



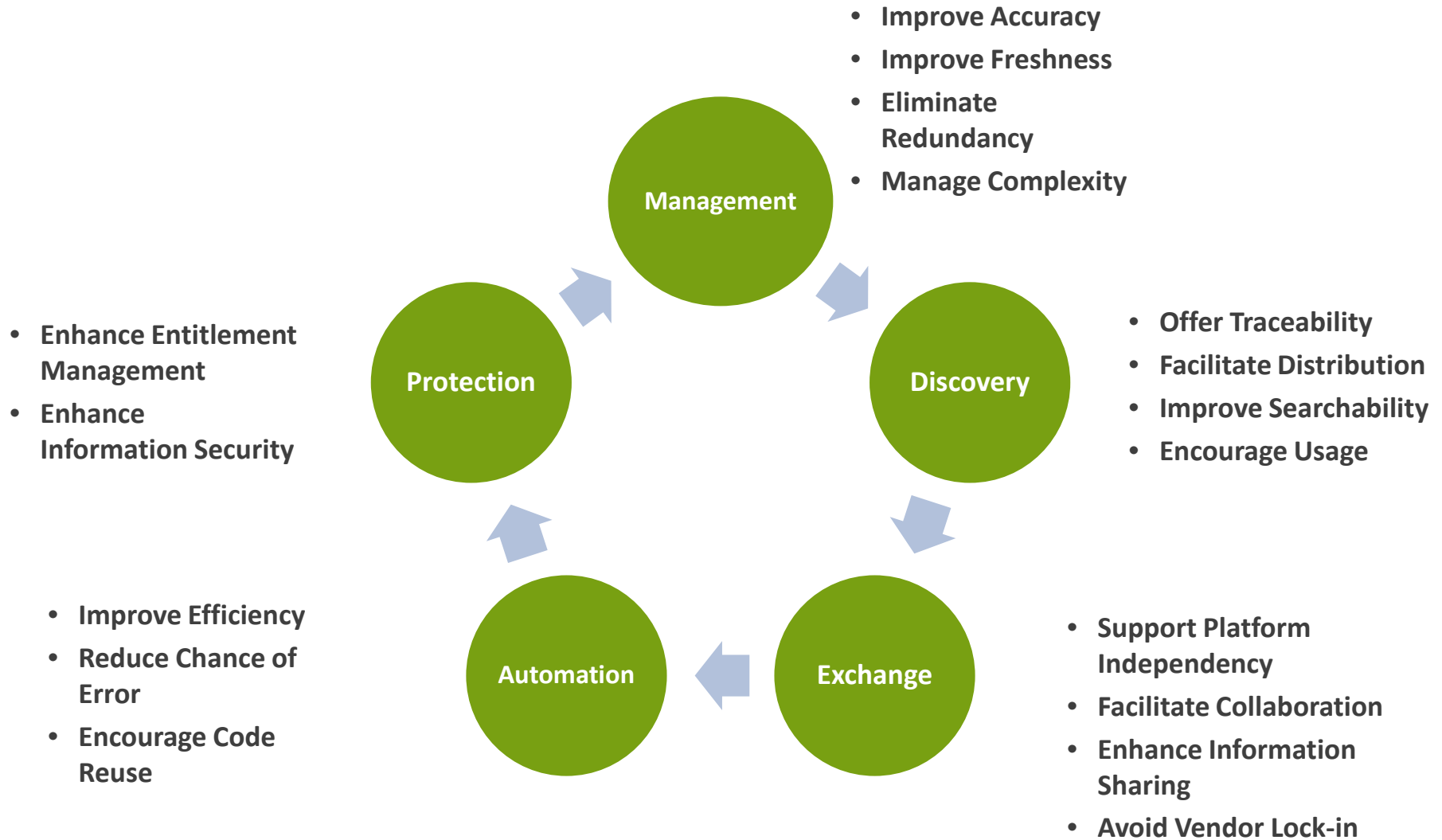
Using Study Metadata from Protocol and SAP to Drive Study Data Collection

PhUSE Single Day Event - Indianapolis

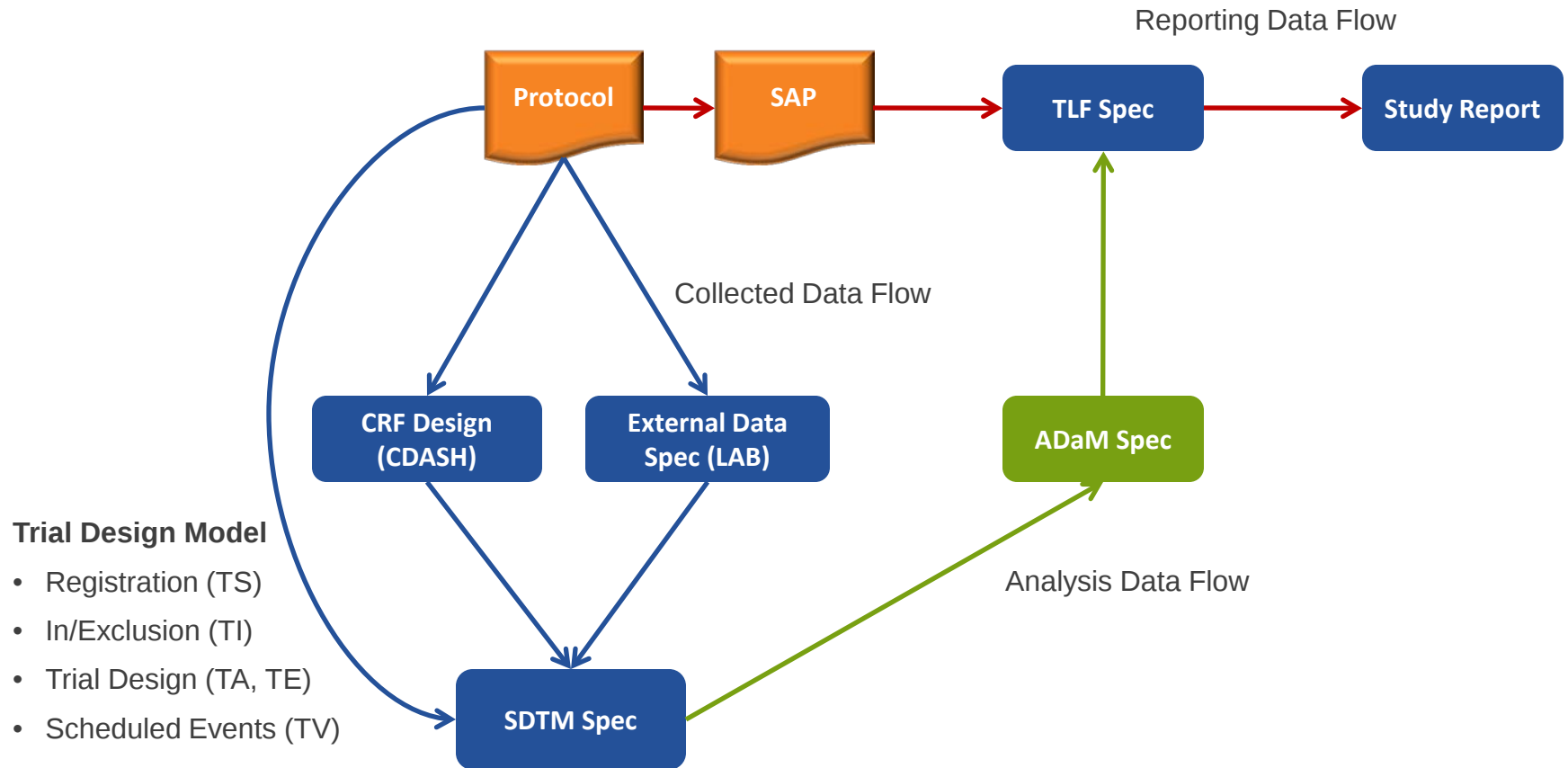
- **Traceability and Metadata**
- **HCL & EDETEK's Metadata Driven Approach for E2E Data Management**
- **Discussion**

- Traceability is a major concern for the FDA. Traceability means understanding the relationship between the results and the source. It facilitates transparency and is an essential component for building confidence in all stages of clinical information processing.
- Building traceability is establishing the path between an element and its immediate predecessor.
- **Metadata Traceability:** provides an explanation of how any downstream information has been created from its predecessors by describing **the algorithms and deviations used**.
- **Data Point Traceability:** provides **data lineage** of how down-stream records are related to the predecessor records(s): are the records originating from source or are these derived records.

IMPORTANCE OF METADATA



METADATA FLOW IN CLINICAL TRIAL DATA PROCESSING



Standards

- Establish Conceptual Models
- Establish CRF, SDTM, and ADaM Standards
- Store in MDR
- Governed and Maintained

Library

- Establish Libraries at Different Levels in Different Areas
- Library would grow to incorporate new scenarios
- CRF Libraries, SDTM Templates, ADaM Templates, TLF Templates

Design

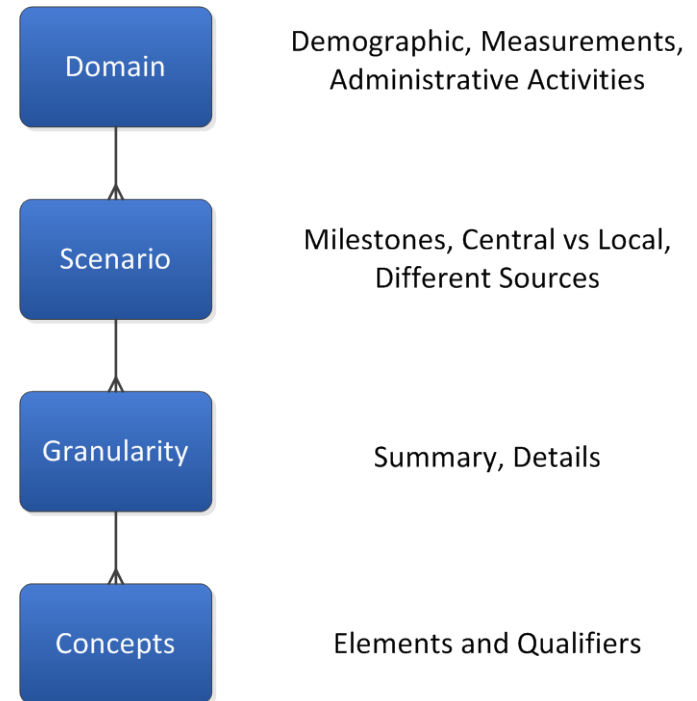
- Extract Metadata from Protocol and SAP
- Study Activities drive CRF Design
- Study Endpoints drive Variable Derivation
- Study Analysis drives TLF Design, ADaM Structure

Customize

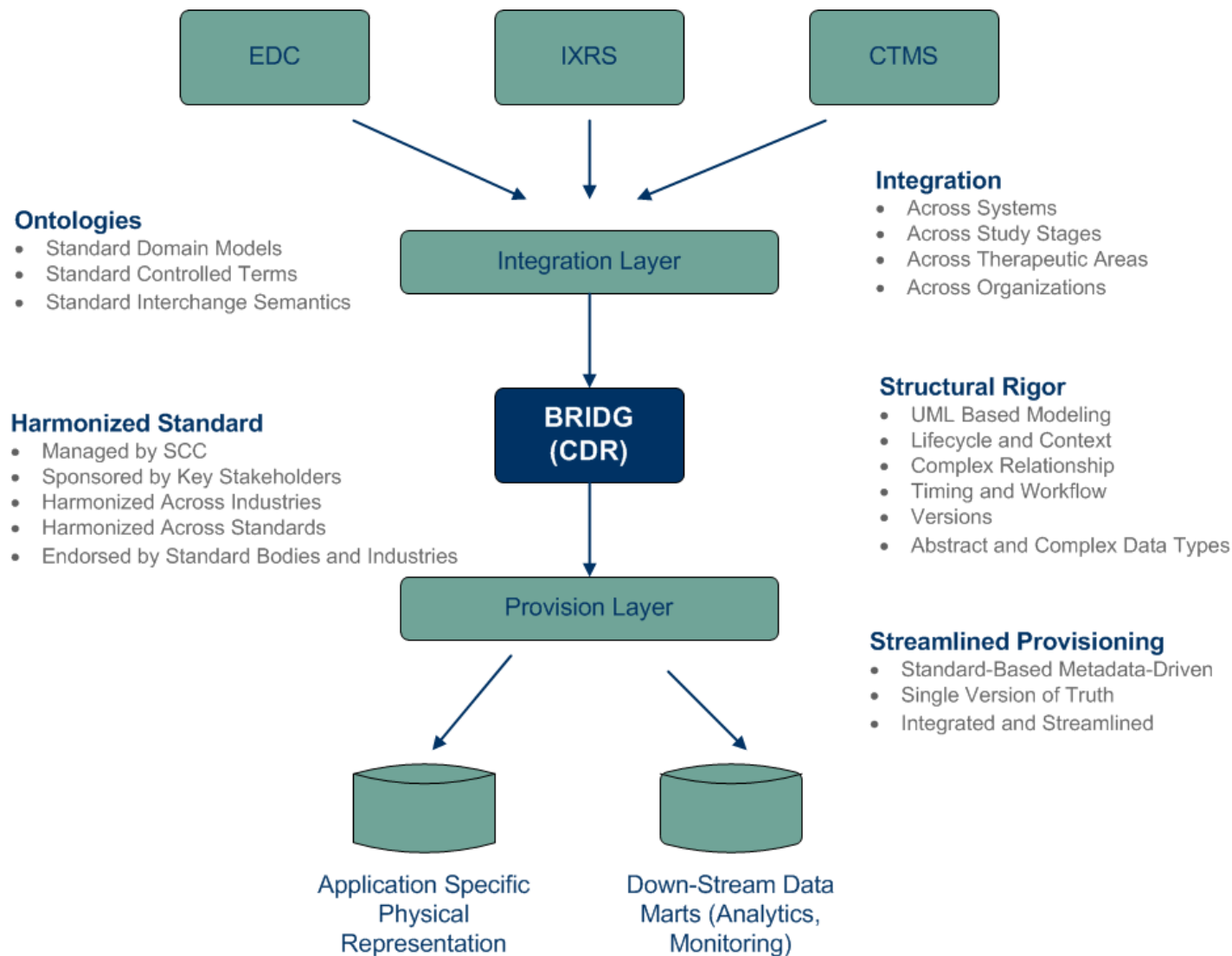
- Study Specific Configuration
- Conformance and Deviation can be monitored
- Mechanism exists to publish to Library

STEP 1 – ESTABLISH STANDARDS

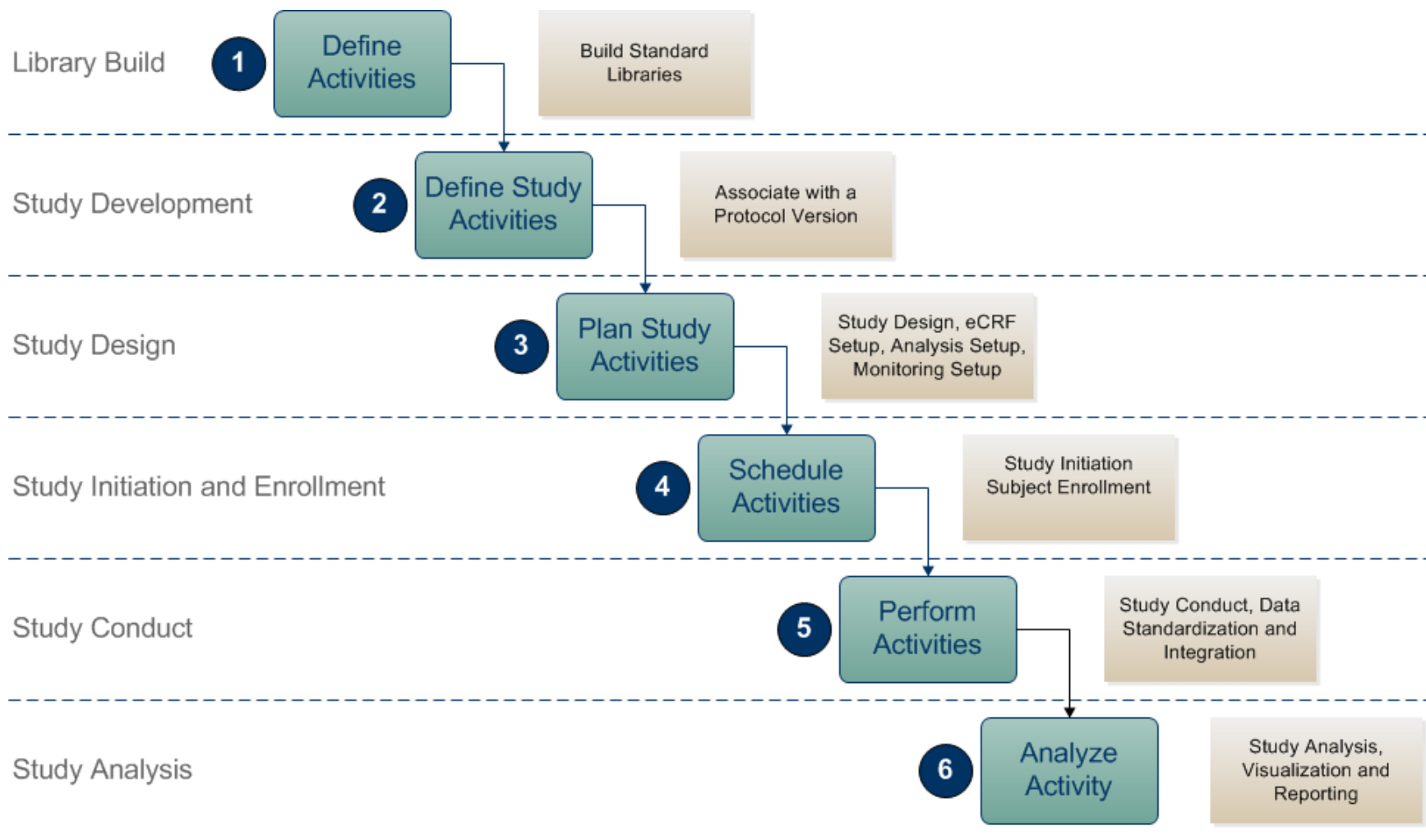
- **Define Conceptual Model**
 - Leverage BRIDG concept
 - Cover all data origins
 - Trial Design
 - Protocol
 - Collected Subject Data
 - Derived Subject Data
 - Cover all trial stages
 - Study Design (TDM -> SDTM)
 - Study Conduct (CDASH -> SDTM)
 - Study Analysis (ADaM)
 - Study Reporting (TLF)
 - Use Controlled Term to add semantics
- **Define CRF, SDTM, and ADaM Standards**



WHY BRIDG?



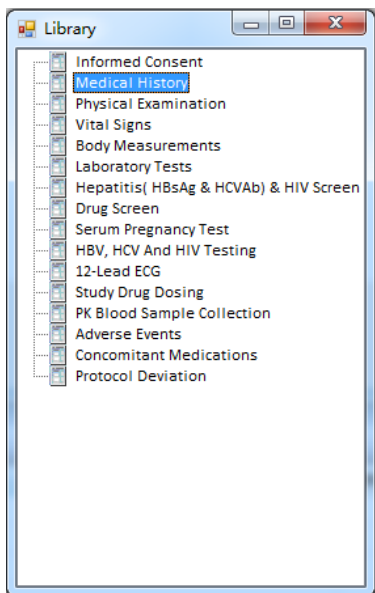
MANAGING STUDY USING BRIDG



STEP 2 – DEVELOP LIBRARIES

- **Libraries at Different Levels**
 - Company
 - Therapeutic Area
 - Program
- **CRF Library**
 - Standard CRFs
 - Cover different scenarios
- **SDTM Template**
 - Standard Structure
 - Standard raw -> SDTM mapping
- **ADaM Template**
 - Standard Structure
 - Standard Derivations
- **TLF Template**
 - Break-down to Standard Units

STEP 2 : CREATE COMMON LIBRARIES – eCRF & TLF



eCRF:

- Common Activities
- Associated Forms
- BRIDG or CDASH Mappings

MEDICAL HISTORY

Does the subject have any past clinically significant medical conditions?

- ☐ Yes
☐ No

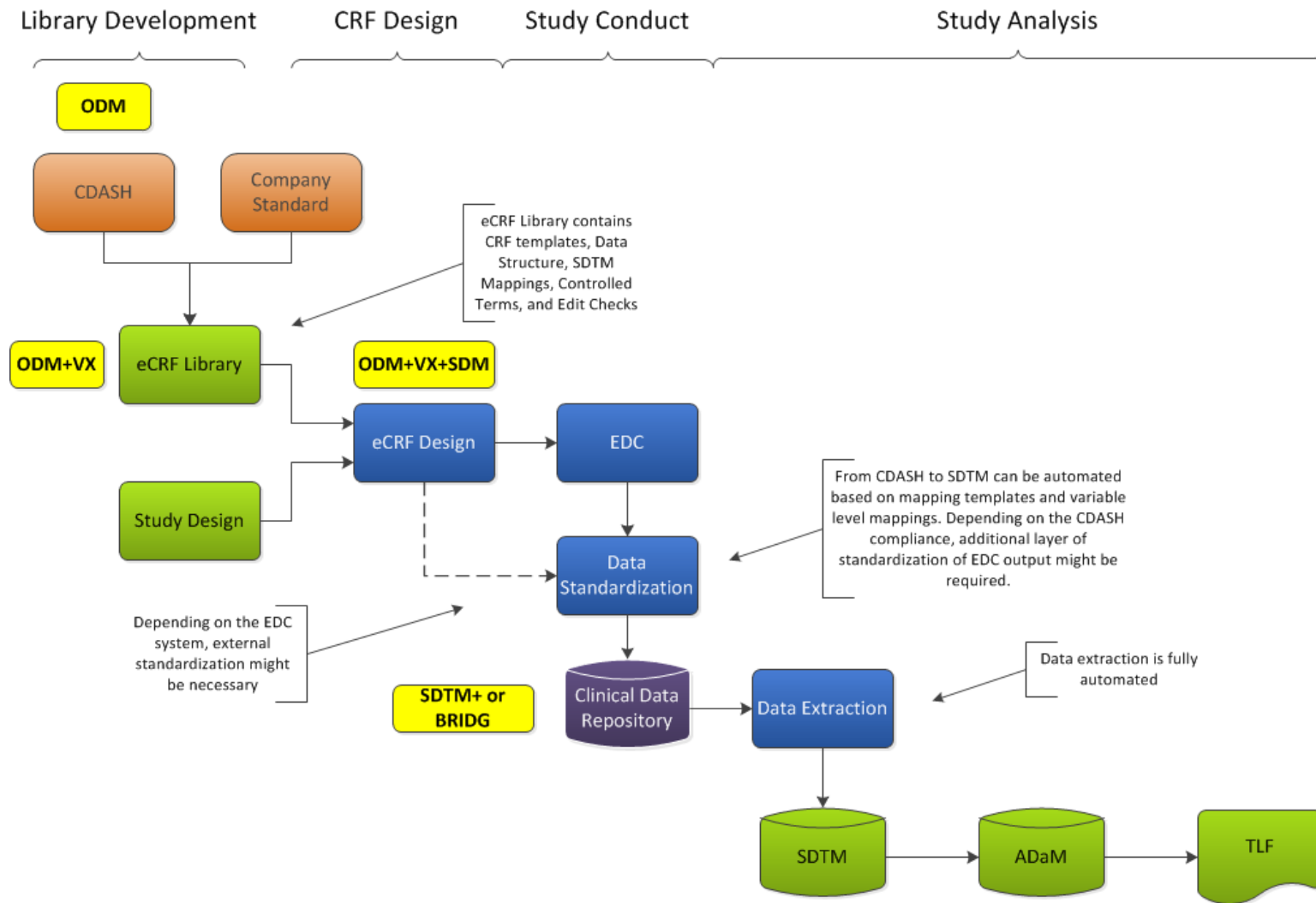
MH Number	Diagnosis/Procedure	Start Date (DD/MM/YYYY)	End Date (DD/MM/YYYY)	Ongoing
				☐

The screenshots show the SAS interface for creating TLFs. The 'Select template' window shows a list of templates on the left and a preview of a table on the right. The 'Statistic Formats Setting' window shows options for 'Simple Statistics' and 'Complex Statistics'. The 'TLF Professional' window shows a list of variables on the left and a preview of a table on the right.

TLFs

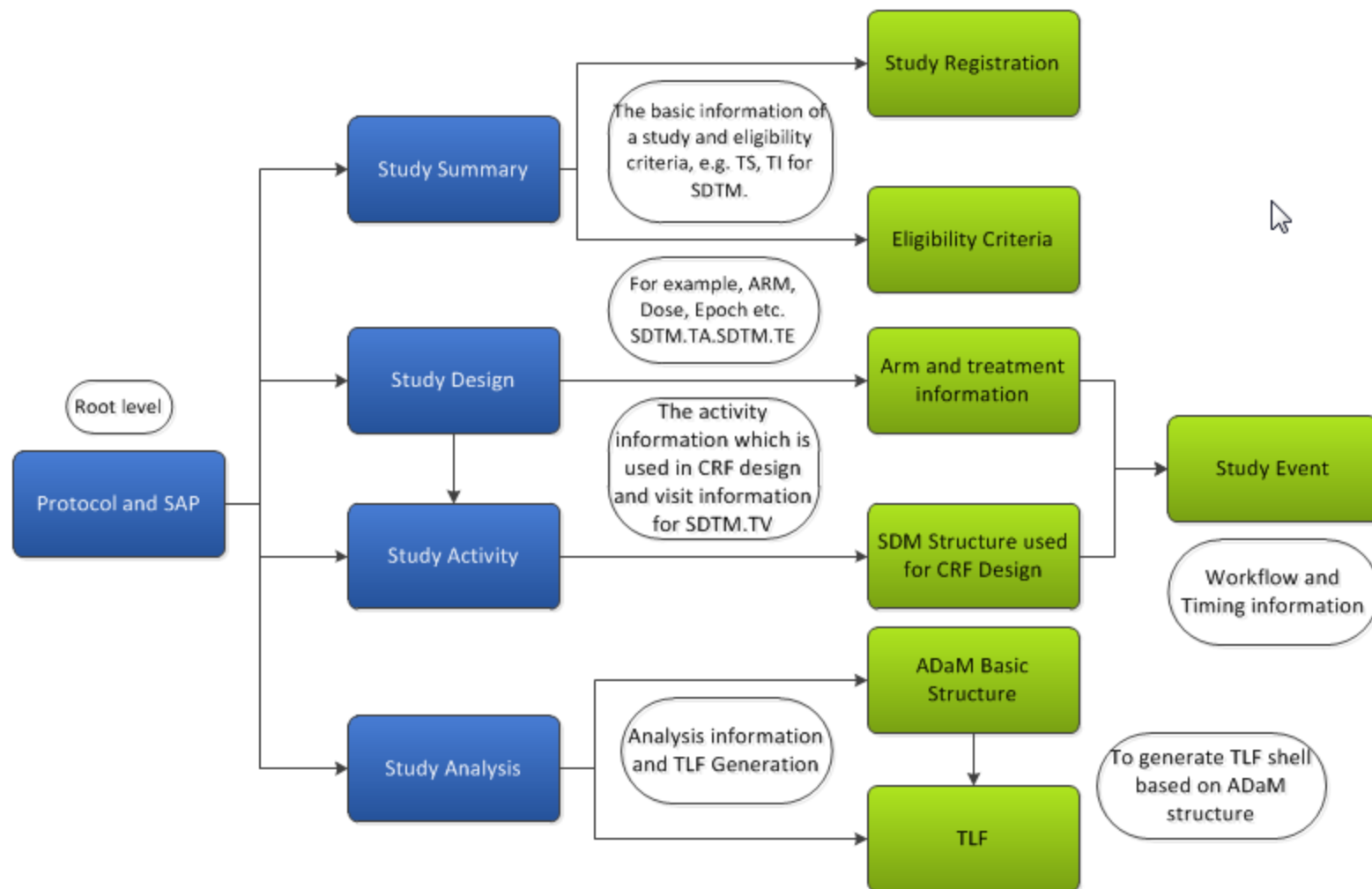
- Common Endpoints
- Reporting Templates
- BRIDG or ADaM Mappings

FLOW FROM DATA COLLECTION TO STUDY ANALYSIS



STEP 3 – STUDY DESIGN USING STUDY ARCHITECT

Study Architect Structure



STEP 3 : STUDY DEVELOPMENT - PROTOCOL

1.0 PROTOCOL SYNOPSIS

Study Title:	A Phase 1, Open-Label, Two-Period, Crossover Study to Evaluate the Relative Bioavailability of a New Tablet Formulation Versus the Current Capsule Formulation of PPI-668 Administered in Healthy Adult Volunteers
Protocol Number:	PPI-668-103
Study Phase:	Phase 1
Study Objectives:	Primary Objective: - To compare the relative bioavailability of PPI-668 when administered as a single oral dose in a tablet formulation vs. in a capsule formulation Secondary Objective: - To evaluate the safety and tolerability of single oral doses of PPI-668 in healthy subjects
Study Design and Duration of Treatment:	This will be an open-label, single-center, single-dose, crossover study. A total of 24 healthy subjects will be enrolled. On Day 1, under a 1:1 randomization scheme eligible volunteers, 12 subjects will receive a single oral dose of PPI-668 in the form of the current capsule formulation and 12 subjects will receive a single oral dose of the test tablet formulation of PPI-668, under fasted conditions. After a one week wash period, on Day 8, subjects will receive under fasted conditions the alternate PPI-668 formulation to the one they received on Day 1. Study medication will be administered approximately 0800 hours on Days 1 and 8 after an overnight fast of at least 10 hours. Screening will take place within 28 days before Study Day 1. Eligible subjects will be screened for eligibility criteria and will be enrolled into the study.

Registration

- Titles and Background Info
- Sponsors
- Study Design

Titles and Background Info

Organization's Unique Protocol ID * FDAAA XXX-XXX-XXX

Brief Title * FDAAA A Phase 1, Open-Label, Two-Period, Crossover Study to Evaluate the Relative Bioavailability of a New Tablet

SAP Name

Rulebook Name

Study Type * FDAAA **Interventional**

Brief Summary * FDAAA

Detailed Description

Previous Next OK Cancel

Table 7-1 Schedule of Study Procedures

	Screening ^a	Day -1	Day 1	Day 2	Day 3	Wash-out	Day 7	Day 8	Day 9	Day 10	Day 15 ± 3 days Follow-up
Written Informed Consent	X										
Medical History	X										
Complete Physical Examination	X										X
Brief Physical Examination		X					X				
Vital Signs ^b	X	X			X		X			X	X
Height	X										
Weight	X	X					X				
Hematology ^c	X	X					X				X
Chemistry ^d	X	X					X				X
Urinalysis	X	X					X				X
Urine Drug and Breath Alcohol Screening	X	X					X				
Pregnancy	X	X					X				X

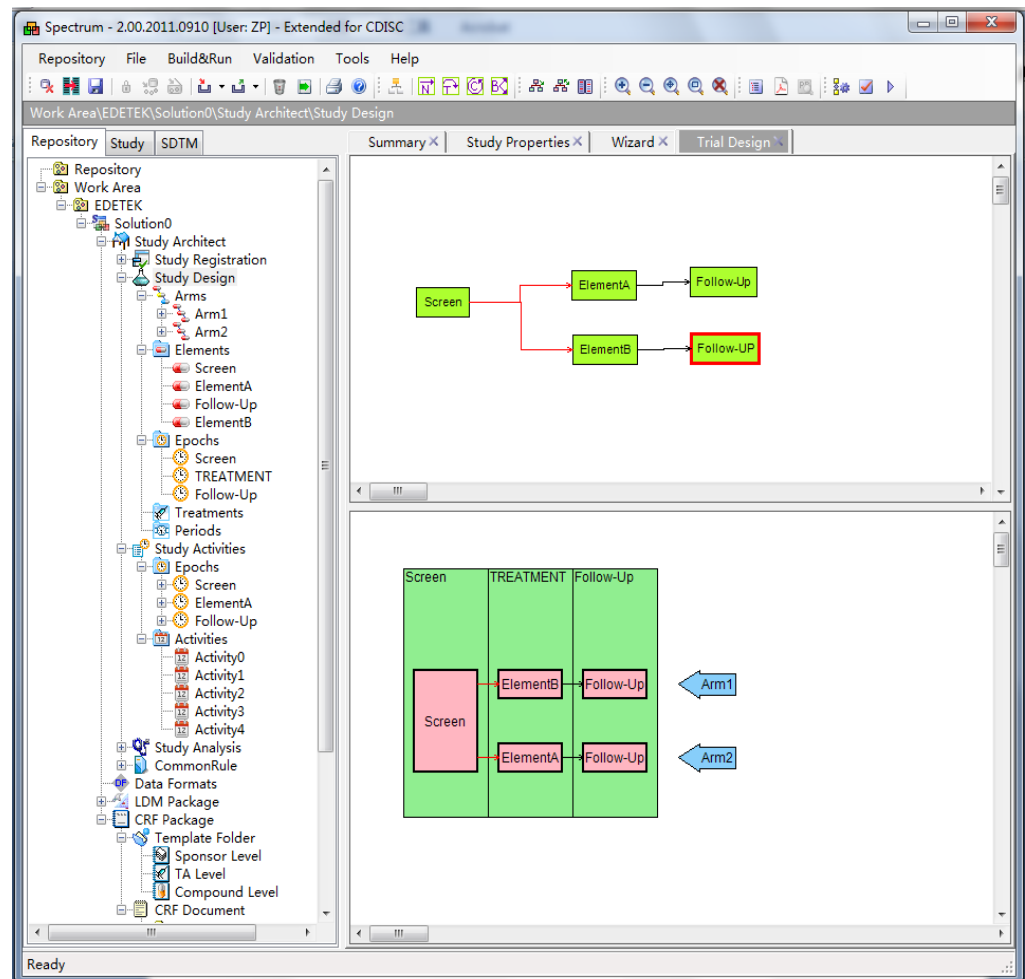
Study Protocol Versions are developed, and are associated with Defined Activities

STEP 3 : STUDY DESIGN – DATA COLLECTION

- Define Arms
- Define Epochs
- Define Segments
- Define Cells
- Define Study Events
- Define Workflow

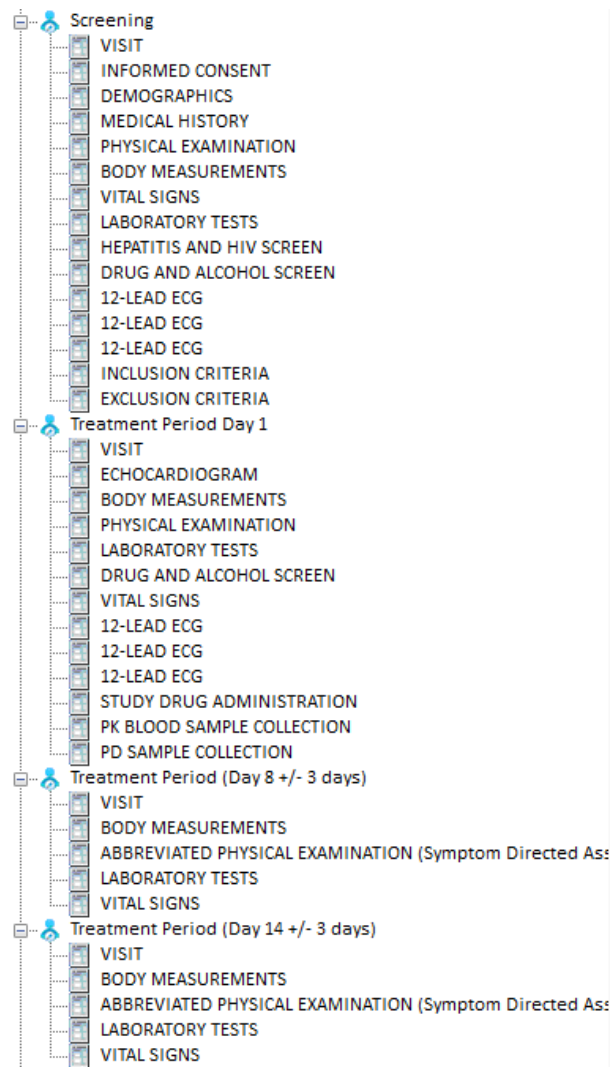


Study Design Model (SDM)

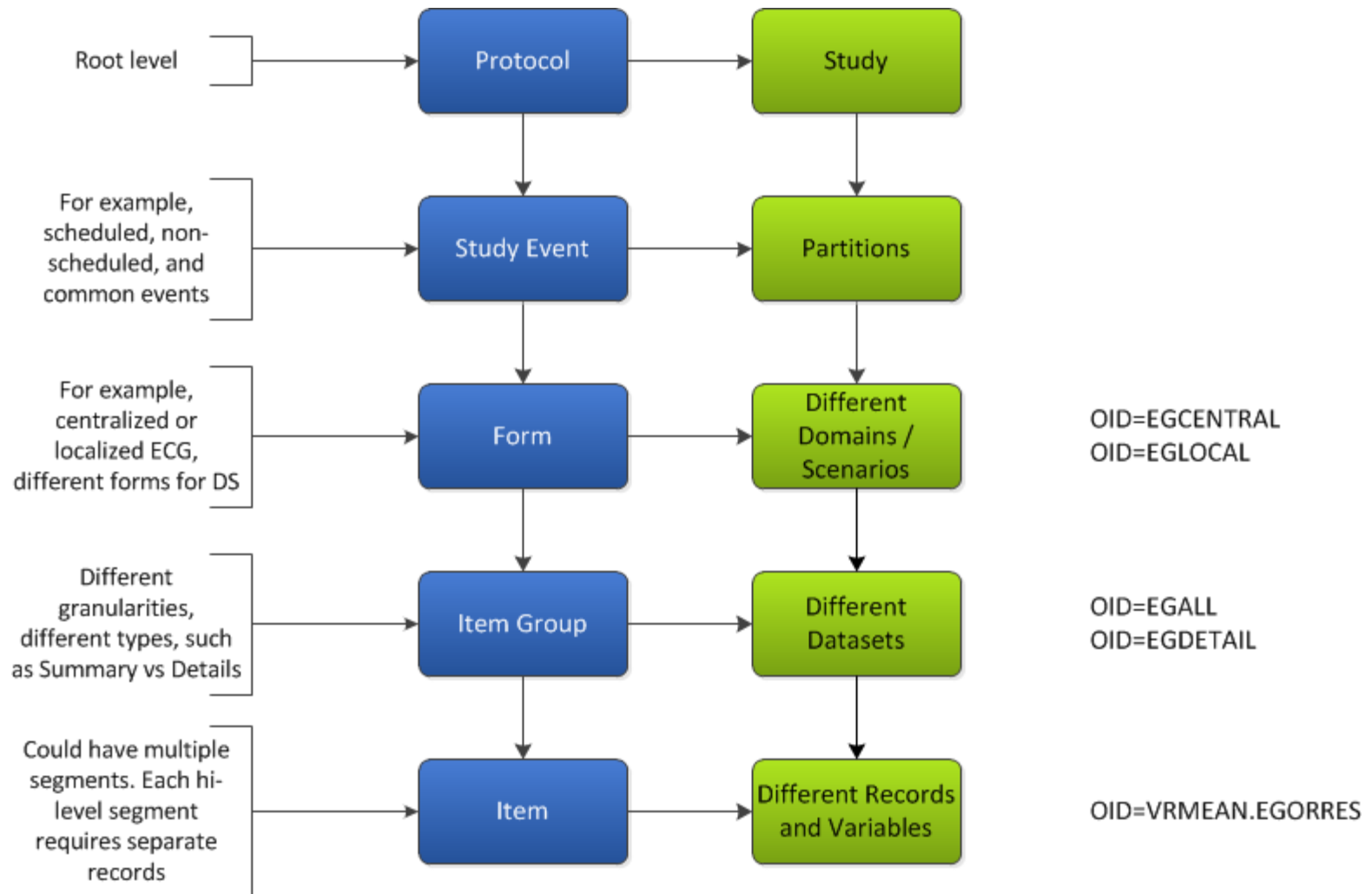


STUDY ACTIVITY FROM SA TO CRF

Epoch	Screening						
Activity/Event	Screening	Treatment Period Day 1	Treatment Period (Day 8 +/- 3 days)	Treatment Period (Day 8 +/- 3 days)	Treatment Period (Day 14 +/- 3)	Treatment Period (Day 21 +/-3)	Treatment Period (Day 28 +/- 3 day)
INFORMED CONSENT	X						
DEMOGRAPHICS	X						
MEDICAL HISTORY	X						
PHYSICAL EXAMINATION	X	X					
BODY MEASUREMENTS	X	X	X	X	X	X	X
VITAL SIGNS	X	X	X	X	X	X	X
LABORATORY TESTS	X	X	X	X	X	X	
HEPATITIS AND HIV SCREEN	X						
DRUG AND ALCOHOL SCREEN	X	X					X
12-LEAD ECG	X	X					
INCLUSION CRITERIA	X						
EXCLUSION CRITERIA	X						
ECHOCARDIOGRAM		X					
STUDY DRUG ADMINISTRATION		X					
PK BLOOD SAMPLE COLLECTION		X					X
ABBREVIATED PHYSICAL EXAMINATION (Symptom Directed Assessment)		X	X	X	X	X	X
PD SAMPLE COLLECTION		X					X
STUDY COMPLETION							
ADVERSE EVENTS							
CONCOMITANT MEDICATIONS							



STANDARDIZED CDASH OUTPUT STRUCTURE



- **Map to unique scientific concepts (logical model element)**
- **Item Group (IG) defines datasets**
 - Develop standard naming convention to cover different scenarios
 - Each name should represent unique set of contents
- **Item defines variable names and hierarchies**
 - Leveraging multiple segment names
 - Category, Sub-category, Test Code, Variable Name
 - Use CDISC Controlled Terminology to convey semantic information

DM CRF DESIGN

DEMOGRAPHICS

OID=F.DM

OID=IG.DM

Date of Birth

 - -

Day

Month

Year

OID=I.BRTHDAT

Sex

☐

Male

OID=I.SEX

☐

Female

Race (Check all that apply)

☐

White

☐

Black/African-American

OID=I.RACE

☐

Asian

☐

Native Hawaiian/Other Pacific Islander

☐

American Indian/Alaska Native

☐

Other Specify

OID=I.RACEOTH

Ethnicity

☐

Hispanic

OID=I.ETHNIC

☐

Not Hispanic

CDASH DM DATASET OBTAINED

BIRTHDAT	ETHNIC	RACE1	RACE2	RACE3	RACE4	RACE5	RACE6	RACEOTH	SEX
23-MAR-1968	2					1			M
27-DEC-1982	2					1			M
15-DEC-1987	2			1					M
06-JAN-1976	2			1					M
14-AUG-1990	2					1			M
13-JAN-1986	1					1			M
20-SEP-1994	2			1					M
25-JUL-1976	2			1					F

EG CRF DESIGN

12-LEAD ECG

OID=F.EG

In triplicate after 5 minutes of resting in the supine position.

OID=IG.EGPERF

☐ Not Done

OID=I.EGPERF

OID=IG.EGDETL Date of ECG	- - Day Month Year	OID=I.EGDAT
Actual Time	: Hour Minute	OID=I.EGTIM
Ventricular Rate	(beats/min)	OID=I.VRMEAN.EGORRES
PR Interval	(msec)	OID=I.VRMEAN.EGORRES
RR Interval	(msec)	OID=I.RRMEAN.EGORRES
QRS Interval	(msec)	OID=I.QRSDUR.EGORRES
QT Interval	(msec)	OID=I.QTC.EGORRES
QTcF Interval	(msec)	OID=I.QTcF.EGORRES

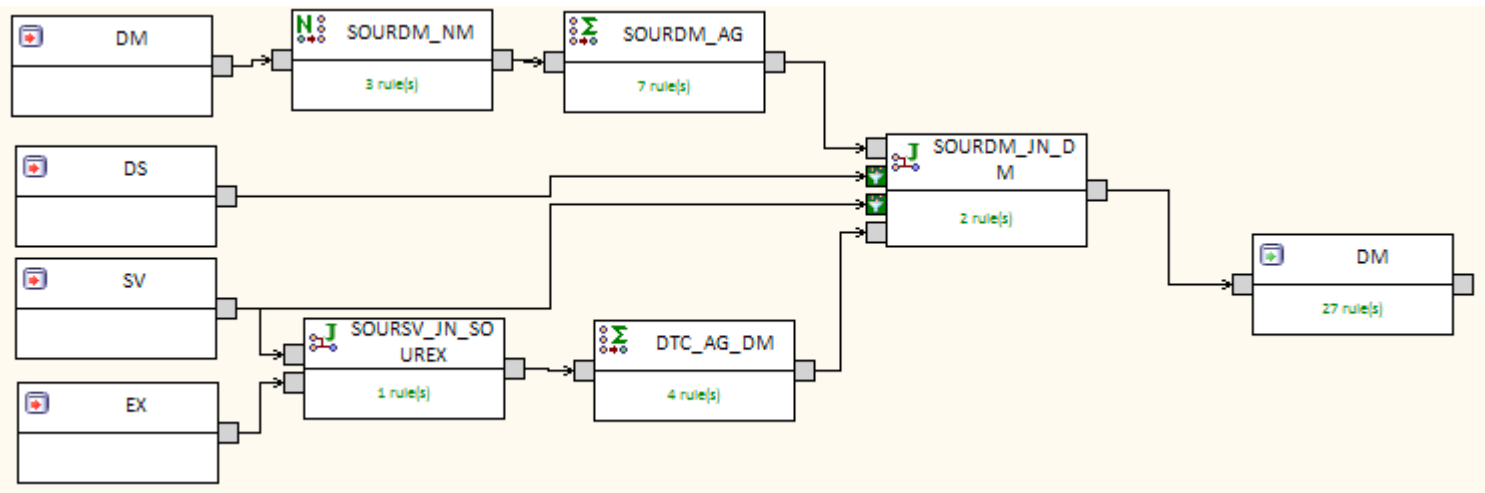
CDASH EG DATASET OBTAINED

EGDAT	EGORRES	EGORRESN	EGORRESU	EGTEST	EGTIM
28-MAR-2014		165	(msec)	PR Interval	0918
28-MAR-2014		985	(msec)	RR Interval	0918
28-MAR-2014		90	(msec)	QRS Interval	0918
28-MAR-2014		410	(msec)	QT Interval	0918
28-MAR-2014		412	(msec)	QTcF Interval	0918
28-MAR-2014	1	.			0918
28-MAR-2014		60	(beats/min)	Ventricular Ra	0918
28-MAR-2014		178	(msec)	PR Interval	0920
28-MAR-2014		966	(msec)	RR Interval	0920
28-MAR-2014		88	(msec)	QRS Interval	0920
28-MAR-2014		411	(msec)	QT Interval	0920
28-MAR-2014		415	(msec)	QTcF Interval	0920
28-MAR-2014	1	.			0920
28-MAR-2014		62	(beats/min)	Ventricular Ra	0920
28-MAR-2014		170	(msec)	PR Interval	0922
28-MAR-2014		979	(msec)	RR Interval	0922
28-MAR-2014		86	(msec)	QRS Interval	0922
28-MAR-2014		407	(msec)	QT Interval	0922
28-MAR-2014		409	(msec)	QTcF Interval	0922
28-MAR-2014	1	.			0922
28-MAR-2014		61	(beats/min)	Ventricular Ra	0922

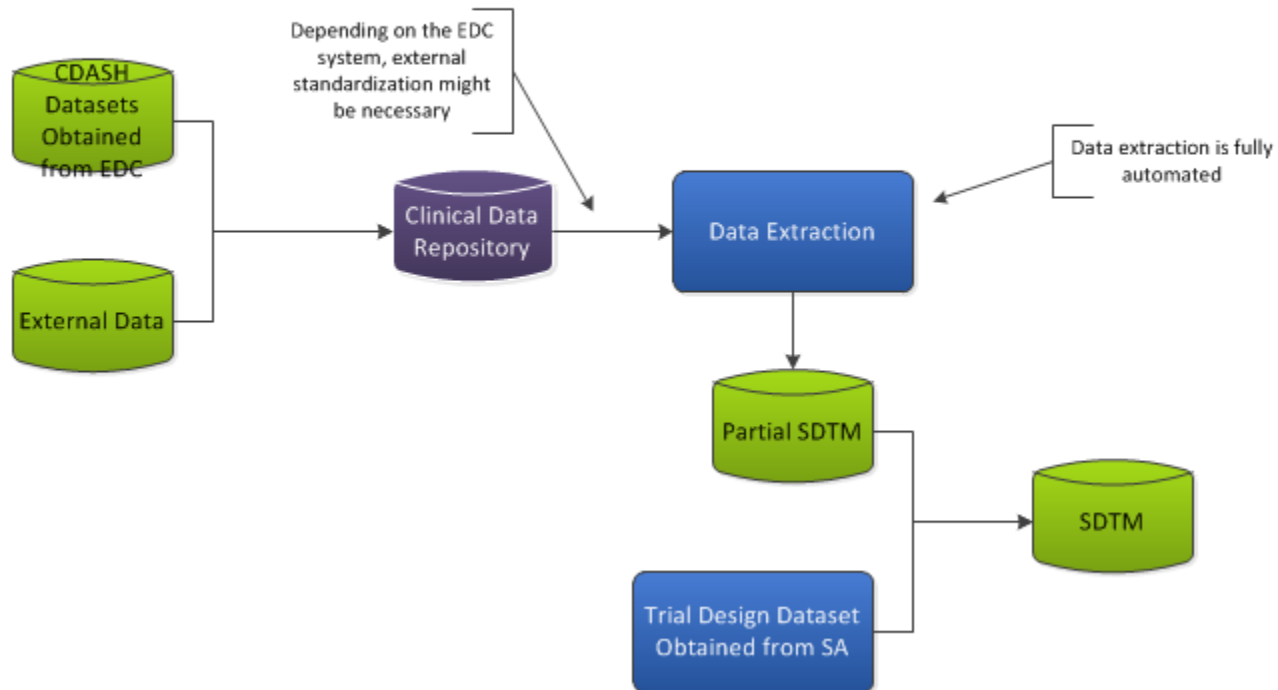
- **Match IG-based EDC Output with Partition-Level Templates**
 - Scenario based templates
 - Auto-matching based on available IG-based outputs
 - Auto-Pruning of extra template variables
- **Match Domain-Level Templates**
 - All partitions are merged together
 - Merged results are matched against domain-level templates
 - Finalization stage is applied based on domain class
 - Baseline calculation
 - --SEQ assignment
 - SUPPQUAL creation

- **Non-standard derived variables have to be defined based on SAP or mapping standards.**
 - Baseline Definition
 - Default: Last non-missing observation before dosing
 - Average of repeated measurements
 - Reference Dates
 - First Dose/Last Dose Dates
 - Treatment Emergent AE
 - Last Participation Date
 - Unit Systems for Lab Standardization
- **Variables of Protocol Origin have to be defined**
 - Trial Design
 - Other protocol specified information not available elsewhere
- **These are specified at study design**

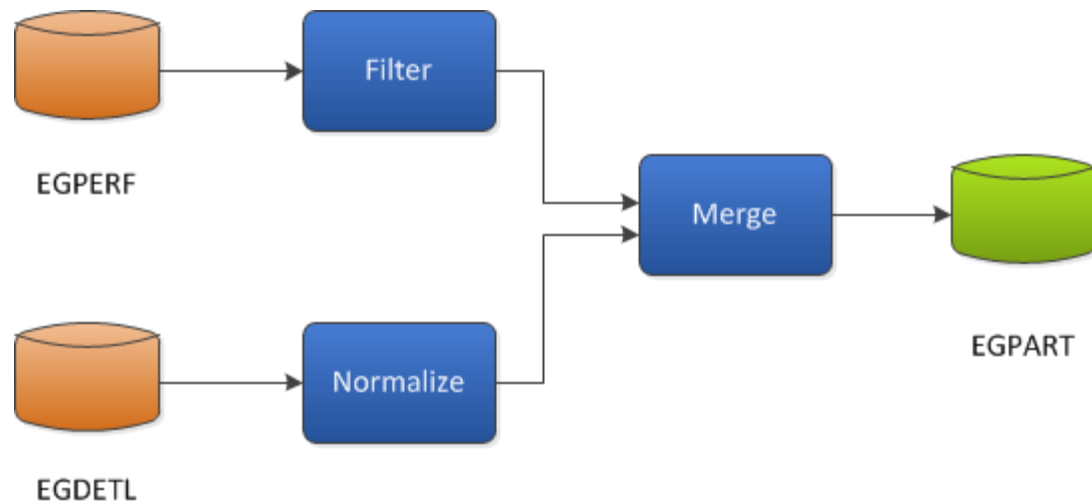
CREATING SDTM/ADAM TEMPLATES



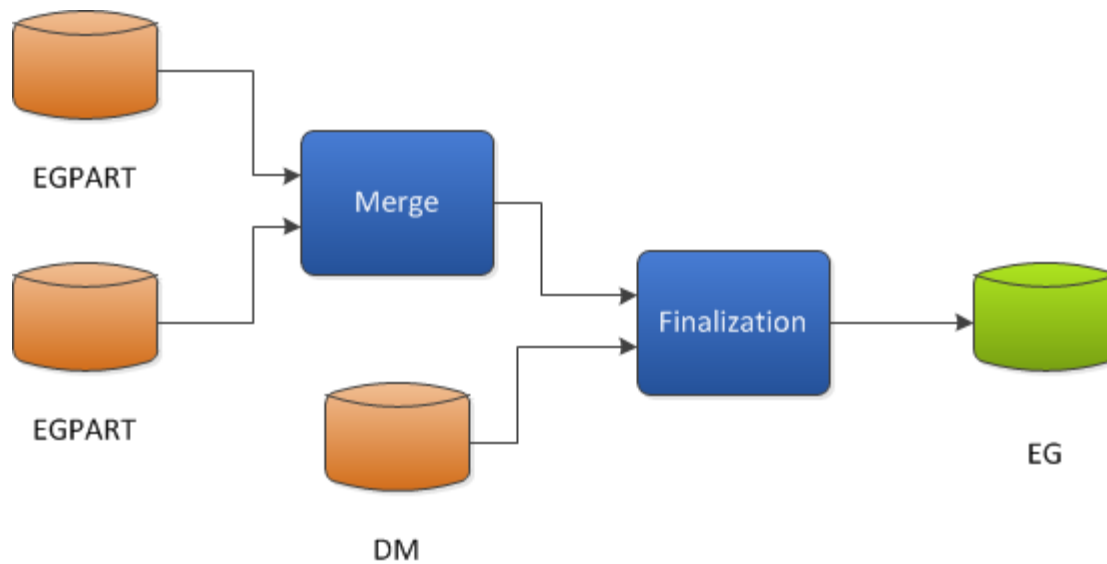
CDASH - SDTM Dataset Generation



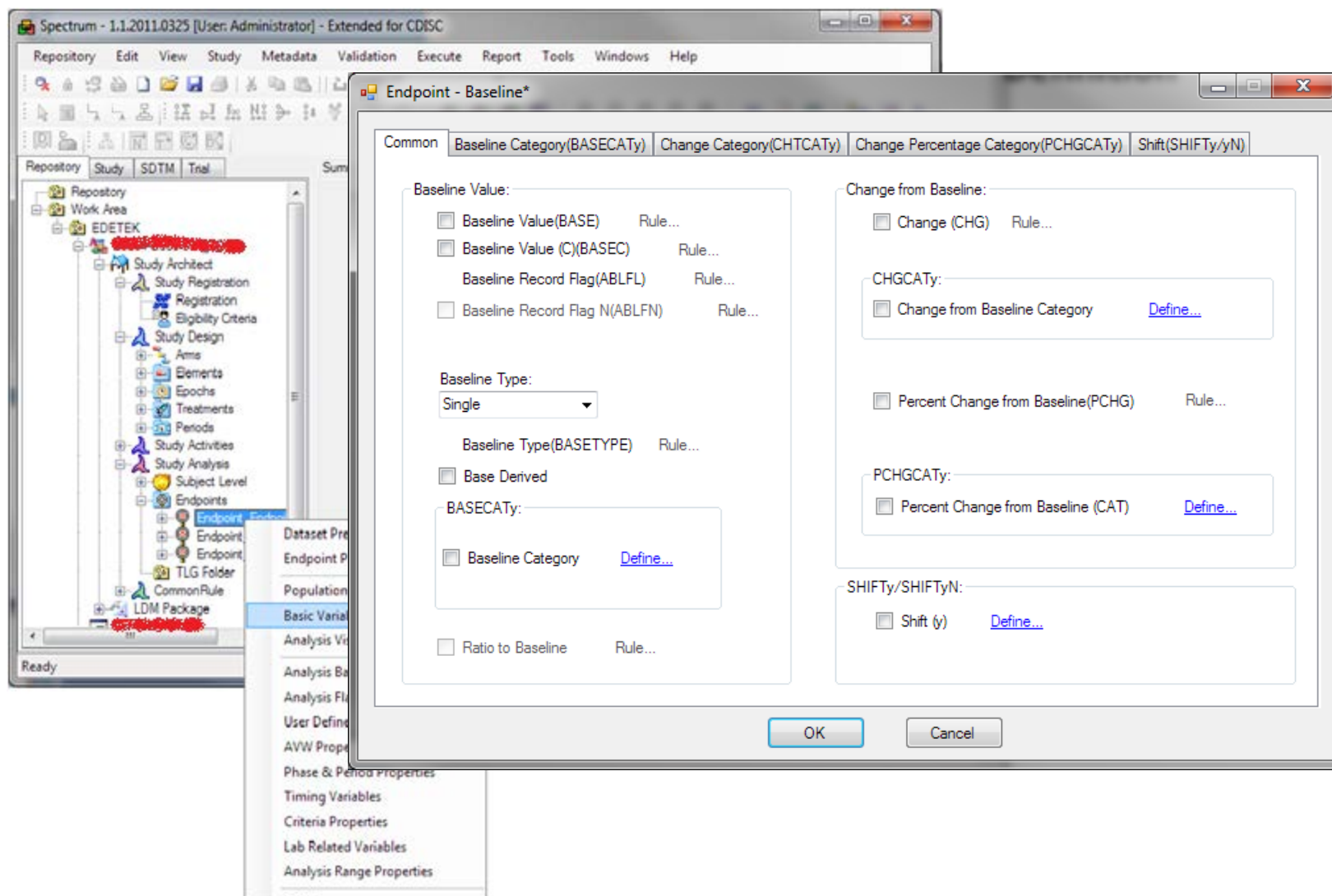
PARTITION LEVEL TEMPLATES



DOMAIN LEVEL TEMPLATES



STEP 3 : STUDY DESIGN - ANALYSIS



STUDY 3 : STUDY DESIGN - REPORTING

The screenshot illustrates the workflow for creating a new table in the software. The top toolbar includes buttons for New, Import1, Import2, Export1, Export2, Spec, Shell, Macros, and SAS Script. A yellow arrow points to the 'New' button, which triggers a dropdown menu with 'Demo Spec', 'Table', and 'Shell'. Another yellow arrow points to the 'Table' option, which opens the 'New Table' dialog.

The 'New Table' dialog shows the following fields and options:

- Table Name:** Table_0
- Components:** A list of components with checkboxes: ID, Title, Population, Data Source, Columns, Reporting Item, and Program Name. All are checked. Buttons for 'All' and 'Reverse' are present.
- Buttons:** OK and Cancel.

The 'Main Panel' window shows the 'Demo Characteristics' tab. It contains the following fields and tables:

- ID:** 34.1.2
- Title:** Demographics
- Population:** Safety Population
- Data Source:** ADSL
- Columns:** A table with 3 columns: Column1, Column2, and Column3. Each column has a 'label' and a 'type'.
- Reporting Item:** A table with 3 columns: name, label, and stats. Each row has a 'name', 'label', and 'stats'.
- Program Name:** A text field.
- Buttons:** Add, Delete, and Update.

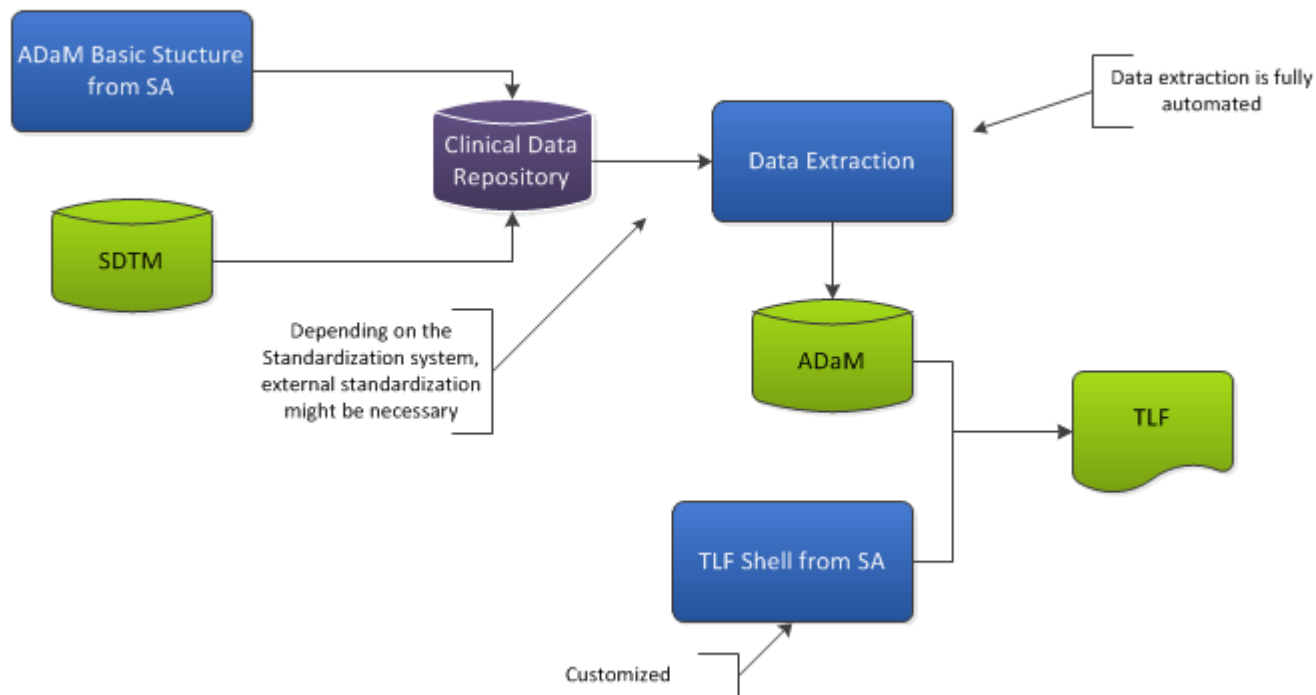
Column	label	type
Column1	Demographic	rawlabel
Column2	Statistic	statistics
Column3		

name	label	stats
AGE	Age(years)	n_mean_sd_median...
GENDER		
DISP_GENDER		

Table 14.1.2			
Demographics			
Safety Population			
Demographic	GROUP A	GROUP B	GROUP C
Age (years)			
n	xxx	xxx	xxx
Mean (Sd)	xx.x (xx.x)	xx.x (xx.x)	xx.x (xx.x)
Min, Max	xx, xx	xx, xx	xx, xx
Median	xx	xx	xx
Gender			
Female	xx(x.xx)	xx(x.xx)	xx(x.xx)
Male	xx(x.xx)	xx(x.xx)	xx(x.xx)
Race			
American Indian or A	xx(x.xx)	xx(x.xx)	xx(x.xx)
Black	xx(x.xx)	xx(x.xx)	xx(x.xx)
White	xx(x.xx)	xx(x.xx)	xx(x.xx)
Asian	xx(x.xx)	xx(x.xx)	xx(x.xx)
Native Hawaiian or o	xx(x.xx)	xx(x.xx)	xx(x.xx)

AUTOMATED ADaM & TLF CREATION

Study Architect-ADaM-TLF



After the clinical data repository which contains rule template as the SDTM repository in page 14-16, ADaM datasets are generated finally.

- **ADaM structure is generated to support TLF creation**
 - Analysis parameter variables
 - Analysis group variables
 - Analysis population variables
 - Analysis visit windowing variables
 - Analysis flag variables
 - Analysis enabling variables
- **Traceability to SDTM is established through ADaM templates**
 - Derivations
 - Macros – with metadata wrapper
- **Traceability to data collection is established through SDTM templates**

Table 14.1.2+
Demographics+
Safety Population+

Demographic+	Statistic+	KD025+ (N=8)+	Total+ (N=8)+
Age (years)+	N+	8+	8+
	Mean (SD)+	31.0 (8.88)+	31.0 (8.88)+
	Median+	29.5+	29.5+
	Min, Max+	19.0, 46.0+	19.0, 46.0+
Gender+			
Male+	n (%)+	7 (87.50%)+	7 (87.50%)+
Female+	n (%)+	1 (12.50%)+	1 (12.50%)+
Race+			
American Indian or Alaska Native+	n (%)+	0 (0.00%)+	0 (0.00%)+
Asian+	n (%)+	0 (0.00%)+	0 (0.00%)+
Black+	n (%)+	4 (50.00%)+	4 (50.00%)+
Native Hawaiian or other Pacific Islander+	n (%)+	0 (0.00%)+	0 (0.00%)+
White+	n (%)+	4 (50.00%)+	4 (50.00%)+
Other+	n (%)+	0 (0.00%)+	0 (0.00%)+
Ethnicity+			
Hispanic or Latino+	n (%)+	1 (12.50%)+	1 (12.50%)+
Not Hispanic or Latino+	n (%)+	7 (87.50%)+	7 (87.50%)+
Weight (kg)+	N+	8+	8+
	Mean (SD)+	78.8 (12.84)+	78.8 (12.84)+
	Median+	80.6+	80.6+
	Min, Max+	58.2, 96.4+	58.2, 96.4+