



Cancer Imaging Phenomics Toolkit (CaPTk):

A Radio(geno)mics Platform for Quantitative Image Analytics

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Primary Aim: To enable swift and efficient translation of cutting-edge academic research into clinically useful tools^[1].

Target Audience:

- **Clinical experts:** facilitating use of complex algorithms for clinically relevant studies through a user-friendly interface.
- **Computational experts:** allowing batch-processing of multiple subjects and integration of new algorithms.



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(HPC-shared resources for computationally demanding pipelines)

Web-accessible

Cross-Platform

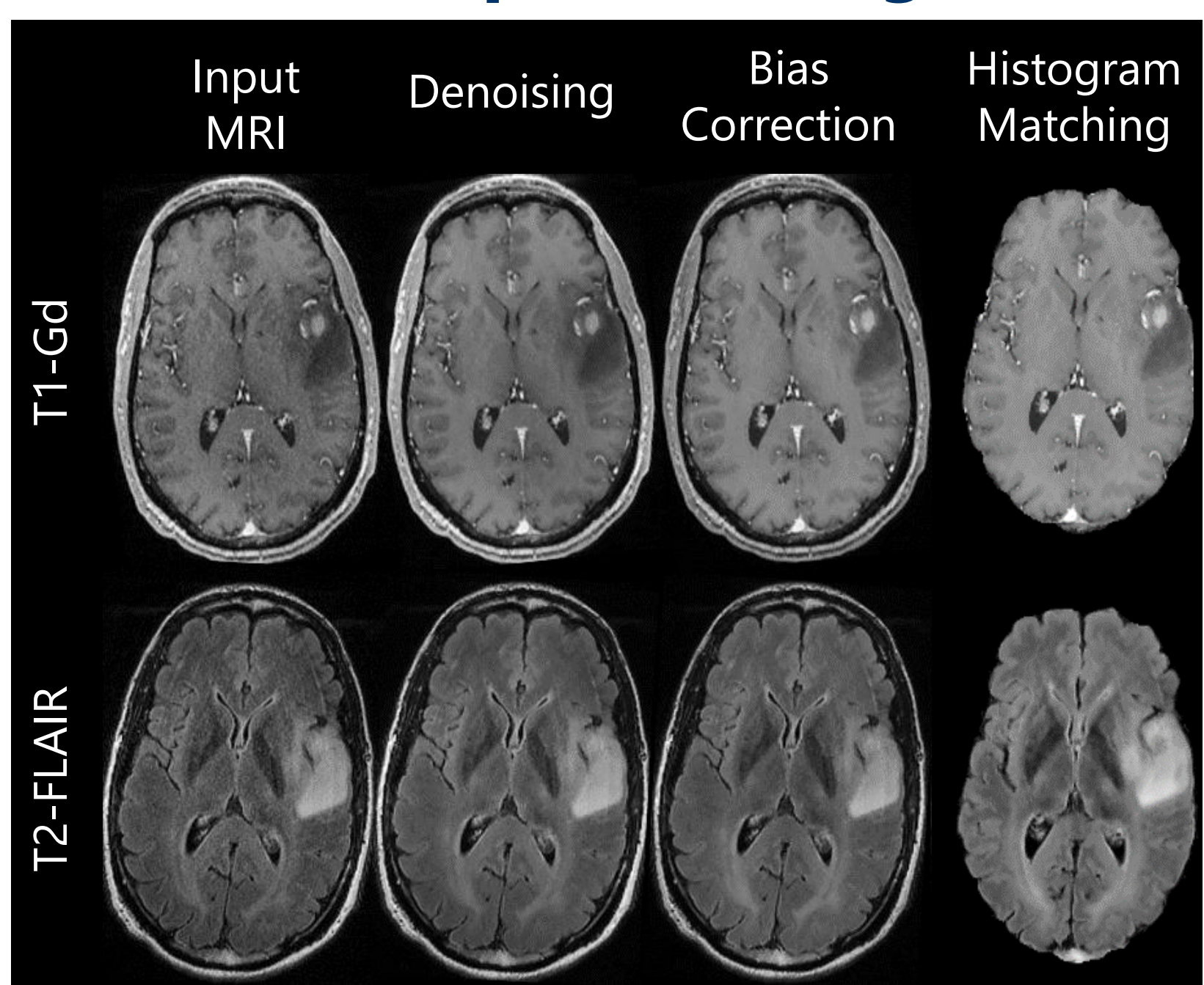
Open-Source

Easy to use

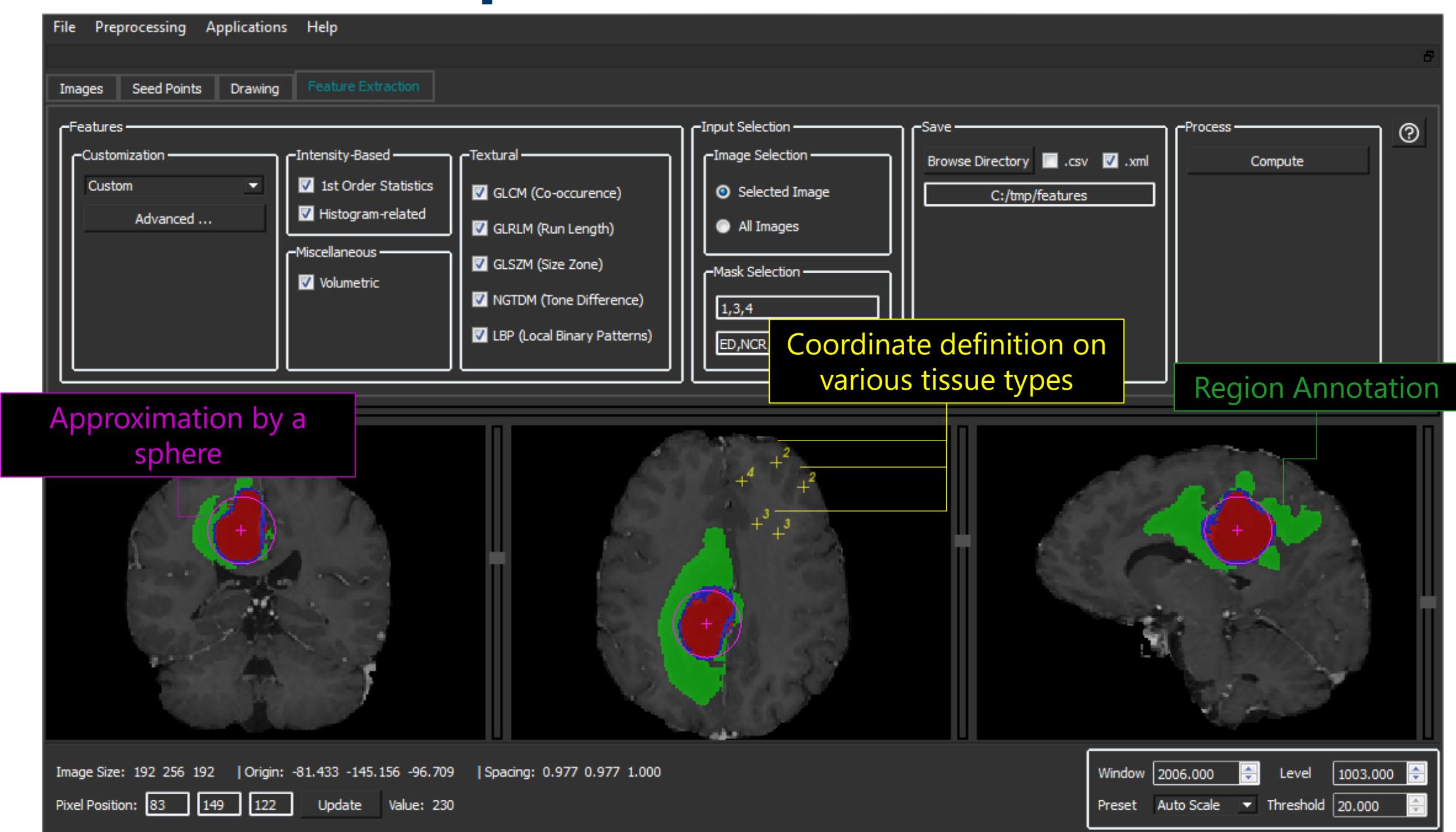
Extendable

– General Purpose Tools –

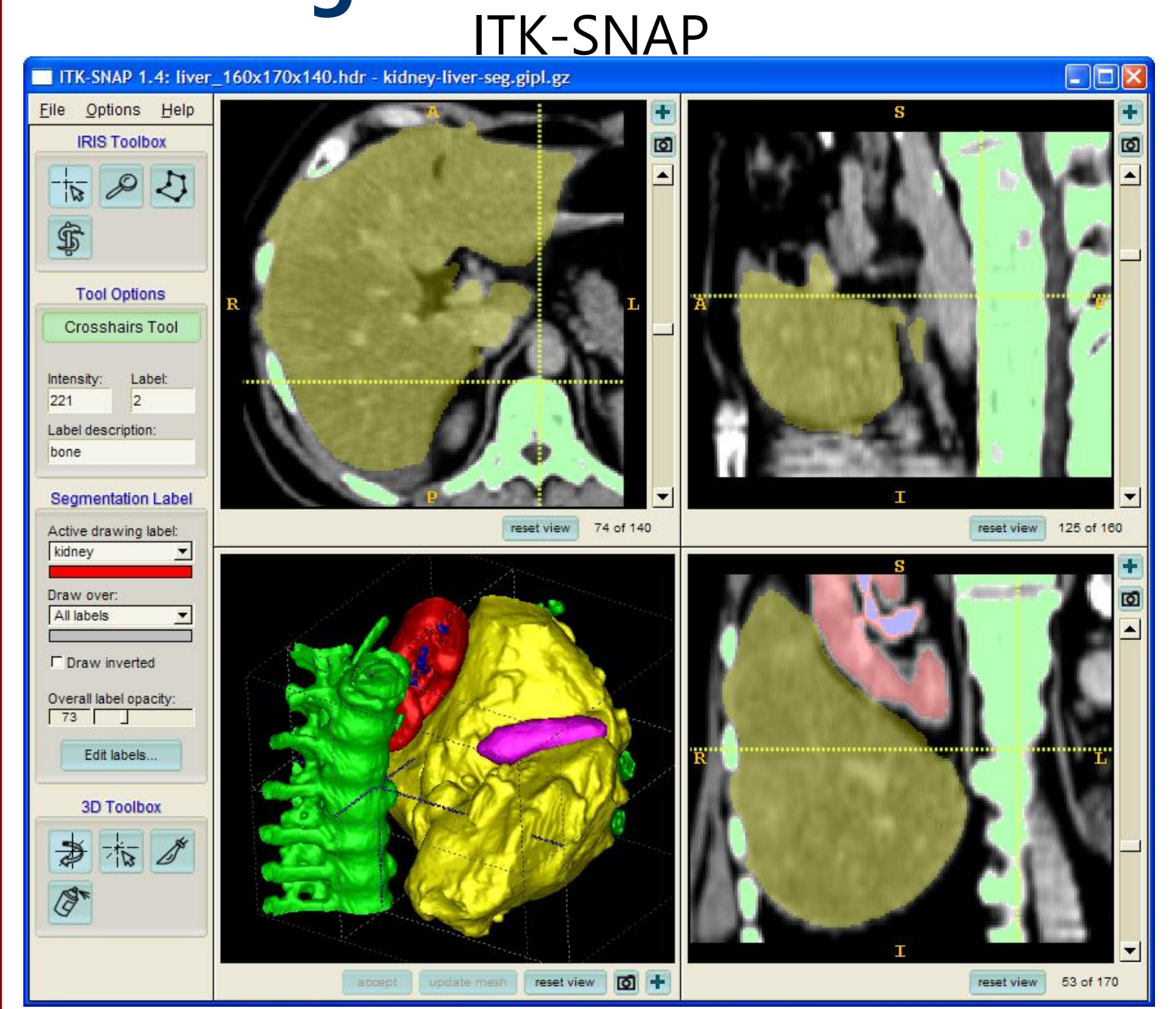
Pre-processing^[2]



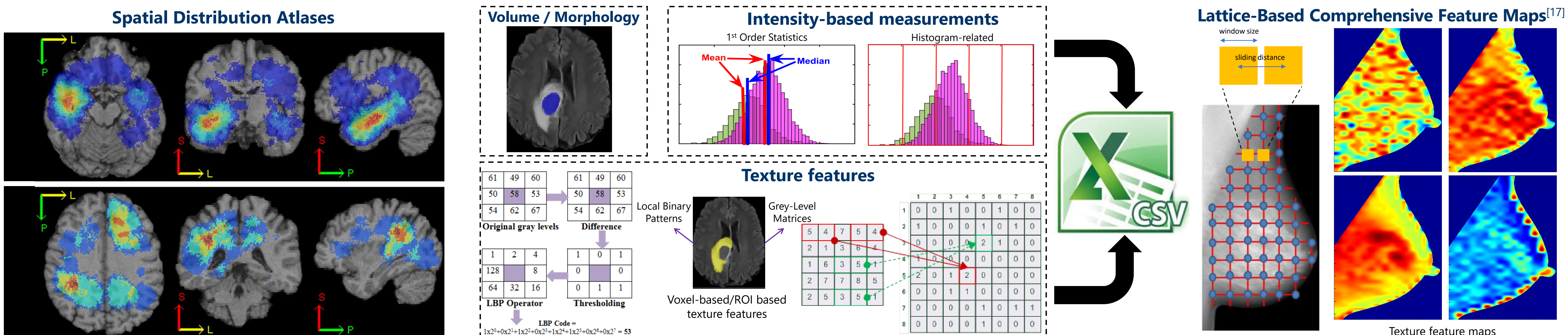
Graphical Interaction



Segmentation^[3,4]

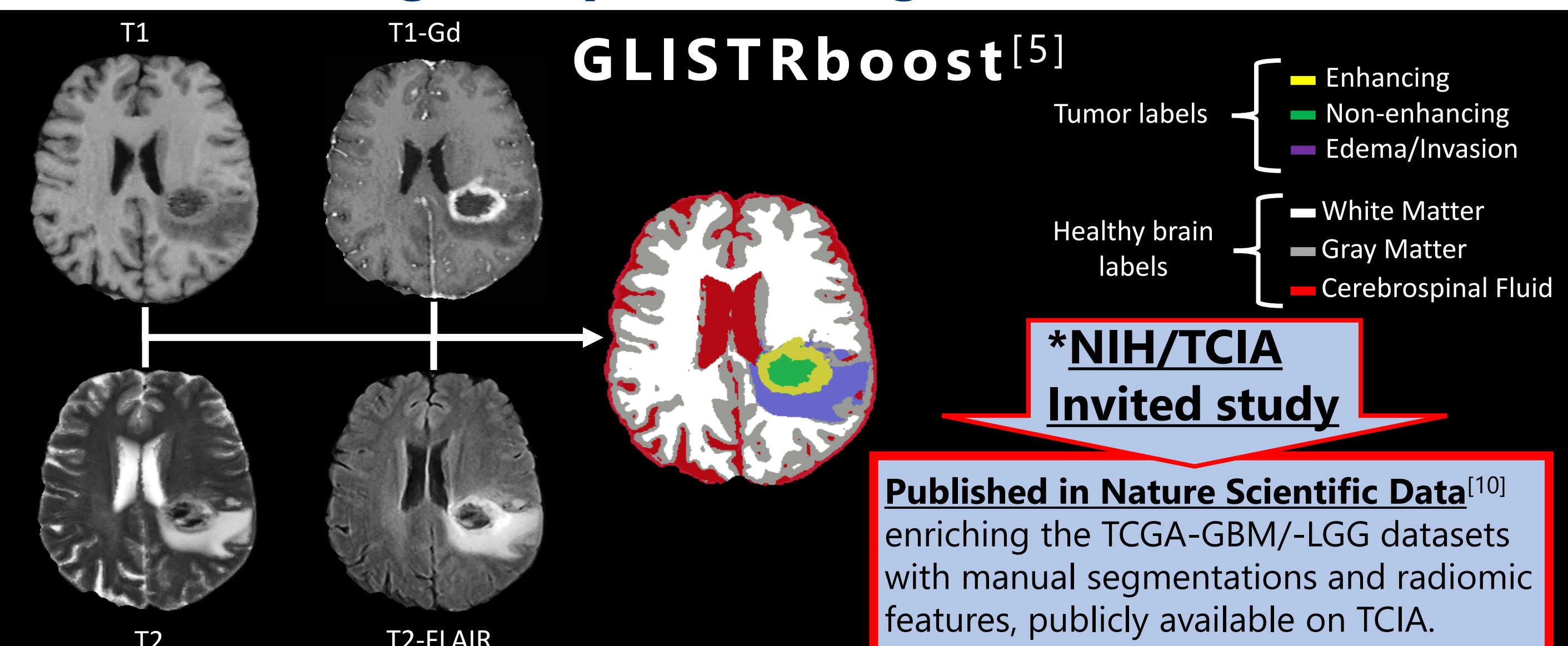


Quantitative Feature Extraction

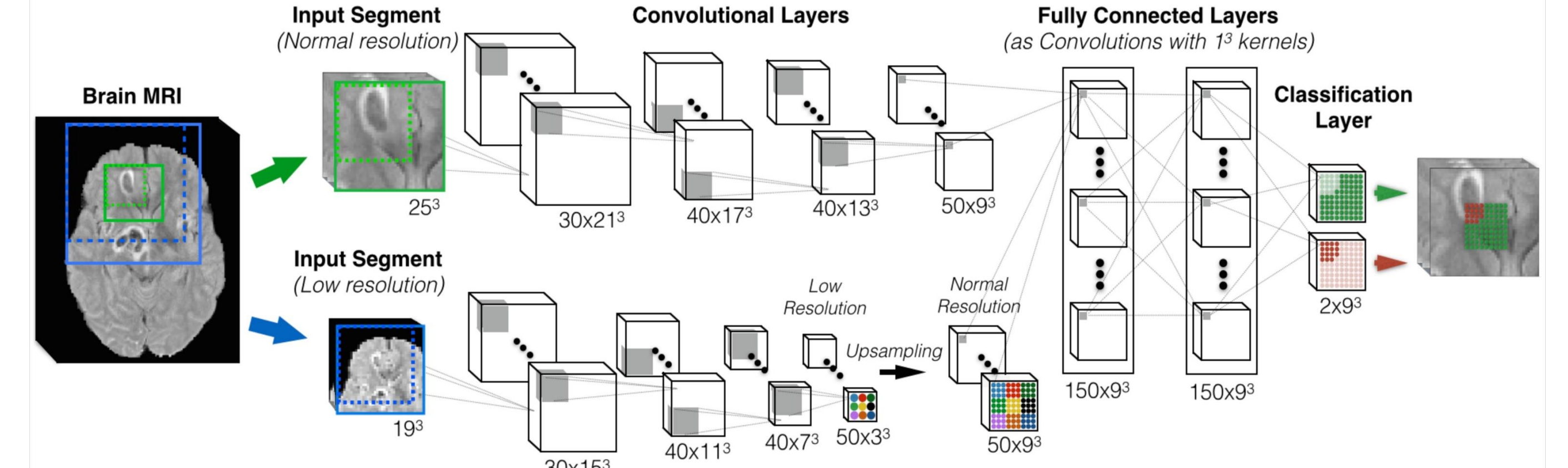


– Specialized Applications –

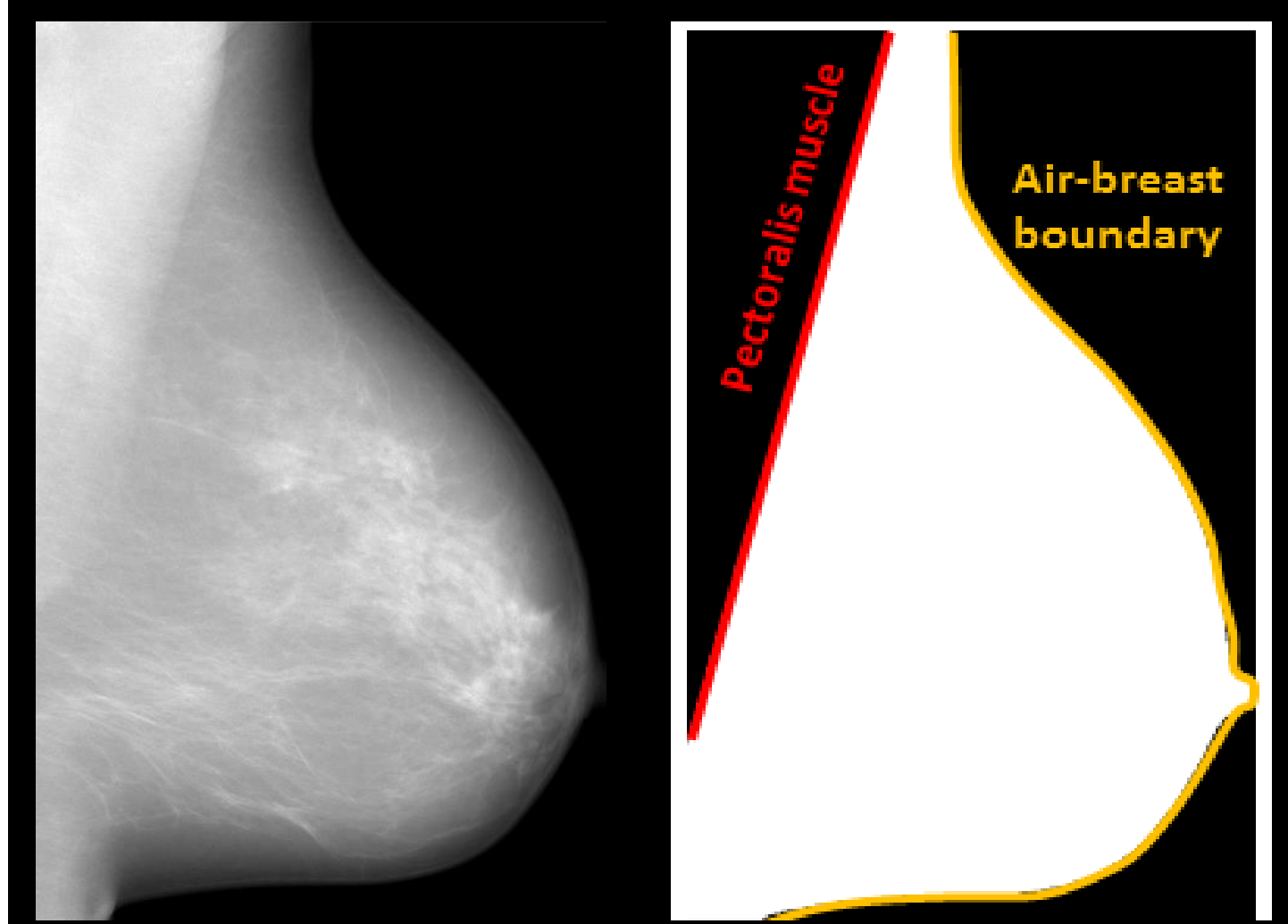
Organ-Specific Segmentation



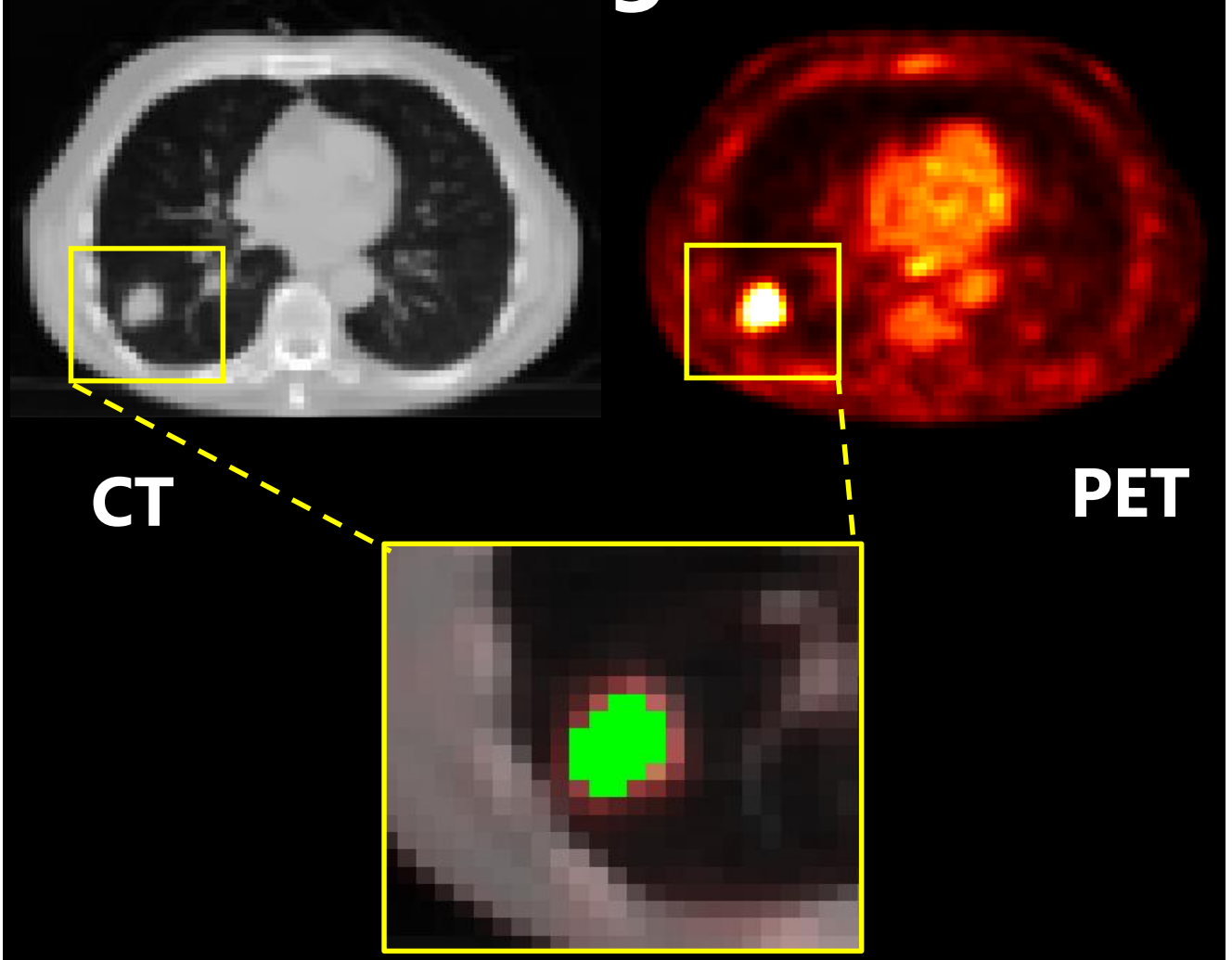
DeepMedic^[6]



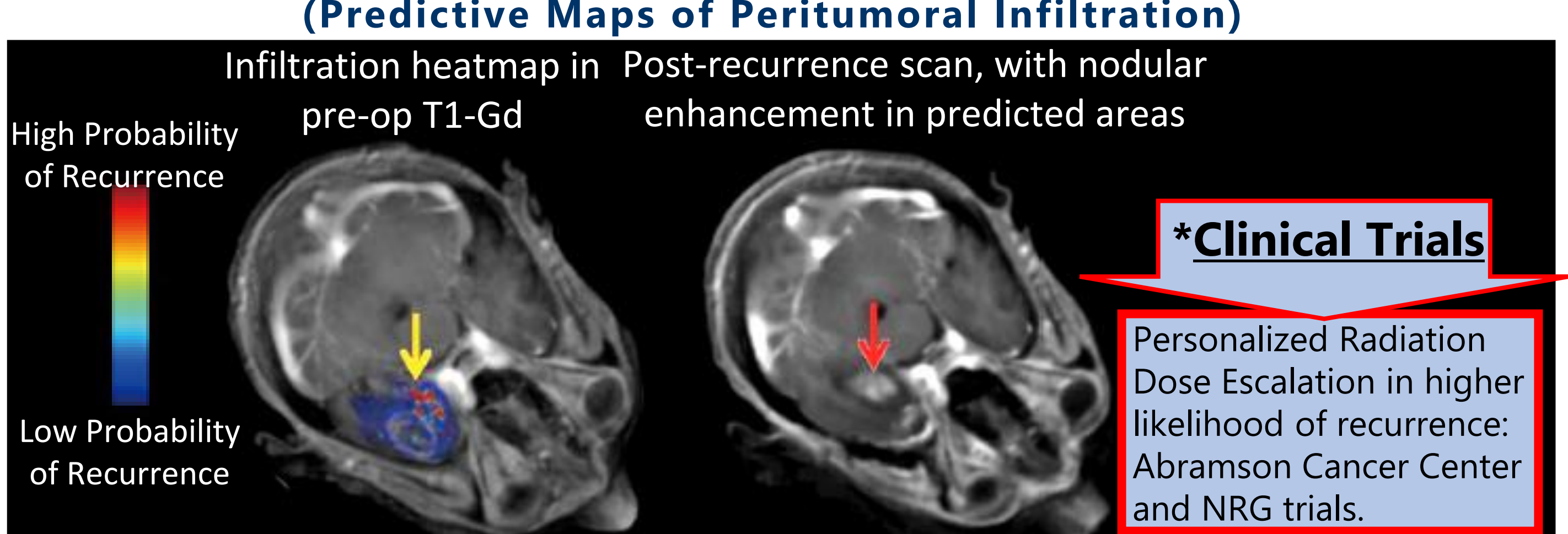
LIBRA^[7,8]



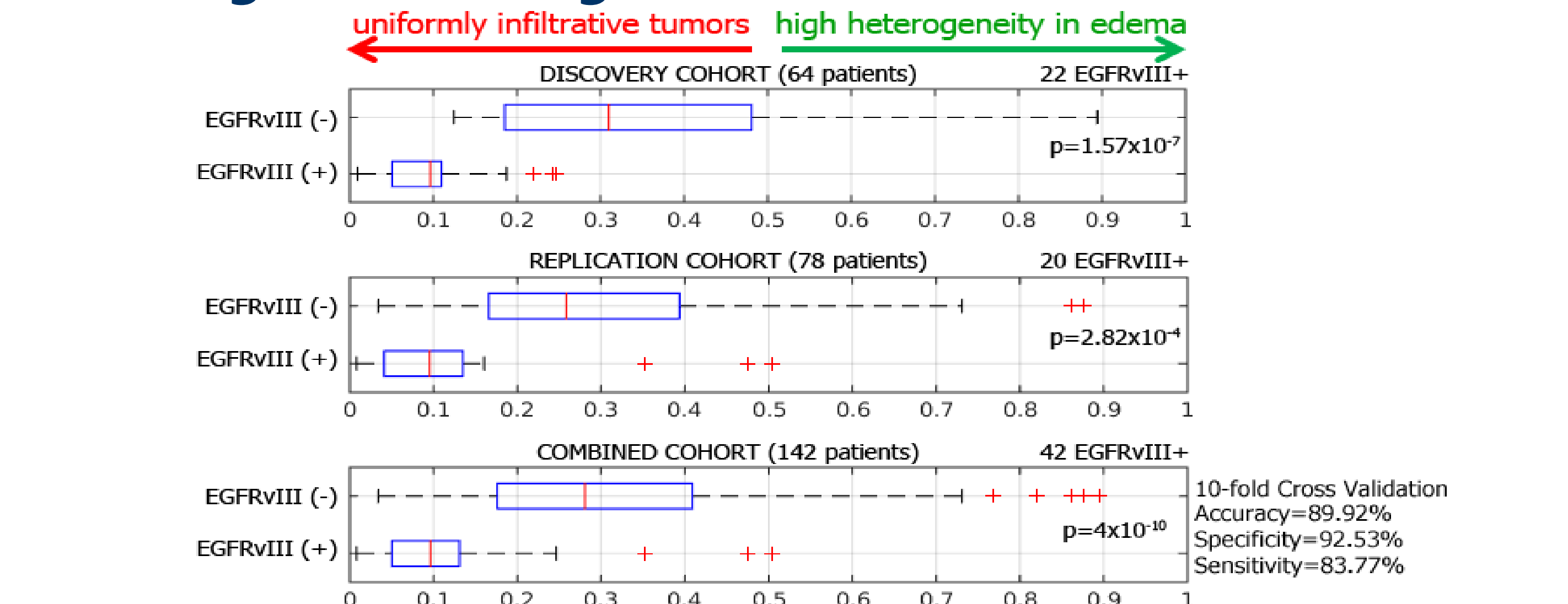
SBRT Segment^[9]



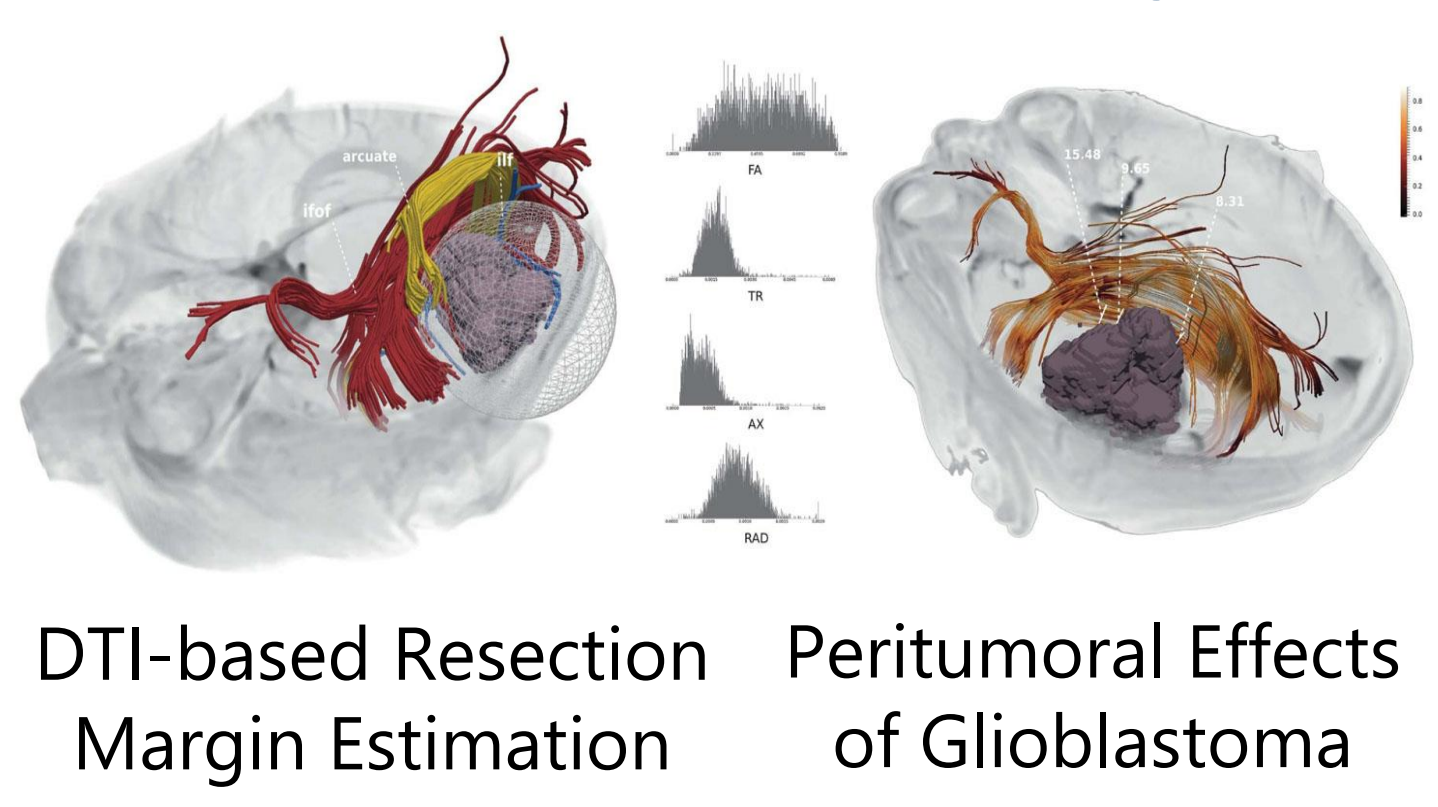
GBM Recurrence Prediction^[11, 12]



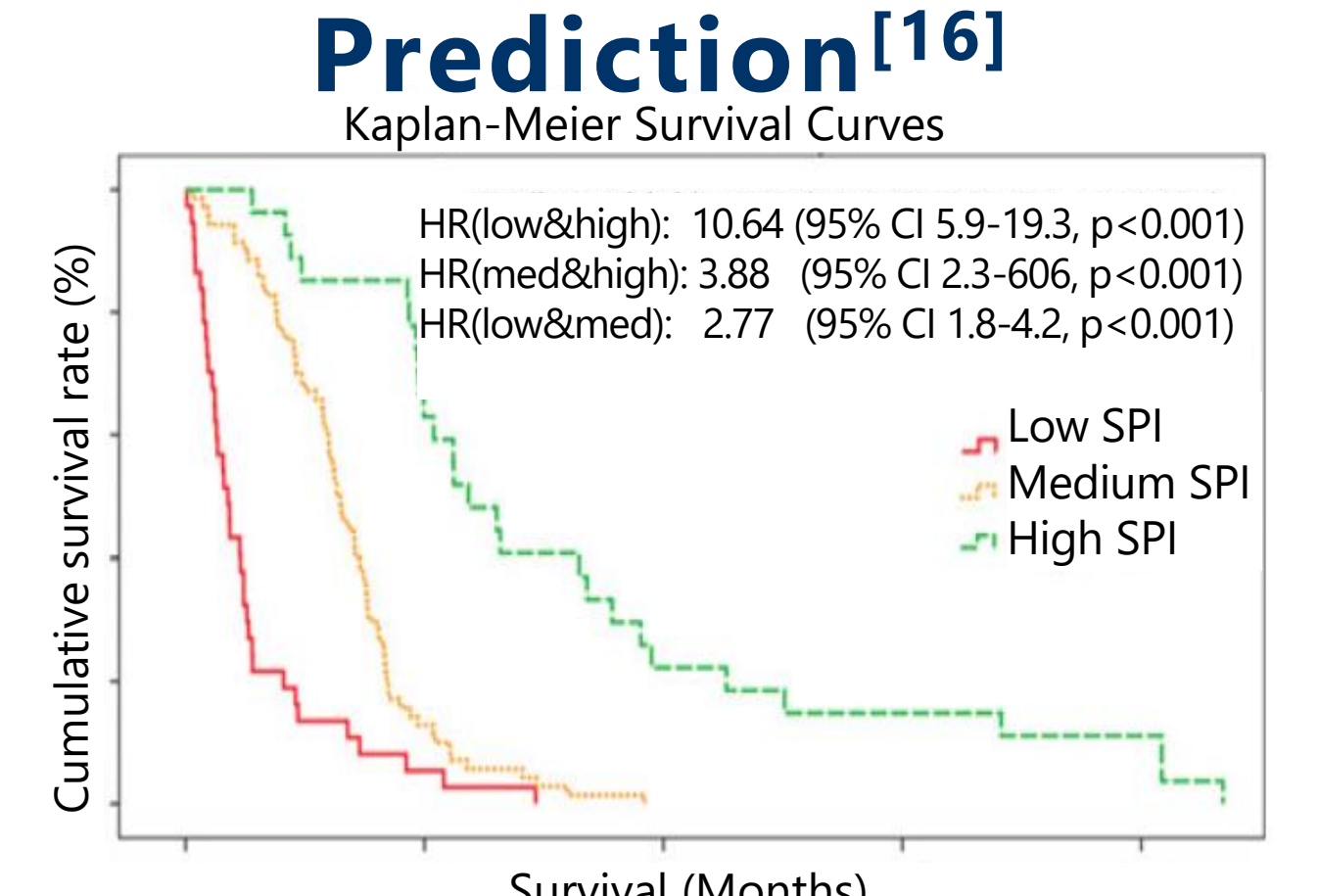
Radiogenomic Signature of EGFRvIII in GBM^[13]



Computational Study of Brain Connectivity^[14,15]



GBM Survival Prediction^[16]



– Future Directions –

Cohort-Based Interface | Traditional Machine Learning Training Module | Multiscale Feature Extraction

References

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