

# MRI predicts later outcomes

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- Quantitative analysis of tissue volume from MRI at term equivalent age has been shown to predict:
  - Impaired visual function in VLBW infants at age 2 (Shah et al. 2006)
  - Object working memory deficits at age 2 (Woodward et al. 2005)
  - PDI and MDI at age 2 (Thompson et al. 2008)

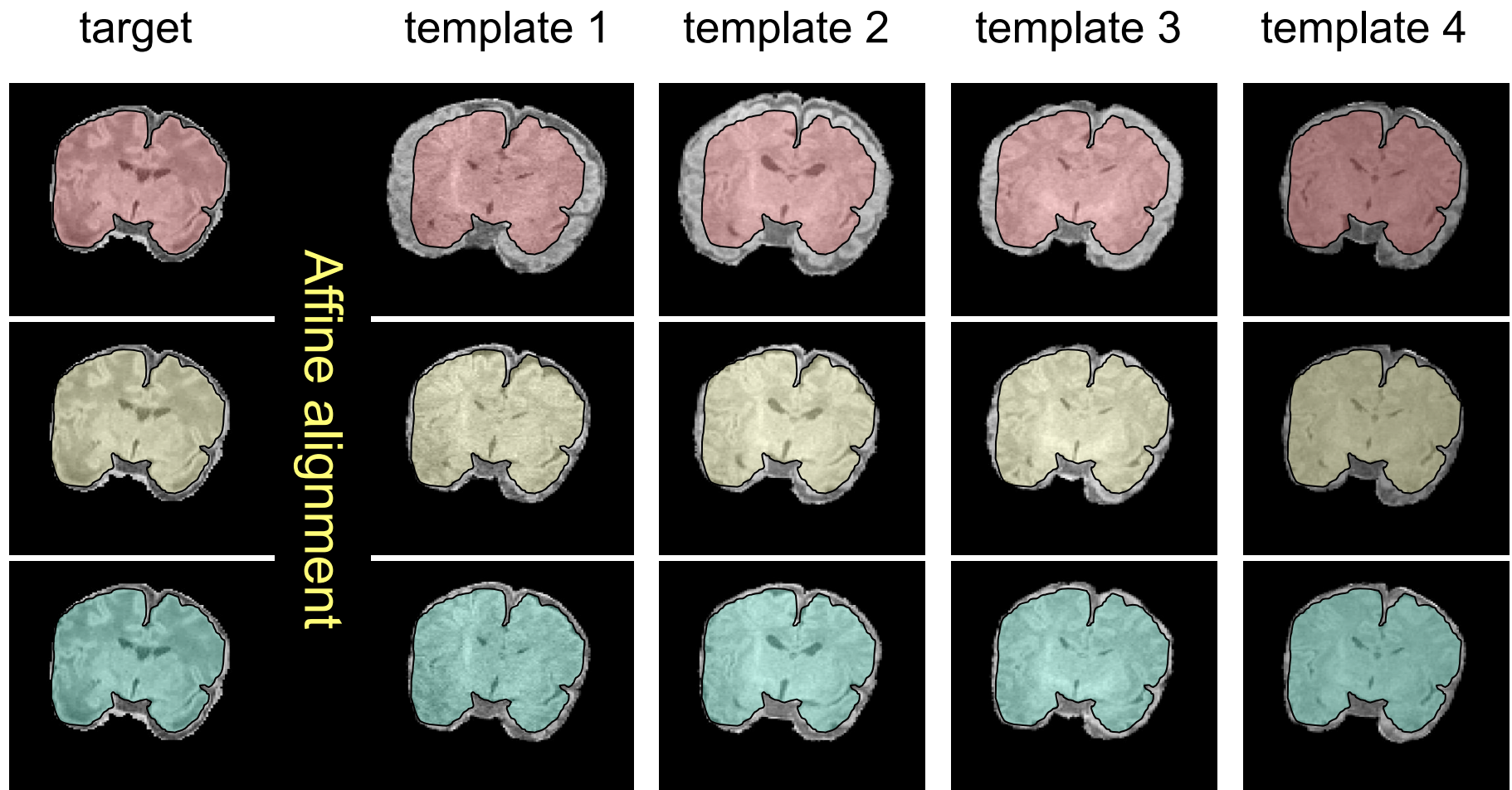
# Tissue Class Training Data

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- Our previous work has utilized interactive selection of per-subject training data:
  - Time consuming,
  - Subject to intra-rater and inter-rater variability,
  - Enabled identification of subtle contrast between different tissue types.
- We sought to develop an algorithm that avoids per-subject interaction, while maintaining excellent performance.
  - Weisenfeld and Warfield, NeuroImage, 2009.

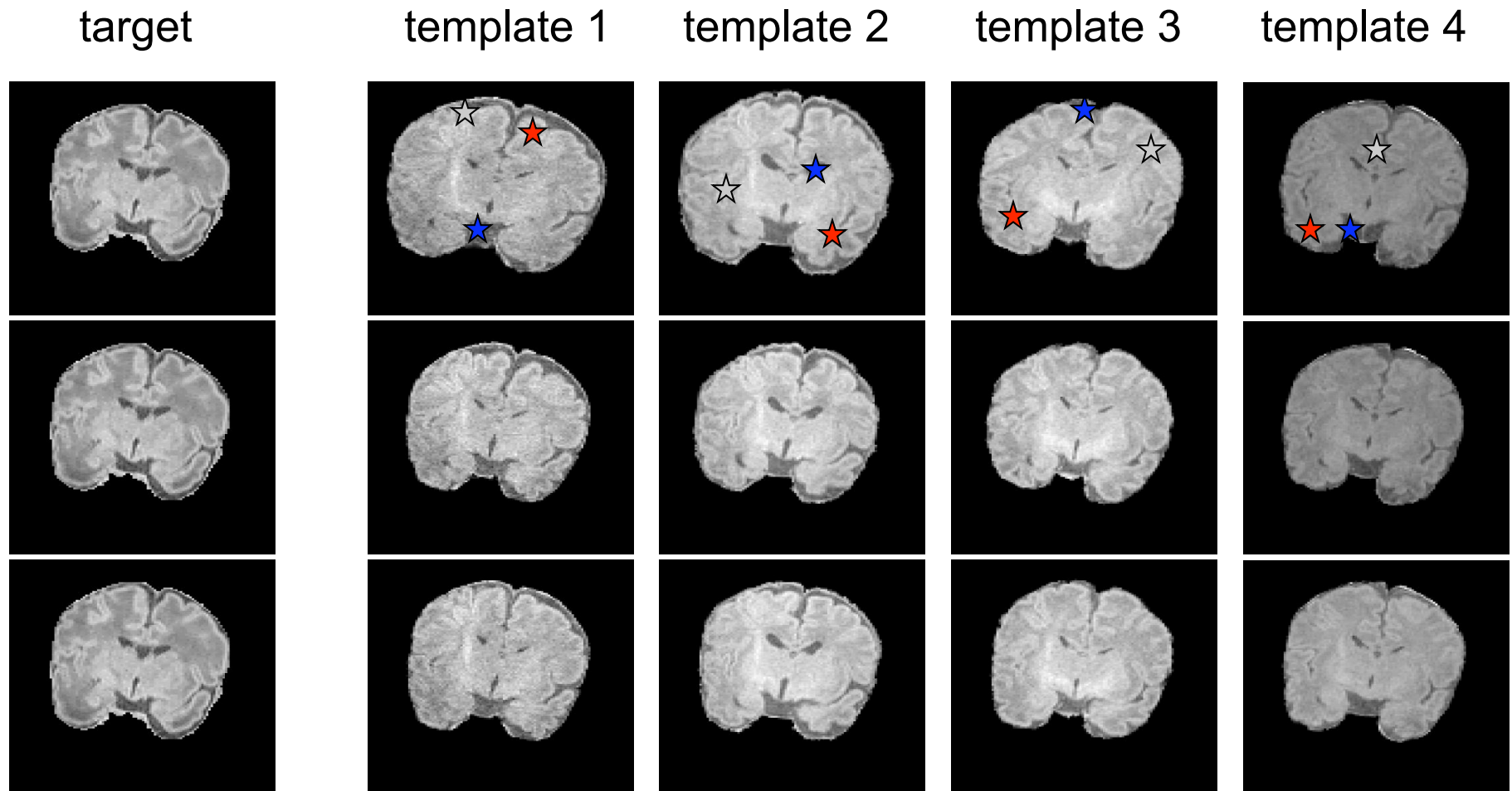
# Template to Target Registration

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# Tissue prototypes manually identified

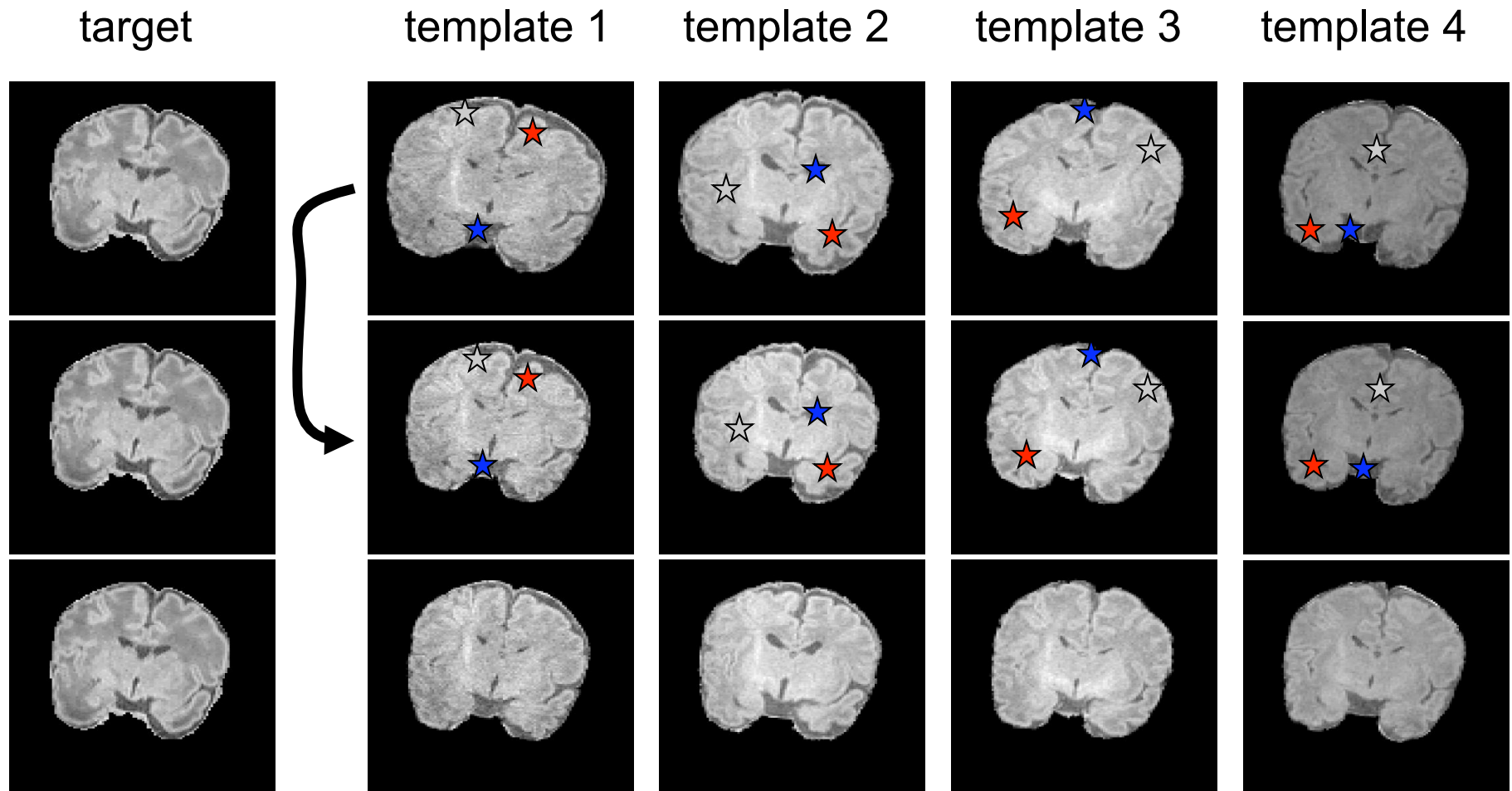
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tissue class samples selected once on the original template images.

# Tissue prototypes transferred

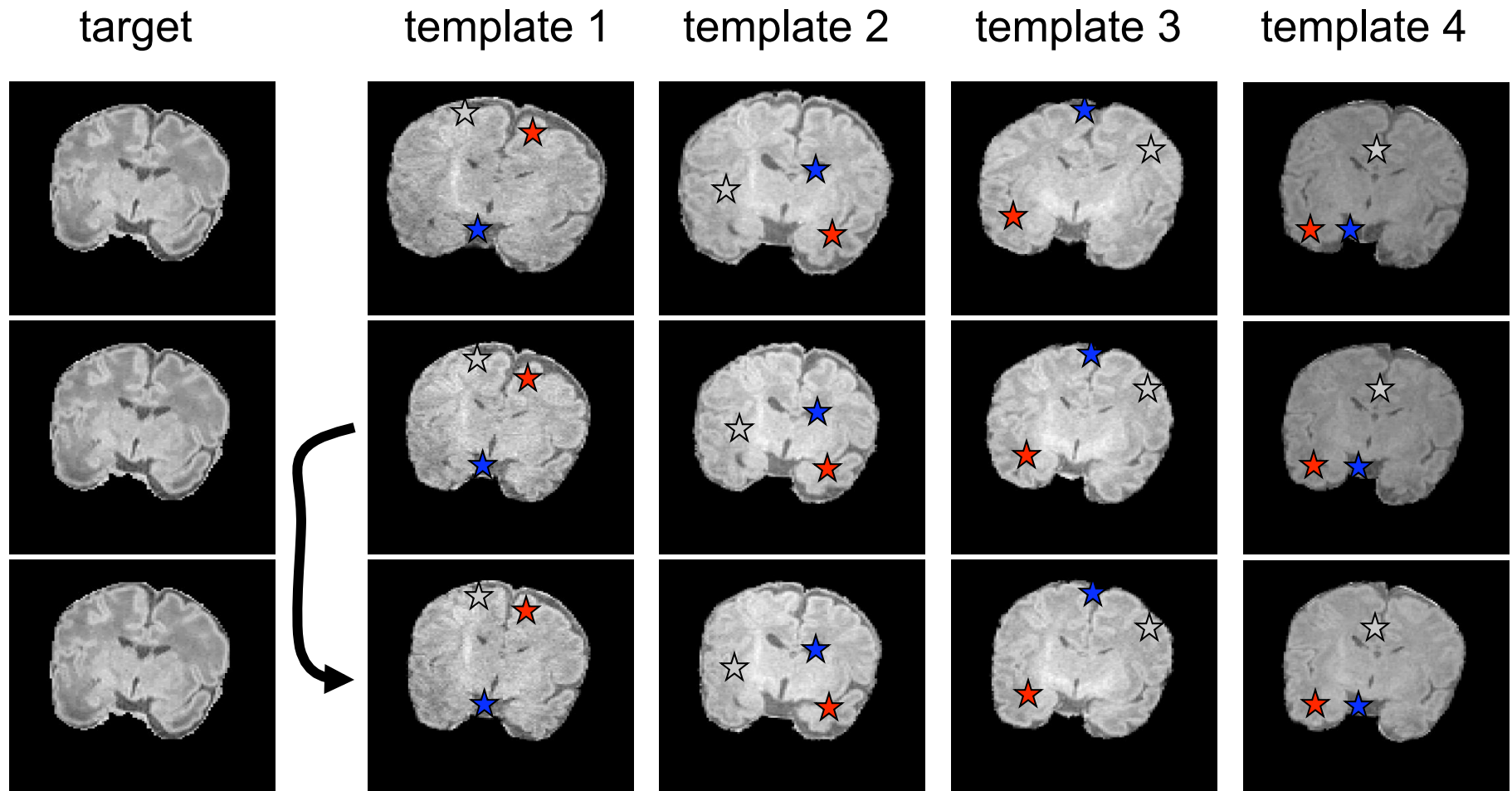
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and then projected through the affine transform...

# Tissue prototypes transferred

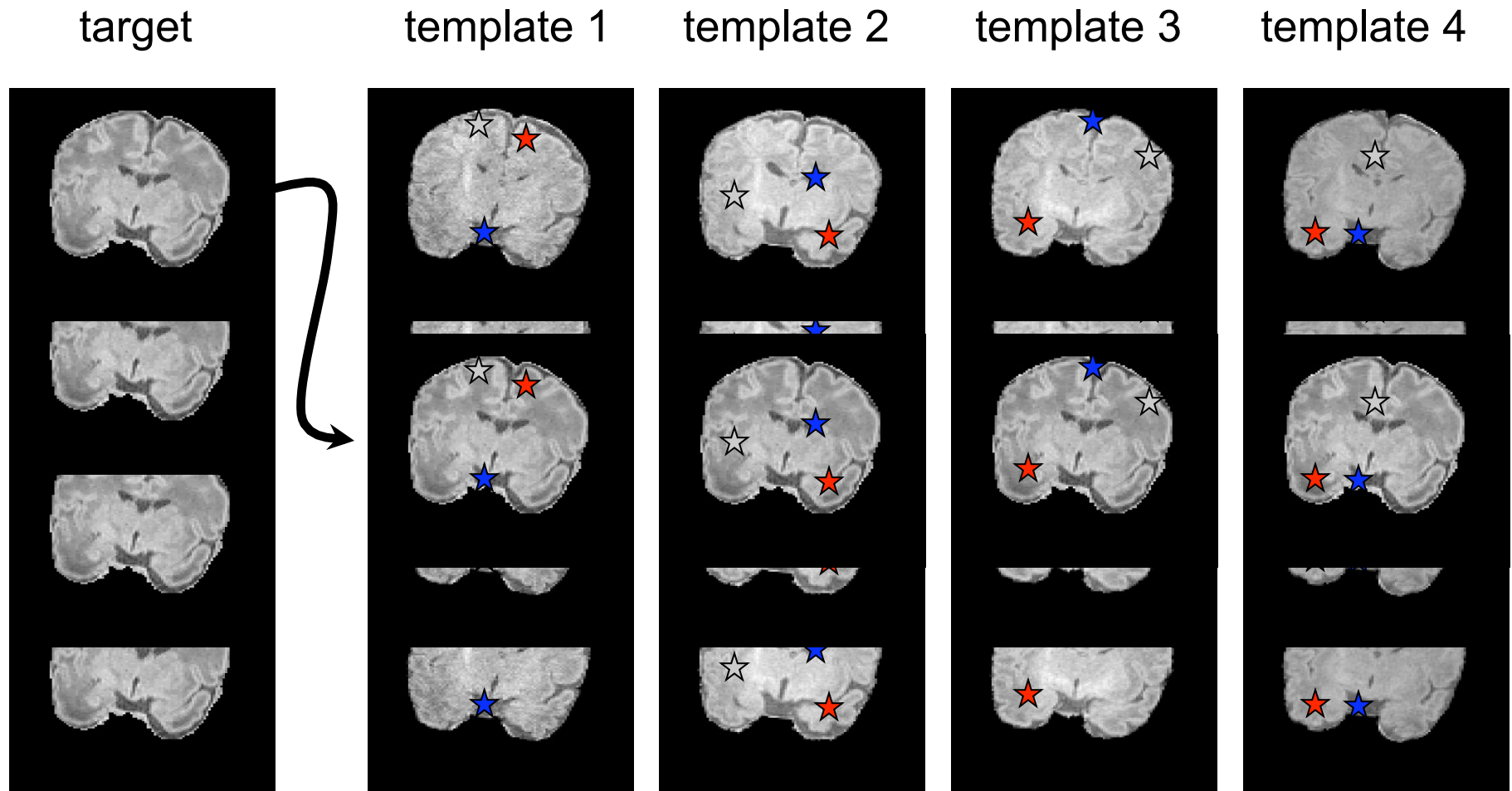
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and then projected through the b-spline *non-linear* transform...

# Tissue prototypes transferred

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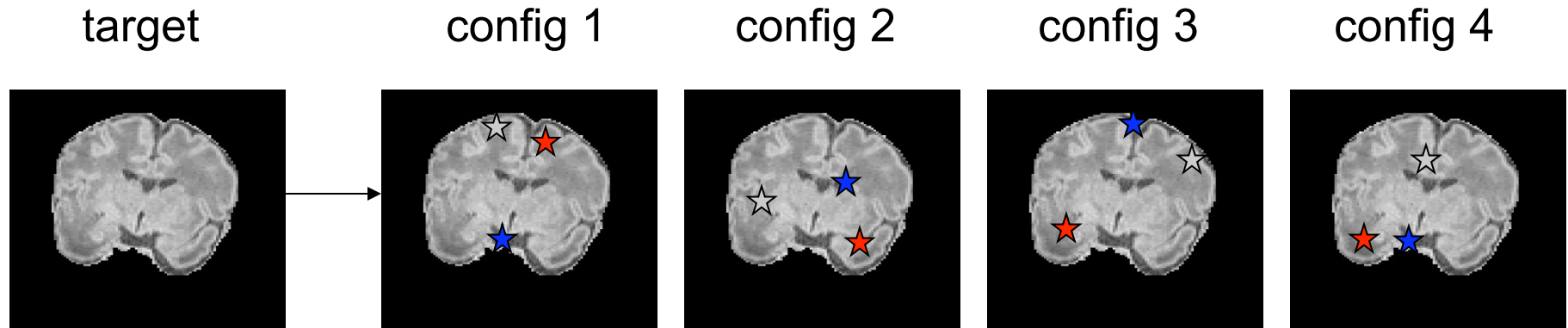


Different prototype configurations are projected onto the target subject

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# Multiple Configurations on the Target

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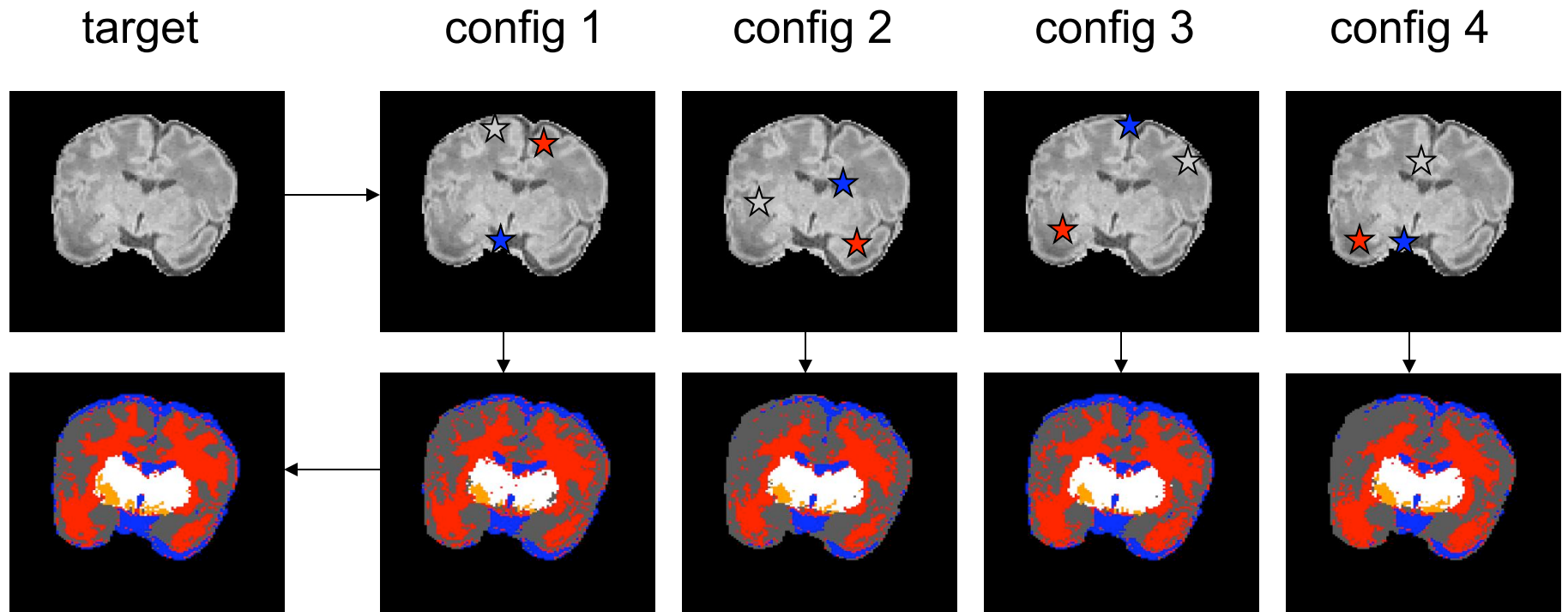


The different prototype configurations represent the physical variation among the template subjects. By adding template subjects, and choosing prototypes by hand *only once*, a wider range of physical variation can be accommodated. Once a template subject is added, it is re-used without further human intervention.

The image *intensity* data used is *only* from the individual under study (the target).

# Multiple Configurations on the Target

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Each configuration of sample coordinates leads to a different candidate segmentation of the target subject.

STAPLE is used to combined candidate segmentations.

# Configurations are Edited

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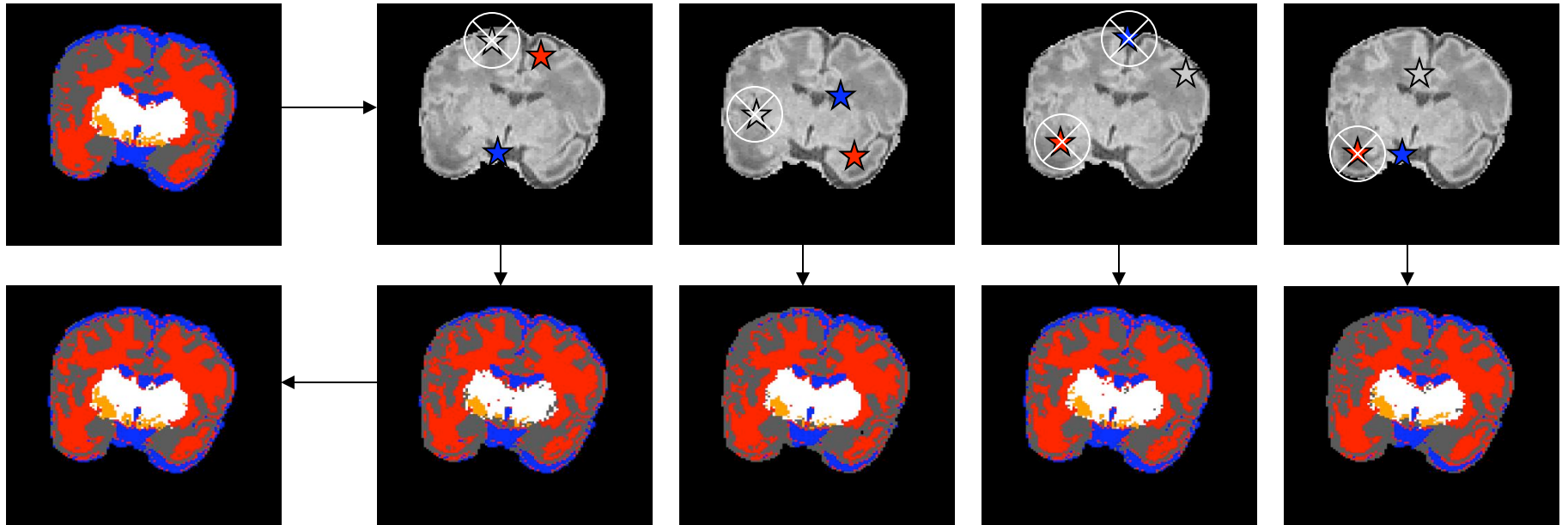
estimated truth

config 1

config 2

config 3

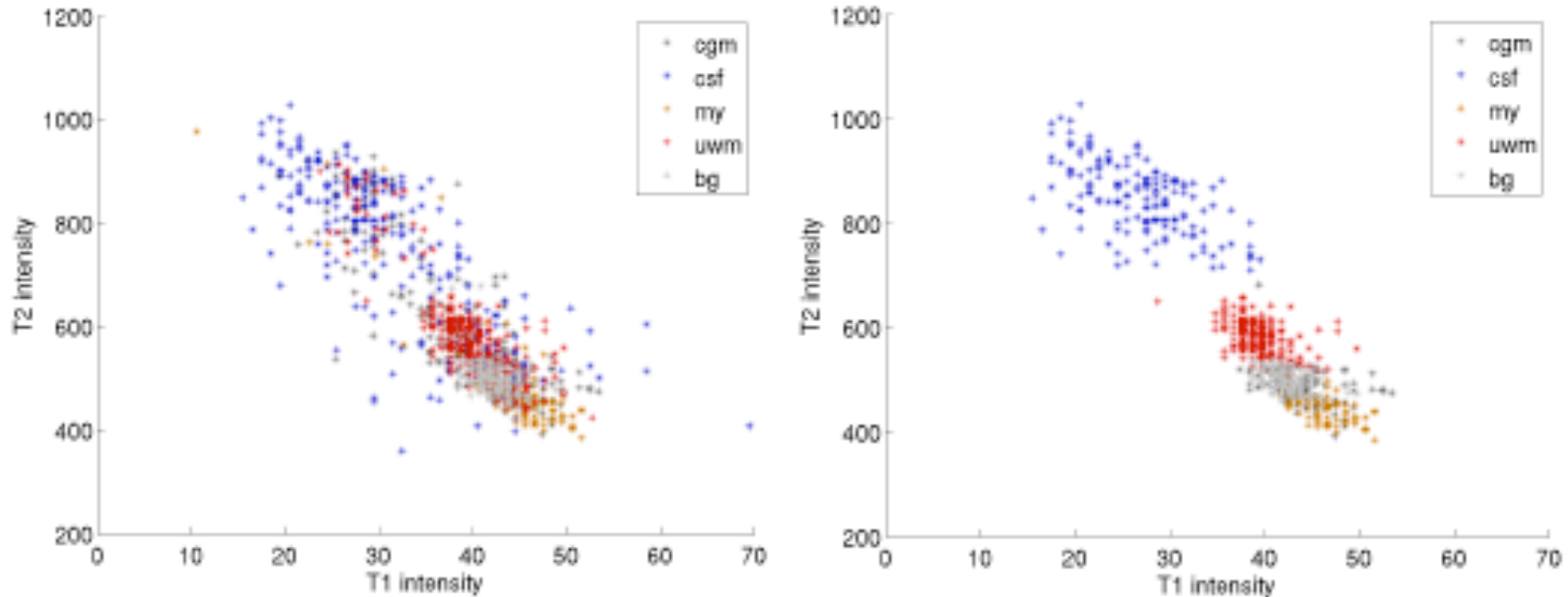
config 4



The previous iteration's STAPLE output (top left) is used to identify and eliminate prototypes which are inconsistent with the data.

# Adaptation of training data

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Evolution of feature space of training data through the automated projection and editing process.

Tissue class boundaries in feature space are identified.

# Evaluation of training data

		train	
		$\mathcal{M}$	$\mathcal{A}$
test	$\mathcal{M}$	$0.95 \pm 0.02$	$0.93 \pm 0.02$
	$\mathcal{A}$	$0.95 \pm 0.01$	$0.97 \pm 0.01$

Posterior probability of correct classification with manually and automatically generated training data.

Subject	$\mathcal{M}$	test				
		$\mathcal{A}^0$	$\mathcal{A}^1$	$\mathcal{A}^2$	$\mathcal{A}^3$	$\mathcal{A}^{\text{FINAL}}$
1	0.98	0.66	0.77	0.86	0.91	0.93
2	0.96	0.65	0.77	0.87	0.92	0.94
3	0.96	0.66	0.79	0.89	0.94	0.95
4	0.96	0.65	0.78	0.87	0.92	0.94
5	0.94	0.66	0.78	0.87	0.93	0.95
6	0.94	0.67	0.80	0.90	0.94	0.95
7	0.94	0.66	0.79	0.89	0.94	0.96
8	0.94	0.70	0.82	0.91	0.96	0.97
9	0.94	0.67	0.79	0.89	0.93	0.95
10	0.96	0.69	0.80	0.89	0.94	0.96
mean±sd	$0.95 \pm 0.02$	$0.67 \pm 0.02$	$0.79 \pm 0.02$	$0.88 \pm 0.02$	$0.93 \pm 0.01$	$0.95 \pm 0.01$

Improved consistency of GM, UWM and CSF over iterations of editing of training data.

# Segmentation comparison

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Subject	CGM	CSF	myelin	UMWM	SCGM
1	0.95	0.82	0.63	0.95	0.92
2	0.89	0.94	0.80	0.89	0.86
3	0.89	0.93	0.71	0.91	0.89
4	0.84	0.95	0.70	0.85	0.81
5	0.92	0.93	0.77	0.93	0.88
6	0.89	0.98	0.77	0.93	0.85
7	0.93	0.96	0.70	0.96	0.89
8	0.91	0.97	0.79	0.93	0.87
9	0.87	0.94	0.66	0.91	0.80
10	0.94	0.80	0.67	0.95	0.88
mean±sd	0.90 ± 0.03	0.92 ± 0.06	0.72 ± 0.06	0.92 ± 0.03	0.86 ± 0.04

Dice coefficient comparing interactive to automated tissue classification.

# Segmentation comparison

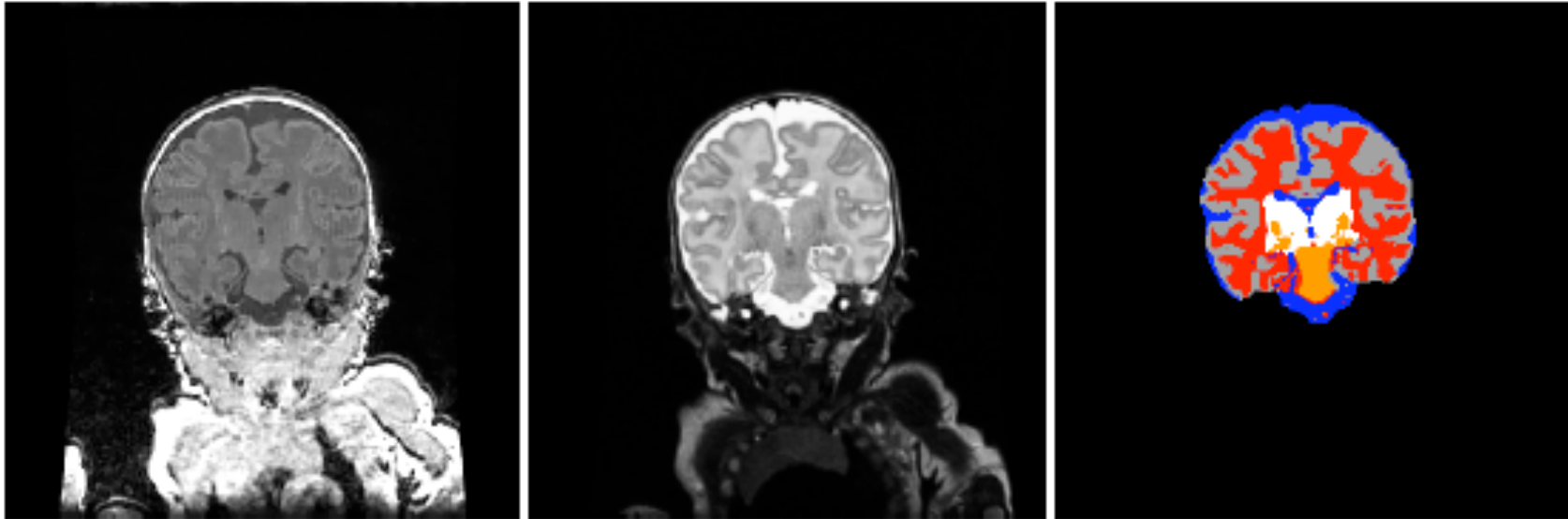
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Subject		CGM	CSF	myelin	UMWM	SCGM
1	experts	$0.86 \pm 0.06$	$0.89 \pm 0.05$	$0.81 \pm 0.11$	$0.85 \pm 0.05$	$0.86 \pm 0.08$
	automatic	0.75	0.96	0.86	0.79	0.96
2	experts	$0.87 \pm 0.06$	$0.93 \pm 0.02$	$0.96 \pm 0.05$	$0.87 \pm 0.06$	$0.90 \pm 0.12$
	automatic	0.77	0.98	0.96	0.72	0.74
3	experts	$0.90 \pm 0.04$	$0.91 \pm 0.02$	$0.77 \pm 0.06$	$0.88 \pm 0.03$	$0.91 \pm 0.03$
	automatic	0.77	0.97	0.81	0.78	0.95
4	expert	$0.87 \pm 0.08$	$0.91 \pm 0.02$	$0.81 \pm 0.06$	$0.87 \pm 0.04$	$0.94 \pm 0.04$
	automatic	0.84	0.95	0.69	0.70	0.94

Comparison of predictive values of tissue segmentations obtained by interactive drawing and by automated tissue classification.

# Newborn brain segmentation

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# Segmentation Algorithm

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- Weisenfeld and Warfield, NeuroImage 2009
- Automatic estimation of training data is comparable to interactive selection by an expert.
- Automated segmentation compares well to hand-drawn segmentations.
- Software for pediatric MRI analysis, CRKit, supported by NIH.

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