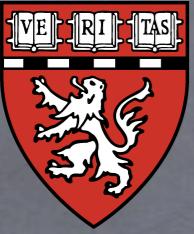


Interpolating Multi-Fiber Models by Gaussian Mixture Simplification

ISBI 2012, Barcelona, Spain

Maxime Taquet,
Benoît Scherrer,
Christopher Benjamin,
Sanjay Prabhu,
Benoît Macq,
Simon K. Warfield

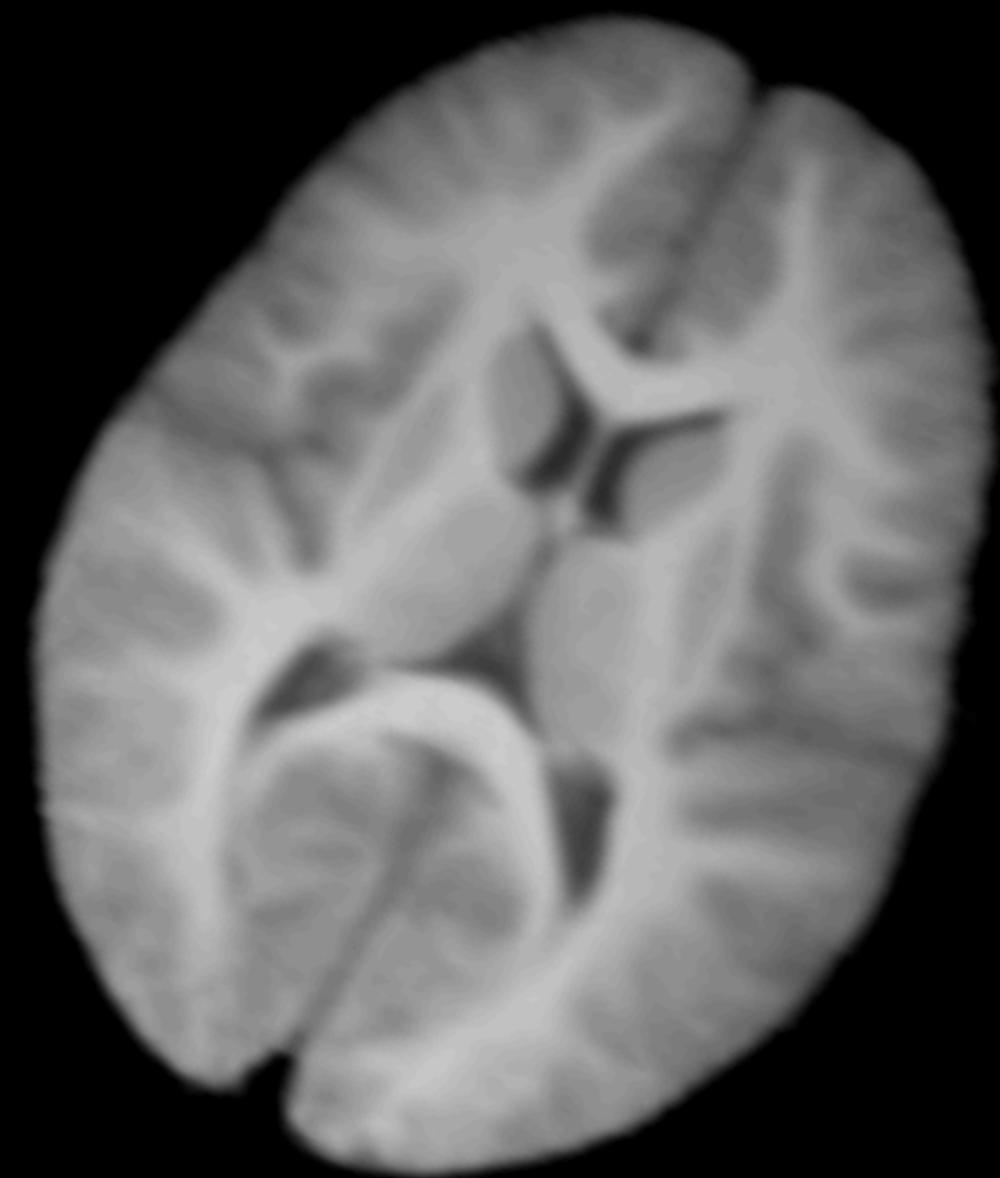
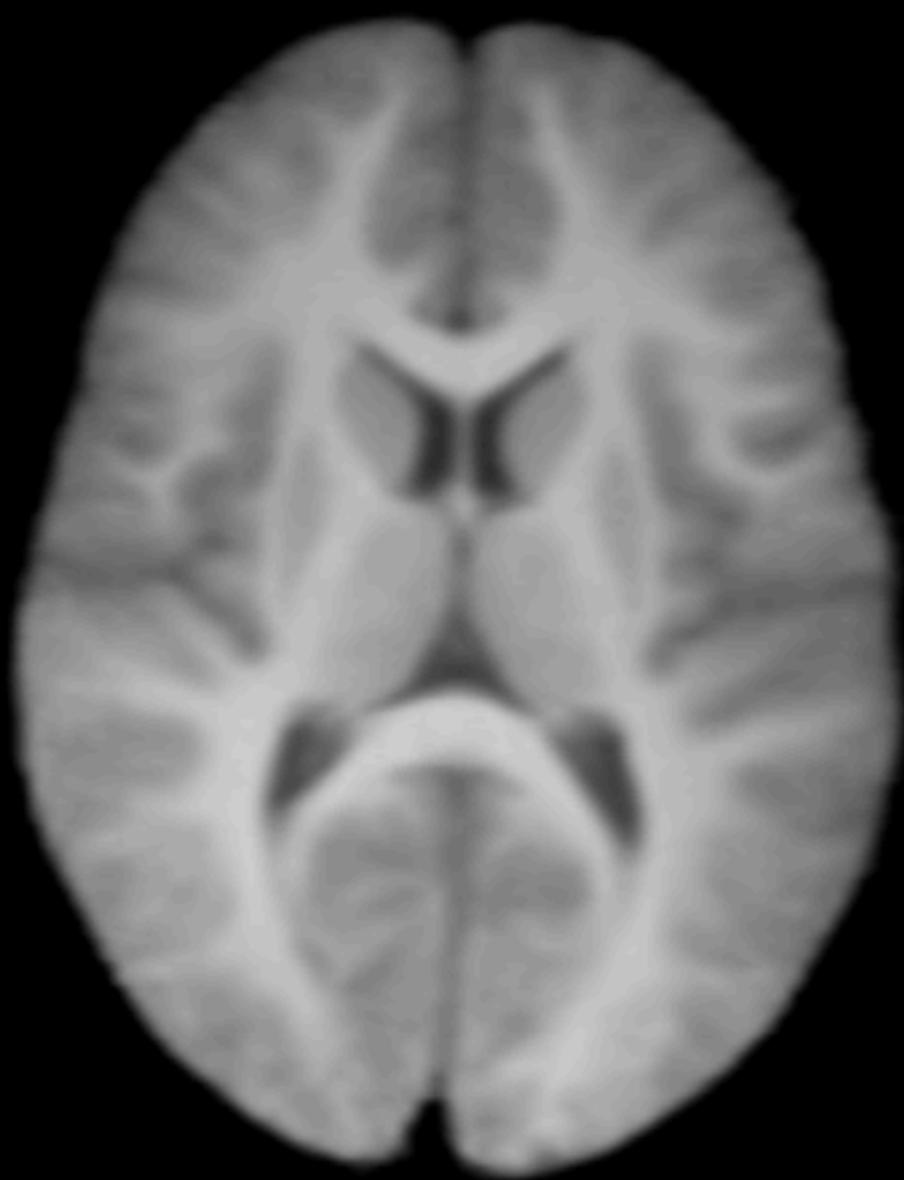


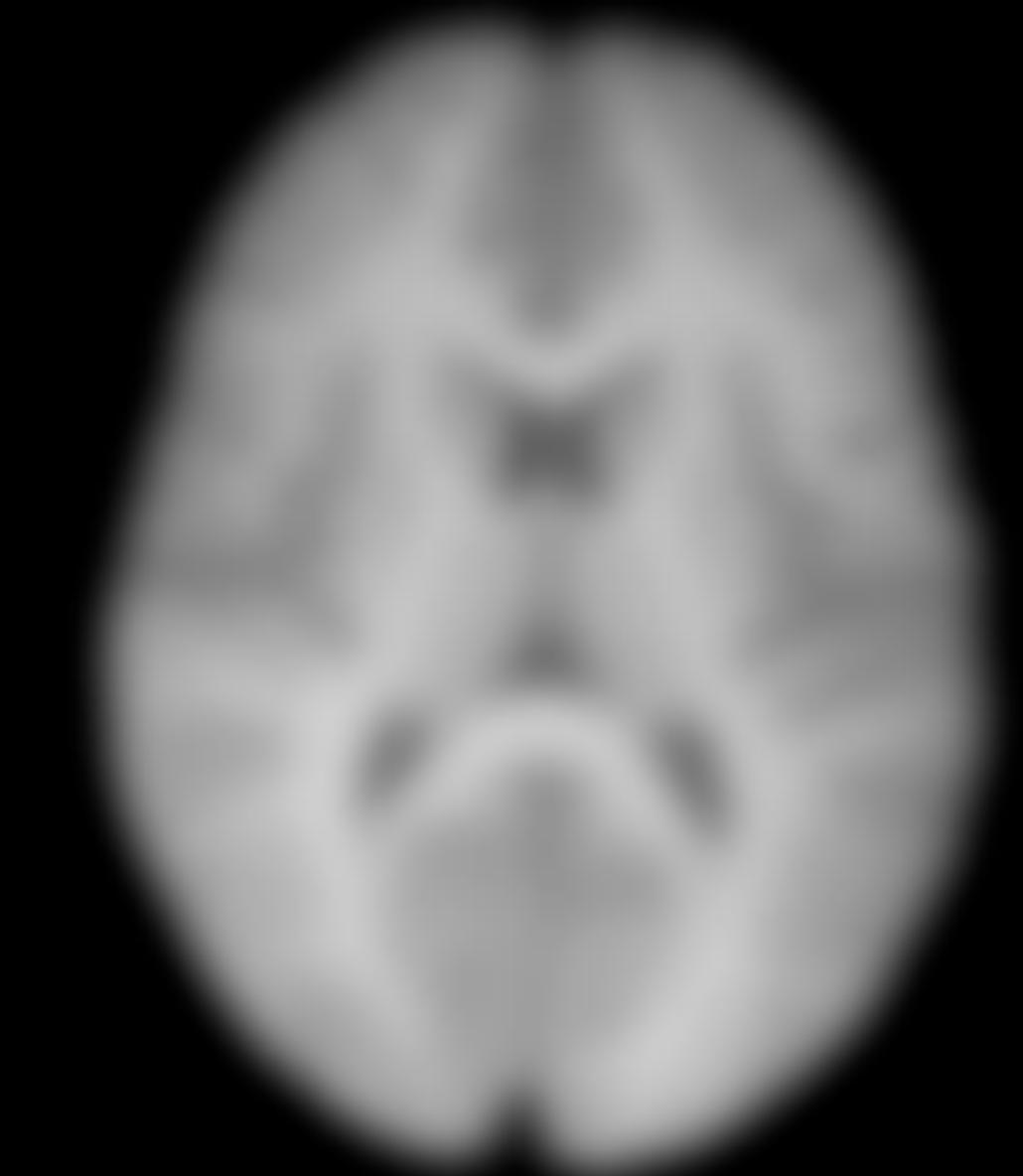
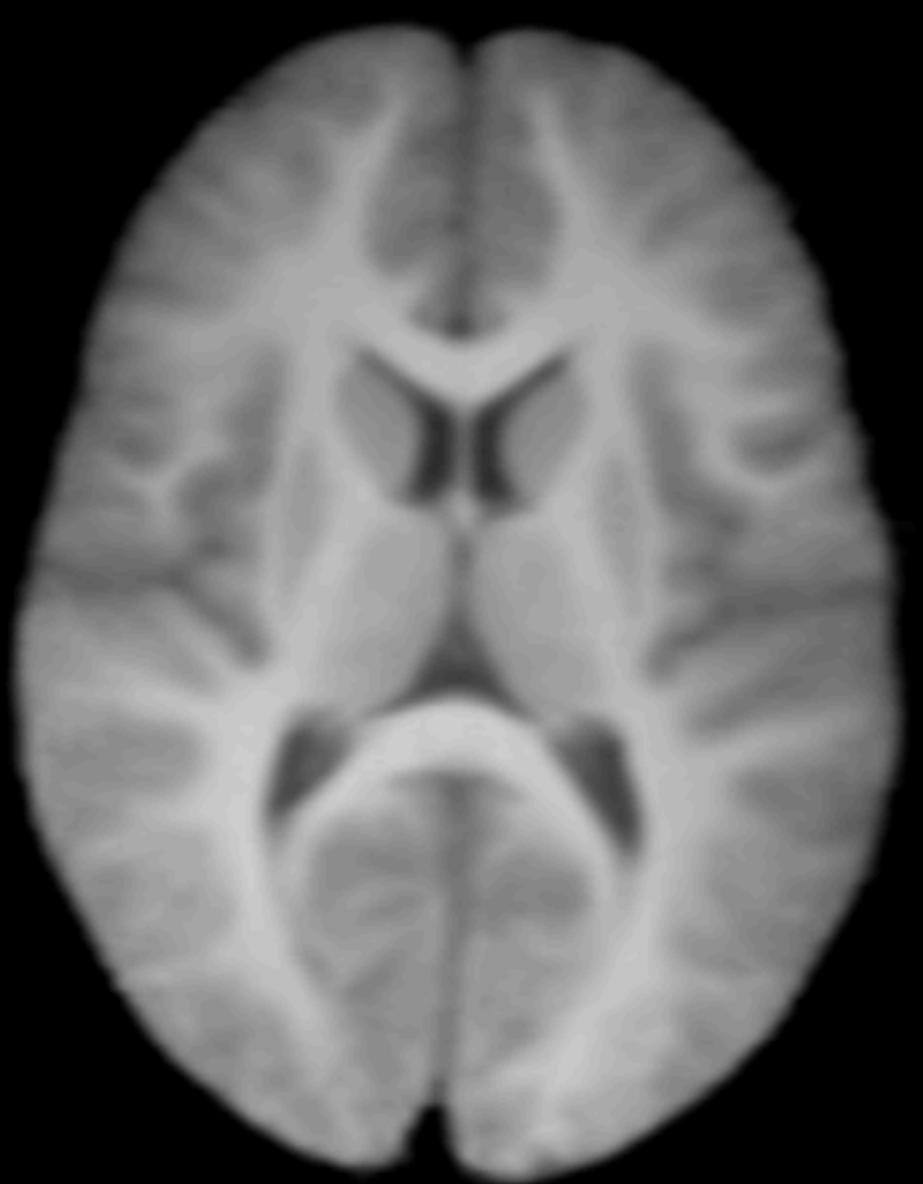


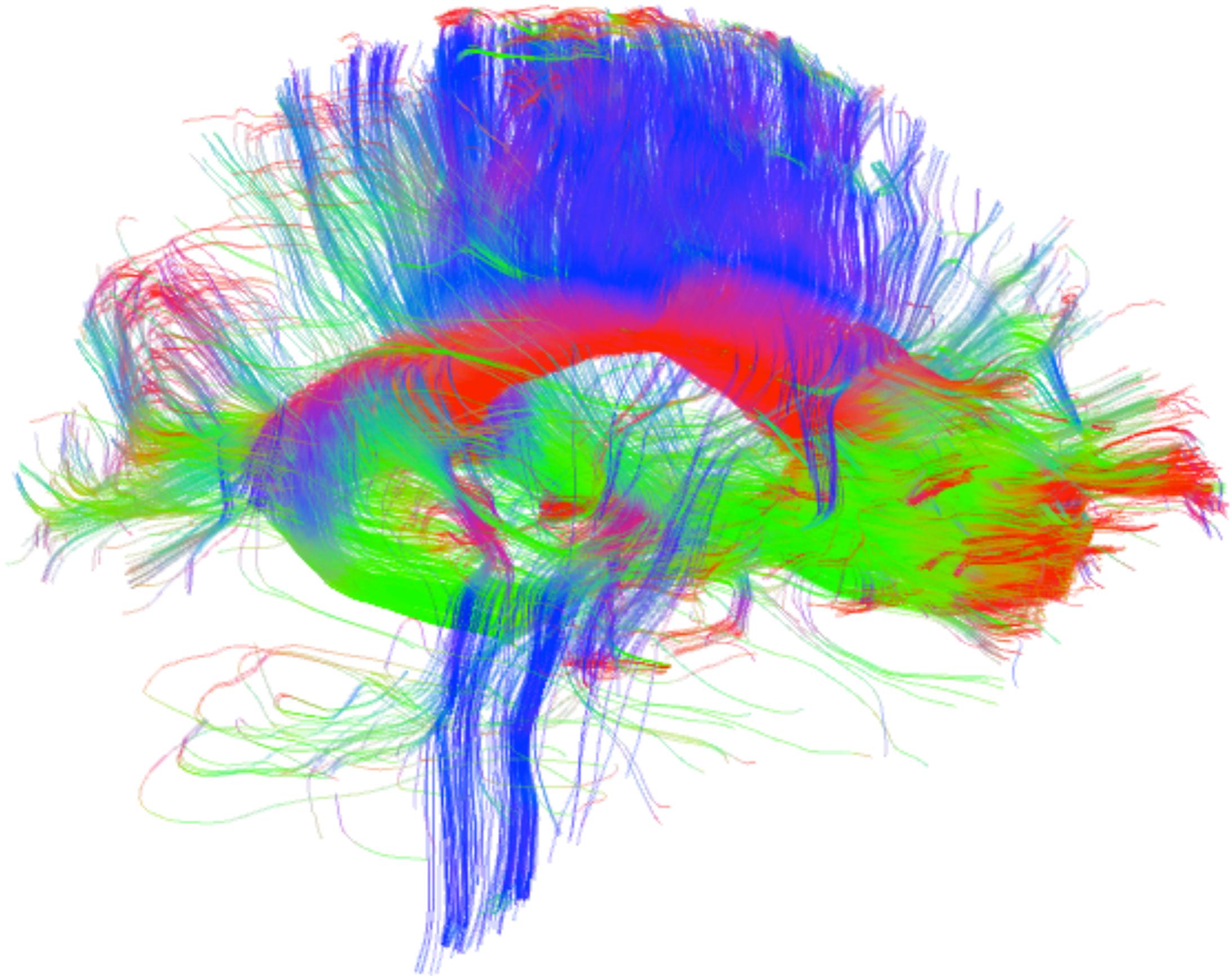
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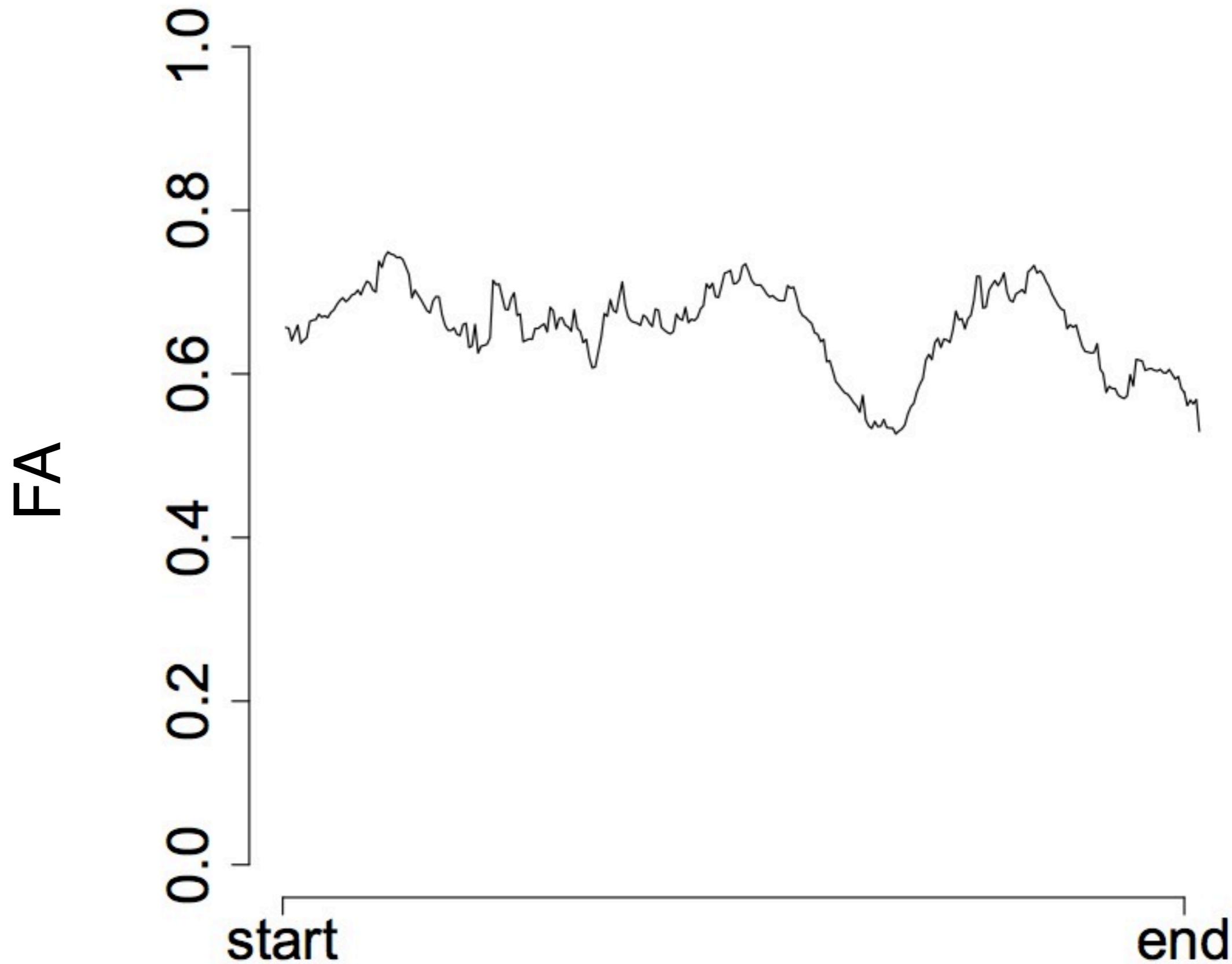
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Scalar characteristic along a tract



Heuristics fail
a global approach is required

The problem is well known
as a Gaussian Mixture Model Simplification

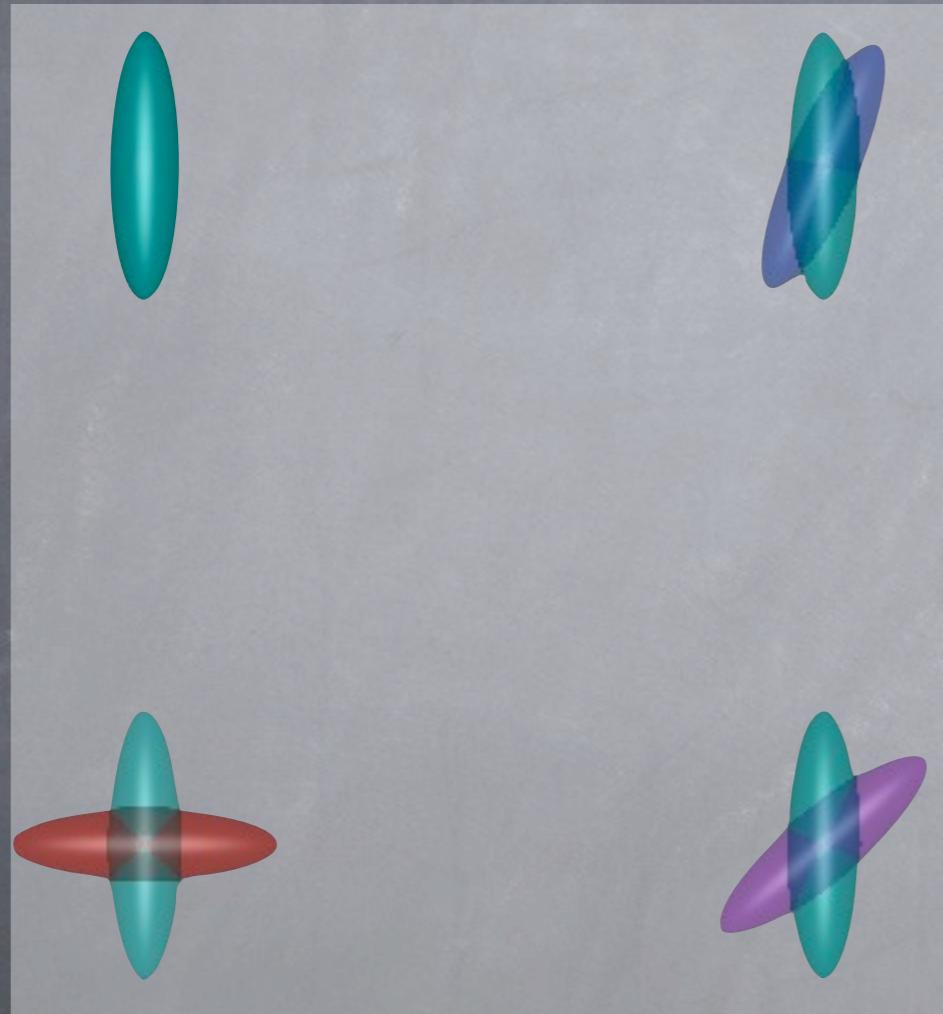
It can be efficiently solved
in an expectation-maximization scheme

Heuristics fail
a global approach is required

The problem is well known
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It can be efficiently solved
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A multi-fiber model represents fibers independently



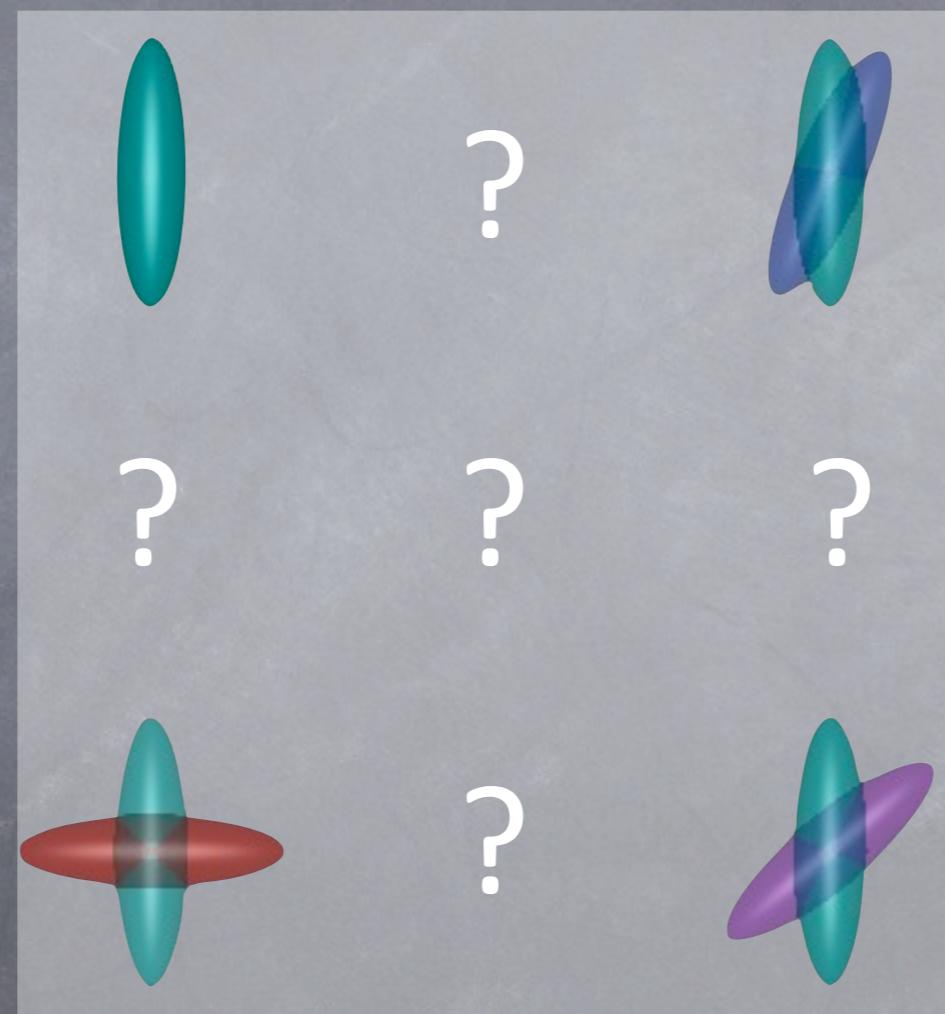
$$S = \sum_{i=1}^N f_i S_i$$

Single-fiber model
Volumetric fractions

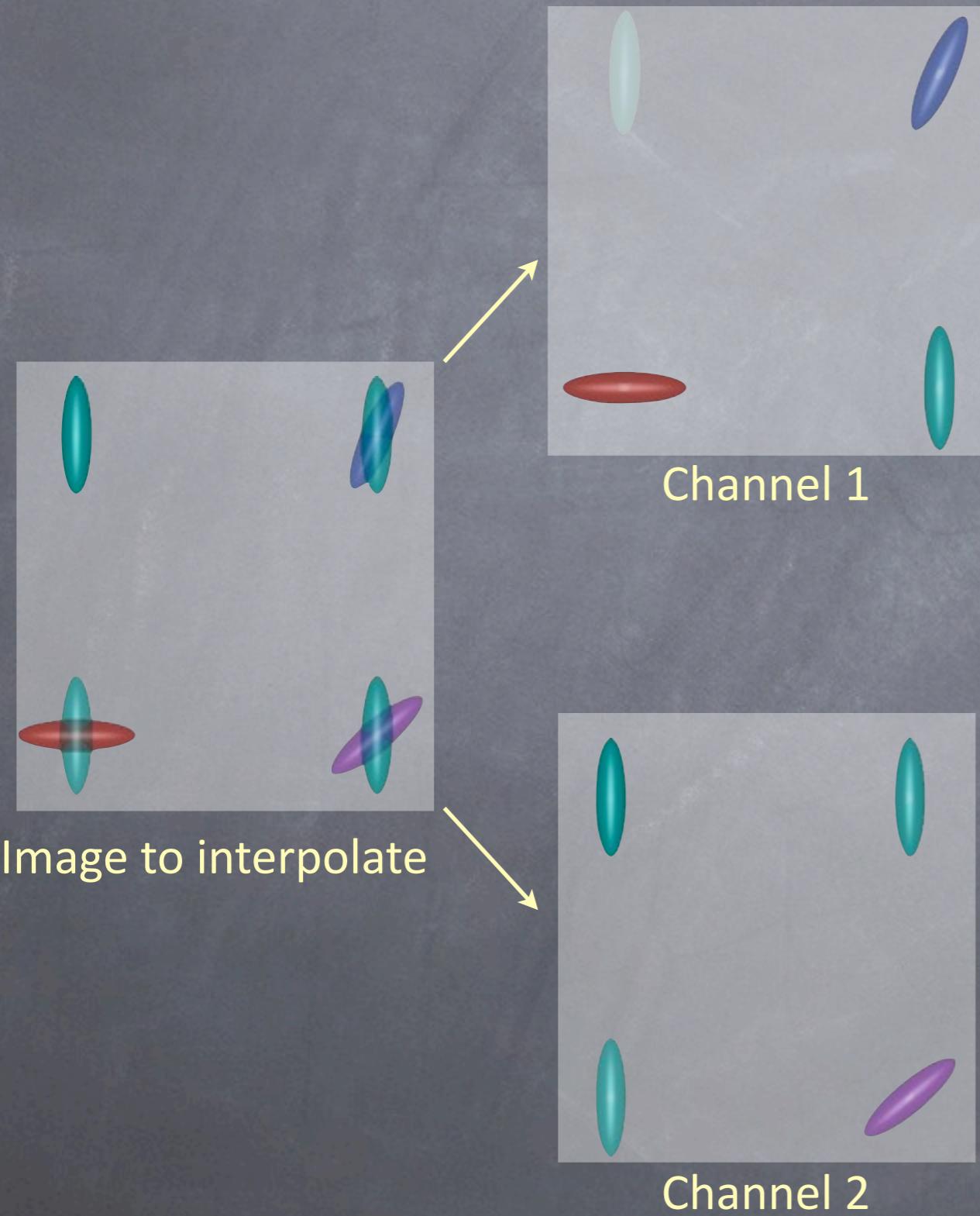
Particular case: the multi-tensor model

$$S = S_0 \sum_{i=1}^N f_i e^{-b\mathbf{g}^T D_i \mathbf{g}}$$

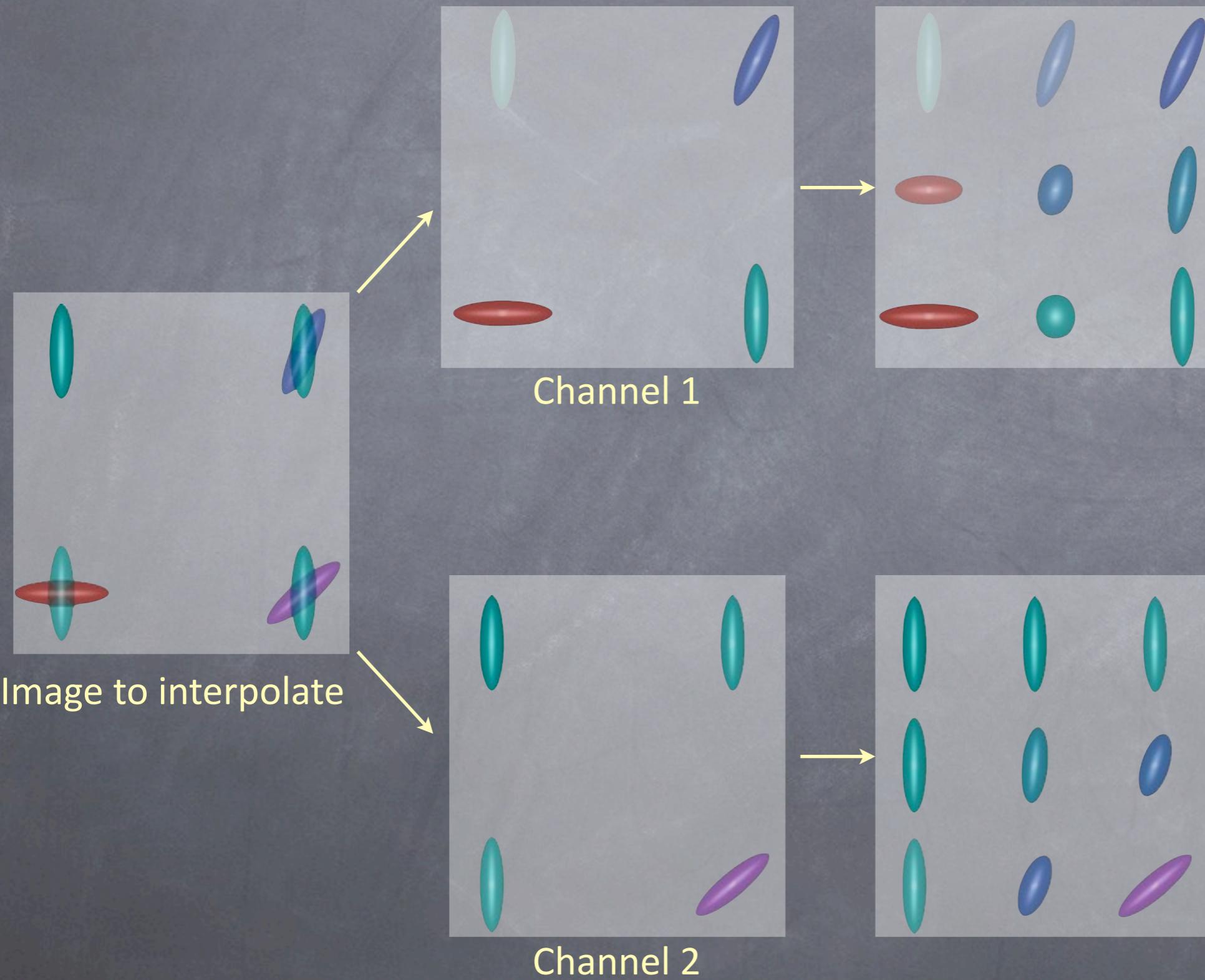
Interpolating a multi-fiber model is challenging



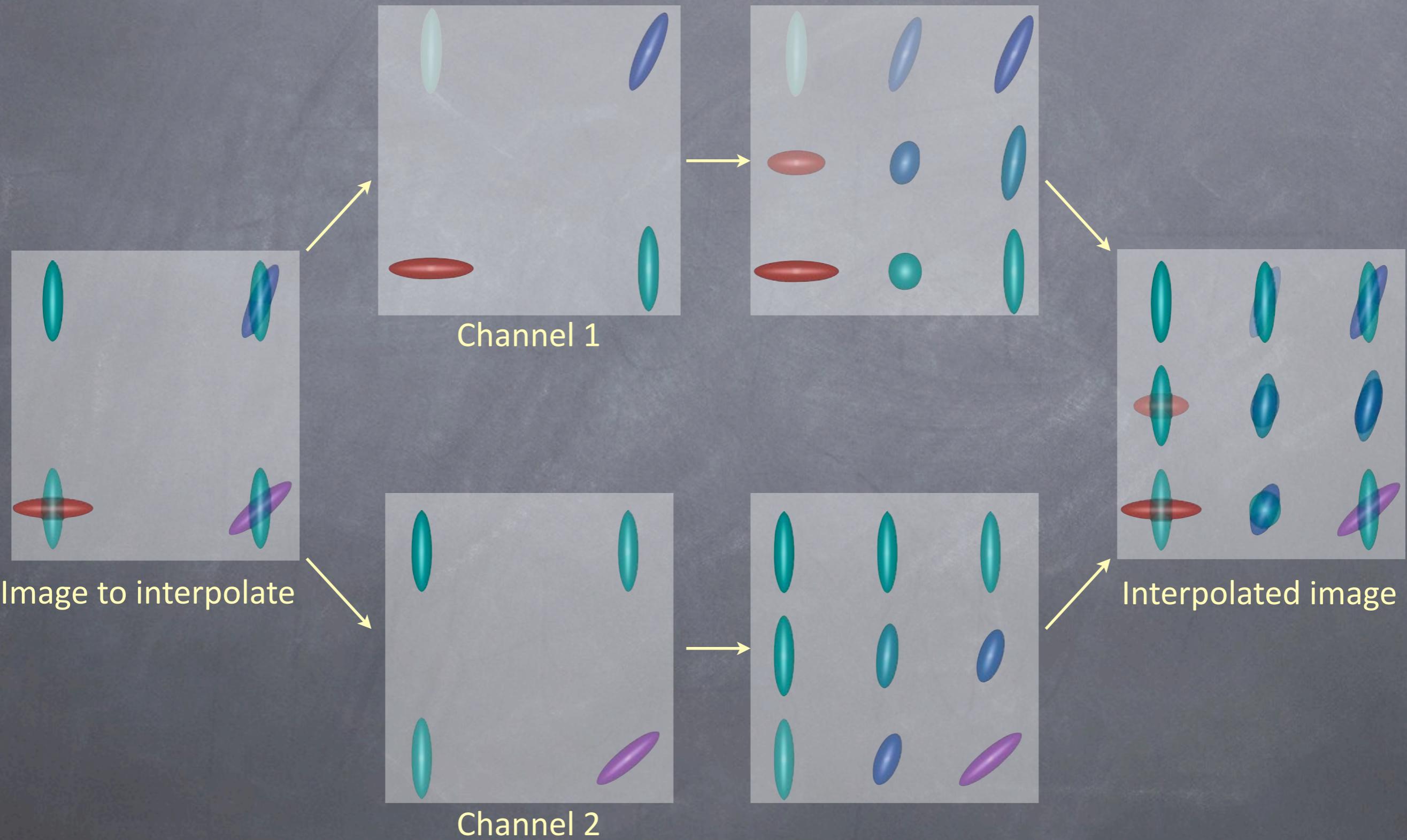
Channel based interpolation does not work



Channel based interpolation does not work



Channel based interpolation does not work



Channel based interpolation does not work

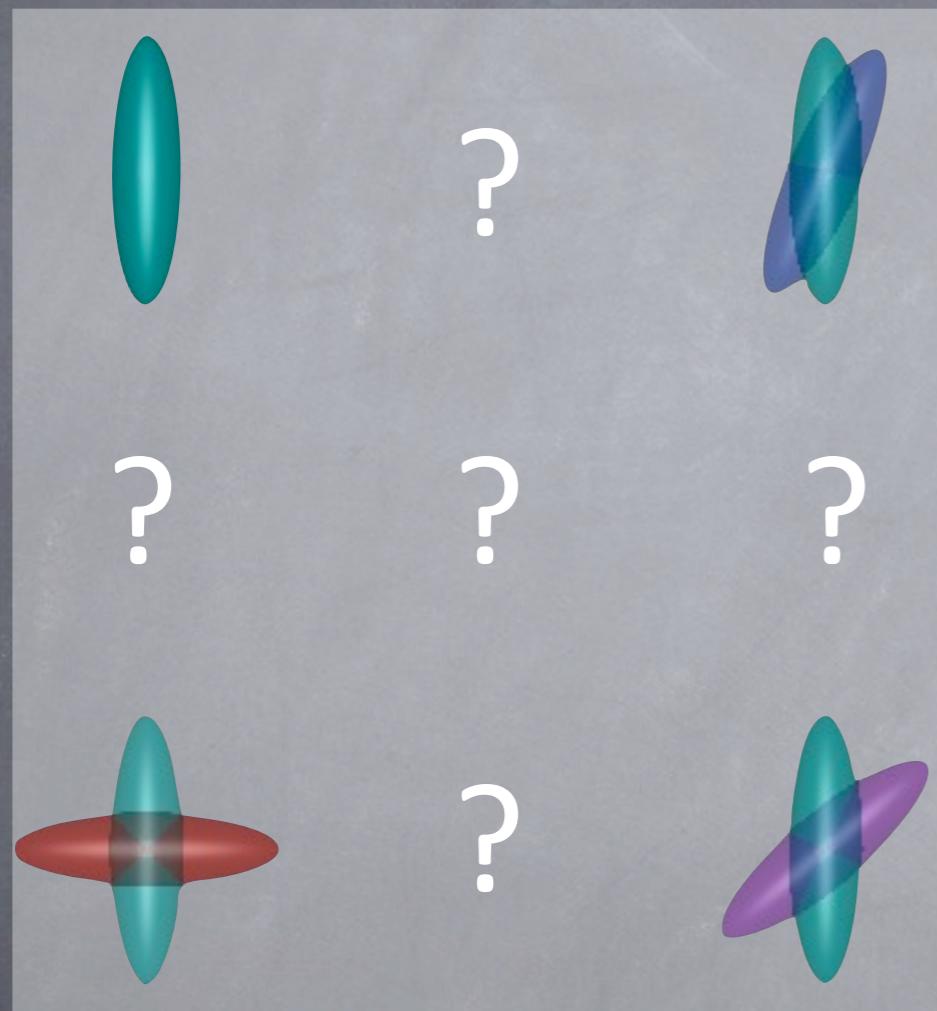
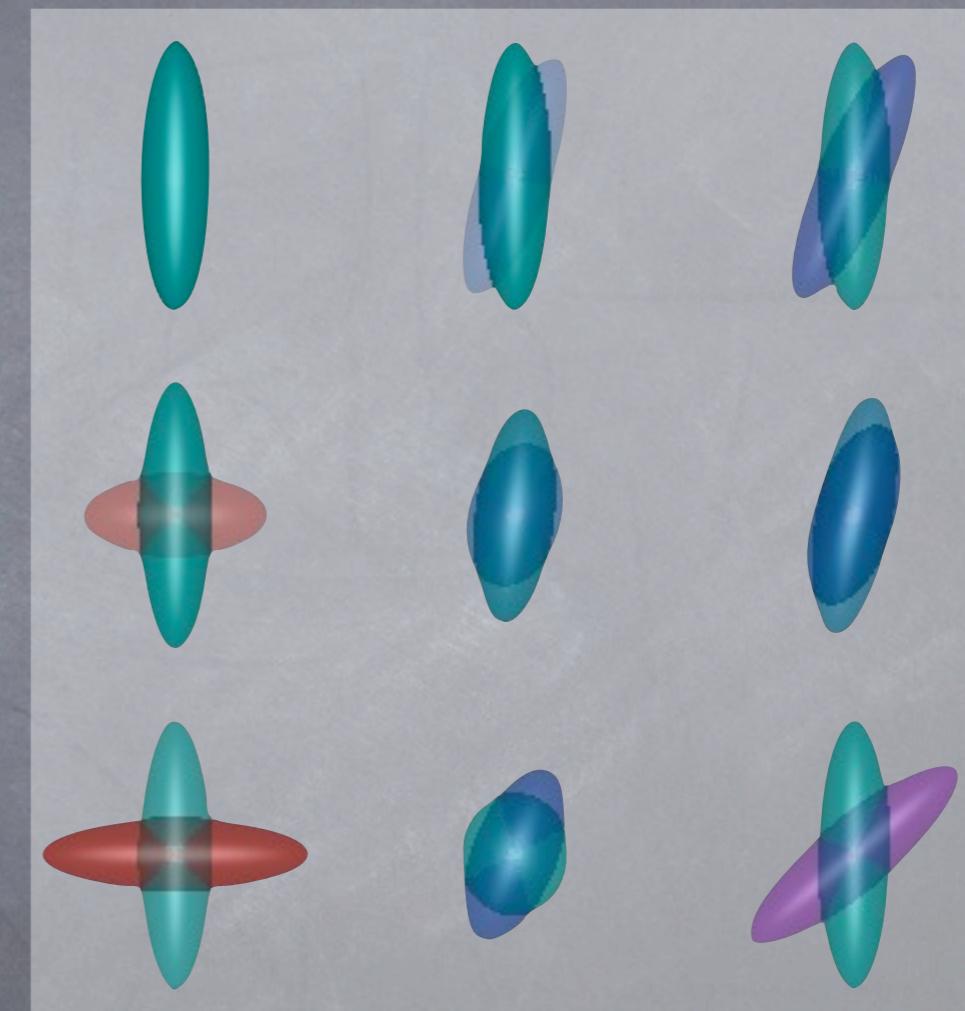


Image to interpolate



channel-based interpolation

Orientation-based interpolation does not work

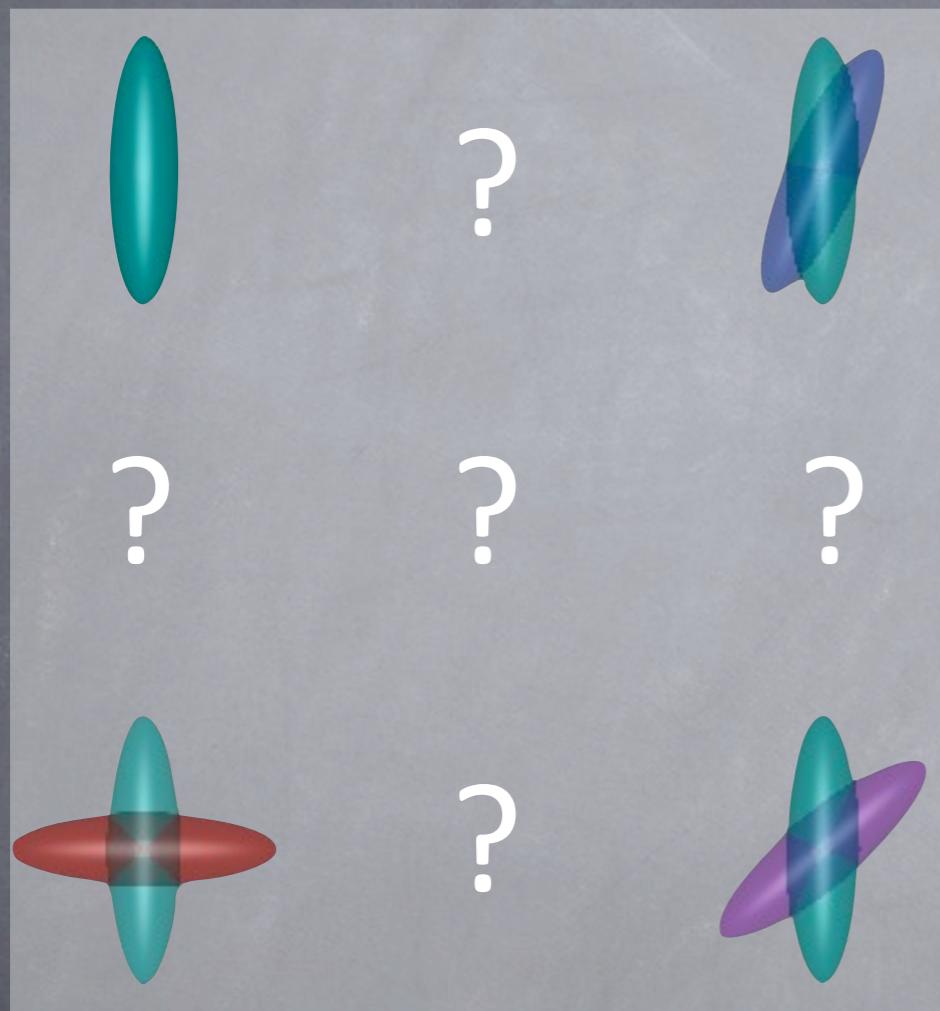
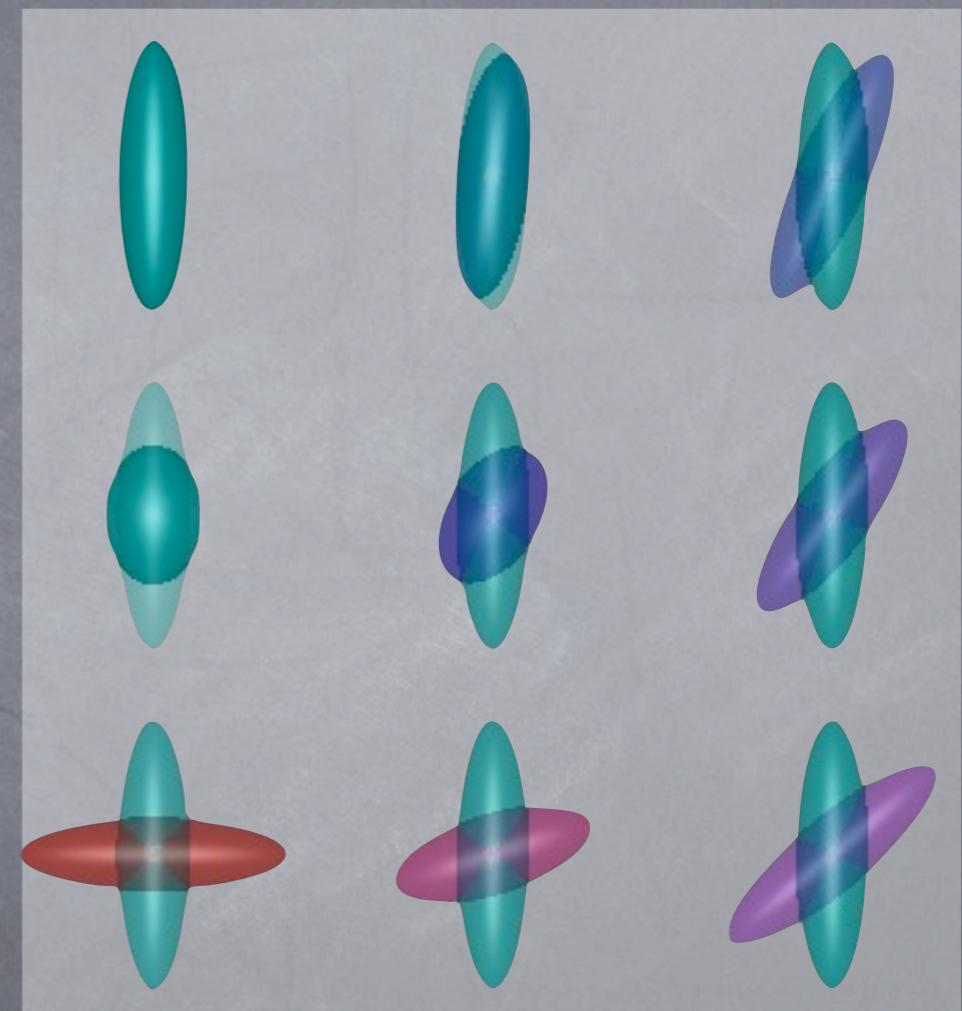


Image to interpolate



orientation-based interpolation

The multiple fibers are paired in two groups based on their orientation.

A global approach is required

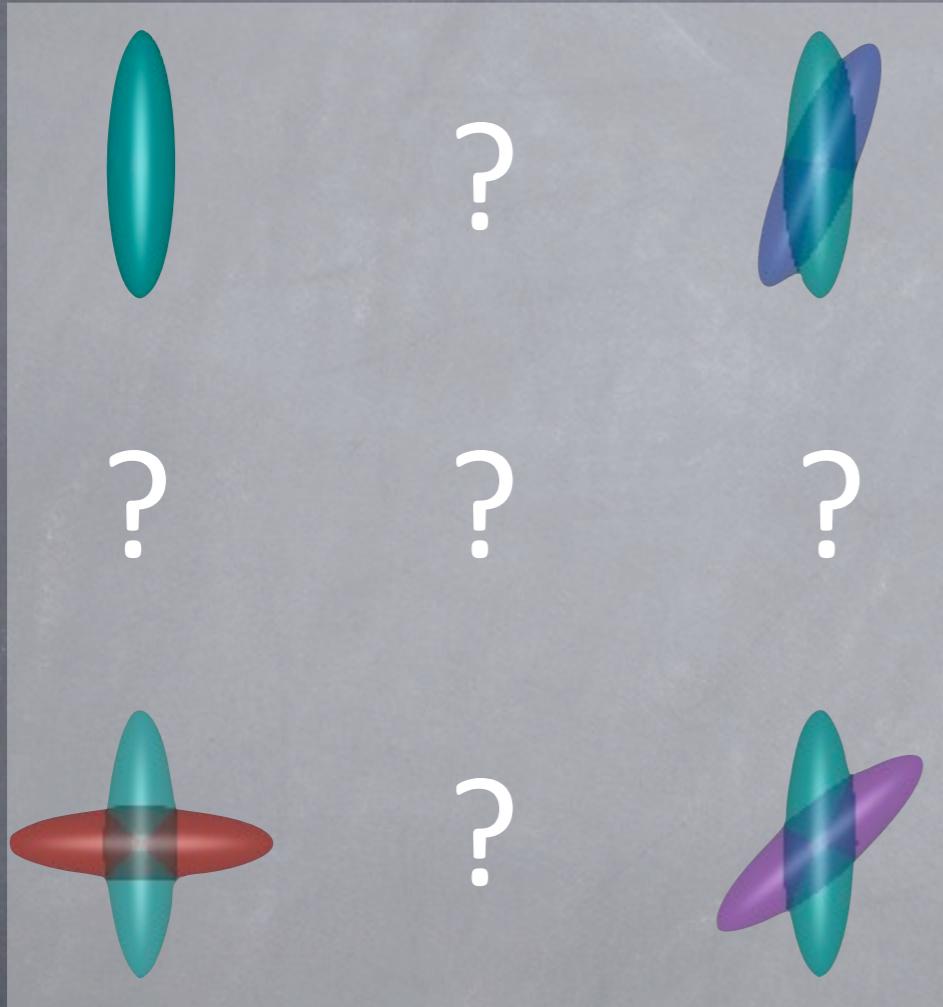
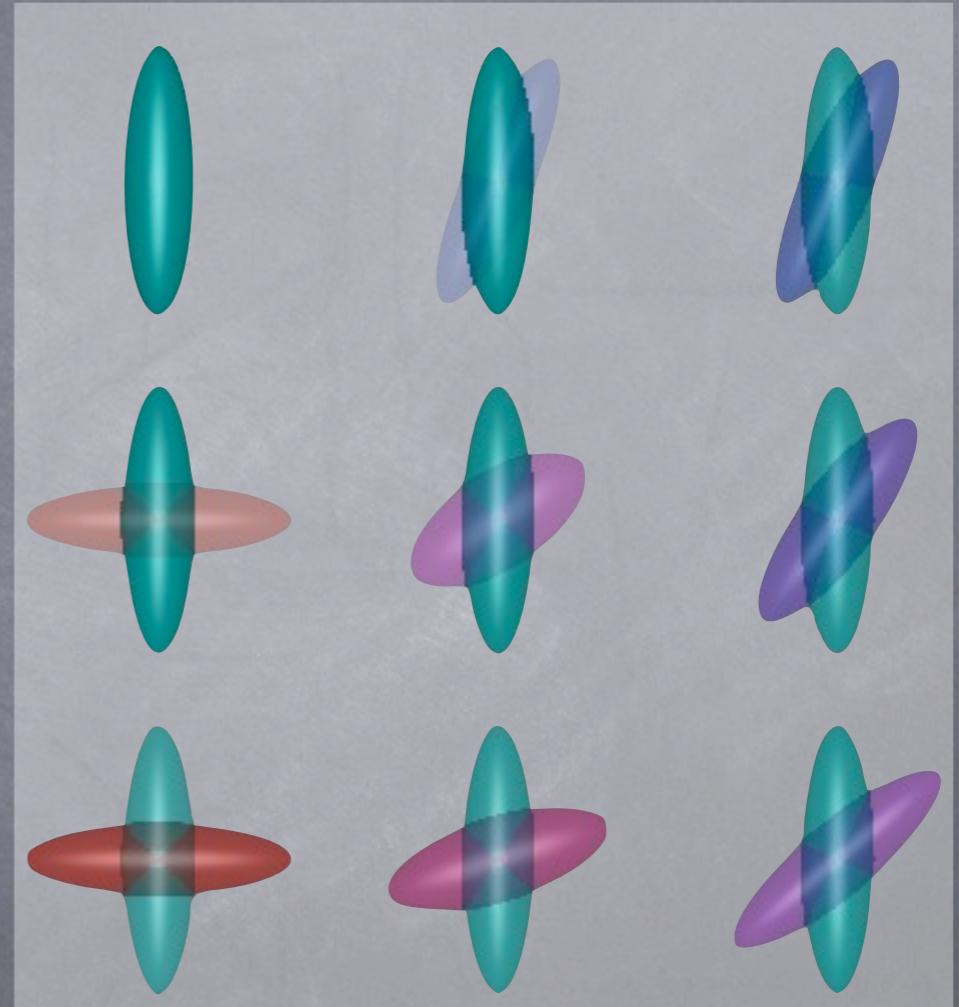
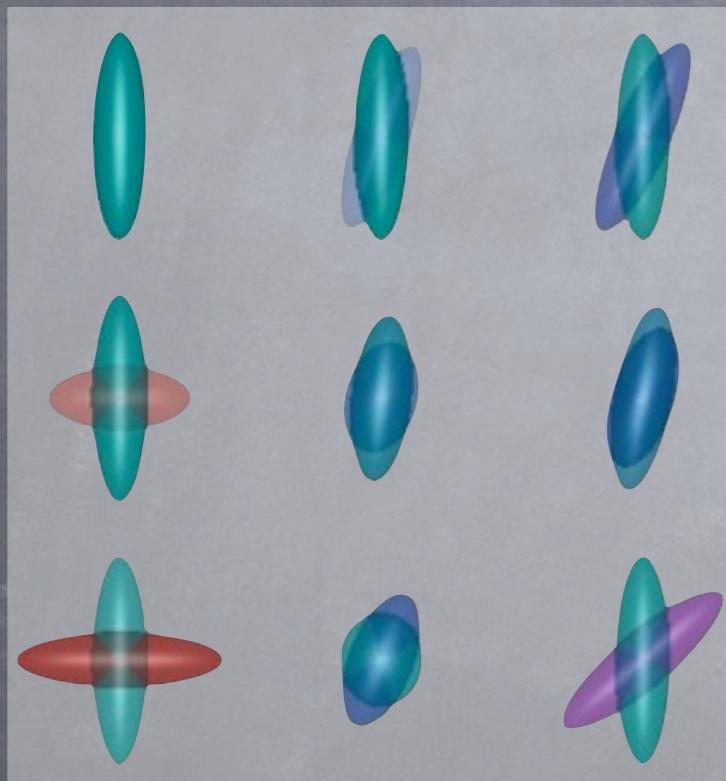


Image to interpolate

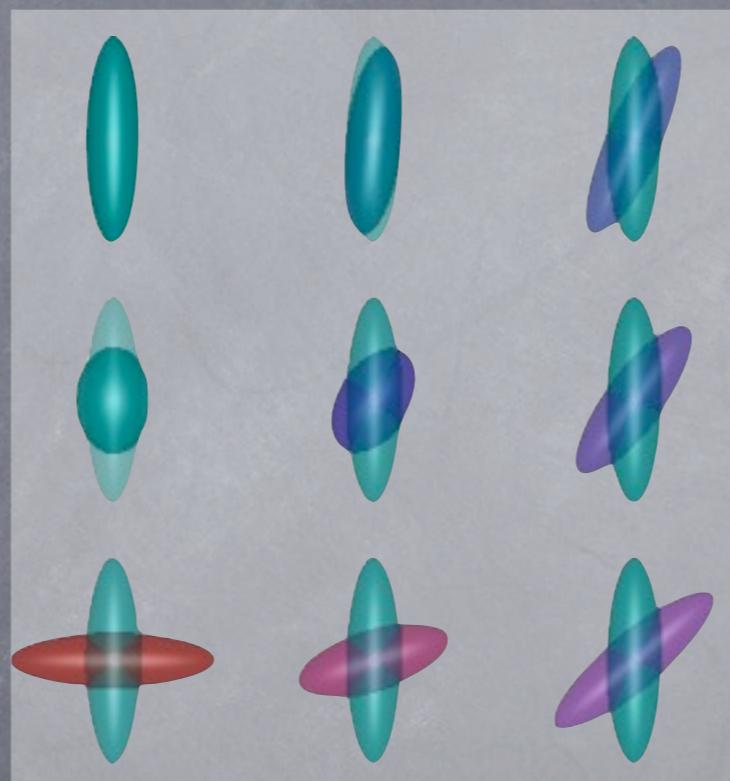


Proposed method

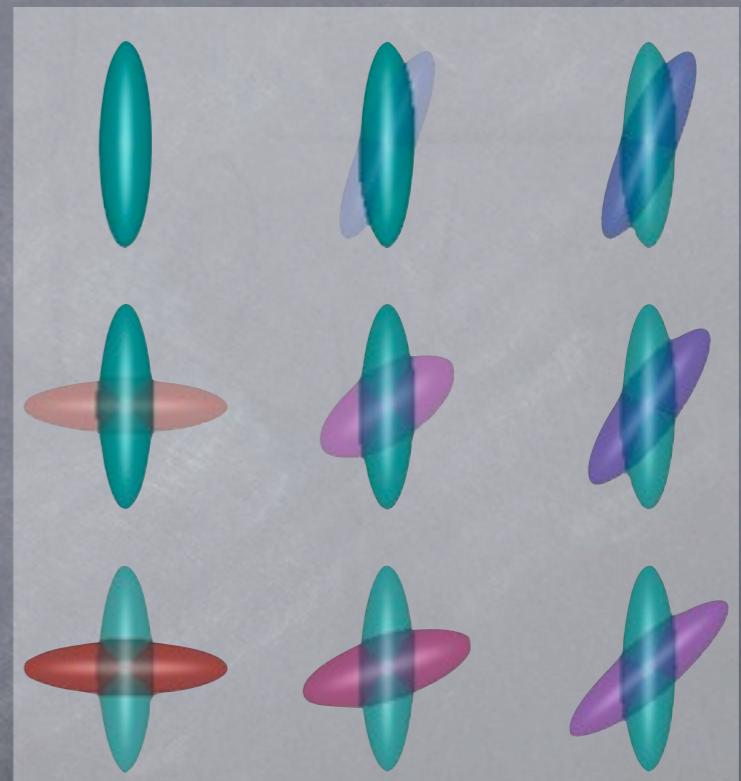
A global approach is required



channel-based



orientation-based



proposed method

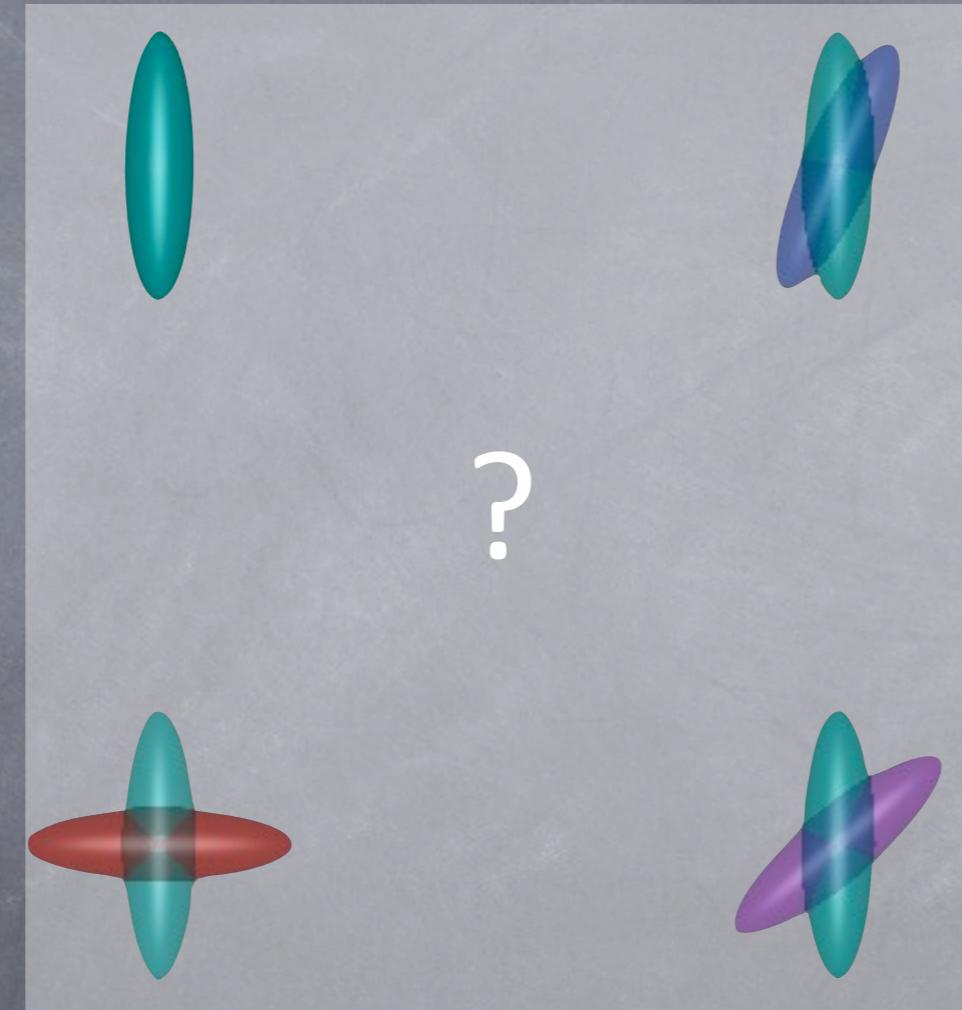
Heuristics fail
a global approach is required

The problem is well known
as a Gaussian Mixture Model Simplification

It can be efficiently solved
in an expectation-maximization scheme

Mathematically, we mix mixtures

$$p_1(\mathbf{x}) = \sum_i \alpha_{1,i} G_{1,i}$$



$$p_2(\mathbf{x}) = \sum_i \alpha_{2,i} G_{2,i}$$

$$p_3(\mathbf{x}) = \sum_i \alpha_{3,i} G_{3,i}$$

$$p_4(\mathbf{x}) = \sum_i \alpha_{4,i} G_{4,i}$$

Mathematically, we mix mixtures

$$p_1(\mathbf{x}) = \sum_i \alpha_{1,i} G_{1,i}$$

$$p_2(\mathbf{x}) = \sum_i \alpha_{2,i} G_{2,i}$$

Scalar interpolation
coefficients

$$p(\mathbf{x}) = \sum_{j=1}^4 a_j \sum_i \alpha_{j,i} G_{j,i}$$

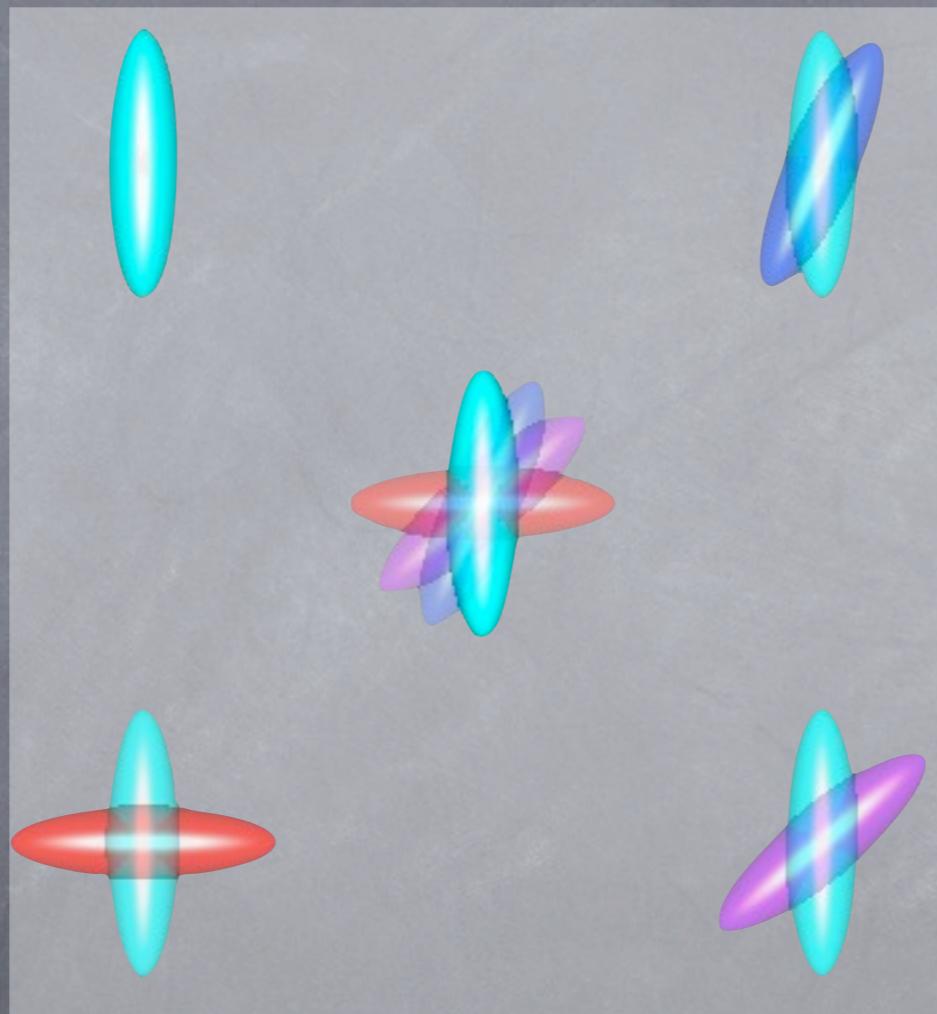
All mixtures

$$p_3(\mathbf{x}) = \sum_i \alpha_{3,i} G_{3,i}$$

$$p_4(\mathbf{x}) = \sum_i \alpha_{4,i} G_{4,i}$$

The result is a high order mixture model

$$p_1(\mathbf{x}) = \sum_i \alpha_{1,i} G_{1,i}$$



$$p_2(\mathbf{x}) = \sum_i \alpha_{2,i} G_{2,i}$$

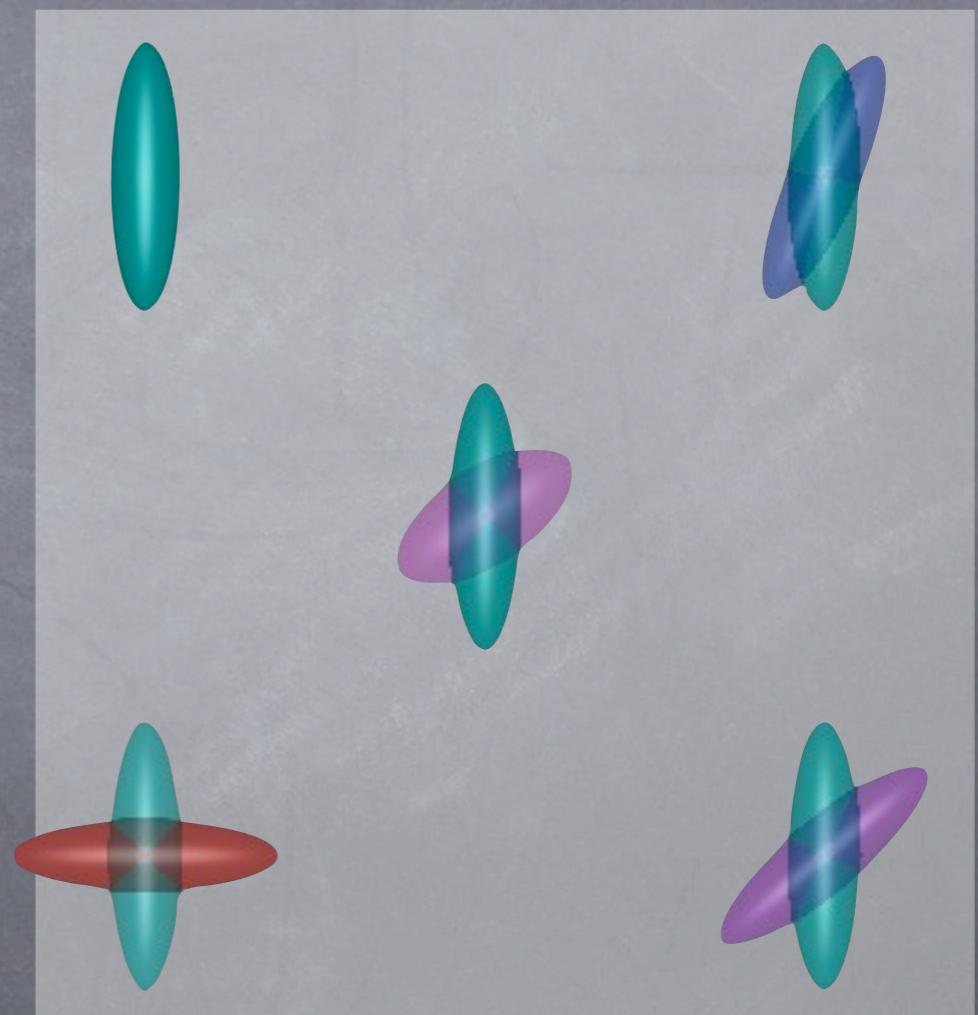
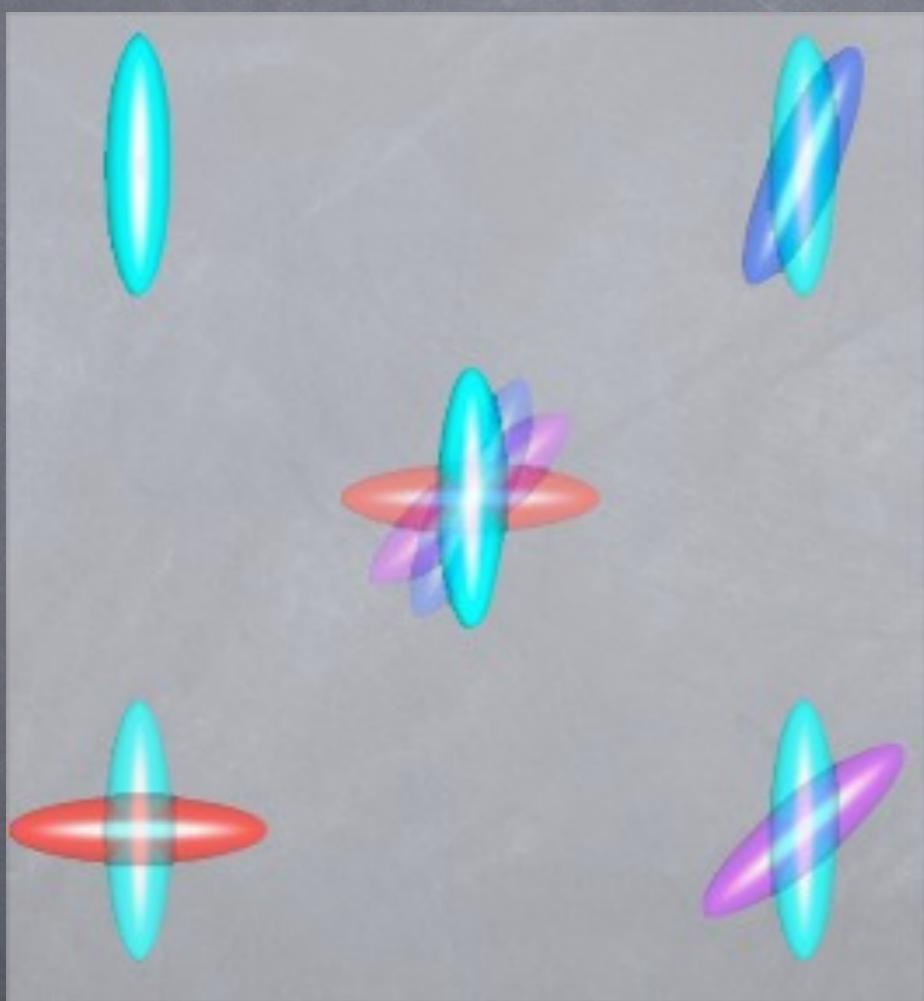
$$p_3(\mathbf{x}) = \sum_i \alpha_{3,i} G_{3,i}$$

$$p_4(\mathbf{x}) = \sum_i \alpha_{4,i} G_{4,i}$$

The complete mixture needs be simplified

$$p(\mathbf{x}) = \sum_{j=1}^4 a_j \sum_i \alpha_{j,i} G_{j,i}$$

$$p_S(\mathbf{x}) = \sum_i f_i R_i$$



The complete mixture needs to be simplified

$$p(\mathbf{x}) = \sum_{i=1}^N \alpha_i G_i \quad \xrightarrow{N > K} \quad p_S(\mathbf{x}) = \sum_{j=1}^K f_j R_j$$

The complete mixture needs to be simplified

$$p(\mathbf{x}) = \sum_{i=1}^N \alpha_i G_i \quad \xrightarrow{N > K} \quad p_S(\mathbf{x}) = \sum_{j=1}^K f_j R_j$$

$$p_S^* = \arg \min_{p_S} D(p, p_S)$$

Minimize some discrepancy measure

The complete mixture needs to be simplified

$$p(\mathbf{x}) = \sum_{i=1}^N \alpha_i G_i \quad \xrightarrow{N > K} \quad p_S(\mathbf{x}) = \sum_{j=1}^K f_j R_j$$

$$p_S^* = \arg \min_{p_S} D(p, p_S)$$

Minimize some discrepancy measure

$$D(p, p_S) = \sum_j \sum_{i:\pi_i=j} \alpha_i D(G_i || R_j) \quad \text{Cumulative Differential Relative Entropy}$$



Latent variables indicating the grouping of components

Heuristics fail
a global approach is required

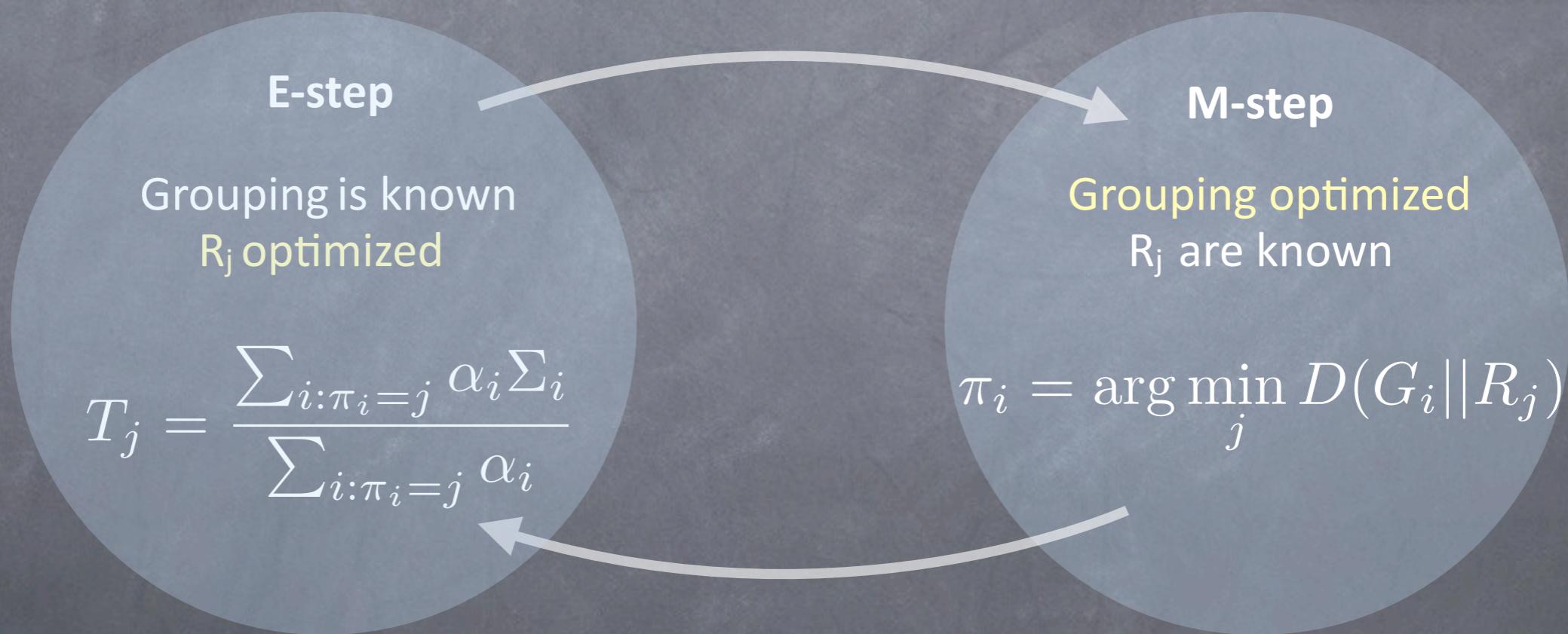
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Simplifying Gaussian mixtures is guaranteed to converge

Computation time:
100μs/voxel
(< 10 iterations)

$$\mathbf{R}^* = \arg \min_{\mathbf{R}} \sum_j \sum_{i:\pi_i=j} \alpha_i D(G_i \parallel R_j)$$



Both steps can be carried out in closed form (Davis and Dihillion, 2007)

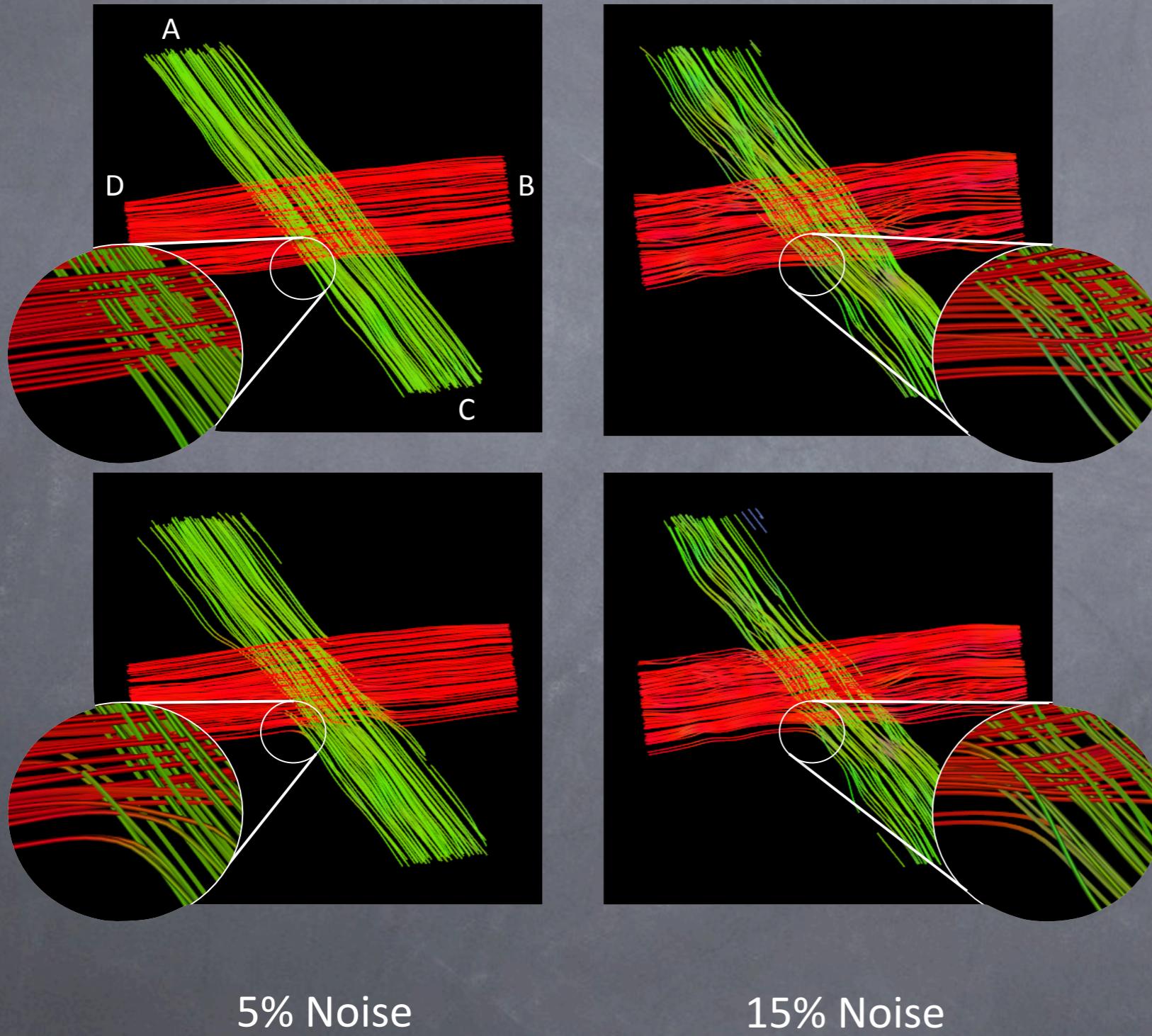
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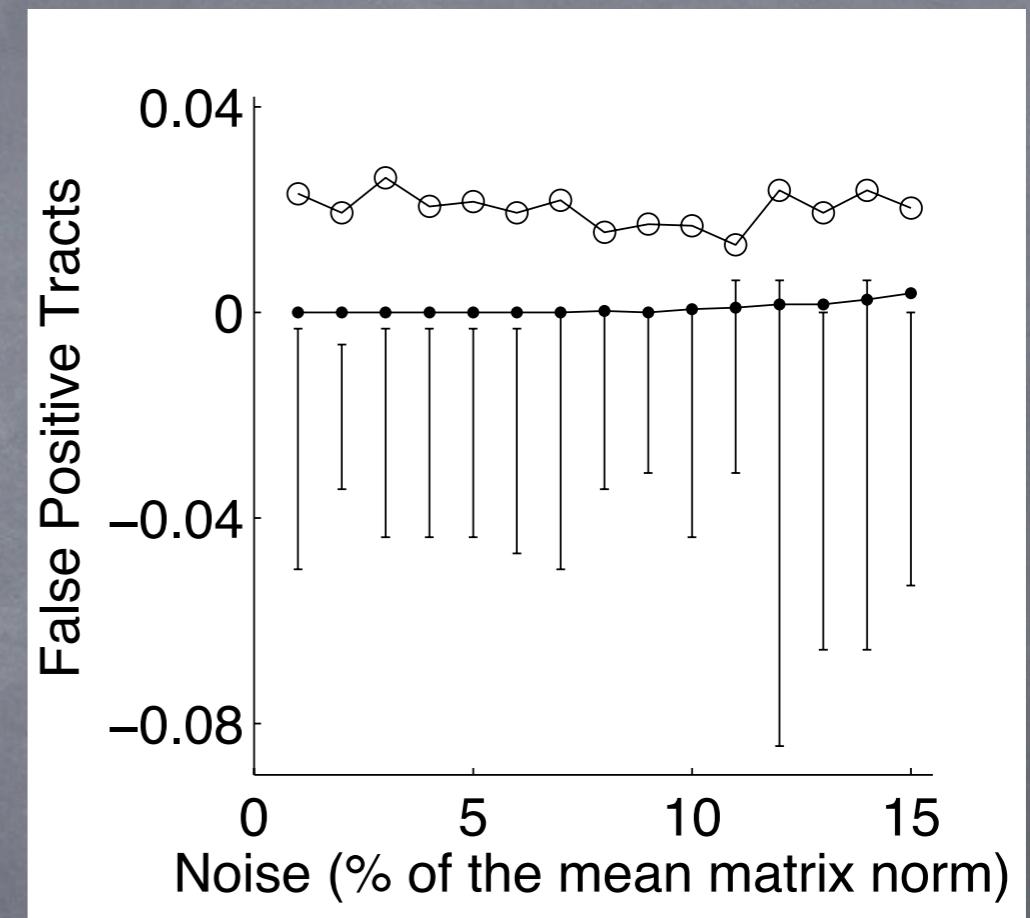
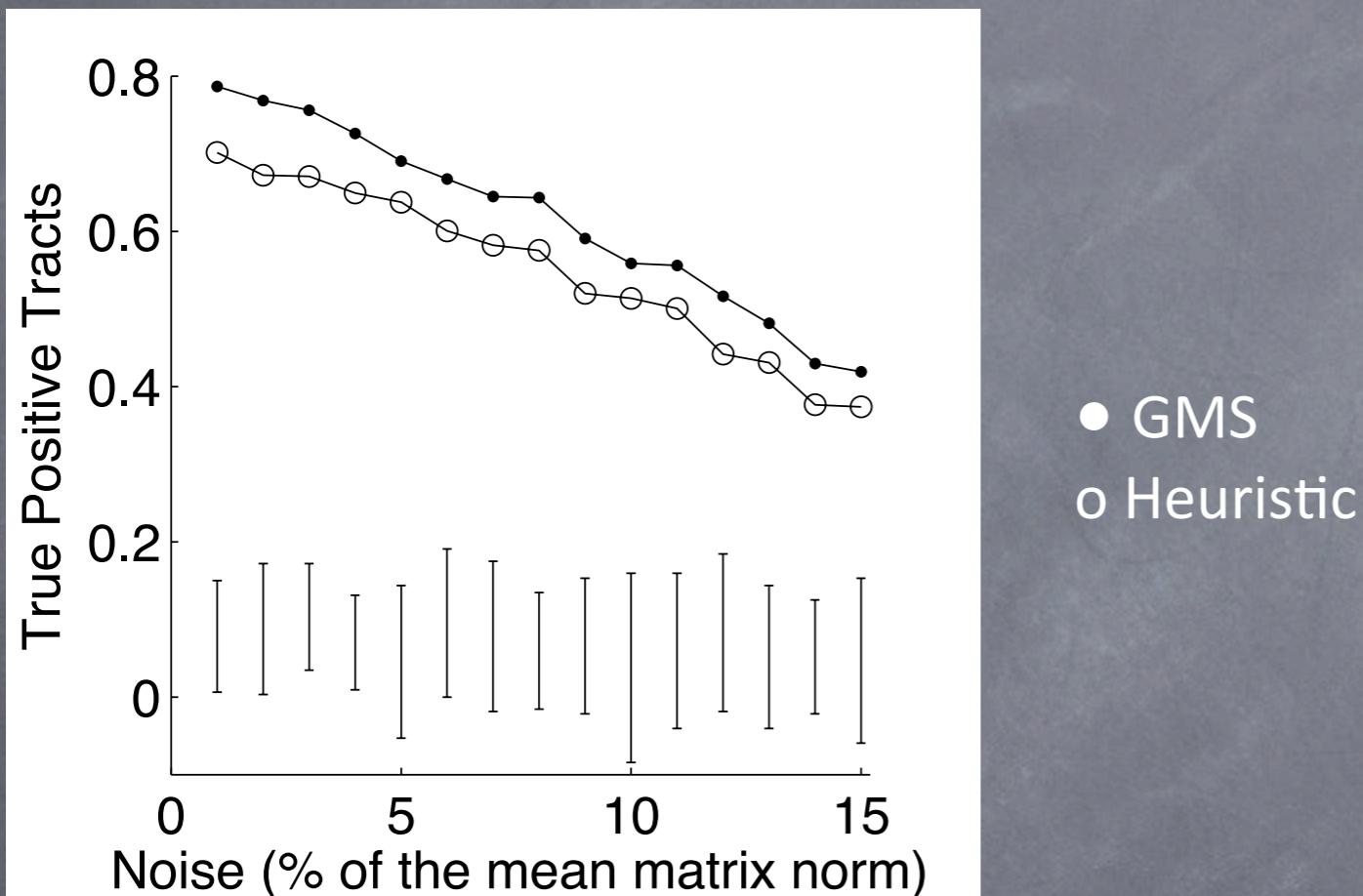
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How useful is it in practice?

Using GMS prevents confounding white matter fascicles



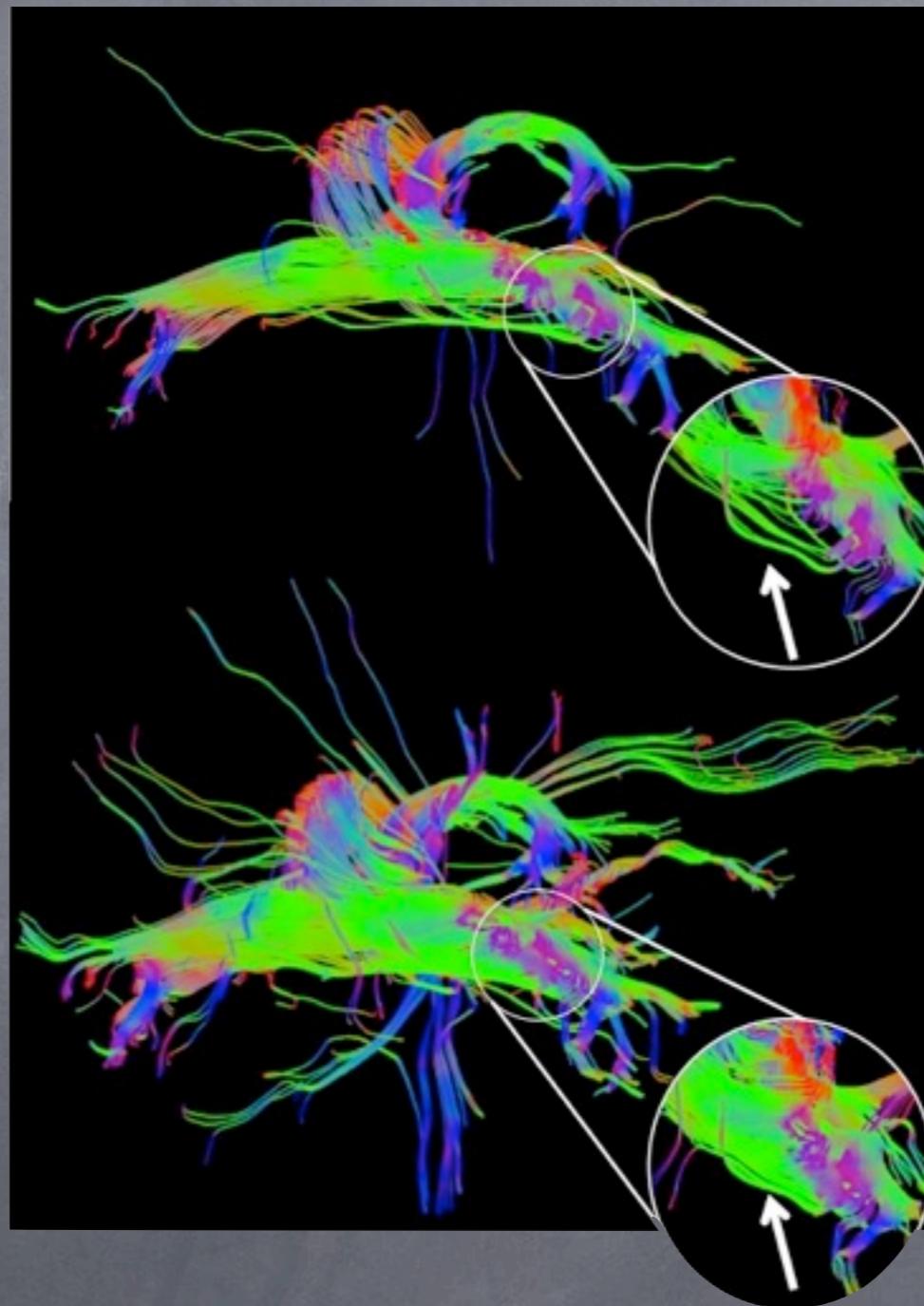
Using GMS prevents confounding white matter fascicles



GMS consistently

- increases the true positive rate
- decreases the false positive rate

This advantage may have clinical implications

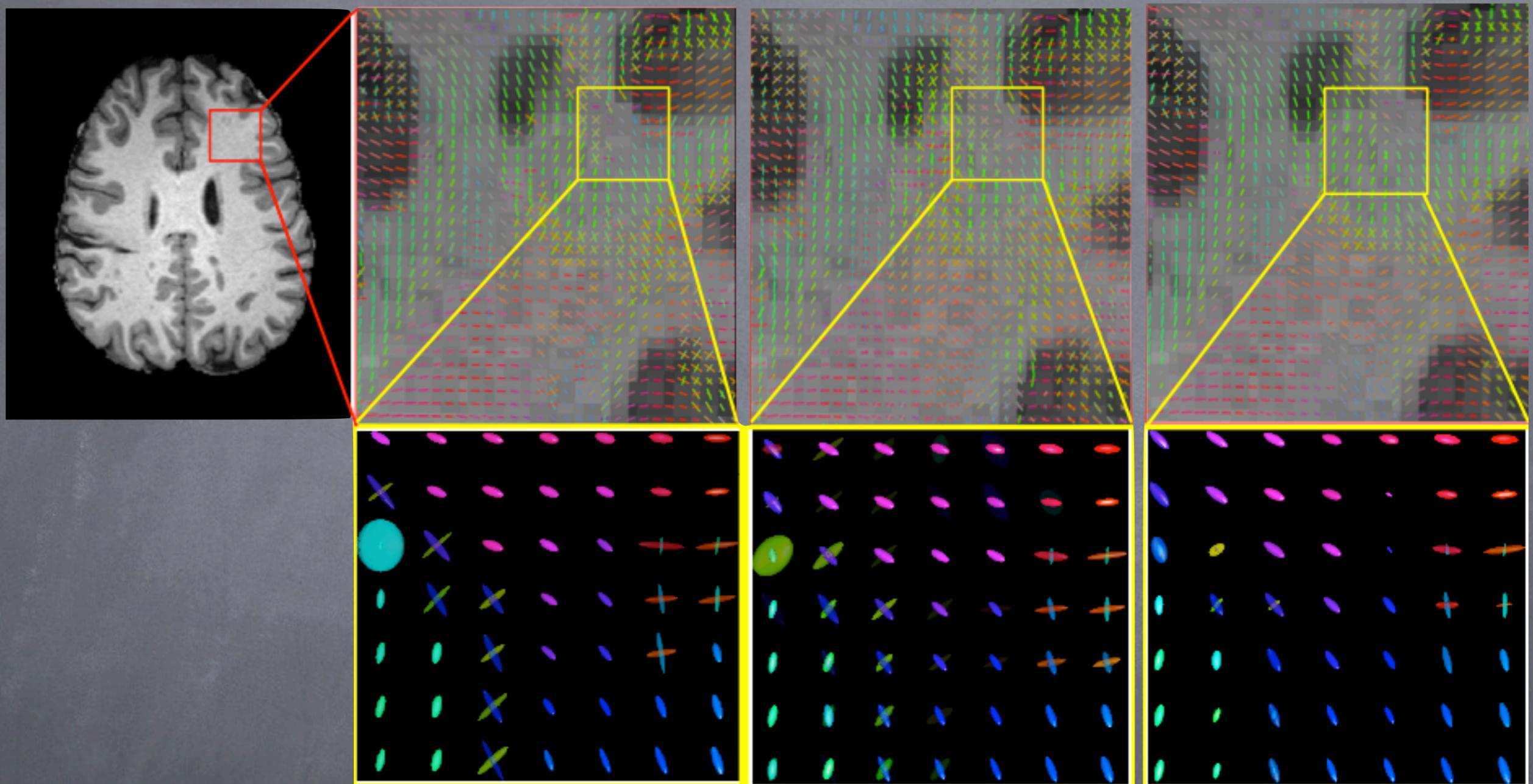


GMS (our method)

Heuristic (orientation-based)

The Meyer's loop is more properly identified
There are fewer spurious tracts

In spatial normalization, GMS preserves more information



I_0

$T^{-1} \circ (T \circ I_0)$

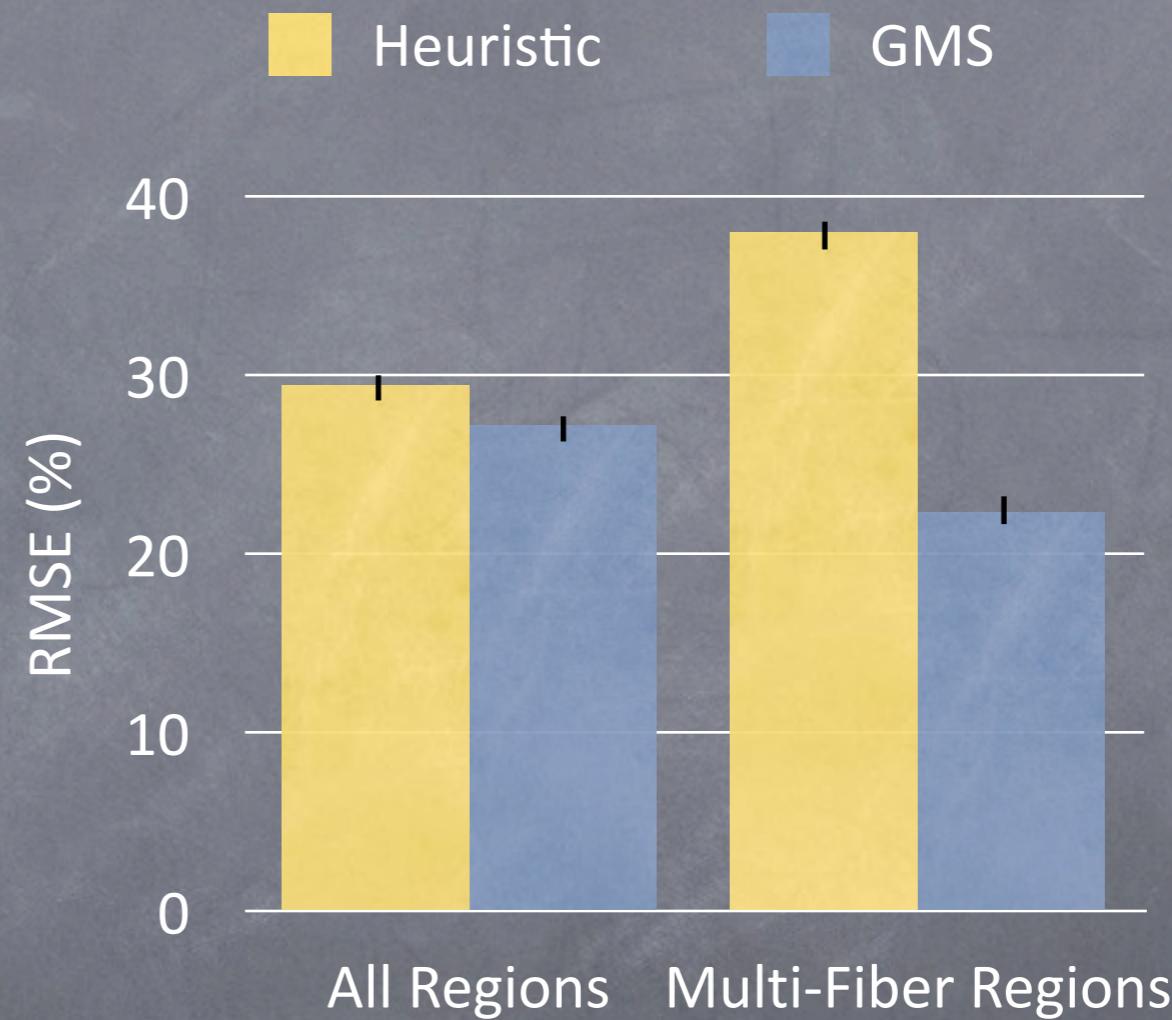
GMS

$T^{-1} \circ (T \circ I_0)$

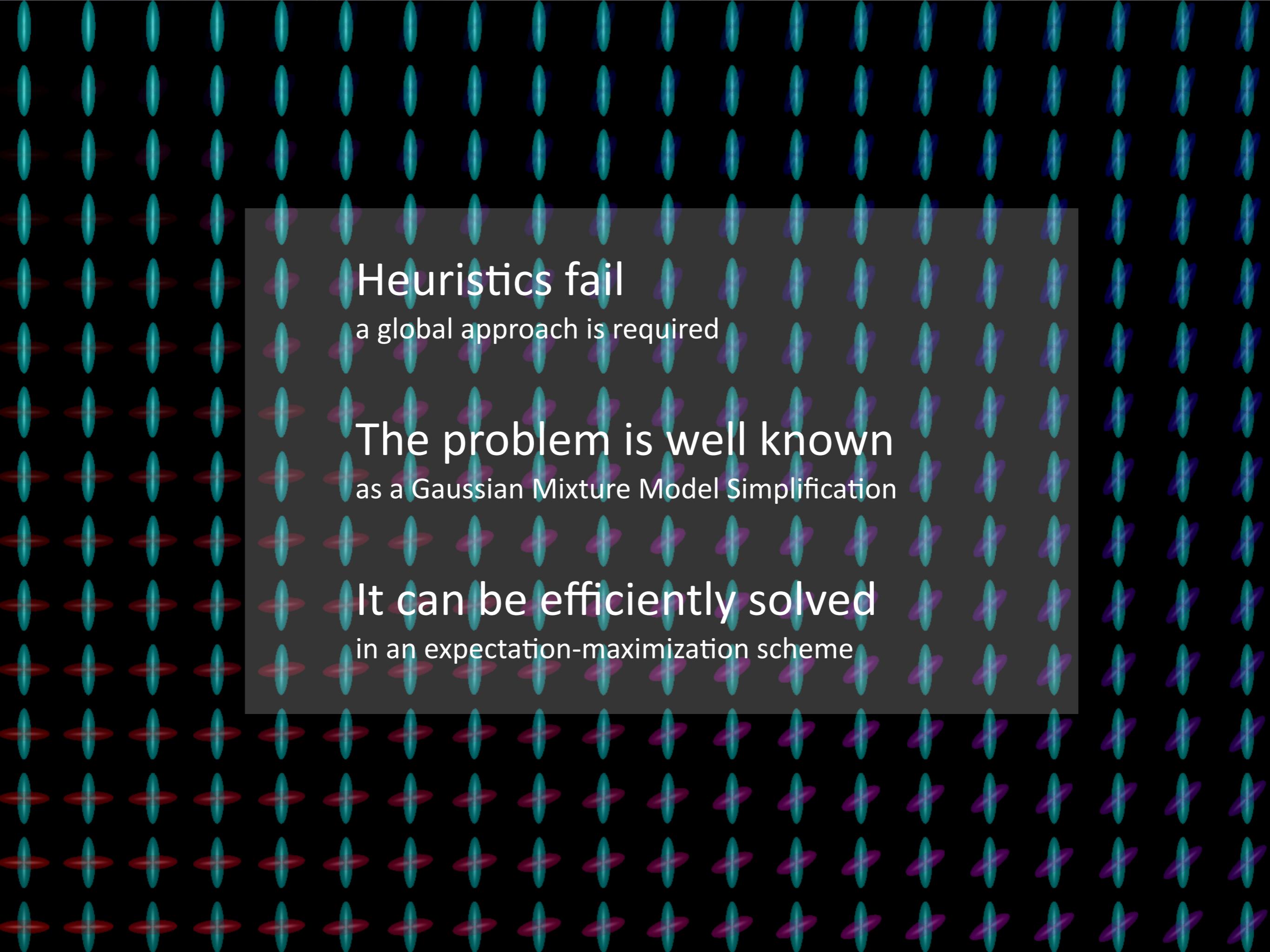
Heuristic

In spatial normalization, GMS introduces less error

- 10 multi-fiber DTI
- $1.8 \times 1.8 \times 2.4 \text{ mm}^3$
- CUSP-45 Acquisition (ISBI 2011)



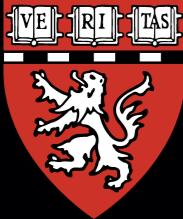
GMS reduces interpolation errors in all regions
and more strikingly in multi-fiber regions



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Thank you

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Benoît Scherrer,
Christopher Benjamin,
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Benoît Macq,
Simon K. Warfield**