



Building Genomic Medicine Capability

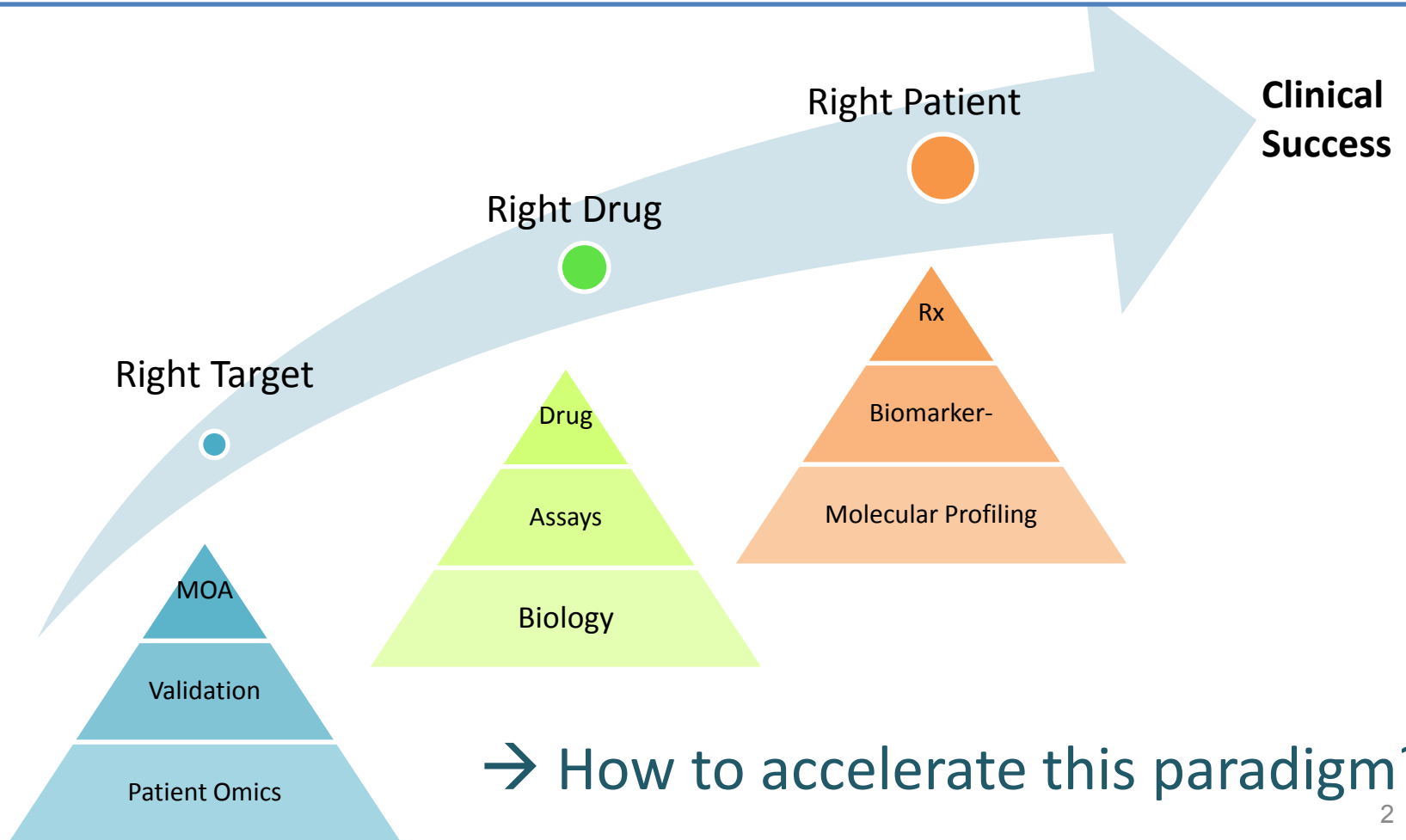
Challenges and opportunities of big data



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Personalised/Stratified/Precision Medicine for Cancer

Personalised medicine will enable the much needed paradigm shift in clinical care delivery, but we will need appropriate tools & know-how to realize the model and implement the vision



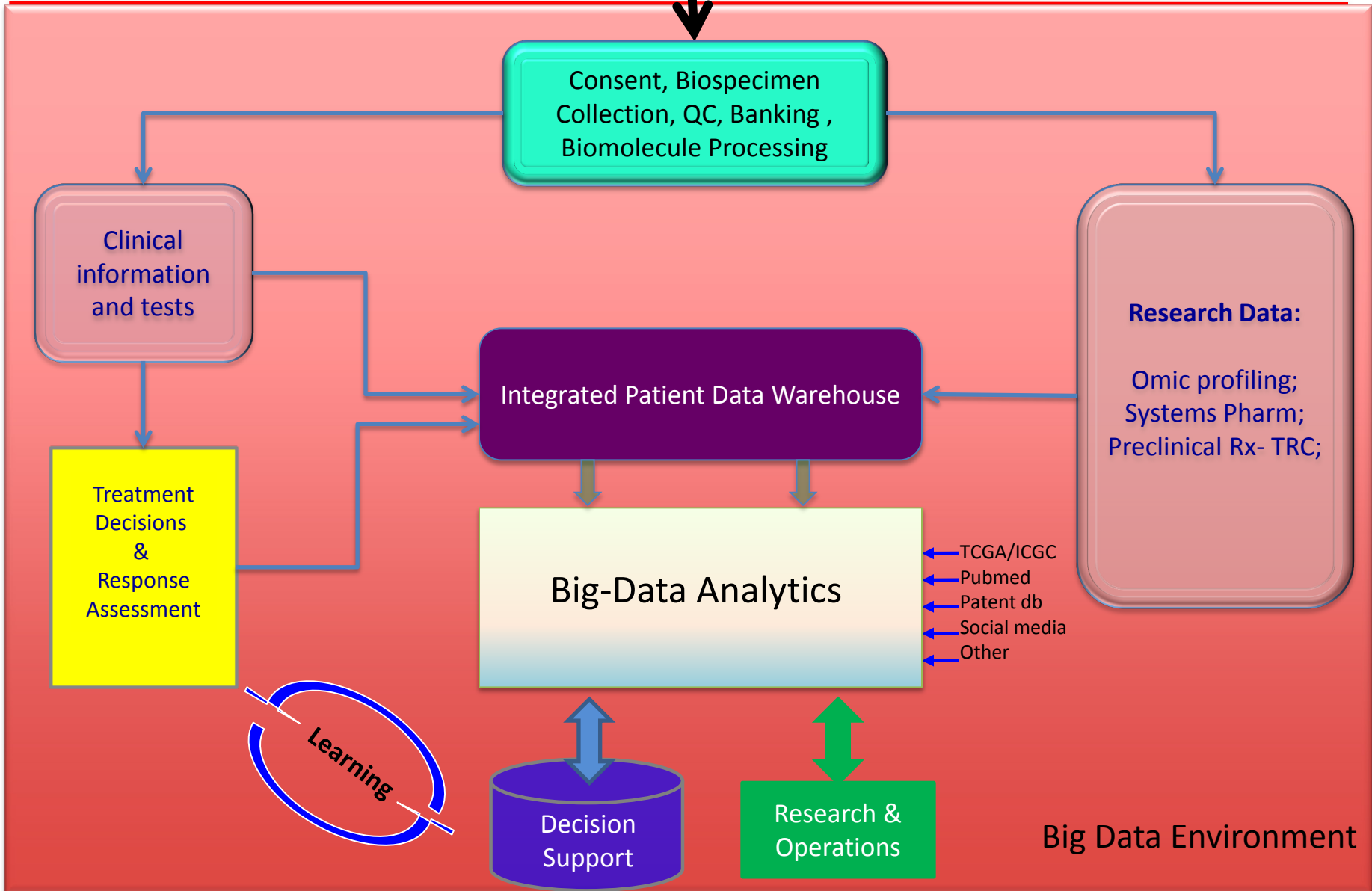
- The selected cancers are:
- [Triple Negative Breast Cancer](#)
- [High-grade Serous Ovarian Cancer](#)
- [Leukemia \(AML/MDS\)](#)
- [Leukemia \(CLL\)](#)
- [Lung](#)
- [Melanoma](#)
- [Prostate](#)

- Focus on patient impact and reduction in mortality world-wide
- Comprehensive, spanning the cancer care continuum
- Collaborative, internal and external
- Innovative, in organizational constructs and technology

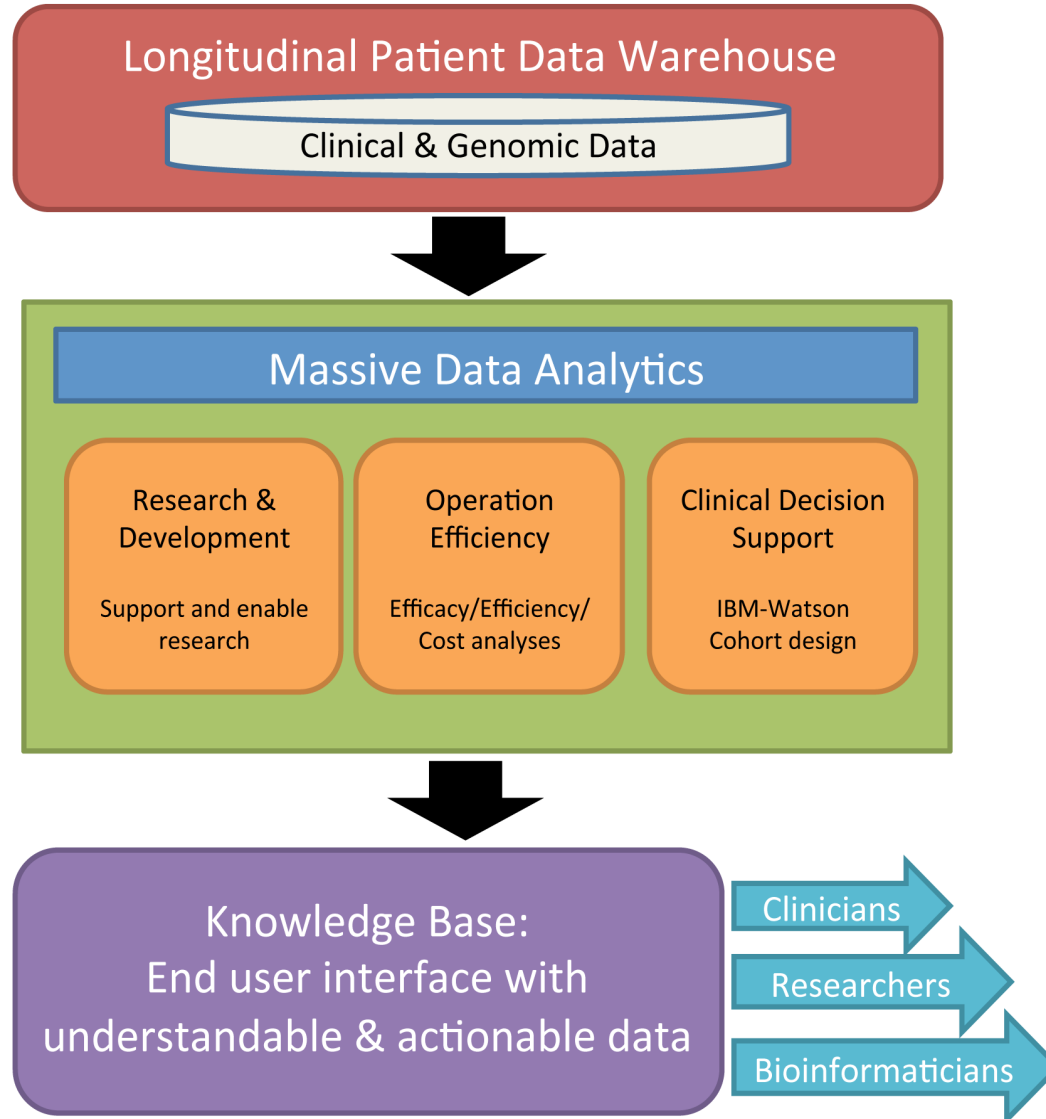
Moonshot Platforms

- Center for Co-clinical trials
- Institute for Personalised Cancer therapy
- Cancer Control
- Early detection/Diagnostics
- Clinical Genomics
- Immunology
- Institute for Applied Cancer Sciences
- Translational Research Continuum
 - Research Genomics/Informatics
 - Big Data
 - Adaptive Learning

Adaptive Learning in Genomic Medicine



Big (well, it is Texas after all) Data Analytics



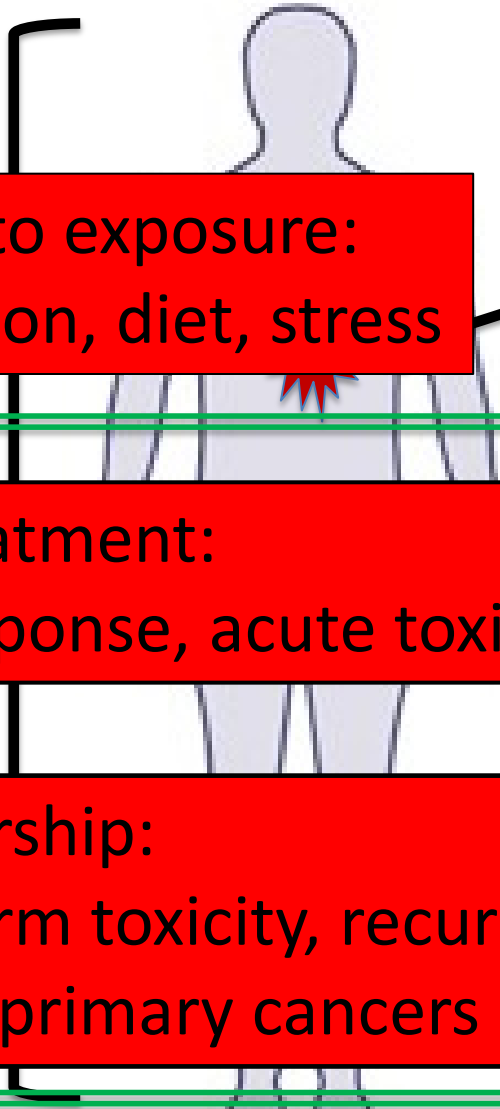
- 1000 leukemia patients by fall 2013– MDS/AML/CLL focus
- Focused on but not limited to newly diagnosed patients
- Samples taken at diagnoses/presentation and thereafter at each patient visit.
- Saliva/buccal for normal, bone marrow and/or peripheral blood
- Bone marrow/bloods accessed in context of normal clinical workup/care
- All samples collected and held in CLIA compliant chain of custody

- Exome sequencing, low-pass WGS
- Data generated on normal/tumor (presentation) and from relapse sample(s)
- All clinical data currently collected in Departmental database plus extraction from patient records
- A few early potential questions –
 - MDS to AML progression
 - risk of death during induction chemotherapy
 - subclonality and risk of relapse/progression

- Other Opportunities (some of them)
 - Genetic/genomic heterogeneity
 - Comprehensive cancer patient genomics –
 - Interplay of germline and somatic genomics in the same patient
 - Impact of genomics on outcomes
 - adverse events
 - survivorship

- Genetic heterogeneity is a key determinate of variation in outcomes
 - What are the cancer genes operative?
 - What is the level of intra-tumor heterogeneity?
 - What are the germline/somatic sequence variants that are influencing factors including:
 - Drug metabolism
 - Immune response
 - Cancer susceptibility
 - Toxicity
 - How do these factors interact and influence outcomes?

Comprehensive Cancer Patient genomics a tale of (at least!) two genomes



Risk and response to exposure:
Tobacco, UV radiation, diet, stress

Somatic

Germline

Treatment:
Response, acute toxicity, resistance

Survivorship:
Long term toxicity, recurrence,
second primary cancers

Adaptive Learning/Leukemia Team

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