

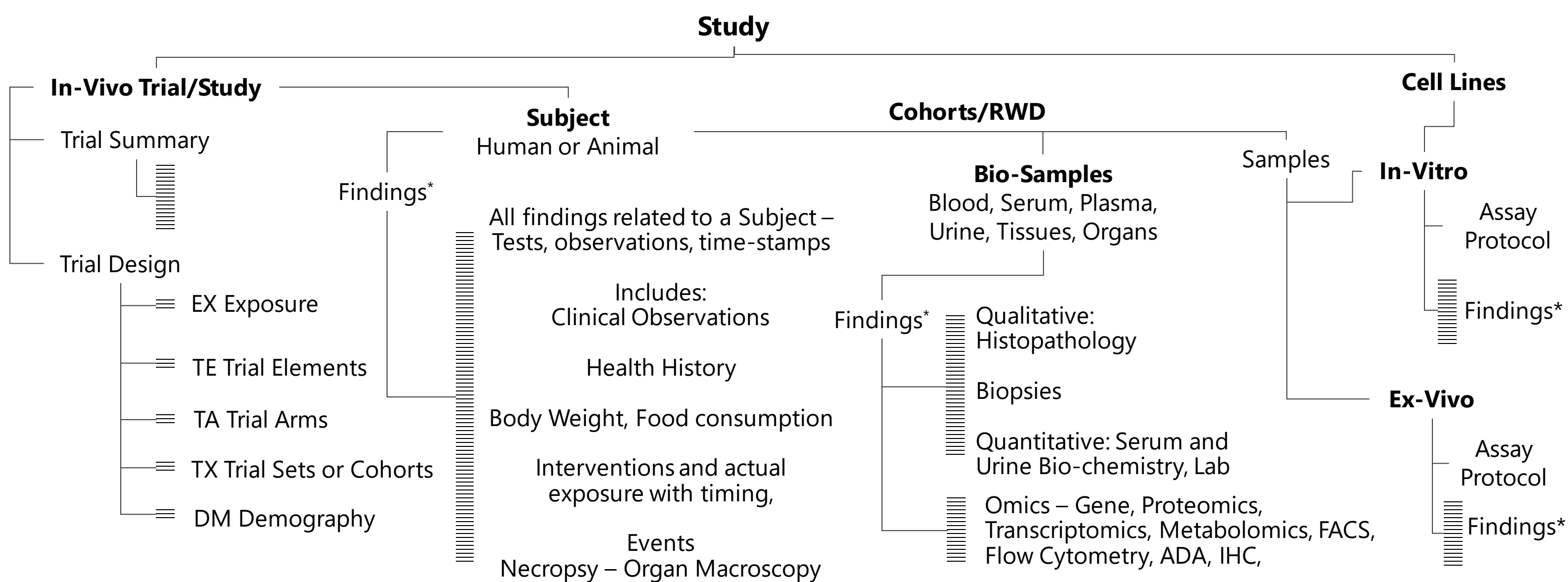
## Input: Metadata & Raw Clinical Data

### Create End-to-Start Specification

- ☐ Produce a standards-based, machine readable specification

### Generate Start-to-End Metadata

- ☐ Use standards specification to generate study metadata artifacts
- ☐ Demonstrate the ability to generate study metadata given a specification

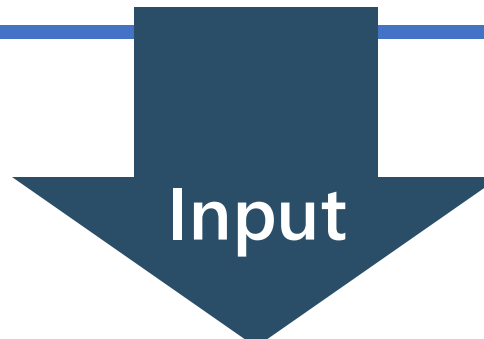
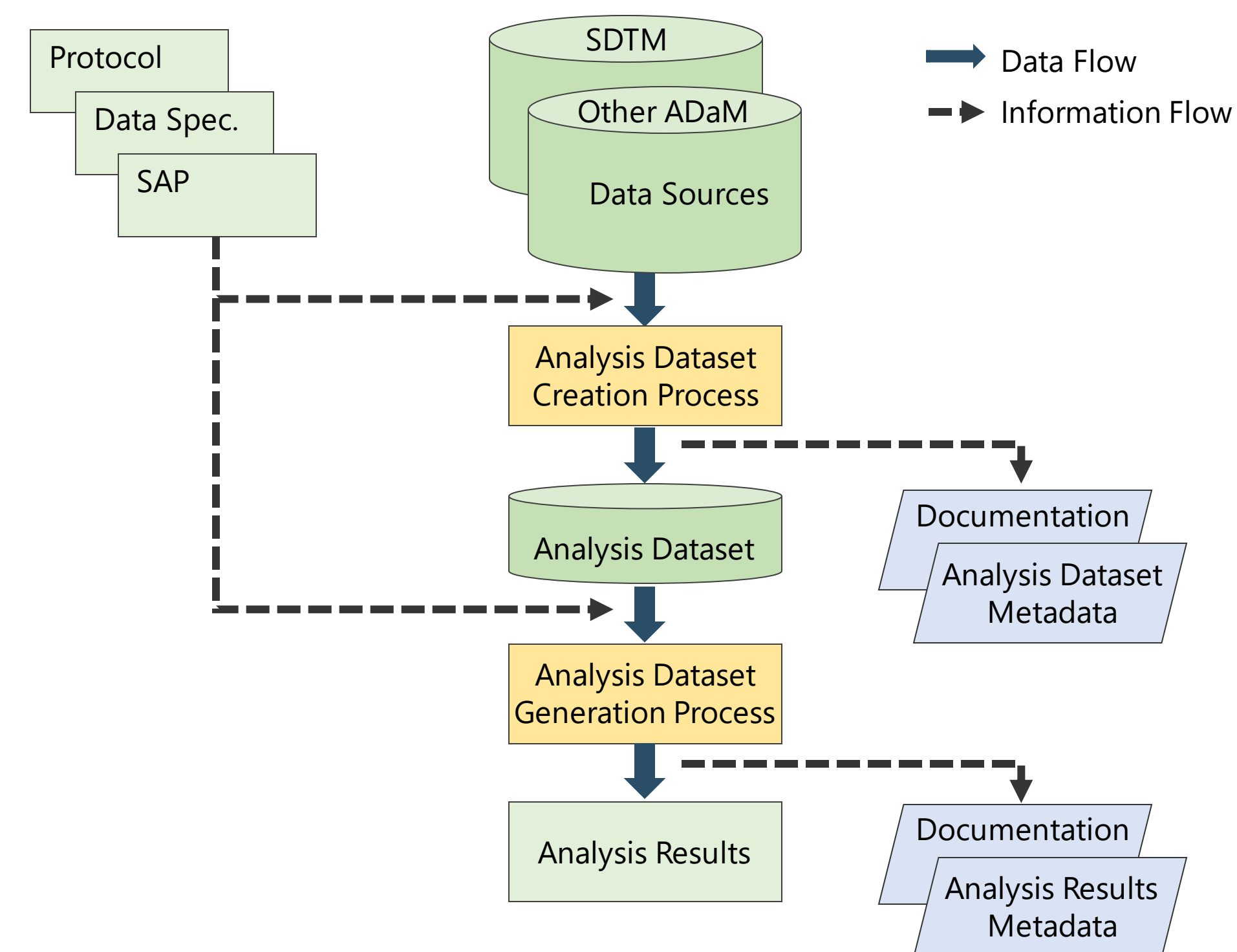


\* Findings are held in a common storage model with Observation/Testname, date-Time stamp, Units and other attributes

## Output: SDTMs, ADaMs, Define.xml & TFLs

### Data Curation

- ☐ Repetitive Process to Optimize Data and Metadata to ensure Valuable use of Data



## Transformation and Automation: Reusability & Repeatability

### Transformation Data Start-to-End

- ☐ Use machine-readable metadata to generate study data artifacts
- ☐ Demonstrate the ability execute data transformations given the study

### Replication and automation are the focuses

- Use or create utilities to replicate the process: Project Set Up, Mapping Specification, Mapping Creation
- Use analytics tool to identify the areas for replication and automation: Data Profiling & Data Rules for Source Data Review / Edit Checks

### Metadata-driven process is the key for automation

- Metadata makes data meaningful
- Metadata is machine readable
- Metadata is the base for automation

### Standard adoption is the key for code reusability

- Train people to understand the standards
- Define standard templates
- Build public libraries for code snippets and public transformation: Custom functions, procedures and packages; public data rules; and public Experts
- Group code snippets and functional transformation into modular mapping and transformation: pluggable maps
- Define workflow to govern the process: Workflow Manager and Process Flows

