



Data Deluge: Challenges and Solutions in Implementing a Precision Medicine Approach to Create the TRACK-TBI **Information Commons**

Tatiana Khasanova¹; Mary Vassar²; Mónica Coelho¹; Sabrina Taylor²; Laura Brovold¹; Geoffrey Manley² and the TRACK-TBI Investigators³

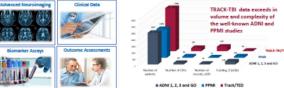
- ¹ Rancho BioSciences, LLC, 16955 Via Del Campo #220, San Diego, CA
- ² University of California San Francisco, Department of Neurosurgery, Brain and Spinal Injury Center

TRACK-TBI Background

- Traumatic Brain Injury (TBI) is a serious condition, affecting an estimated 5.3 million people living with the longe, and psychological health disabilities, with annual direct and indirect costs estimated a term physical, cognitive, and psychological health disabilities, with annual direct and indirect costs estimated at over 576 billion (II). Each year in the U.S. there are approximately 2,8 million TBF-taled emergency department (ED) visits, hospitalizations, and deaths (2). TBI is a contributing factor in a third of all injury-related US deaths. Despite promising preclinical research on neuroprotective agents these advances have failed to translate into a single successful clinical trial or treatment (3). The heterogeneity of TBI has post challenges to the design of clinical trials and approaches to effective treatment of TBI represents a great unmet need in public health.
- The NINDS and DoD funded² multicenter Transforming Research and Clinical Knowledge in Traumatic Brain Injury (TRACK-TBI) study aims to improve TBI classification and identify new diagnostic and prognostic markers for future
- Since 2013, in collaboration with expert public-private partners and industry, we are collecting and analyzing since 2025, in consolution with expert point-private granters and industry, we are contenting and analyzing comprehensive acute and long-term data on subjects at 19 U.S. Trauma centers, across the spectrum of fininy. This includes physiology measures, advanced neuroimaging, blood biospecimens, and outcome assessments. TRACK-TBI has currently enrolled 3,264 persons with TBI and will complete follow-ups for the first 2700 TBI and another 300 Orthopedic Control subjects in 2019.
- * TRACK-TBI network has established a robust research infrastructure involving hundreds of multidisciplinary experts collaborating in a concerted effort to improve the diagnosis, treatment and outcomes for persons with TBI.

Data Types

- Clinical, standardized high resolution neuroimaging, proteomic, genomic and outcome biomarkers
- Injury spectrum ranges from concussion to coma for patients enrolled within the first 24 hours fol
- Subjects are followed during their acute care phase and at four follow-up visits over 12 months:
- Clinical and outcomes core data are collected in a web-based electronic case report form that has over 200 forms and 15,000
- months are submitted electronically to a central imaging repository



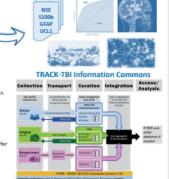
Precision Medicine

The ultimate goal for this study is to enable development of new diagnostic and prognostic biomarkers for future treatment clinical studies. Ideally, all data types should be analyzed holistically to arrive at the credible biomarke

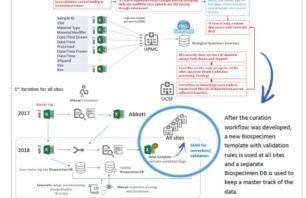


Commons is being built to harmonize disparate datasets across multiple domains and applications

- · Our teams have been developing curation datasets for analytic teams
- Data has been harmonized with the NIH TBI Common Data Elements and submitted to the Federated Interagency TBI Repository (FITBIR) for future sharing with the research community
- Examples are presented for the workflows, challenges and solutions



At the time the study was launched biospecimen data collection consisted of Excel sheets logs used by the study sites with no validation controls to flag and prevent errors.



Biospecimen Validation Process

men logs are maintained for the inventory of over 90,000 bio



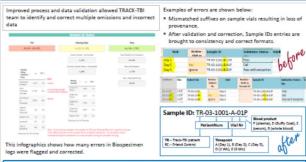
Challenges and Solutions

 In examples, we harmonized three primary sources of data: clinical and outcomes (via CRFs), Biospecimen samples, and neuroimaging findings. For clinical data, we started by implementing OpenClinica solution to track data entry errors and omissions. This work silowed us to de-lineate sources of mistakes (Dingram 1), However, a more errors and omissions. This work allowed us to de-lineate sources of mistakes (Diagram 1). However, a more succinct solution was developed using an R Shiny app (Diagram 2).



- ling to consistency; and (3) identify omissions in collect sample collection is ongoing; (2) bring timestamp cool and request corrective actions.
- We solved these issues by developing a uniform template for biospecimen log entries (with multiple rules for autodata checking) and a biospecimen validation process.





Future Directions

- or the first 3000 subjects for ha repositories and creation of locked datasets for sharing.
- Develop analytic pipelines for biomarker-based pathoana approach in an exploratory clinical trial utilizing the established TRACK-TBI infrastructure.

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References