

Using a Standards Library to support end-to-end CDISC automation

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Meet the Speaker

Stuart Malcolm

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Stuart Malcolm is Head of Standards, Efficiency and Automation at Veremed where he is responsible for the delivery of software platforms, tools, and techniques to optimise the delivery of clinical trial analysis projects

Stuart has over 25 years experience delivering software-based solutions in clinical trials, and has previously worked in telecoms, finance and media.

Stuart is co-chair of PHUSE Emerging Trends & Technology Working Group

"To define the future, one must study the past"

TFL Automation using CDISC Analysis Results Metatadata (ARM) PharmaSUG 2019

A Language-Orientated Approach to CDISC Metadata-Driven Automation CDISC Interchange 2023

PharmaBUG 2019 - Paper AD-203 Large-scale TFL Automation for regulated Pharmaceutical trials using CDISC Analysis Results Metatadata (ARM) Stuart Macion, Frontier Scienco (Scotland) Ltd.

BSTRACT

The creation of a Clinical Study Report (CSR) for Phase IIIII Pharmaceutical clinical trial involves the production of several hundred Tables, Figures and Listings (TFL). This can be a time-consuming activity when each TFL is programmed manually.

While some TFL are common for many studies, there is always a requirement to create study-specific TFL in addition, CDISC Analysis Results Metadata (ARM) Define.xml is often requested at the end of the rial.

his paper outlines an approach to TFL automation that involved creation of the CDISC Analysis Results fetadata at the start of the process, not the end, and uses this metadata to generate the TFL. SAS program structure is described that allows standard TFL to be created while also providing explicit to easily incorporte study-specific analyses.

INTRODUCTION

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The development of TFL is primarily a collaboration between Clinical Statisticians and Statistical Programmers, as shown in Figure 1 below. The Programmer's primary role is to create and QC TFL, and the Statistican's role is to specify the TFL – using a combination of a SAP (Statistical Analysis Plan) and Mox't Table Stote.

igure 1 Users and uses of an automated TFL system



In an automated TFL system, the Metadata performs a similar role as programming specification for traditional measurements of the TL Typically the metadata is created by the Programmer and reviewed by the Statistician. The development of the Metadata is the main area of collaboration between Programmers and Statisticians, and one of the benefits of using the COIISC APM statisticand for the Programmers and Statisticians, and one of the benefits of using the COIISC APM statisticand for the





Automation of TFL Generation using CDISC 360 Enriched Metadata CDISC EU Interchange 2020



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• The views and opinions expressed in this presentation are those of the author(s) and do not necessarily reflect the official policy or position of CDISC.

• The author(s) have no real or apparent conflicts of interest to report.



End-to-End Vision



What is 360?

360 is the CDISC initiative aimed at implementing standards as linked metadata with a conceptual foundation providing the additional semantics needed to support metadata driven-automation across the end-to-end clinical research data lifecycle.

New software tools will be able to consume this new metadata to ease standards implementations while increasing data processing efficiencies.

Standards-based metadata-driven automation is a critical component in realizing the primary benefits expected of the CDISC standards: substantially improved efficiency, consistency, and re-usability across the clinical research data lifecycle.

These benefits drive the return on investment in the CDISC standards implementations expected by CDISC stakeholders.

"SUBSTANTIALLY IMPROVED

- EFFICIENCY
- GNSISTENCY
- REUSABILITY

ACROSS THE LUNICAL RESEARCH LIFECYCLE



Death Stars are expensive...



...take a lot of time and effort to build, and..



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... are static and hard to change!





What if we could build trials out of components?..













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It's not metadata... it's a model

It's not metadata... it's a model

Model

2 Model Co the SDTN



The SDTM provides a gene animal studies. The model is

animal studies. The model is built around the concept of *observations*, which consist of discrete pieces of information collected during a study. Observations normally correspond to rows in a dataset. A *domain* is

a collection of ol adverse event of clinical trial.

The primary pur medical devices. Within those clas observation class <u>Observation Clas</u> purpose Domain

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Standards Library





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An agile approach to standards management



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https://imranqureshi.co.uk/devops-for-dummies-3rd-ed-notes/

A New Hope? Agile, Decentralised and Automated





Thank you!

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