Fast HARDI Uncertainty Quantification and Visualization with Spherical Sampling

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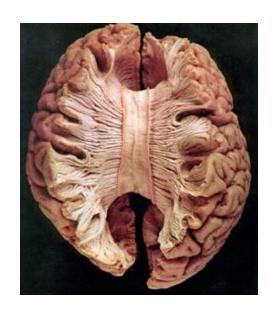




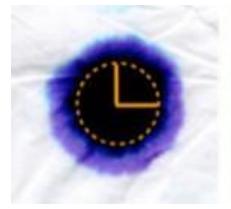


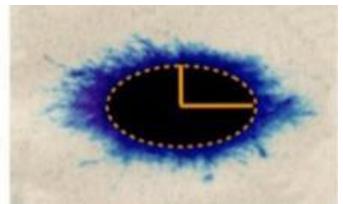


Introduction



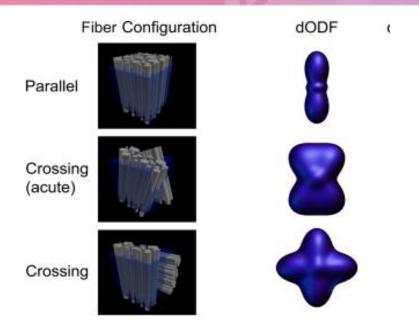




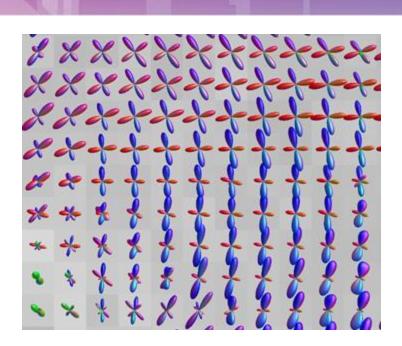


Drops of dye in Kleenex and newspaper show the diffusivity through Brownian Motion

Diffusion MRI



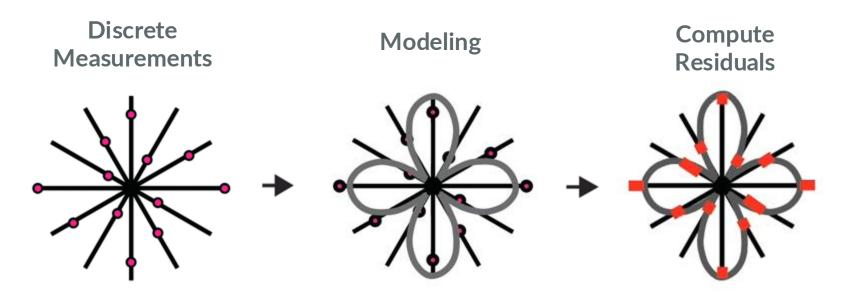
Diffusion MRI measures the diffusivity function per voxel



Directly visualized as glyphs

[Sotiropoulos & Zalesky 2017]

Measuring and Modeling

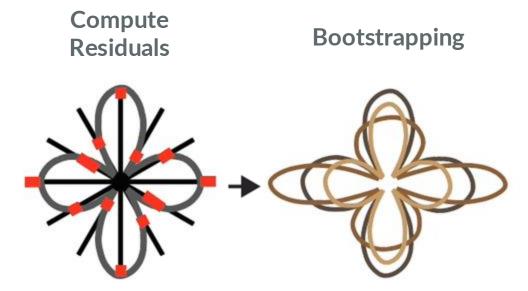


Diffusion magnitude along gradient vectors

Estimate function from samples

Red bars denote residual magnitude

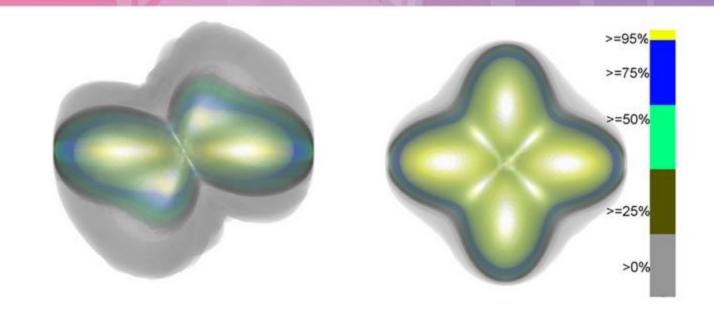
Bootstrapping



Red bars denote residual magnitude

- Bootstrapping simulates multiple acquisitions
- Each simulation randomly perturbed by residuals
- The ensemble encodes uncertainty

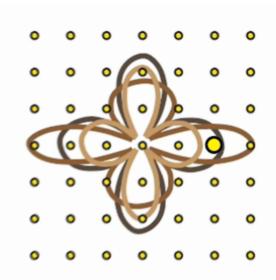
Shape Inclusion Probability



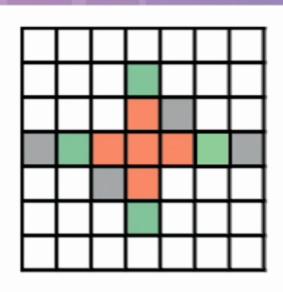
- Directly shows the variation of the bootstrap ensemble
- High cost to compute, store, and render

Volume Sampling

- Samples SIP function on structured grid
- Entire ensemble is evaluated to compute one voxel



Yellow dots denote sampling points



Volume rendered with transfer function

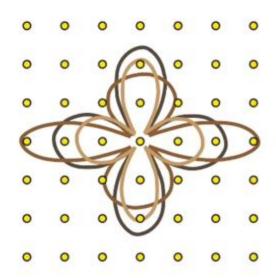
Our Contributions



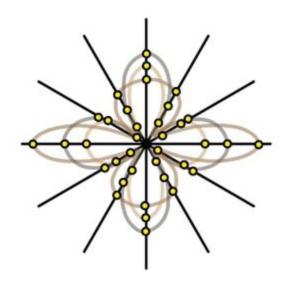
Volume Sampling vs Spherical Sampling

- Evaluates SIP values directly at boundaries
- Needs much less samples

Volume Sampling

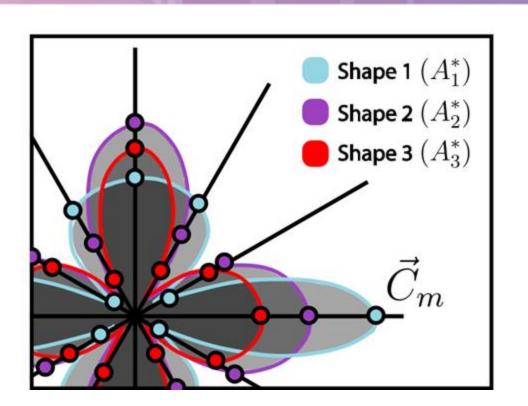


Spherical Sampling



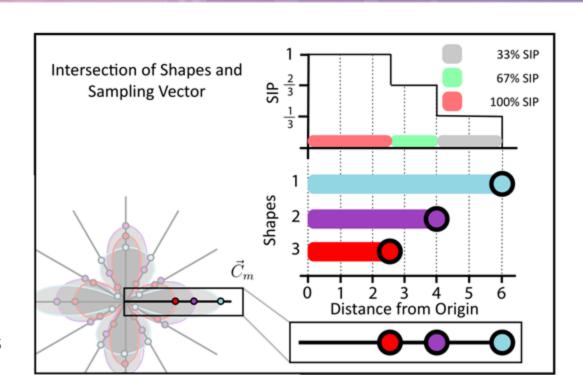
Spherical Sampling

- Sampling vectors are roughly distributed evenly
- Each shape intersects a vector exactly twice
- Each shape is evaluated once per sampling vector



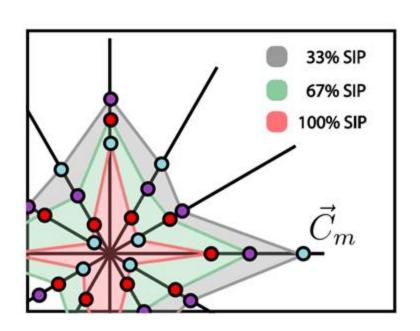
Spherical Sampling

- SIP function changes at vertices
- Sorting each vector's samples gives isolevels in order
- Isosurfaces easily extracted from array access (Ex. 2nd element of length 3 array corresponds to 67% SIP)

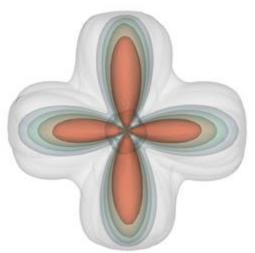


SIP Isosurface Extraction

- SIP isosurfaces are derived from the shapes' vertices
- All isolevels are computed simultaneously

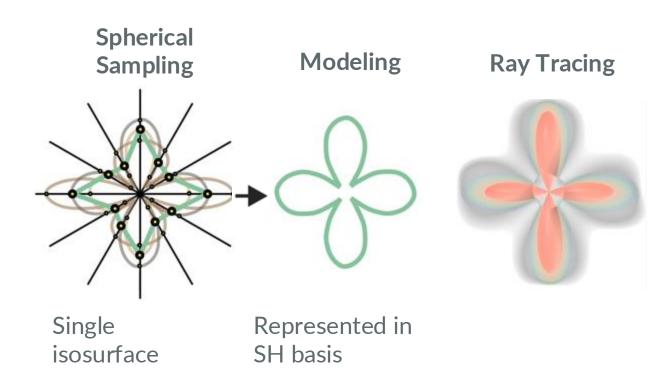






SIP Modeling

- Each isosurface is modeled
- Modeling requires much less samples
- Reduces memory
- Enables ray tracing



Ray Tracing

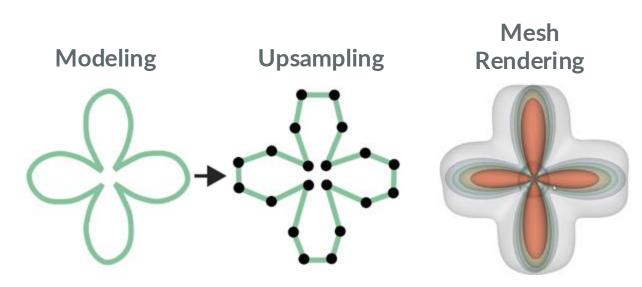
Low memory cost

Multiple volumetric regions



SIP Upsampling

- Modeling allows efficient upsampling
- Quickly creates high resolution meshes
- Triangle meshes are easy to visualize



Black dots are vertices of upsampled isosurface

Evaluation

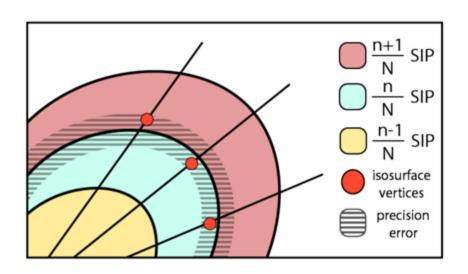
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Speed

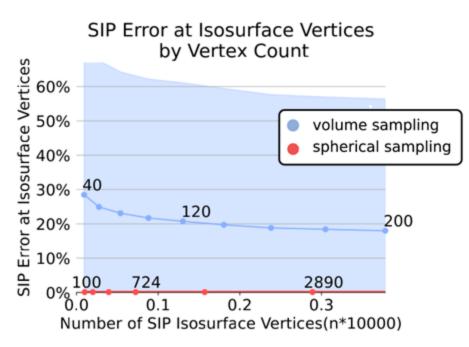
- Up to 8164X faster
- Previous method can take days to months
- New methods are usable in clinically feasible times

SIP Interval Modeling	Single Voxel	Data Slice (81,106)
Volume (R=100³)	2 min	12 days
Volume (R=200³)	12 min	2 months
Spherical (M = 500)	114 ms	12 min
Spherical (M = 1,000)	215 ms	24 min
Upsampling (SH = 4)	31 ms	5 min
Upsampling (SH = 8)	85 ms	13 min

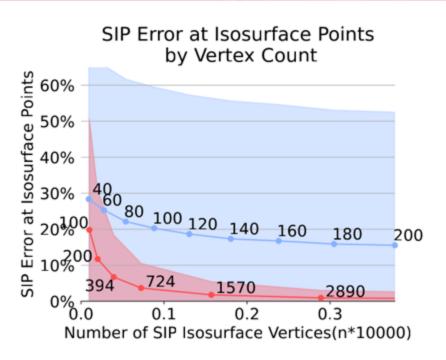
Accuracy

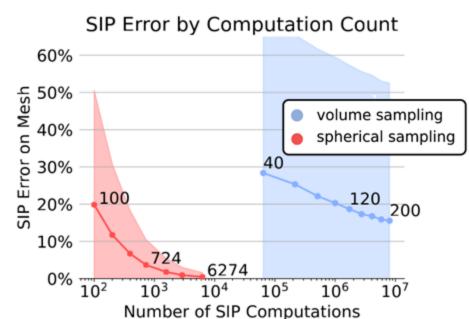


- Error is minimal at SIP boundaries
- Good choice for modeling



Accuracy





 Error of the mesh converges as resolution increases with respect to resolution (left) and compute time

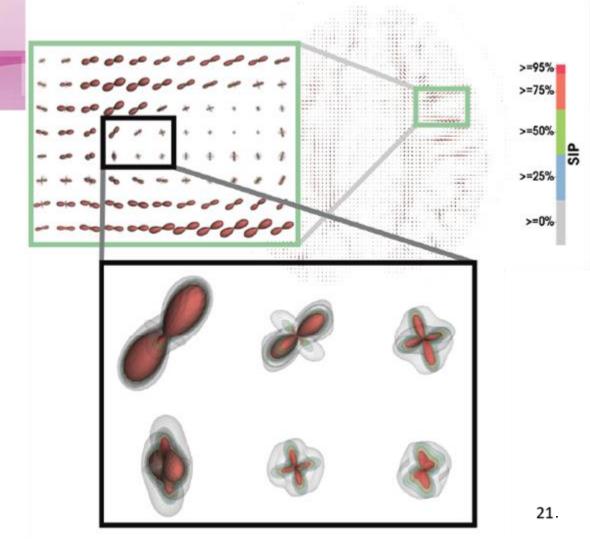
Memory

- Up to 37282x memory reduction
- Polygon mesh and model representation may alleviate storage constraints

Technique	Single Voxel	Data Slice (81,106)
Volume (R=100 ³)	4 MB	34 GB
Volume (R=200³)	32 MB	268 GB
Polygon Mesh (M=500)	10 KB	84 MB
Polygon Mesh (M=1,000)	20 KB	168 MB
Polygon Mesh (M=10,000)	200 KB	2 GB
Model (SH=4)	300 B	3 MB
Model (SH=8)	900 B	8 MB

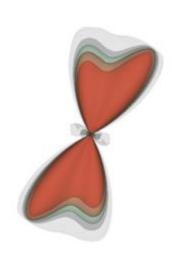
Brain Data

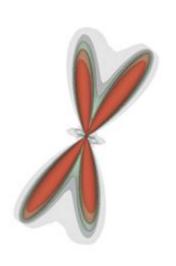
- The example is a slice (81x106) of the Stanford HARDI data set
- Computed in 13 minutes with upsampling
- Clinically feasible time



SH Degree & Discernibility







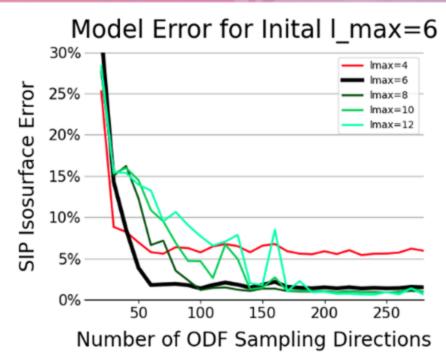
(a)
$$l_{max} = 4$$

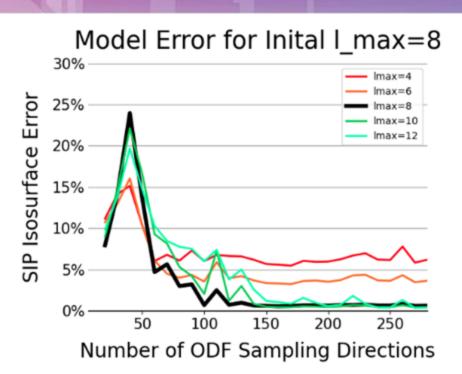
(b)
$$l_{max} = 8$$

(c)
$$l_{max} = 12$$

• Higher SH complexity of initial model discerns smaller angles

SIP Modeling Accuracy





- SIP Modeling error converges as samping vectors increase
- SIP model complexity must be greater or equal to initial model complexity

Future Work

- GPU acceleration
- Uncertainty-aware tractography
- Adaptive modeling complexity

Thanks for your attention



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