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Breakout Session II: Predictive Oncology Algorithms and Software – Challenges, Opportunities and Paths Forward

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Current Resources

- 1. ITCR
- 2. Data Commons
- 3. CANDLE
- 4. QIN (quantitative imaging network)
- 5. Dockstore.org genomic related containers
 - 1. Workflow containers
- 6. Openslide
- 7. TCIA
- 8. NCI cloud resources
- 9. List of medical imaging resources
- 10. Other open domain Machine Learning tools / frameworks

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Key Opportunities

- 1. Maintain a curated list of available tools
 - (as currently done by ITCR)
 - What's the incentive to participate: Publication required as term of grant
- 2. Leverage DOE non-cancer tools
 - system biology, CFD and other scaled code from national labs
- 3. Tie tool development to clinical outcomes
 - 1. Bring tools into clinical workflow
- 4. Improve Standards
- 5. Improve quality and awareness of workflows already available in the cloud
- 6. Map hospital EMR data into a standard form to facilitate input into CANDLE
 - 1. Can we use ML to control variability in hospital specific implementations
- 7. Move from desktop to cloud based tools
- 8. Reference/Benchmark datasets
- 9. Can we generate enough synthetic data to use for training
- 10. Quality assurance, scalability, supportability of the tools
 - Need a standards body



Challenges and Roadblocks 1. Have to put tools into clinical workflows to validate them

- - This dos not necessarily mean clinical trials
 - Start with quality settings?
- Clarify what CANDLE can be leveraged to help parallel the clinical process/workflow
 - What can and cannot be ingested from EMRs
- Interoperability is the key challenge
- Transitioning informatics into something clearly actionable
- Improving digital literacy of physicians Usability of tools for plhysicians
- Visualization
- Data assurance, validation and views
- What is sharable? When? How do we best manage?
- Managing X-domain collaborations
 - 1. Role of tools
 - 2. Impact/convergence for tools
- 10. Still constrained by access to data
- 11. Access to computational power required to train algorithms

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Next Steps

- 1. Improve participation from EMR players
- 2. Ensure expertise in:
 - 1. Complex software systems and integrations
- 3. Clarify and communicate TCIA
- 4. Clarify and communicate Cancer Cloud Resources
- 5. Clarify and communicate availability of Data Commons
- 6. Improve V&V of software
- 7. Broad teams of pathologists and EMR Vendor players