Business & Decision Life Sciences

ADaM Conversions: The Good, The Bad and the Ugly

Jessica Minkue Mi Edou

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Introduction

Business & Decision Life Sciences' experiences in ADaM conversion projects applying:

- Linear approach for a legacy study
- Parallel approach for Pooled datasets

Objective

Describing the good, bad and ugly experiences gained during the execution of these projects



ADaM Conversions: The Good, the Bad and the Ugly





Good understanding of the relationship between

- Analysis results
- Analysis datasets
- SDTM domains

The core of any ADaM conversion is the SDTM/ADaM traceability, this can be found at two levels:

- Metadata traceability: finding a relationship between an analysis result and analysis dataset(s), or a relationship of the analysis variable to its source dataset(s) and variable(s)
- Data point traceability: finding the predecessor record(s)



Table 1 Demographic Data - Per-Protocol Treatment 1 Treatment 2 Baseline body mass index (BMI) [kg/m**2] 167 167 Mean 29.08 29.04 4.84 4.80 SD Min 20.3 16.0 Median 28.69 28.47 Max 40 1 41 2 Baseline BMI (categorical) [N (%)] 41 (24.6%) 71 (21.1%) <25 ka/m**2 25-<30 kg/m**2 60 (35.9%) 130 (38.7%) Patient Demographics – Part I >=30 kg/m**2 66 (39.5%) 135 (40.2%) atient X5 Page 4 Visit Date 11-Dec-2007 Blank Comment CAT = "PROTOCOL MILESTONE PATIENT DEMOGRAPHICS - Part I
 BMI
 BMIGR1
 BMIGR1N
 BMIGR2
 BMIGR2N

 22.777777778
 <30 kg/m**2</td>
 1
 25<<30 kg/m**2</td>
 2
 STUDYID USURIO SUBJID Informed consent was obtained 9999-0001 9999-0001-000001 000001 1 25<30 kg/m**2 9999-0001 9999-0001-000002 000002 25 503615702 <30 ko/m**2 1.25<30kg/m**2 9999-0001 9999-0001-000003 26.175194521 <30 kg/m**2 Gender 1 = male, 2 = female 000003 1 25-<30 kg/m²² 9999-0001 9999-0001 9999-0001-000004 000004 35.15625 >=30 kg/m**2 2 >=30 kg/m**2 Date of birth AGE years AGE Age 30.968858131 >=30 kg/m**2 9999-0001-000005 000005 2 >=30 kg/m**2 (Age is automatically 9999-0001 99999-0001-000006 000000 39.697163916 >=30 kg/m** calculated when ecreen in 8 9999-0001 9999-0001-000007 000007 25.826446281 <30 kg/m**2 1 25-<30 kg/m**2 9 9999-0001 10 9999-0001 2 >=30 kg/m^{**}2 2 >=30 kg/m^{**}2 saved and closed) 9999.0001.000009 000008 30.103806228 >=30 kg/m**2 9999-0001-000009 000009 32.280962683 >=30 kg/m**2 Height or 9999-0001 9999-0001 9999-0001-000010 000010 28.876133787 <30 kg/m~2 1 25-c30 kg/m**2 Weight kg 9999-0001-000011 000011 29.372397383 <30 kg/m**2 1 25-<30 kg/m**2 Waist circumference 9999-0001 9999-0001 9999-0001-000012 000012 26.714852608 <30 kg/m**2 1 25-<30 kg/m**2 14 9999-0001-000013 000013 32.718619869 >=30 kg/m**2 2 >=30 kg/m²2 15 9999-0001 16 9999-0001 9999-0001-000014 000014 28.719723183 <30 kg/m**2 1 25-(30 kg/m**2 ORRES / VSORRESU where STESTCD = "HEIGHT" 9999-0001-000015 000015 32 270420377 >=30 kg/m**2 2 >=30 ko/m¹¹2 WEIGHT", "WAIST" Subject-Level Analysis Dataset Dataset (ADSL) AnalysisladsLapt Vital Signo Dataset (VS Pole The STUDYID variable has a fixed former 'XXXX-IDENTIFIER YYYY', where 'XXXN indicates the 4-digit compound code and the 'Y'YY' the 4-digit study code IDENTIFIER The STUDYID vanishes as fixed former XXXX-YYYY, where XXXX indicates the 4-digit compound code and the YYYY the 4-digit mady code STUDVID Snuty Identifier lest DM STUDYD Study Mentiler test Protocol CRJ Page 1 The USUBJID variable has a fixed format 'XXXX-IDENTIFIER YYYY-ZZZZZ, where 'XXXX' indicates the 4-digit compound code, 'YYY' the 4-digit study code and 'ZZZZZ the 5-digit patient code USUBJID Unique Subject Identifier text Domain Abbreviation text DOMAIN IDENTIFIER Assigned USUBID Usige Subject best Pronocid DENTIFIER The USUBID variable has a fixed format XXXX-YYYY-ZZZZZ, where XXXX indicates the 4-digit compound code, "YTYY" th 4-digit study code and "ZZZZZZ" the 6-digit patient code SUBJID DM Subject Identifier for the text IDENTIFIE HEIGHT Baseline Height (cm) integer Deriver ANALYSIS IDENTIFIES equence mather (nationatically provented) ADSL HEIGHT ANALYSIS EIGHT Baseline Weight (kg) integer Derived STESTCD TOPIC Vital Sime Test Short tent Assigned ADSL WEIGHT Baseline BMI intege Derive ANALYSIS Derived Vital Sizes Test Name tent ADSL BMI BMIGRI Category 1 of Baseline text BMIGR11 Derived ANALYSIS Vital Signs Position of text POSITION CRF Page 13 RECORD ADSL BMIGR1 ANALYSIS BMIGRIN Category 1 of Baseline integer BMIGR1N Derived Derived, CRF Page 9, 13 Rends or Finding in dent. RESULT ADSL BMIGRIN AND ROD Category 2 of Baseline text BMIGR2L Derived ANALYSIS CRF Page 2.12 VSREAL VARIABLE AUSL BMIGR ANALYSIS Category 2 of Baseline integer BMIGR2N Derived STRESC Character Result Finding Inst Derived RESULT QUALIFIER ADSL BMIGR2N in Std Format Category 3 of Baseline text BMIGR3L BMIGRI Derived ANALYSIS s & Decision SSTRESS Numeric Result Finding Bost 3.1 in Standard Units Derived RESULT QUALIFIER ADSL BMIGR3 Category 3 of Baseline integer BMIGR3N BMI, (N) ANALYSIS BMIGRIN Derived ADSL BMIGREN

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Table 1 Demographic Data - Per-Protocol

	Treatment 1	Treatment 2
Baseline body mass index (BMI) [kg/m**2]		
N	167	167
Mean	29.08	29.04
SD	4.84	4.80
Min	20.3	16.0
Median	28.69	28.47
Max	40.1	41.2
Baseline BMI (categorical) [N (%)]		
<25 kg/m**2	41 (24.6%)	71 (21.1%)
25-<30 kg/m**2	60 (35.9%)	130 (38.7%)
>=30 kg/m**2	66 (́39.5%)	135 (40.2%)





	STUDYID	USUBJID	SUBJID	BMI	BMIGR1	BMIGR1N	BMIGR2	BMIGR2N
2	9999-0001	9999-0001-000001	000001	27.77777778	<30 kg/m**2	1	25-<30 kg/m**2	2
3	9999-0001	9999-0001-000002	000002	25.503615702	<30 kg/m**2	1	25-<30 kg/m**2	2
4	9999-0001	9999-0001-000003	000003	26.175194521	<30 kg/m**2	1	25-<30 kg/m**2	2
5	9999-0001	9999-0001-000004	000004	35.15625	>=30 kg/m**2	2	>=30 kg/m**2	3
6	9999-0001	9999-0001-000005	000005	30.968858131	>=30 kg/m**2	2	>=30 kg/m**2	3
7	9999-0001	9999-0001-000006	000006	39.697163916	>=30 kg/m**2	2	>=30 kg/m**2	3
8	9999-0001	9999-0001-000007	000007	25.826446281	<30 kg/m**2	1	25-<30 kg/m**2	2
9	9999-0001	9999-0001-000008	000008	30.103806228	>=30 kg/m**2	2	>=30 kg/m**2	3
10	9999-0001	9999-0001-000009	000009	32.280962683	>=30 kg/m**2	2	>=30 kg/m**2	3
11	9999-0001	9999-0001-000010	000010	28.876133787	<30 kg/m**2	1	25-<30 kg/m**2	2
12	9999-0001	9999-0001-000011	000011	29.372397383	<30 kg/m**2	1	25-<30 kg/m**2	2
13	9999-0001	9999-0001-000012	000012	26.714852608	<30 kg/m**2	1	25-<30 kg/m**2	2
14	9999-0001	9999-0001-000013	000013	32.718619869	>=30 kg/m**2	2	>=30 kg/m**2	3
15	9999-0001	9999-0001-000014	000014	28.719723183	<30 kg/m**2	1	25-<30 kg/m**2	2
16	9999-0001	9999-0001-000015	000015	32.270420377	>=30 kg/m**2	2	>=30 kg/m**2	3





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• ADaM define.xml

omputational Alg	orithms (A	DSL.BMI)										
Reference Nam	e	Computation Method										
SL.BMI	Con rand	tinuous variable, calculated using ADSL.WEIGHT/(ADSL.HEIGHT*0,01)**2 value at visit 3 if visit 3 data not available, the last data collected before Iomisation										
omputational Alg	gorithms (ADSL.HEIGHT)											
Reference Nam	e	Computation Method										
SL.HEIGHT	equa	al to VS.VSS?	IRESN v	when VS.VSTES	TCD="HEIGHT"							
omputational Alg	orithms (A	DSL.WEIGH	HT)									
Reference Nam	e				Co	mputation Meth	od					
SL.WEIGHT	equ	al to VS.VSS	TRESN v	vhen VS.VSTES	TCD="WEIGHT" and VS.V	VISITNUM=30 if	visit 3 data not available,	the last data collected befo	ore randomisation			
					ADSL HF GHT							
WEIGHT	Baseline W	Veight (kg)	integer		ADSLIVEIGHT	Derived	ANALYSIS					
ВМІ	Baseline E (kg/m**2)	ВМП)	integer		ADSL BMI	Derived	ANALYSIS					
BMIGR1	Category BMI	1 of Baseline	text	BMIGR1L	ADSL BMIGR1	Derived	ANALYSIS					
BMIGRIN	Category BMI, (N)	1 of Baseline	integer	BMIGR1N	ADSL BMIGRIN	Derived	ANALYSIS					
BMIGR2	Category BMI	2 of Baseline	text	BMIGR2L	ADSL BMIGR2	Derived	ANALYSIS					
BMIGR2N	Category BMI, (N)	2 of Baseline	integer	BMIGR2N	ADSL BMIGR2N	Derived	ANALYSIS					
BMIGR3	Category BMI	3 of Baseline	text	BMIGR3L	ADSL.BMIGR3	Derived	ANALYSIS					
BMIGR3N	Category BMI, (N)	3 of Baseline	integer	BMIGR3N	ADSL BMIGR3N	Derived	ANALYSIS					
								BUS	siness &			



• SDTM define.xml and aCRF

Value Le	vel Metadat:	a (ValueList.VS.VST	ESTCD)									
Source Variable	e Valu	e		Label	Туре	Controlled Terminology		Origin	Role	Comment		
VSTESTC	D DIABP	DIASTOLIC B	LOOD PRESS	SURE		text		CRI	F Page <u>13</u>			
VSTESTC	D HEIGHT	HEIGHT				text		CRI	F Page <u>9</u>			
VSTESTC	D PULSE	PULSE RATE				text		CRI	F Page <u>13</u>			
VSTESTC	D SYSBP	SYSTOLIC BL	OOD PRESS	URE		text		CRI	F Page <u>13</u>			
VSTESTC	D WAIST	WAIST CIRCU	MFERENCE			text		CRI	F Page <mark>9</mark>			
VSTESTC	D WEIGH	T WEIGHT				text		CRI	F Page <u>9</u>			
									patient code		Ŭ	I
	VSSEQ	Sequence Number	Patient D Patient X5 Page	Demographics ge 4 (Demo_V1a fo	– Part I or Visit 1a) Page 1 of 1.			<u>ER</u>	Sequence number ensure uniqueness	(automa within a	tically generated) to dataset for a subject	
	VSTESTCD	Vital Signs Test Short Name	Visit Date 11-E)ec-2007	Blank 🦳 Comment 🛛							
	VSTEST	Vital Signs Test Name	PATIENT DEMO	OGRAPHICS - Part I	DSCAT = "PROTOCO	STDTC	ONE"	M				
	VSPOS	Vital Signs Position of Subject	DSTERM/ Gender SEX			1 = male, 2 =	female	2				
	VSORRES	Result or Finding in Original Units	Date of birth	BRIHDIC		Age (Age i:	s automatically	Ł				
	VSORRESU	Original Units	Height			saved	and closed)	E N				
	VSSTRESC	Character Result/Findir in Std Format	Weight Waist circumfe	erence		kg cm		2				
	VSSTRESN	Numeric Result/Finding in Standard Units	VSORRES	/ VSORRESU	where			ŝ				þ
Restricted	© Business	& Decision Life Sci	"WEIGHT"	, "WAIST"							Lite Sc	ļ

ADaM Conversions Prerequisites: Linear Approach



ADaM Conversions Prerequisites: Linear Approach



ADaM Conversions Prerequisites: Parallel Approach



ADaM Conversions Prerequisites: Parallel Approach



ADaM Conversions: The Good, the Bad and the Ugly









- E Good Traceability between the statistical outputs, the ADaM datasets and the SDTM domains
- G Since BDLS was responsible for the **SDTM Conversion**, issues with SDTM domains were resolved rapidly
 - Having the ADS in ADaM structure facilitated the addition of this legacy study as part of an Analysis Pooled Database



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E • Hidden differences with SDTM found during validation:

- Different Coding Dictionary Versions in SDTM database
- For consistency with newer studies, some categorical variables were updated with new combination of values
- A Due to SDTM standards restrictions some values used for analysis were not available in the SDTM datasets
 - Key variables in SDTM domains cannot be empty

 \rightarrow If the initial analysis missing values were summarized in a table, these values were not available anymore in the SDTM datasets





- E ADaM specifications created from SAP and protocol
 - Rules specified in Protocol and SAP were differently interpreted in the same way as in the original analysis
- $\mathcal{U} \rightarrow$ Multiple solutions were tried before having the proper result
- G <u>QC: manual comparisons were needed</u>
 - **ADS** not available

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- Listings in pdf format
 - \rightarrow Resolving all these issues was time consuming



- Parrallel Approach
- Creation of a **pooled ADaM mapping**
 - From Sponsor's specifications
- Creation of **pooled ADaM datasets**
 - From Sponsor's Analysis datasets
- Extraction of the **individual ADaM mapping and datasets**











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- E BDLS was in charge of the SDTM and ADaM conversion, this facilitated the communication between the two teams and accelerated the problem solving
- With a mature Pooled ADaM conversion process, data extraction for each study is fast, and consistency checks across studies are no longer needed
 - Codelist were re-usable in ADaM due to its consistency across studies
- D A **comment file** with all questions from and to the sponsor, facilitates the tracking of all answers and decisions





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- E Not well documented **Derivation Rules** caused difficulties to define the traceability
 - Studies were added into the pooled database at different time points, while derivations were not properly updated
- A **SDTM limitations** affected the ADaM conversion
 - some raw variables were not available in SDTM, but were mentioned in ADS specifications
 - Specifications had to be redefined in cooperation with the sponsor
 - Due to traceability between SDTM/ADaM, **timelines** for ADaM datasets were influenced by SDTM timelines





E • Structure of the pooled original ADS datasets changed with the inclusion of new studies

 \rightarrow Impact on the pooled mapping, pooled datasets and extractions

- G Source variables which had some manipulation during the conversion (e.g. variable derived, with a format or transposed) were sensitive to changes in its content at each transfer
- $L \rightarrow$ New compare tools to check changes in dataset structure between the original ADS received



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- BDLS developed **new compare tools** to check changes in dataset structure between the original ADS received, like number of records, empty, new, and missing variables that were present before
- **Specifications** were **compared**: results of a compare were included in the communication file to the sponsor

	В		С	D	E		F		Н		I	J
1	VARIABLE	-	DESCRIPTION 🔹 🔻	TYI 👻	compare_fla 💌	new_do	omain_flag	۳	B&DLS comments	-	Sponsor comments	Updates
2	SOURCE_VAR	₹_1	AESI flag-Angioedem	num	ADDED	NO						
3	SOURCE_VAR	2_2	AESI flag-Embolic an	num	ADDED	NO						
4	SOURCE_VAR	₹_3	AESI flag-Hypoglycae	num	ADDED	NO						
5	SOURCE_VAR	₹_4	AESI flag-Hypersensi	num	MODIFIED	YES						
6	SOURCE_VAR	₹_5	AESI flag-Increased u	num	ADDED	YES						
7	SOURCE_VAR	₹_6	Investigator Special In	num	REMOVED	NO						
8	SOURCE_VAR	₹_7	Investigator Special In	num	REMOVED	NO						
9	SOURCE_VAR	₹_8	Investigator Special In	num	REMOVED	NO						
10	SOURCE_VAR	₹_9	Investigator Special In	num	ADDED	NO						
11	SOURCE_VAR	2_10	Investigator Special In	num	ADDED	NO						
12	SOURCE_VAR	₹ <u>11</u>	Investigator Special In	num	ADDED	NO						
13	SOURCE_VAR	₹_12	Investigator Special In	num	ADDED	NO						
14	SOURCE_VAR	₹_13	Investigator Special In	num	ADDED	NO						
	• •	Issue	es & updates post-rel	ease	Source data is	ssues	specs_com	р	Compare_tool	SD	FM Source not mapped	Review cycl



• ADS content was compared:

Results from content compare for variables being "derived", "with a CL applied", "transposed" during ADaM conversion, were included in the communication file to the sponsor

.1	 Image: Second sec							er against the ne oplies a change i	w transfe n ADaM (er. This compare is don derivation rule, e.g. tra	e ONLY for va Inspose rule,	riables that are 'derived' or code list.
Α	В	С	D			E			F	G	Н	
studyid	DOMAIN 👻	variable 🚽 💌	comment		·	content		B&DL	S Comment	Sponsor	 Updates Comments 	
									we will for this	include a new criteria value. Please confirm		L
0001	ADS11	SRC_VAR_5	new variab	le to transpose present	in new databas	e	Creatinine >= 1.5* ref. s	ample and > ULN	I the val	ue is correct	correct	Pooled mapping updated
									we will for this	include a new criteria value. Please confirm		
0001	ADS2	SRC_VAR_7	7 Domain not present in new database			Creatinine >= 2*baselin	e and > ULN	_N the value is correct		correct	not included	
									this is flag, pl	a new set of population ease confirm if we need		
0001	11 ADS9 SRC VAR 5 Value not present in code list metadate but present in cource date								hhe ot	it	Please inno	e not included
 Image: Image: Im	Issues & (updates post-i	release	Source data issues	specs_comp	Compare_tool	SDTM Source not ma	apped Review	v cycles	Consistency checks	Def 🤆): • • •



- To ensure traceability with SDTM, a good knowledge of **what was mapped from RAW into SDTM** and in which study, was essential
- List of RAW variables per study per domain from each SDTM metadata, specifying if the variable was mapped into SDTM, and if it was empty

STUDY	JT VIEWNAME	J VARNAME	LABEL	▼ VARTYPE	FORMATN	🖃 MAPPED	CONFIRMED	EMPTY 💽
RAW_Study1	AE	AEI	AE indicator	num	YN1F.		х	X
RAW_Study1	AE	AEONTM	AE onset time	num	TIME5.	×	х	
RAW_Study1	AE	AEENDC	AE end date continued	char	\$CONT1F.	×	Х	
RAW_Study1	AE	AEOUT	AE outcome	num	AEOUT1F.	×	Х	
RAW_Study1	AE	AEREL	AE drug relationship	num	YN1E.	×	х	
RAW_Study1	AE	AEONDT	AE onset date	num	DATE9.	×	Х	
RAW_Study1	AE	AEENDDT	AE end date	num	DATE9.	×	х	
RAW_Study1	AE	AELLT	AE lowest level term	char	\$200.		х	
RAW_Study1	AE	AELLTCD	AE lowest level term code	char	\$10.		х	
RAW_Study2	AE	AEI	AE indicator	num	YN1F.		Х	X
RAW_Study2	AE	AEONTM	AE onset time	num	TIME5.	×	Х	
RAW_Study2	AE	AEENDC	AE end date continued	char	\$CONT1F.	×	х	
RAW_Study2	AE	AEOUT	AE outcome	num	AEOUT1F.	×	х	
RAW_Study2	AE	AEREL	AE drug relationship	num	YN1F.	×	х	
RAW_Study2	AE	AEONDT	AE onset date	num	DATE9.	×	х	
RAW_Study2	AE	AEENDDT	AE end date	num	DATE9.	×	х	
RAW_Study2	AE	AELLT	AE lowest level term	char	\$200.		X	
RAW_Study2	AE	AELLTCD	AE lowest level term code	char	\$10.		Х	



ADaM Conversions: The Good, the Bad and the Ugly





Conclusions

• Parallel approach:

- ensured reproduction of TFLs
- consistency between studies within a pooled database

• Linear approach:

- SDTM/ADaM traceability is demonstrated
- Bigger risk of not reproducing identical results to the initial analysis
- Source pooled ADS should be stable and specifications should be clear
- SDTM datasets & pooled ADaM datasets should be **finalized** before the individual study extractions can start



Conclusions

- Dictionary update: SDTM and ADaM uses the same dictionary version. Specific ADaM variables are available
- **Good communication** is needed between SDTM, ADaM team members and the sponsor, during the whole process
- **Sponsor's feedback** on draft mapping and draft ADaM datasets is crucial to address all details upfront
- All decisions can be easily tracked if they are properly documented in a **communication file**



Questions?







Business & Decision

Jessica Minkue | Project Manager Statistics | jessica.minkue@businessdecision.com

Business & Decision Life Sciences Sint-Lambertusstraat 141 rue Saint-Lambert B-1200 Brussels T: +32 2 774 11 00 F: +32 2 774 11 99 <u>lifesciences@businessdecision.com</u> http://www.businessdecision-lifesciences.com/