

Data Specification

MetrumRG

2019-01-12 10:32:01

spec

Description: [data set description]

Column	Type	Field	Value
C	character	short-name	commented row indicator
		values	., C
		long-name	commented rows
USUBJID	character	short-name	USUBJID
		long-name	universal subject identifier
ID	numeric	short-name	ID
		long-name	NONMEM ID number
EVID	numeric	short-name	event ID
		long-name	event ID indicator
		comment	per NONMEM specifications
MDV	numeric	short-name	MDV
		long-name	missing DV indicator
		comment	per NONMEM specifications
SEQ	numeric	short-name	SEQ
		values	1 = concentration, 2 = response, 3 = other
		long-name	record type indicators
AMT	numeric	short-name	amount
		unit	mg
		range	0 to 1000
		long-name	dose amount
II	numeric	short-name	II
		unit	hours
		long-name	interdose interval
CMT	numeric	short-name	Compartment
		values	1, 2, 3, 4
		long-name	compartment number
		comment	per NONMEM specifications
TAFD	numeric	short-name	TAFD
		unit	hours
		long-name	time after first dose
WT	numeric	short-name	Weight
		unit	kg
		range	40 to 100
		long-name	baseline weight
EGFR	numeric	short-name	eGFR
		unit	ml/min/1.73m2
		range	10 to 300
		long-name	estimated glomerular filtration rate

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Column	Type	Field	Value
STUDY	character	short-name values	Study 100 = The first phase 1 study, 202 = The first phase 2 study conducted only at sites 1, 2, 3 when the formulation was still oral liquid only, 303 = The renal impairment study, 203 = The second phase 2 study
		long-name	study identifier
SEX	numeric	short-name values	SEX 0 = male, 1 = female
BQL	numeric	short-name values long-name	below limit of quantification 0 = FALSE, 1 = TRUE DV value is below the quantitation limit
DV	numeric	short-name unit long-name comment	DV ng/mL dependent variable see SEQ
DV2	numeric	short-name long-name	DV2 second dependent variable
HAIR	character	short-name values long-name	HAIR 0 = brown, 1 = blonde, 2 = black patient hair color
CLCR	numeric	short-name unit range	creatinine clearance ml/min 40 to 180
ALB	numeric	short-name unit range comment	adjusted albumin g/dL 1 to 5 in the lookup document as ALB2