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#### **USING PARSL IN CREATING MAPPERTRAC**



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# OUTLINE

- Acknowledgements
- Refresher on the science use case
- Evolution of MaPPeRTrac
- Progress since last year
- Future plans



#### ACKNOWLEDGEMENTS











Joseph Moon

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Pratik Mukherji

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# WHY DO WE NEED CONNECTOMES

- Structural connectome is a powerful tool to characterize the network architecture of the human brain
  - Shows great potential for generating important new biomarkers for neurologic and psychiatric disorders like Traumatic Brain Injury
- Structural connectome is based on network-centric view of the brain has the potential to uncover how information is rapidly communicated and integrated across multiple brain areas
  - Uses graph analysis of networks constructed using white matter fiber tractography
  - Resulted in discovery of highly connected regions in brain called "rich-club" nodes
- Structural connectome using Edge density imaging (EDI) maps the number of network edges that pass through every white matter voxel
  - Focus on white matter pathways that constitute the edges of the network





# **EVOLUTION OF THE PIPELINE**

- Bash scripts
  - Hard-coded locations for applications and data
  - Worked well on one cluster at UCSF
- Python scripts
  - Parametrized
  - Ability to submit and run to SLURM
  - Worked well at LLNL
  - Application paths still hard-coded
- MaPPeRTrac
  - Uses Parsl
  - Successfully run at ANL, LLNL, AWS
  - Used Singularity container





# MAPPERTRAC HAS DRASTICALLY SIMPLIFIED AND ACCELERATE THE CONNECTOME RELATED







### **PERFORMANCE IMPROVEMENTS**







### MAPPERTRAC IS READY, ANALYSIS TOOLS ARE BEING REFINED AND VALIDATED

- The TRACK-TBI pipeline is ready to be deployed and to be generalized to other use cases
  - Fast and standardized analysis to support clinical research and application
  - ~1300 TRACK connectomes currently processed
  - Validation and verification is ongoing
- Available in open at https://github.com/LLNL/MaPPeRTrac
- New deep learning-based process is explored
  - CT anatomical segmentation and processing will enable improved multimodal analysis of neuroimaging and clinical features.
  - Accelerated MRI anatomical segmentation will result in faster processing of connectomes.





# WHAT'S NEXT FOR MAPPERTRAC

- MapperTrac
  - Complete testing on Google Cloud
  - Complete testing using docker container
  - Complete the manuscript
- Parsl issues
  - AWS issues
  - It would be nice to have a repository of configurations for known HPC and cloud infrastructures that can be installed as plug-ins





### **THANK YOU**



