

GENERATION OF HIGH-QUALITY TETRAHEDRAL HEAD MESH MODELS FROM MRI SCANS



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OUTLINE

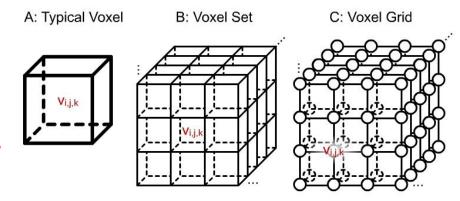
- Background
- Methodology
 - o Segmenting MRI scans
 - o Preprocessing the data
 - O Generating and combining surface meshes for each of the tissue layers
 - o Creating the final volumetric mesh
- Summary



REPRESENTATION OF MEDICAL DATA (1/2)

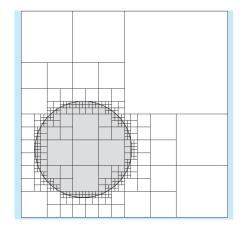
Voxel representation:

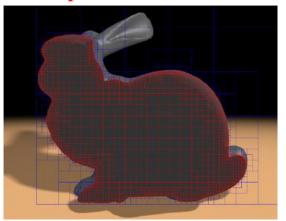
- Most common in medical imaging
- Easiest to generate
- Large number of elements required
- Less accuracy for complex boundaries



Octree representation:

- Similar to voxel, but can adaptively refine where needed
- Less elements than the voxel representation
- Accessing the data is more complicated (recursive access)
- Boundaries still have the staircase pattern





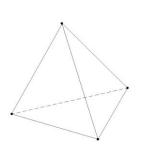
C. Dick et al. (2010)



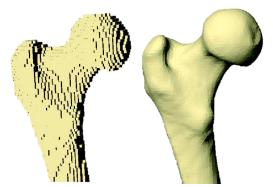
REPRESENTATION OF MEDICAL DATA (2/2)

Tetrahedral meshes

- Can represent complex boundaries
- Can be adaptively refined or coarsened
- Low memory cost
- Suitable for finite element analysis

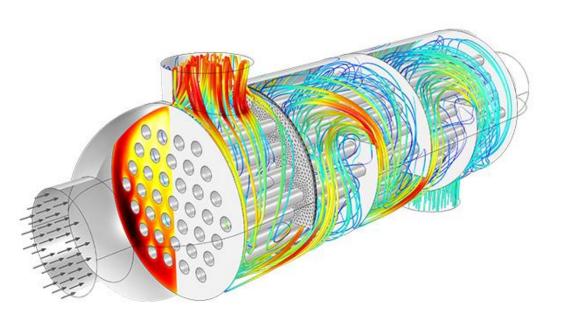


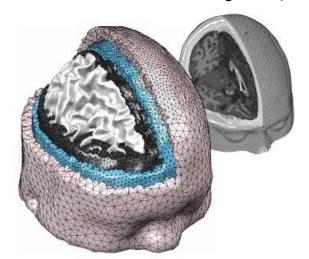
Tetrahedral element



Voxel Tetrahedral mesh representation

P.G. Young et al. (2008)



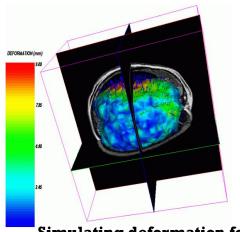


Colin27 head mesh model

Q. Fang (2010)



NEUROLOGICAL APPLICATIONS



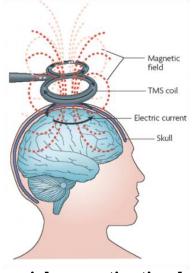
Simulating deformation for neurosurgery (Warfield et al., 2000)

Electric field/current density

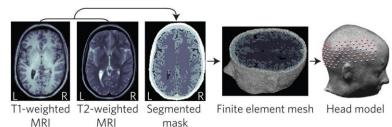
0 33% 66% Peak

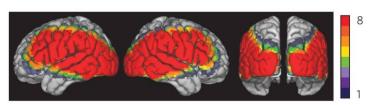
Simulation of transcranial direct current stimulation (tDCS)

(Dayan et al., 2013)



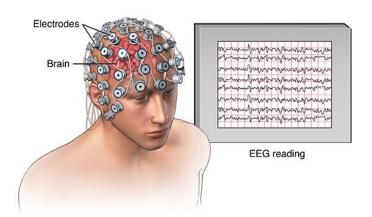
Transcranial magnetic stimulation (TMS)





Mapping brain functions through functional nearinfrared spectroscopy (fNIRS)

(Eggebrecht et al., 2014)



Electroencephalogram (EEG)

(Adebimpe et al., 2016)





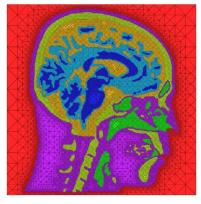
EXISTING MESHING WORKFLOWS



Biomesh3D

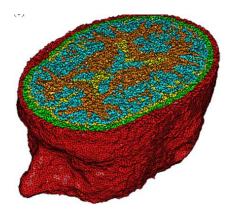
- 8-12 hrs
- Robustness issues

(Callahan et al., 2007)



Cleaver

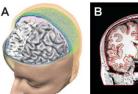
- ~Few minutes
- Large number of elements
- Not much control on element size
- Memory requirements are high (Bronson et al., 2012)



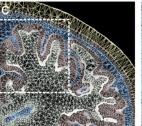
CGAL

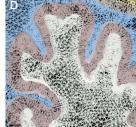
- ~5 min
- Robust
- Accuracy of boundaries

(Fabri & Pion, 2009)





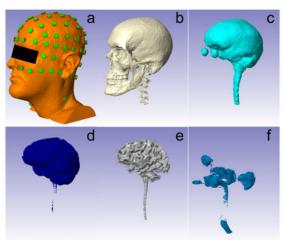




Mri2Mesh

- Accurate boundaries
- ~2-3 hours
 - Input flexibility

(Windhof et al., 2013)



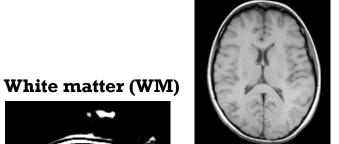
ScanIP

- Large number of elements
- ~2-3 hours
- Not open-source (Huang et al., 2013)





MESH GENERATION FROM A SEGMENTED VOLUME (1/2)





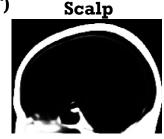
T1-weighted



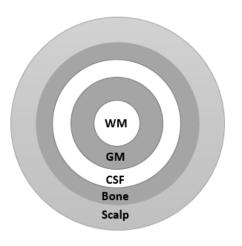




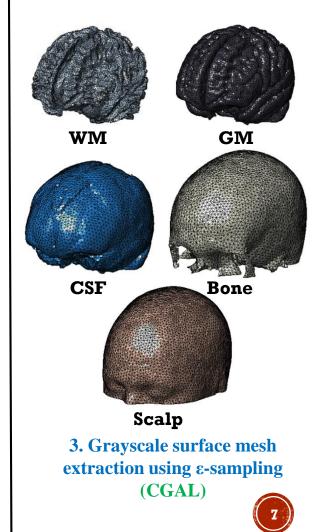




Segment MRI scans (SPM, FSL, BrainSuite etc.)

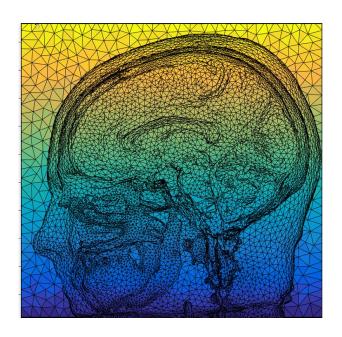


2. Pre-process segmented volumes to ensure brain layers don't intersect (Iso2Mesh)





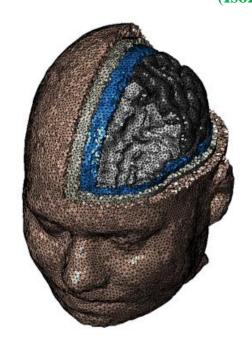
MESH GENERATION FROM A SEGMENTED VOLUME (2/2)

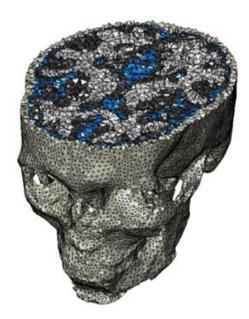


4. Generate a merged surface mesh (Cork)

5. Generate tetrahedral mesh using defined parameters (TetGen)

6. Label different tissue regions (Iso2Mesh)







RESULTS

Can use a wide range of segmentation tools

Head and

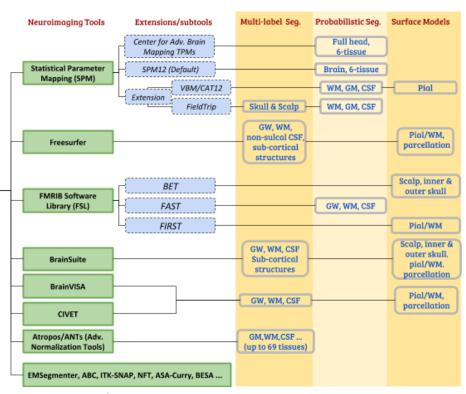
brain

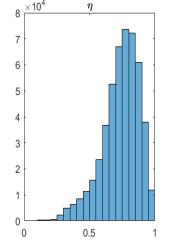
T1/T2 MRI

- High-quality elements
- Open-source
- Processing times of ~1 min
- Fine-grained control on element sizes
- Smooth and accurate boundaries
- Generate accurate brain models with low
 number of elements

(400k – 1M elements vs. 6-7 M in other meshing tools)

· Generalizable to other animal brains



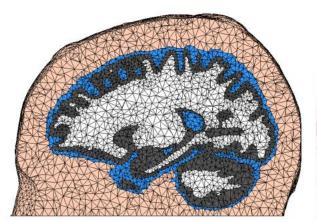


Toe-Liu quality metric



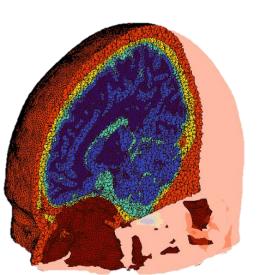


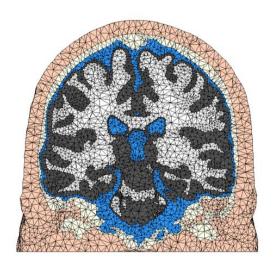
SOME BRAIN MESHING RESULTS

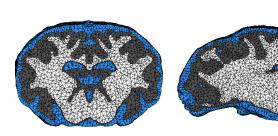




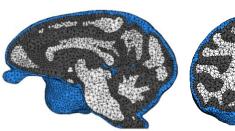
Mouse brain







Ovine brain





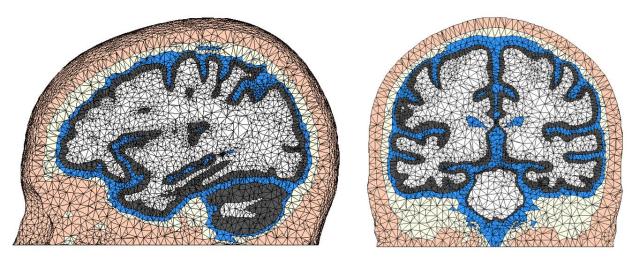
Baboon brain



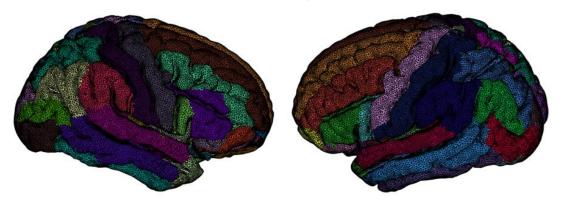
Mesh resulting from a FSL segmentation



WORKFLOW USING CORTICAL SURFACES



Tetrahedral mesh generated from FreeSurfer surfaces and SPM segmentation outputs for a 30-34 years-old average brain.





ACKNOWLEDGEMENTS

Professor Qianqian Fang, primary advisor **Professor Edgard Goluch,** co-advisor

Computational Optics and Translational Imaging lab members:

Xin Sun Xu Sun Morris Vanegas Yaoshen Yuan Yu Shi Edward Xu

Northeastern University, Department of Chemical Engineering and Bioengineering

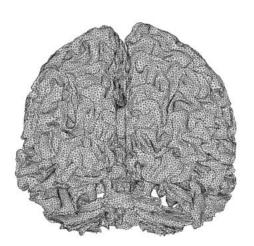
NIH funding: R01GM114365 and R01CA0204443

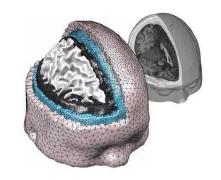




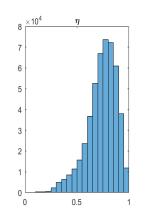
QUESTIONS

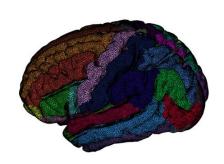




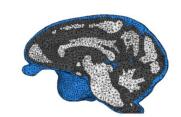








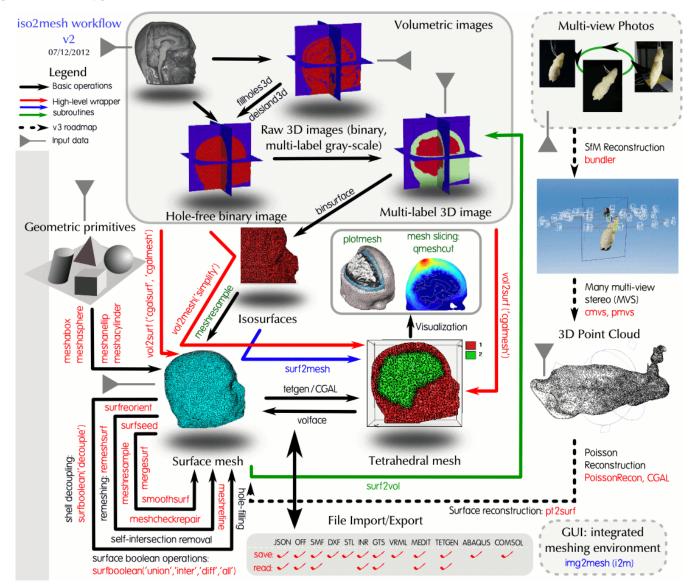






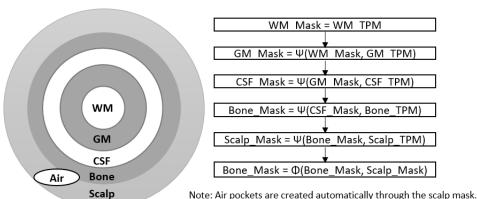


ISO2MESH





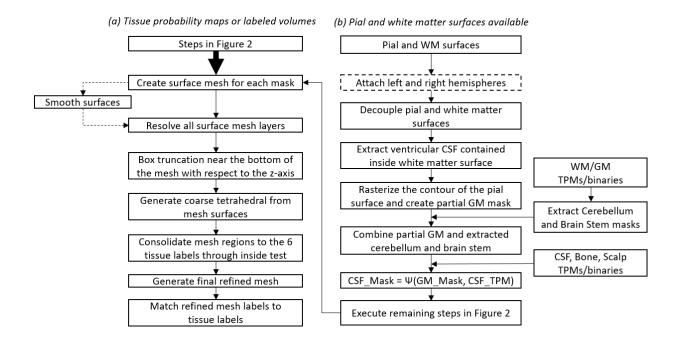
DETAILED WORKFLOW



 $\Psi(A,B) = \max\{A+B, \max_filter(A)\}\$

 $\Phi(A,B) = \min\{A, \min_{} filter(B)\}$

TPM = Tissue probability maps. Same procedure for binary values.





MESH DATABASE

