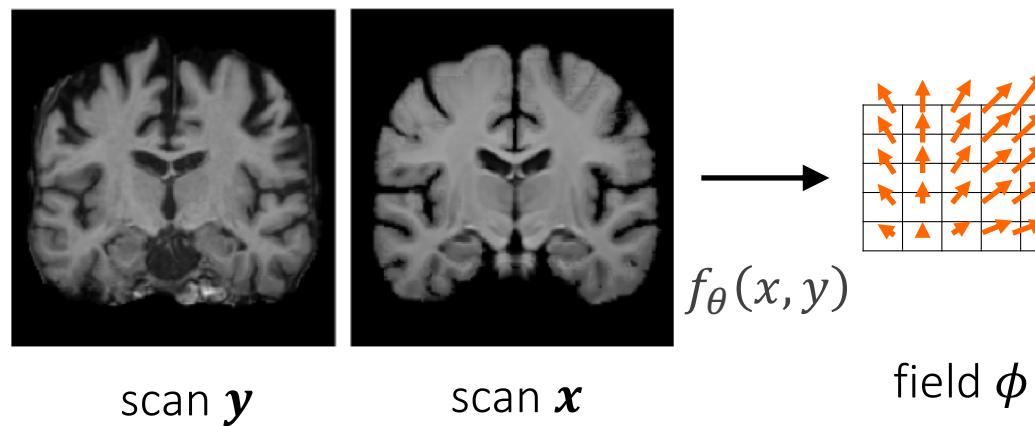
# What kind of architecture?



Ronneberger et al, 2015

# Supervised Learning

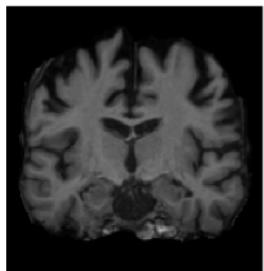
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fast for new image pair!  
need **ground truth** registration  $\phi$

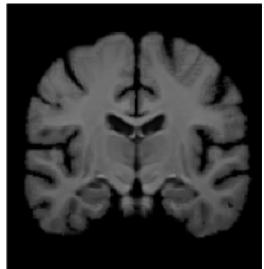
# Unsupervised Learning: VoxelMorph

Moving 3D Image ( $m$ )



[voxelmorph.mit.edu](http://voxelmorph.mit.edu)

Fixed 3D Image ( $f$ )

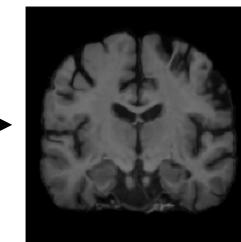


deformation

$\phi$

spatial transform

Moved ( $m \circ \phi$ )



param:  $\psi$

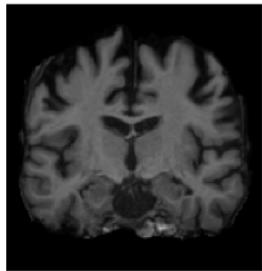
$$\min_{\psi} \|m \circ \phi - f\| + \lambda \|\nabla \phi\|$$

images match

Anatomically smooth

# Unsupervised Learning: VoxelMorph

Moving 3D Image ( $m$ )



[voxelmorph.mit.edu](http://voxelmorph.mit.edu)

Fixed 3D Image ( $f$ )

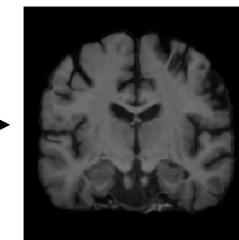


deformation



spatial transform

Moved ( $m \circ \phi$ )



param:  $\psi$

Loss ( $\mathcal{L}$ )

$$\min_{\psi} \|m \circ \phi - f\| + \lambda \|\nabla \phi\|$$

images match

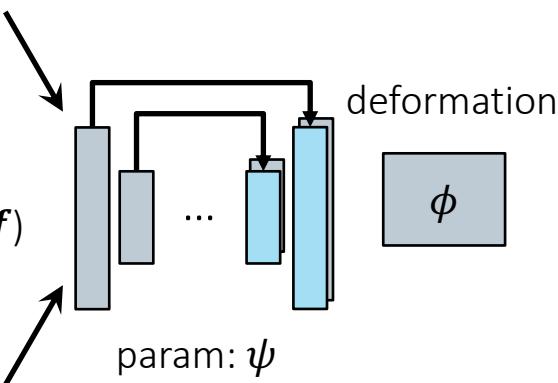
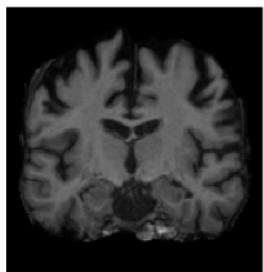
Anatomically smooth

Balakrishnan et al  
CVPR 2018, TMI 2019

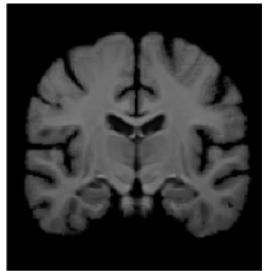
# Registering a new image pair

---

Moving 3D Image ( $\mathbf{m}$ )

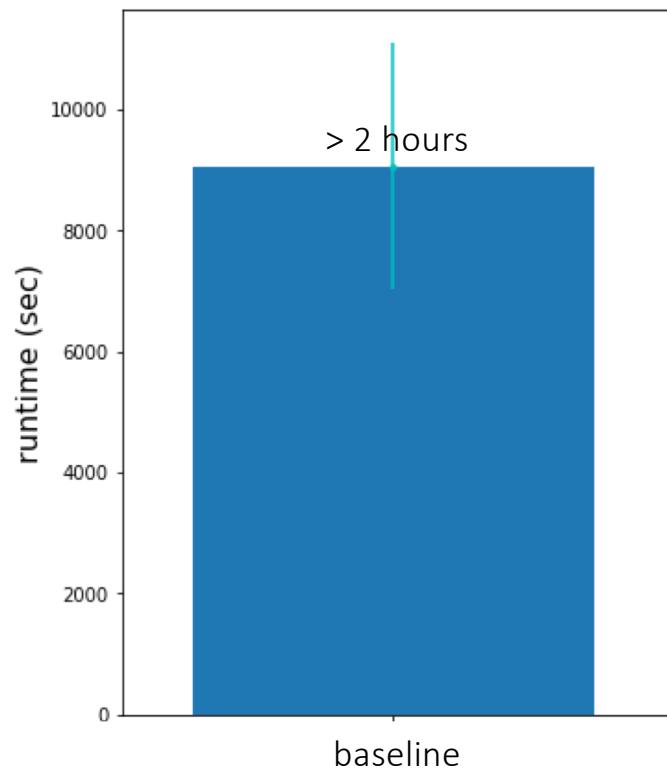


Fixed 3D Image ( $\mathbf{f}$ )



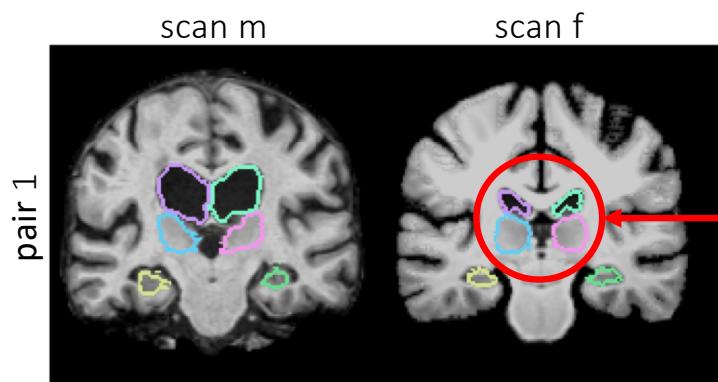
# Runtime for a new 3D image pair

---



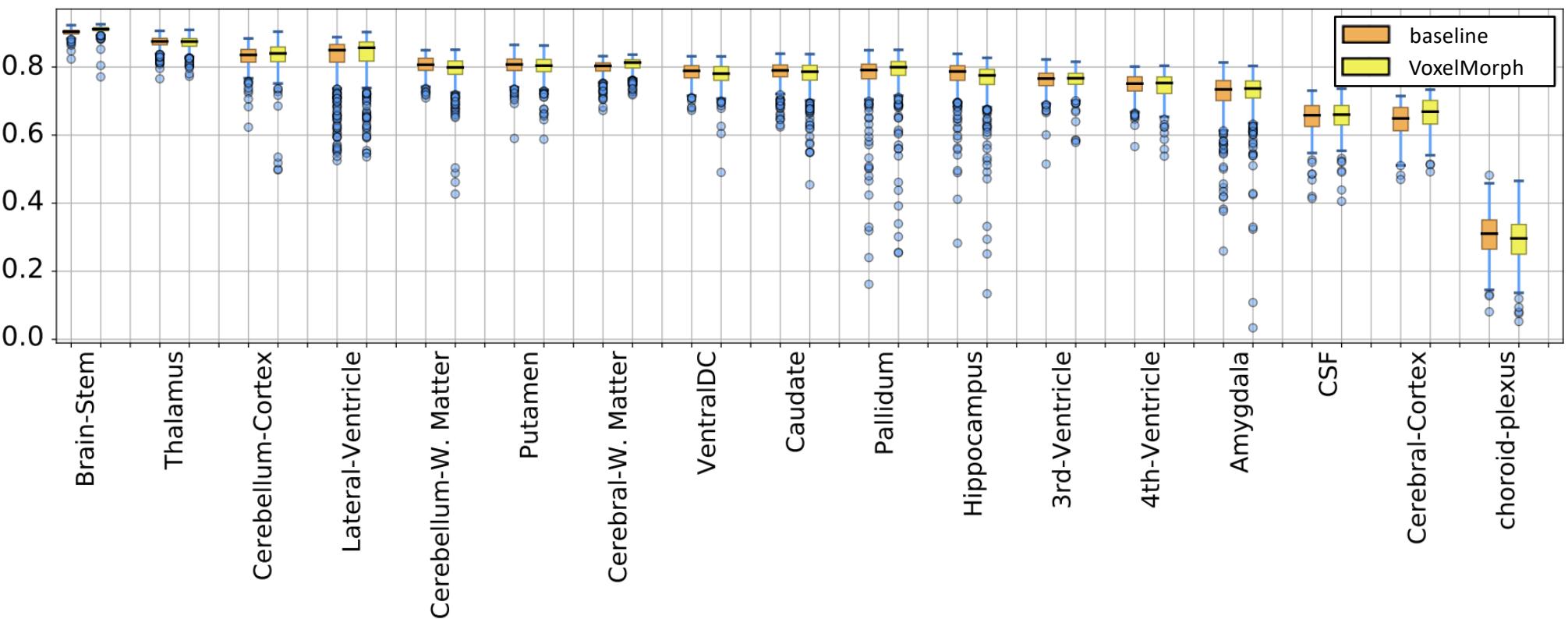
# How to evaluate?

---



\*algorithms only see images, no segmentation maps

# Accuracy via volume overlap (Dice)



# Remarks

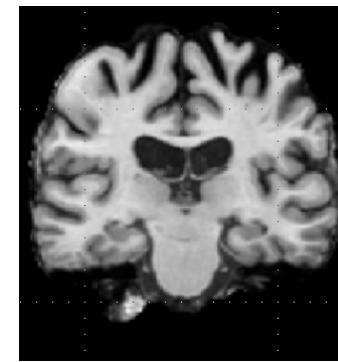
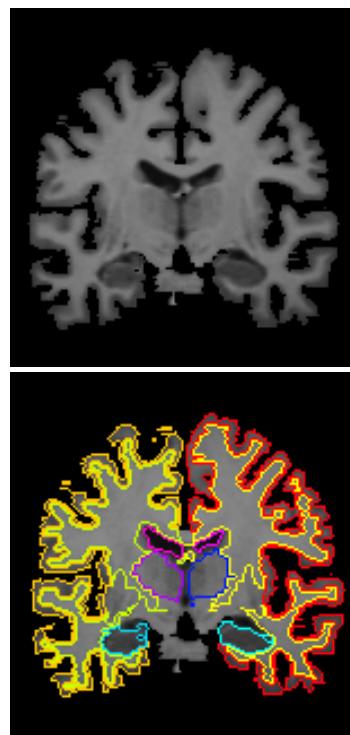
---

- We derive network probabilistically **from probabilistic model**
  - $p(m|\phi; f) * p(\phi) \rightarrow p(\phi|m; f)$
  - Variational approximation to  $p(\phi|m; f)$  leads to network
- Can impose stricter anatomical consistency (**diffeomorphisms**)
  - Provide topological guarantees
- Can use segmentations during training if we have them.

# Going back to segmentation...

---

- Can similarity of brains help?



# Questions?

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## Caveat: registration isn't perfect

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- Supervised segmentation (with 200 training images):  $85 \pm 9$
- Registration-based segmentation (with 1 training image):  $76 \pm 14$

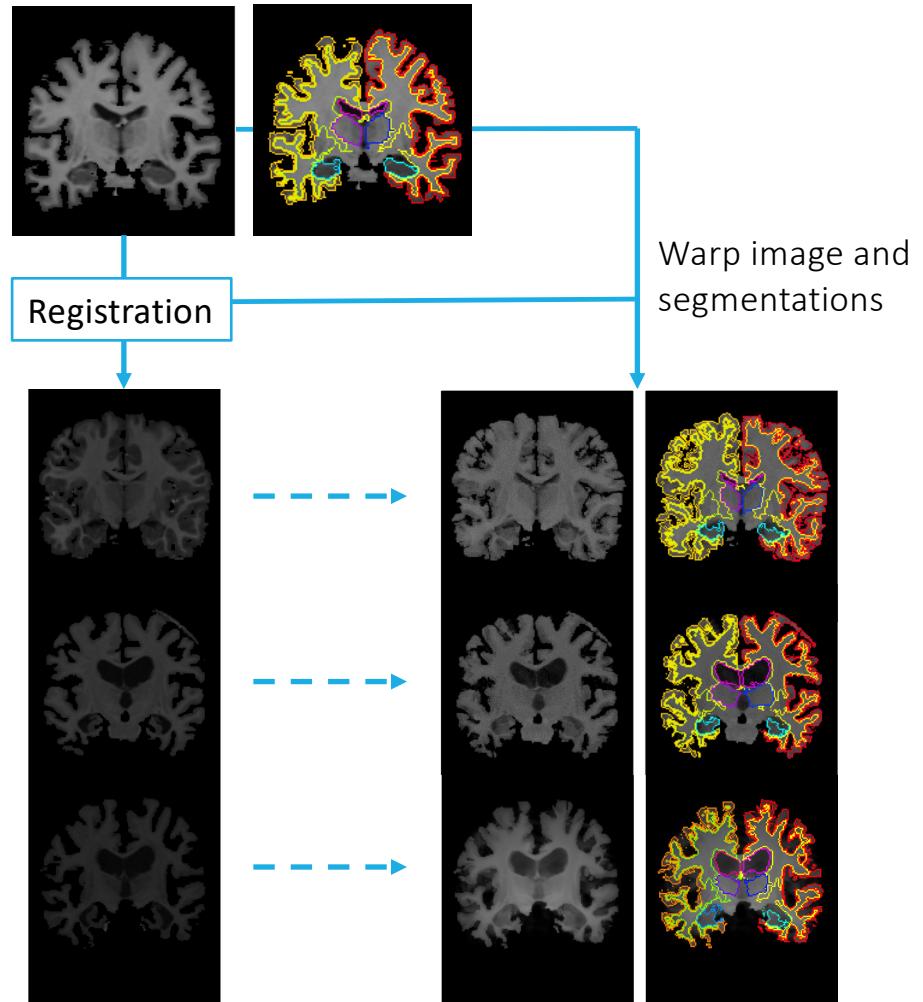
## Caveat: registration isn't perfect

---

- Supervised segmentation (with 200 training images):  $85 \pm 9$
- Registration-based segmentation (with 1 training image):  $76 \pm 14$
- Combine advantages!

# Supervised segmentation & registration

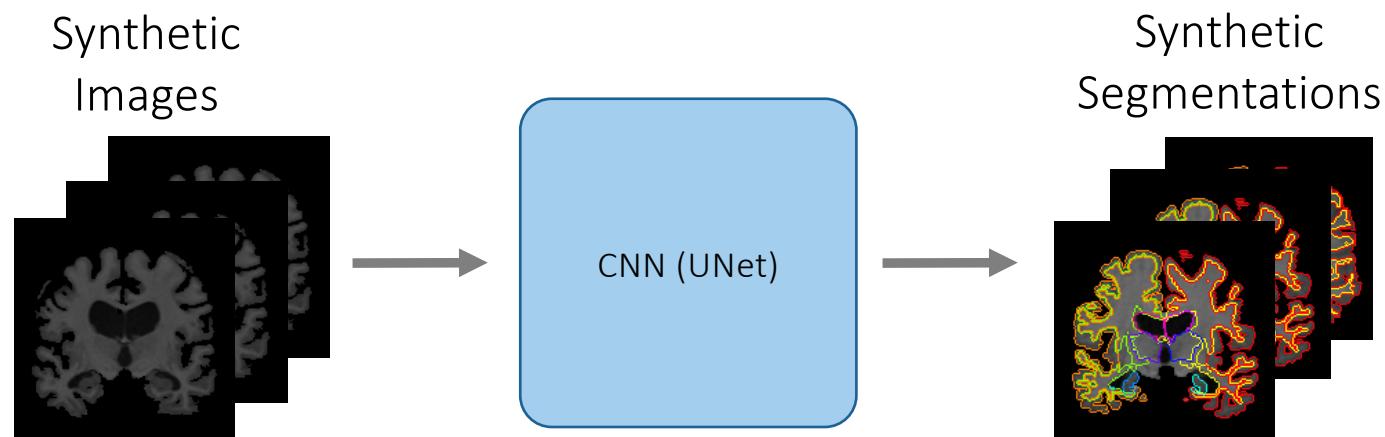
- Register training image to every image in dataset
  - distribution of transforms
- Warp labelled scan and segments to produce *supervised dataset*
  - Span anatomical distribution
  - Accurately segmented



# “Supervised” Network

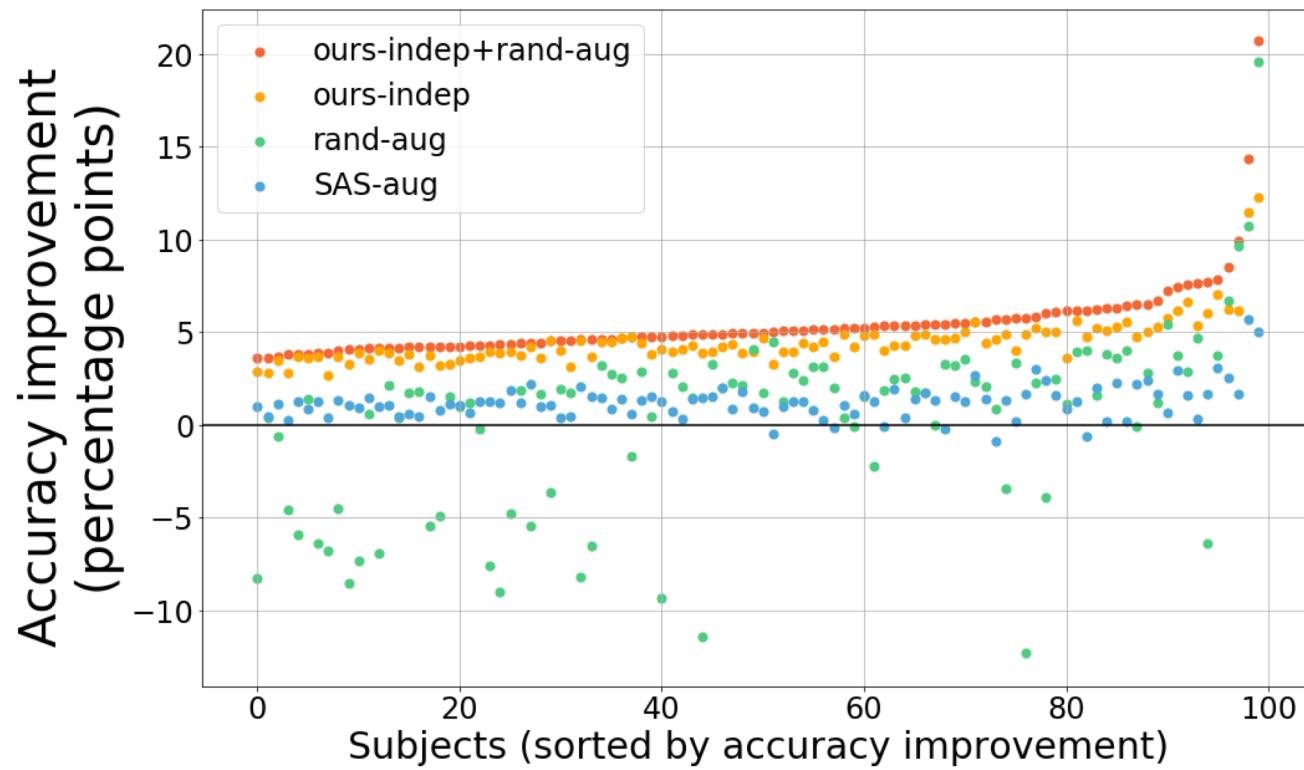
---

- Train supervised segmentation on synthesized “realistic” data  
accuracy increase  $76 (\pm 14) \rightarrow 81.5 (\pm 12)$



# Results

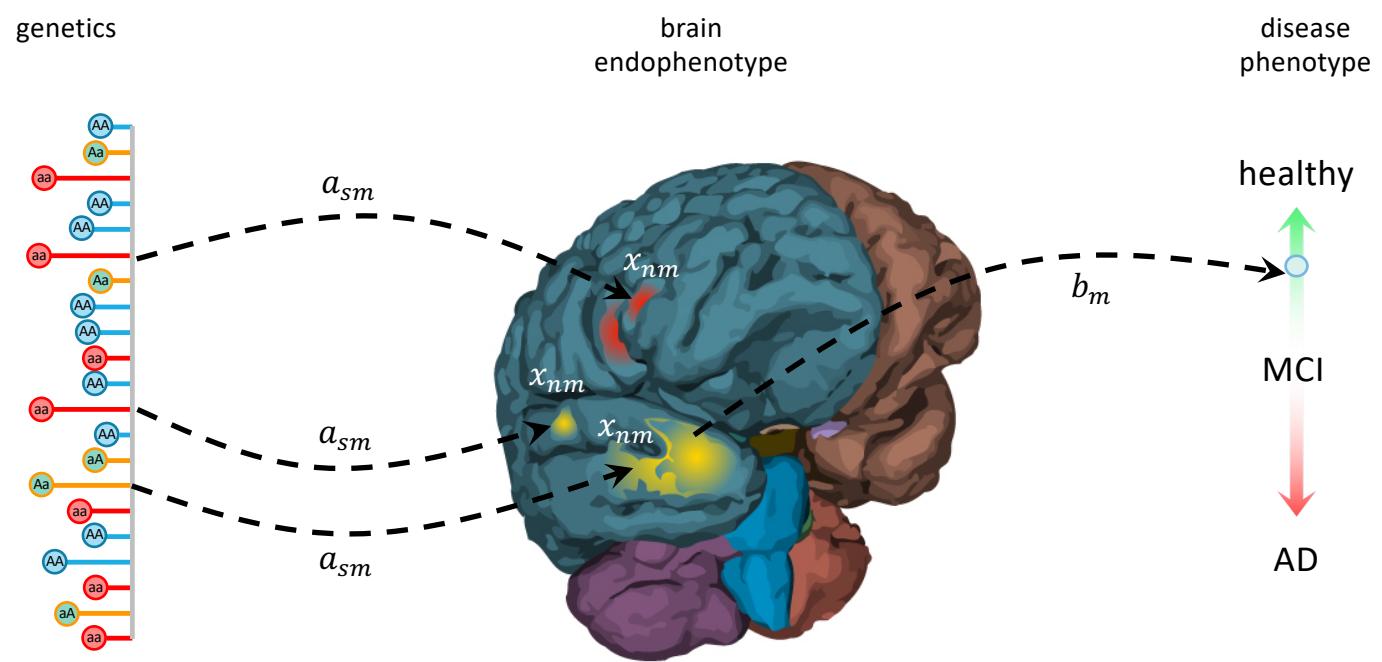
---



# Outline

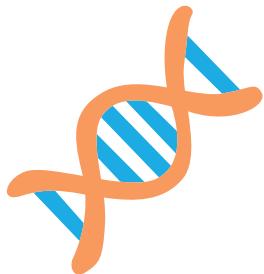
---

- Overview of Medical Imaging
  - Utility and properties
- Example: Segmentation
  - *Classical* and deep learning approaches
- Example: Registration (alignment):
  - Optimization and learning approaches
- **Example: Imaging Genetics**
- Takeaways





Scan at age 50

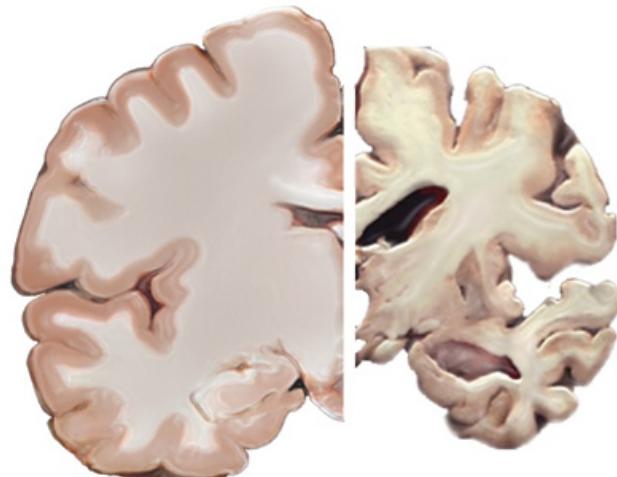


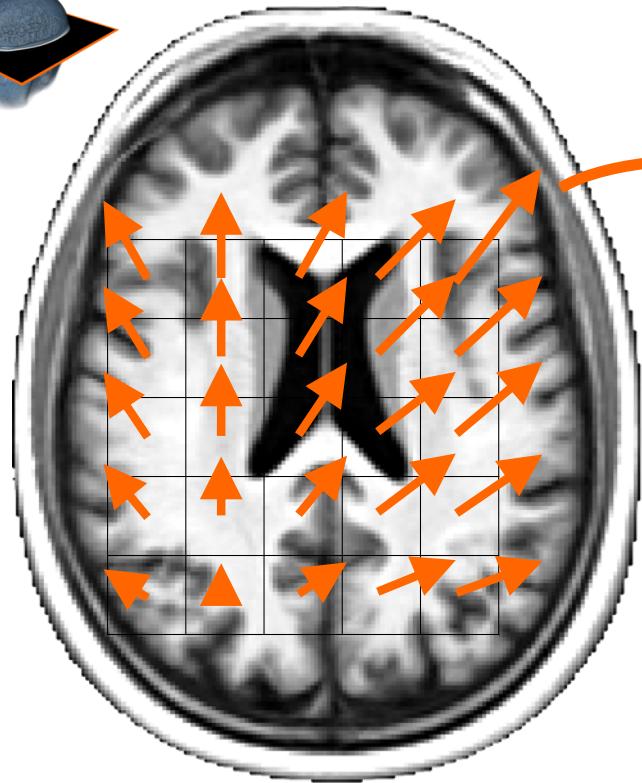
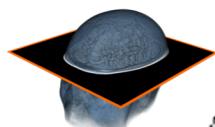
Scan at age 60

# Motivation

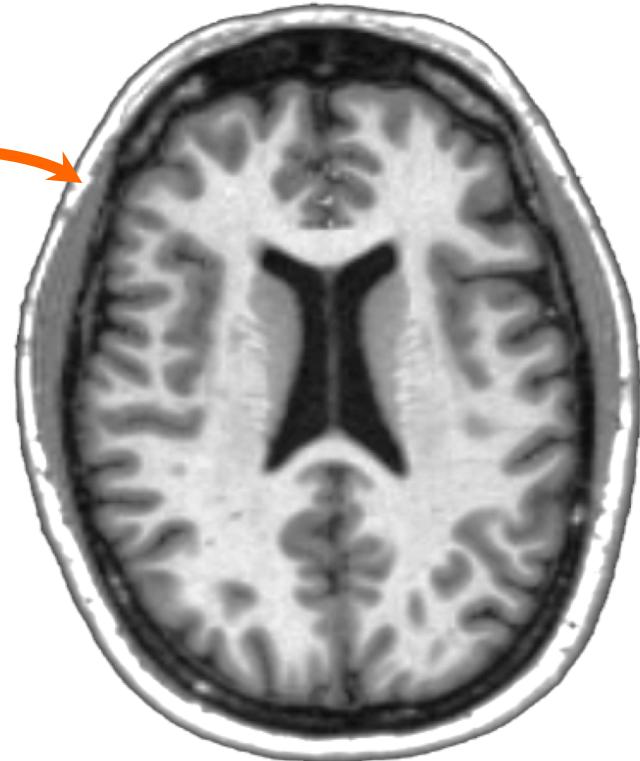
---

Healthy      Severe  
Brain      Alzheimer's

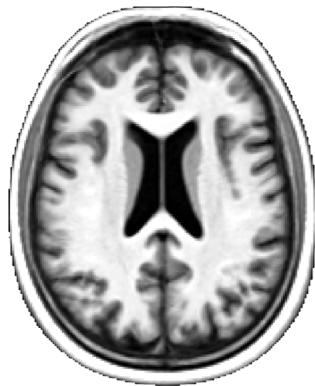




Atlas  
(average brain)



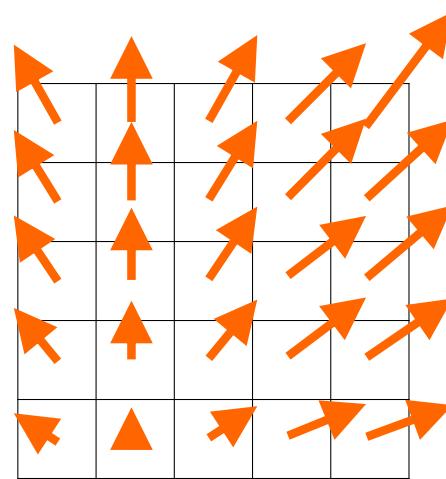
Subject  
medical scan



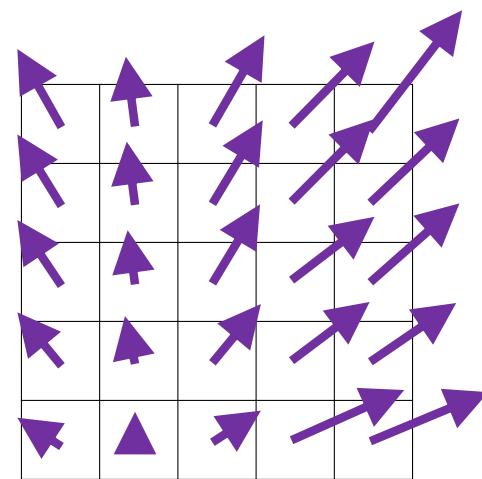
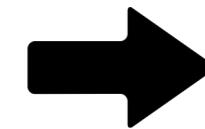
atlas



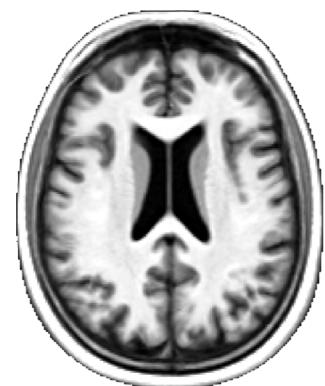
subject  
initial



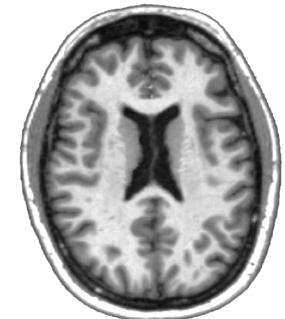
Initial  
displacement  
field



follow-up  
displacement  
field

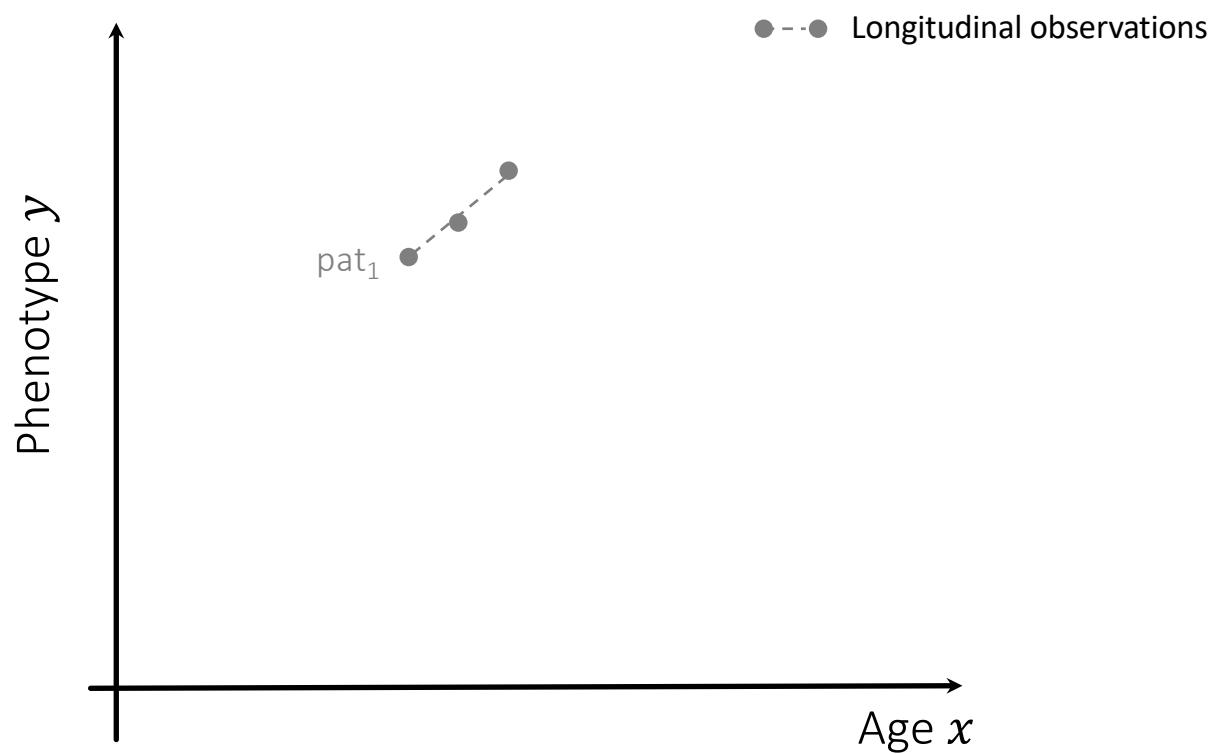


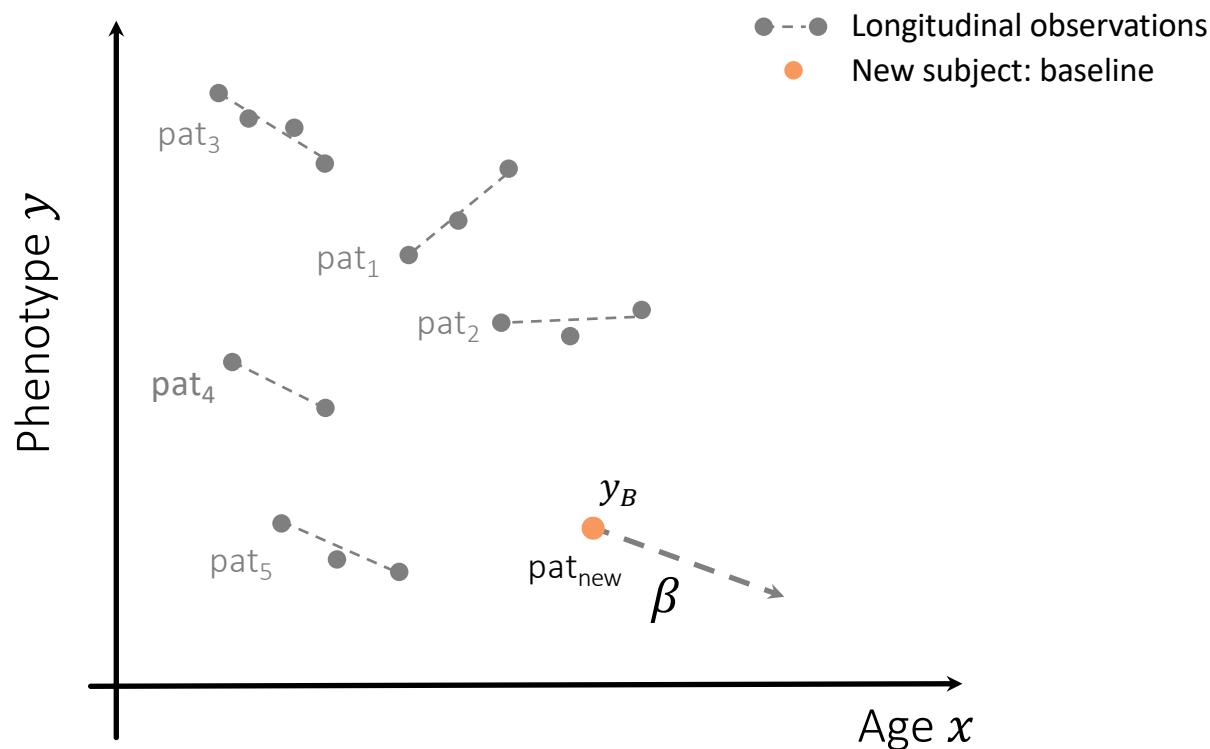
atlas



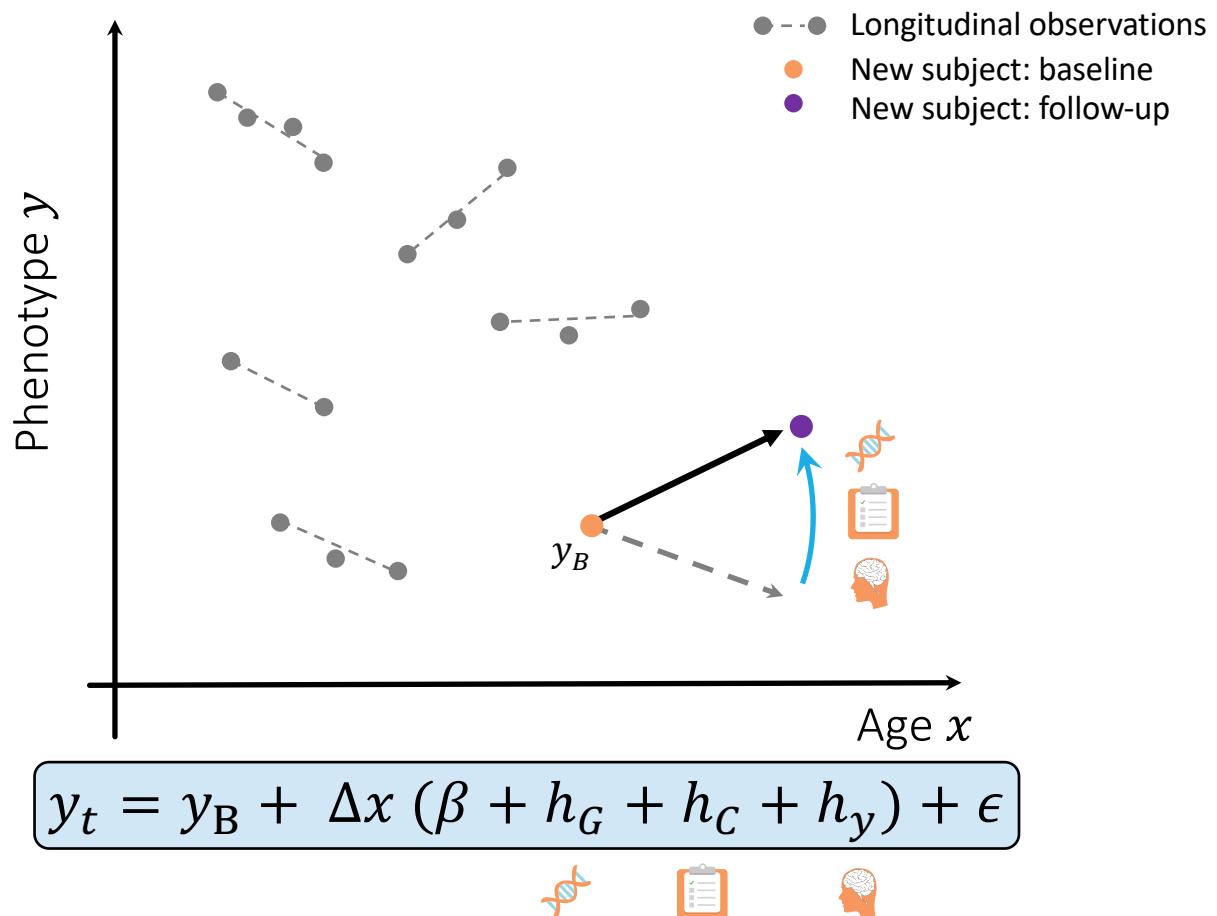
subject  
follow-up

# Scalar Phenotype Change





$$y_t = y_B + \Delta x_t \beta$$



# Model

---

$$y_t = y_B + \Delta x (\beta + h_G + h_C + h_y) + \epsilon$$



subject  $j$  param.

$$h_X(z_i) \sim \sum_j \overbrace{\alpha_j}^{} \underbrace{K_X(z_i, z_j)}_{}$$

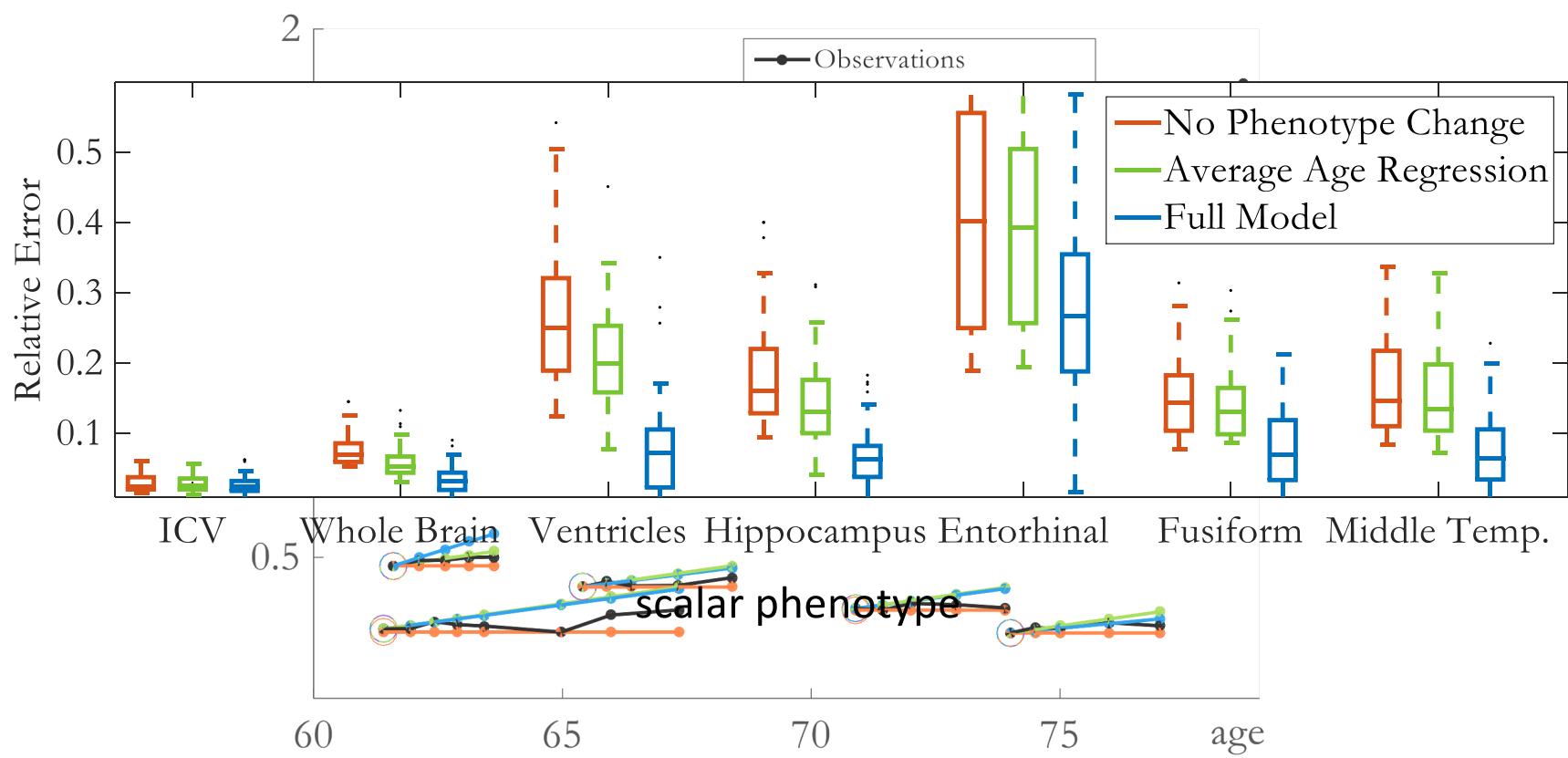
phenotype  $z$  similarity

Genetics Kernel

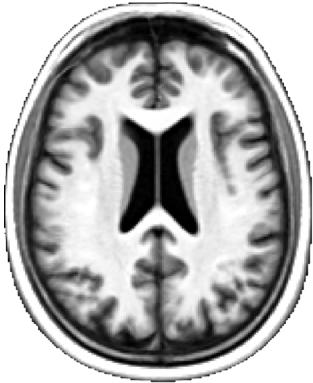
Main parameters:  $\alpha_j$  and  $\beta$

$\frac{1}{s}$

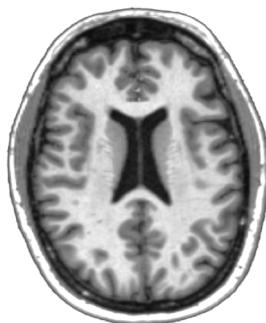
# Scalar Phenotype Results



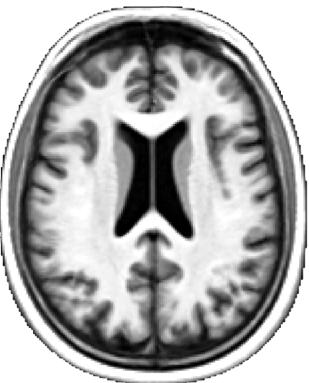
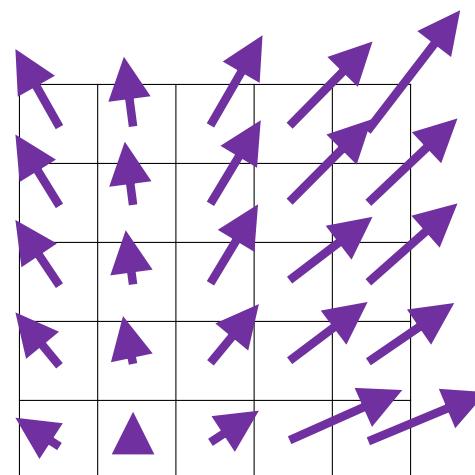
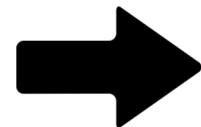
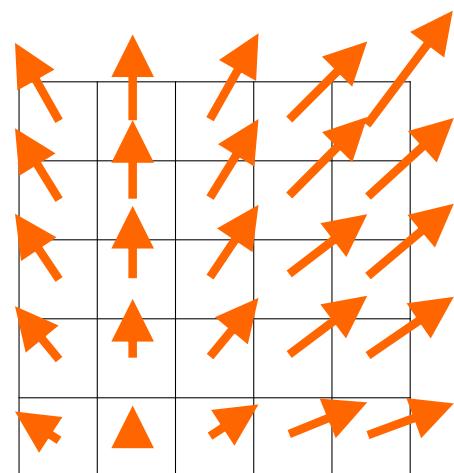
# Predict Anatomical Scans



atlas



baseline



atlas

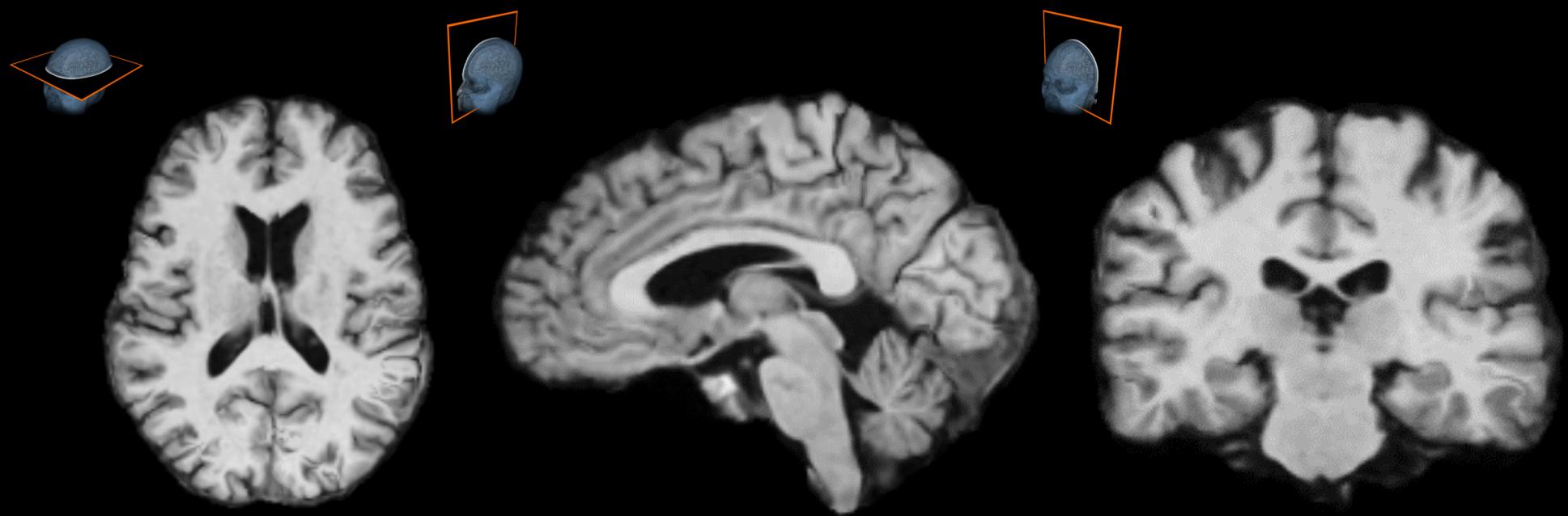


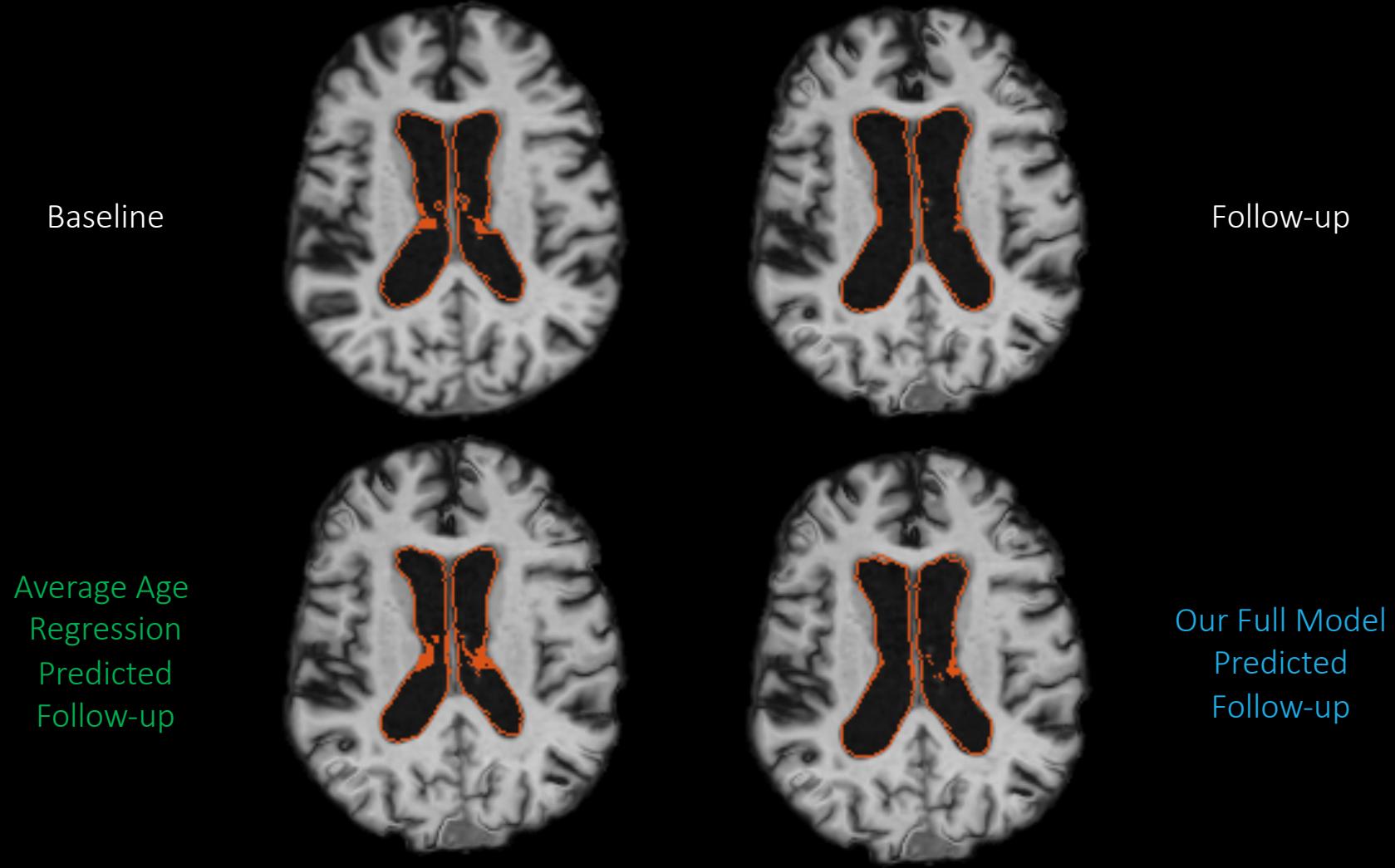
follow-up

$$\text{purple arrow} = \text{orange arrow} + \Delta x (\beta + h_G + h_C + h_y) + \epsilon$$

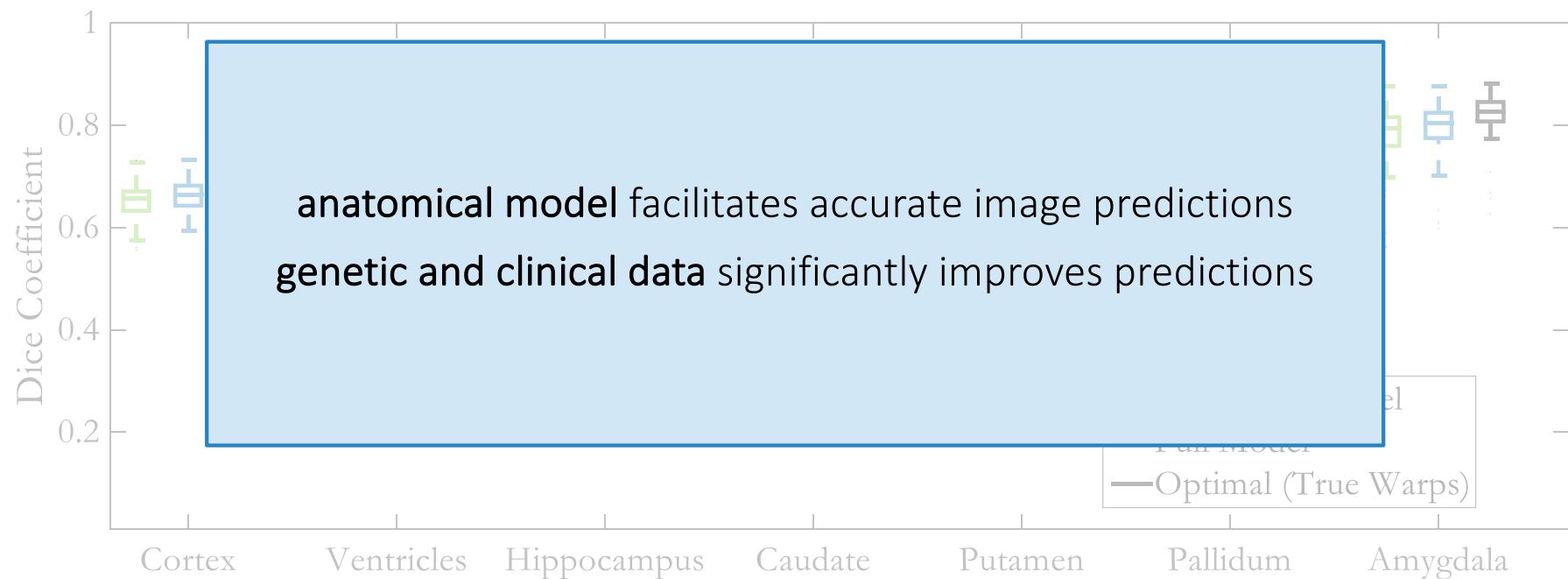


# Predicted follow-up scans



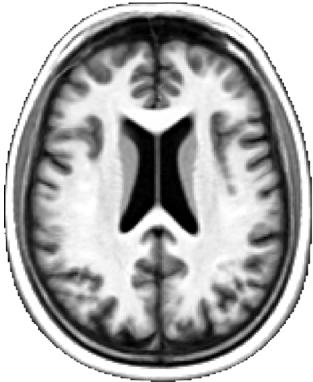


# Volume overlap: predicted vs real structures

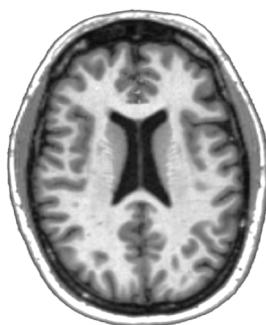


# Questions?

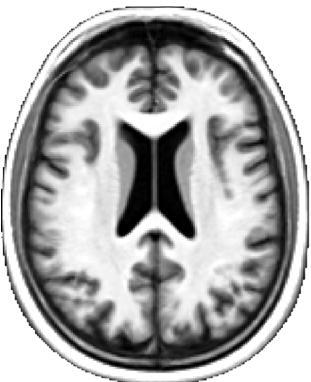
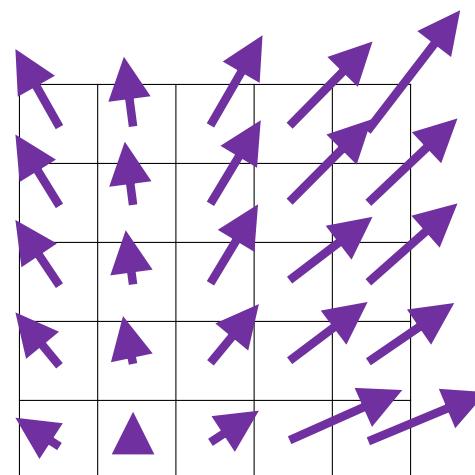
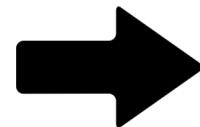
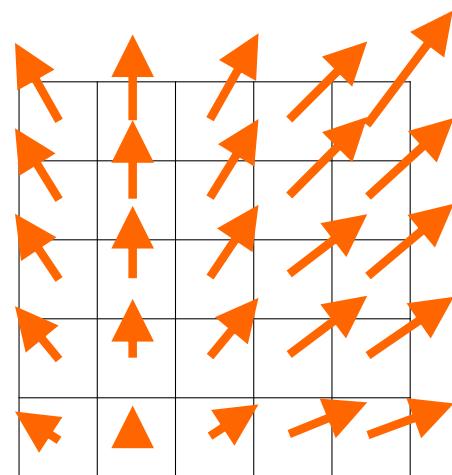
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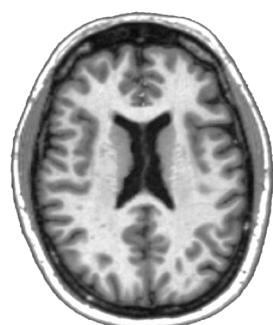
atlas



baseline



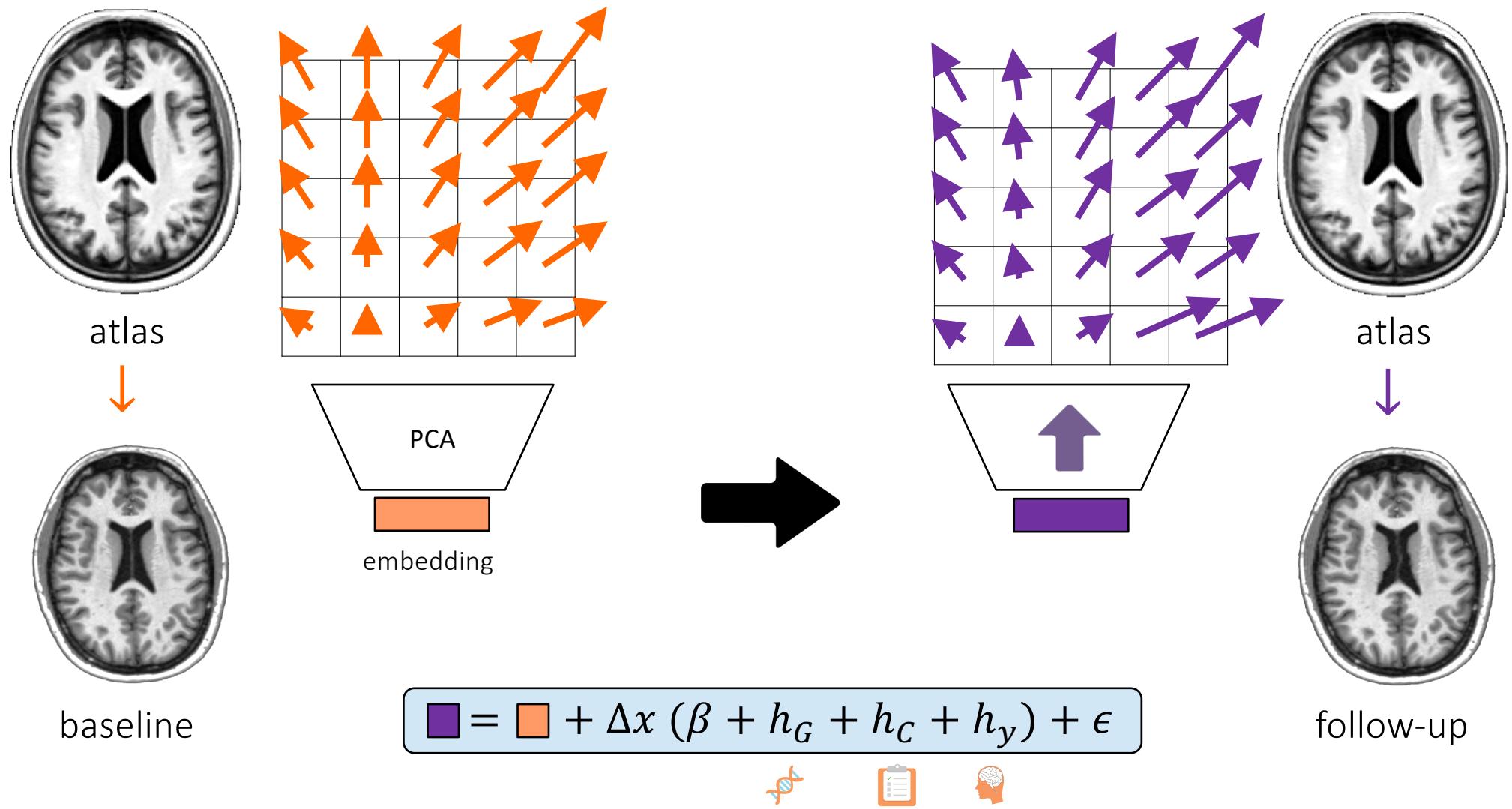
atlas



follow-up

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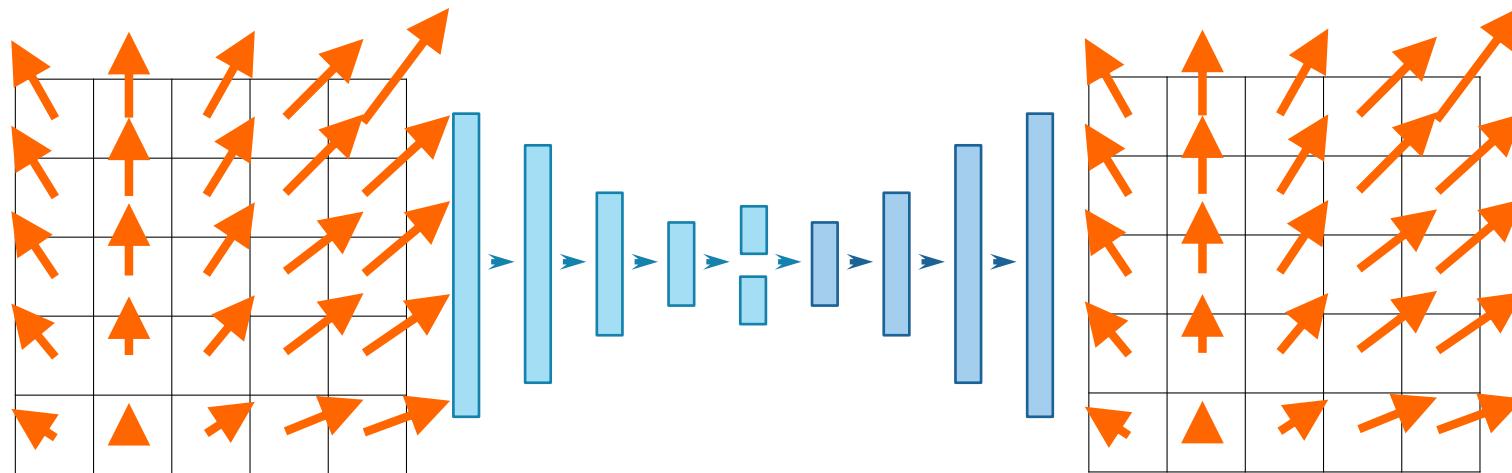




Linear models don't capture imaging or deformations well  
What to do?

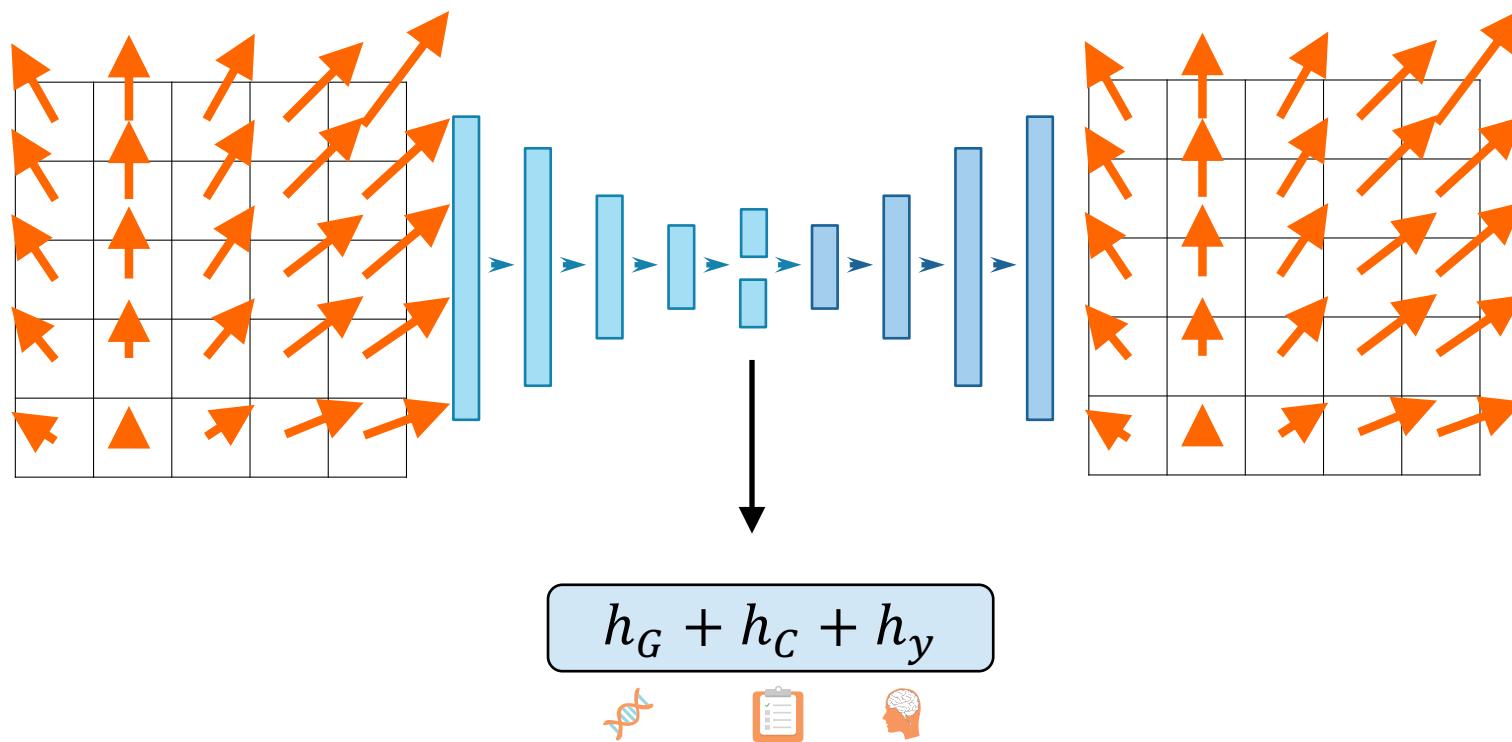
# Gaussian Process Prior VAE

- Capture deformations with a VAE:  $p(z) p(\phi|z)$



# Gaussian Process Prior VAE

- Capture deformations with a VAE:  $p(z) p(\phi|z)$
- Insert external data in the prior  $p(z; G, C, y)$  using Gaussian process!



# Conclusions

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- A lot of medical imaging data
  - Machine (deep) learning enabling fast, successful methods
- In realistic scenarios, usually few **labelled** images
- Combine **learning** concepts and **clinical** knowledge
  - Limited supervised data: leverage unlabeled data
  - Large data: anatomically regularized deep networks
- Measure success if you impact **downstream clinical tasks!**

# Outline

---

- Overview of Medical Imaging
  - Utility and properties
- Example: Segmentation
  - *Classical* and deep learning approaches
- Example: Registration (alignment):
  - Optimization and learning approaches
- Example: Imaging Genetics
- **Takeaways**

# Takeaway Goals

---

- Problems
  - Help the clinicians or scientists (don't replace them)
- Tools and approaches
  - Probabilities, convolutions, and anatomical models
  - Clinical interpretation
- Challenges
  - The systems don't really work (yet)
- Opportunity
  - Impact healthcare (and research)!

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