

MIDUS DNA

MIDUS DNA

Title

MIDUS DNA

Related Materials

Documentation

- [MIDUS Polygenic Risk Score README](#)
- [MIDUS Polygenic Risk Score Documentation](#)

M2MR_P6_PolygenicRiskScores_N2118_20211028

Title	M2MR_P6_PolygenicRiskScores_N2118_20211028
File Name	M2MR1_P6_PolygenicRiskScores_N2118_20211028.sav
Case Quantity	2118
Variable Count	65

M2MRID - M2 AND MR Public IDs combined in a single variable

Type	Numeric (Integer)
Display Values	2
Stats	1

Valid	Invalid	Minimum	Maximum	Mean	StdDev
2118	0	10002	39977	23028.720...	10360.625...

SAMPLMAJ - Major sample identification

Type	Code
Display Values	1
Stats	1

			Frequency	% of total	% of valid
Valid	1	CORE MAIN RDD	640	30.22%	30.22%
	2	CORE SIBLING	6	0.28%	0.28%
	3	CORE TWIN	388	18.32%	18.32%
	4	CORE CITY OVERSAMPLE	20	0.94%	0.94%
	13	CORE MILWAUKEE	201	9.49%	9.49%
	20	REFRESHER MAIN	746	35.22%	35.22%
	21	REFRESHER MILWAUKEE	117	5.52%	5.52%

	Total	2,118	100%	100%
--	-------	-------	------	------

Valid	Invalid	Minimum	Maximum
2118	0	1	21

M2MRCASE - Which sample is R in? 1=M2, 2=MR1

Type	Code
Display Values	1
Stats	1

			Frequency	% of total	% of valid
Valid	1	M2	1255	59.25%	59.25%
	2	MR1	863	40.75%	40.75%
		Total	2,118	100%	100%

Valid	Invalid	Minimum	Maximum
2118	0	1	2

GENCONSENT - Did R consent to genetics 1=Yes, 2=No

Type	Code
Display Values	1
Stats	1

			Frequency	% of total	% of valid
Valid	1	YES	2009	94.85%	94.85%
	2	NO	109	5.15%	5.15%
		Total	2,118	100%	100%

Valid	Invalid	Minimum	Maximum
2118	0	1	2

BRA6DTISSUE - Tissue sample source for DNA

Type	Code
Display Values	1
Stats	1

			Frequency	% of total	% of valid
Valid	1	BLOOD	1647	77.76%	77.76%
	2	SALIVA	471	22.24%	22.24%
		Total	2,118	100%	100%

Valid	Invalid	Minimum	Maximum
2118	0	1	2

BRA6DPAVAIL - Is Polygenic Risk Score (PRS) data available?

Type	Code
ForwardSkip	IF BRA6DPAVAIL = 2, SKIP BRA6DPCEUR TO BRA6DPRSWHR
Display Values	1
Stats	1

			Frequency	% of total	% of valid
Valid	1	YES	1293	61.05%	61.05%
	2	NO	825	38.95%	38.95%
		Total	2,118	100%	100%

Valid	Invalid	Minimum	Maximum
2118	0	1	2

BRA6DPCEUR - European Ancestry Component Score

Type	Numeric (Double)
Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	0.907935	0.99996	0.9919500...	0.0147970...

BRA6DPCEAS - East Asian Ancestry Component Score

Type	Numeric (Double)
Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Display Values	2
Stats	2

	Frequency	% of total	% of valid
--	-----------	------------	------------

Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	1E-5	0.059261	8.2E-4	0.0031256...

BRA6DPCAMR - Ad-Mixed American Ancestry Component Score

Type	Numeric (Double)				
Numeric Details	Decimals: 2				
BackwardSkip	(BRA6DPAVAIL = 2)				
Display Values	2				
Stats	2				

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	1E-5	0.059545	0.0016406...	0.0050313...

BRA6DPCSAS - South Asian Ancestry Component Score

Type	Numeric (Double)				
Numeric Details	Decimals: 2				
BackwardSkip	(BRA6DPAVAIL = 2)				
Display Values	2				
Stats	2				

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	1E-5	0.067207	0.0037339...	0.0099977...

BRA6DPCAfr - African Ancestry Component Score

Type	Numeric (Double)				
Numeric Details	Decimals: 2				
BackwardSkip	(BRA6DPAVAIL = 2)				

Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	1E-5	0.042671	0.0018554...	0.0050552...

BRA6DPC1 - First Principal Component

Type	Numeric (Double)
Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	-0.0127806	-0.00845361	-0.011865...	5.24E-4

BRA6DPC2 - Second Principal Component

Type	Numeric (Double)
Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	0.00574898	0.0112056	0.0100766...	4.34E-4

BRA6DPC3 - Third Principal Component

Type	Numeric (Double)
Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	-0.0154213	0.00976799	-0.003240...	0.0029155...

BRA6DPC4 - Fourth Principal Component

Type	Numeric (Double)
Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	-0.00689364	0.0072943	2.88E-4	0.0020416...

BRA6DPC5 - Fifth Principal Component

Type	Numeric (Double)
Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
-------	---------	---------	---------	------	--------

1293	825	-0.103595	0.0479037	0.0040011...	0.0147113...
------	-----	-----------	-----------	--------------	--------------

BRA6DPRSADHDF - Polygenic Risk Score (ADHD for Females)

Type	Numeric (Double)
Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Notes	Computed Variable, for details see documentation M2MR_P6_PolygenicRiskScore_Documentation_20211129 for PRSice method.
Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	-2.124596...	1.3369620...	-0.236189...	0.5139934...

BRA6DPRSADHDM - Polygenic Risk Score (ADHD for Males)

Type	Numeric (Double)
Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Notes	Computed Variable, for details see documentation M2MR_P6_PolygenicRiskScore_Documentation_20211129 for PRSice method.
Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	-3.731282...	2.9345738...	-0.268295...	1.0011645...

BRA6DPRSADV - Polygenic Risk Score (Adventurousness)

Type	Numeric (Double)
------	------------------

Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Notes	Computed Variable, for details see documentation M2MR_P6_PolygenicRiskScore_Documentation_20211129 for PRSice method.
Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	-3.519634...	3.8361615...	-0.004681...	1.1105288...

BRA6DPRSAOM - Polygenic Risk Score (Age of Menarche)

Type	Numeric (Double)
Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Notes	Computed Variable, for details see documentation M2MR_P6_PolygenicRiskScore_Documentation_20211129 for PRSice method.
Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	-0.898444...	2.2263126...	0.6243924...	0.5682290...

BRA6DPRSZ - Polygenic Risk Score (Alzheimers Disease)

Type	Numeric (Double)
Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Notes	Computed Variable, for details see documentation M2MR_P6_PolygenicRiskScore_Documentation_20211129 for PRSice method.

Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	-1.585654...	0.4085417...	-0.620642...	0.3008868...

BRA6DPRSAN - Polygenic Risk Score (Anorexia Nervosa)

Type	Numeric (Double)
Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Notes	Computed Variable, for details see documentation M2MR_P6_PolygenicRiskScore_Documentation_20211129 for PRSice method.
Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	0.1626018...	1.0442848...	0.6835206...	0.1082258...

BRA6DPRSANX - Polygenic Risk Score (Anxiety)

Type	Numeric (Double)
Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Notes	Computed Variable, for details see documentation M2MR_P6_PolygenicRiskScore_Documentation_20211129 for PRSice method.
Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	-2.976697...	3.7162255...	0.1247699...	1.0060020...

BRA6DPRSASTH - Polygenic Risk Score (Asthma)

Type	Numeric (Double)
Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Notes	Computed Variable, for details see documentation M2MR_P6_PolygenicRiskScore_Documentation_20211129 for PRSice method.
Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	-1.105975...	-0.278311...	-0.651068...	0.1313875...

BRA6DPRSASD - Polygenic Risk Score (Autism Spectrum Disorder)

Type	Numeric (Double)
Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Notes	Computed Variable, for details see documentation M2MR_P6_PolygenicRiskScore_Documentation_20211129 for PRSice method.
Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	-3.049707...	2.5271398...	-0.093738...	0.8791189...

BRA6DPRSAUTO - Polygenic Risk Score (Automobile Speeding Propensity)

Type	Numeric (Double)
------	------------------

Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Notes	Computed Variable, for details see documentation M2MR_P6_PolygenicRiskScore_Documentation_20211129 for PRSice method.
Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	-2.561926...	3.4307402...	0.3769058...	0.8804458...

BRA6DPRSBPD - Polygenic Risk Score (Bipolar Disorder)

Type	Numeric (Double)
Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Notes	Computed Variable, for details see documentation M2MR_P6_PolygenicRiskScore_Documentation_20211129 for PRSice method.
Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	-1.093612...	-0.059585...	-0.566047...	0.1394306...

BRA6DPRSBHT - Polygenic Risk Score (Birth Height)

Type	Numeric (Double)
Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Notes	Computed Variable, for details see documentation M2MR_P6_PolygenicRiskScore_Documentation_20211129 for PRSice method.

Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	-0.632659...	1.7366506...	0.6727856...	0.3632642...

BRA6DPRSBWT - Polygenic Risk Score (Birth Weight)

Type	Numeric (Double)
Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Notes	Computed Variable, for details see documentation M2MR_P6_PolygenicRiskScore_Documentation_20211129 for PRSice method.
Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	-0.211914...	1.5590875...	0.6841271...	0.3218244...

BRA6DPRSBMI - Polygenic Risk Score (Body Mass Index)

Type	Numeric (Double)
Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Notes	Computed Variable, for details see documentation M2MR_P6_PolygenicRiskScore_Documentation_20211129 for PRSice method.
Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	-1.602481...	0.3938214...	-0.693935...	0.3262796...

BRA6DPRSCIQ - Polygenic Risk Score (Child I.Q.)

Type	Numeric (Double)
Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Notes	Computed Variable, for details see documentation M2MR_P6_PolygenicRiskScore_Documentation_20211129 for PRSice method.
Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	-3.809493...	2.3467496...	-0.308782...	0.9364676...

BRA6DPRSCIGS - Polygenic Risk Score (Cigarettes per day)

Type	Numeric (Double)
Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Notes	Computed Variable, for details see documentation M2MR_P6_PolygenicRiskScore_Documentation_20211129 for PRSice method.
Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	-1.224052...	2.4406534...	0.5761155...	0.5915900...

BRA6DPRSCOGP - Polygenic Risk Score (Cognitive Performance)

Type	Numeric (Double)
------	------------------

Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Notes	Computed Variable, for details see documentation M2MR_P6_PolygenicRiskScore_Documentation_20211129 for PRSice method.
Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	-2.969767...	3.8598221...	0.1228462...	0.9041236...

BRA6DPRSEDU - Polygenic Risk Score (College Education)

Type	Numeric (Double)
Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Notes	Computed Variable, for details see documentation M2MR_P6_PolygenicRiskScore_Documentation_20211129 for PRSice method.
Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	-2.771620...	3.1172658...	-0.180453...	0.8760975...

BRA6DPRSCAD - Polygenic Risk Score (Coronary Artery Disease)

Type	Numeric (Double)
Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Notes	Computed Variable, for details see documentation M2MR_P6_PolygenicRiskScore_Documentation_20211129 for PRSice method.

Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	-2.359930...	1.3852684...	-0.500251...	0.6031382...

BRA6DPRSCD - Polygenic Risk Score (Crohns Disease)

Type	Numeric (Double)
Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Notes	Computed Variable, for details see documentation M2MR_P6_PolygenicRiskScore_Documentation_20211129 for PRSice method.
Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	-1.152587...	1.1593308...	-0.572535...	0.2100993...

BRA6DPRSCRD - Polygenic Risk Score (Cross Disorder)

Type	Numeric (Double)
Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Notes	Computed Variable, for details see documentation M2MR_P6_PolygenicRiskScore_Documentation_20211129 for PRSice method.
Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	-1.079917...	1.9285984...	0.5160135...	0.4745363...

BRA6DPRSDEP - Polygenic Risk Score (Depressive Symptoms)

Type	Numeric (Double)
Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Notes	Computed Variable, for details see documentation M2MR_P6_PolygenicRiskScore_Documentation_20211129 for PRSice method.
Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	-3.500963...	2.4216724...	-0.320964...	0.9866357...

BRA6DPRSDISI - Polygenic Risk Score (Disinhibition)

Type	Numeric (Double)
Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Notes	Computed Variable, for details see documentation M2MR_P6_PolygenicRiskScore_Documentation_20211129 for PRSice method.
Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	-3.241073...	2.2887112...	-0.392497...	0.8412533...

BRA6DPRSDPW - Polygenic Risk Score (Number of Drinks per Week)

Type	Numeric (Double)
------	------------------

Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Notes	Computed Variable, for details see documentation M2MR_P6_PolygenicRiskScore_Documentation_20211129 for PRSice method.
Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	-3.718809...	3.3868110...	-0.131657...	1.0329806...

BRA6DPRSEA - Polygenic Risk Score (Educational Attainment)

Type	Numeric (Double)
Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Notes	Computed Variable, for details see documentation M2MR_P6_PolygenicRiskScore_Documentation_20211129 for PRSice method.
Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	-1.797938...	2.3836542...	0.2158835...	0.6096756...

BRA6DPRSES - Polygenic Risk Score (Ever Smoked)

Type	Numeric (Double)
Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Notes	Computed Variable, for details see documentation M2MR_P6_PolygenicRiskScore_Documentation_20211129 for PRSice method.

Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	-2.998216...	2.2755885...	-0.163444...	0.9245886...

BRA6DPRSEXTA - Polygenic Risk Score (Extraversion)

Type	Numeric (Double)
Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Notes	Computed Variable, for details see documentation M2MR_P6_PolygenicRiskScore_Documentation_20211129 for PRSice method.
Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	-1.886540...	2.6241547...	0.4607443...	0.6554421...

BRA6DPRSGRT - Polygenic Risk Score (General Risk Tolerance)

Type	Numeric (Double)
Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Notes	Computed Variable, for details see documentation M2MR_P6_PolygenicRiskScore_Documentation_20211129 for PRSice method.
Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	-2.656831...	2.3024496...	-0.527982...	0.6897536...

BRA6DPRSHDL - Polygenic Risk Score (High-Density Lipoproteins - HDL)

Type	Numeric (Double)
Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Notes	Computed Variable, for details see documentation M2MR_P6_PolygenicRiskScore_Documentation_20211129 for PRSice method.
Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	-1.695465...	2.0598182...	0.0637175...	0.5757995...

BRA6DPRSHT - Polygenic Risk Score (Height)

Type	Numeric (Double)
Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Notes	Computed Variable, for details see documentation M2MR_P6_PolygenicRiskScore_Documentation_20211129 for PRSice method.
Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	-0.111922...	1.2486548...	0.7358602...	0.1764487...

BRA6DPRSIHC - Polygenic Risk Score (Infant Head Circumference)

Type	Numeric (Double)
------	------------------

Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Notes	Computed Variable, for details see documentation M2MR_P6_PolygenicRiskScore_Documentation_20211129 for PRSice method.
Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	-0.071707...	1.2627153...	0.7006454...	0.1962089...

BRA6DPRSICV - Polygenic Risk Score (Intracranial Volume)

Type	Numeric (Double)
Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Notes	Computed Variable, for details see documentation M2MR_P6_PolygenicRiskScore_Documentation_20211129 for PRSice method.
Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	-3.475987...	4.3506785...	-0.002124...	1.0808810...

BRA6DPRSLDL - Polygenic Risk Score (Low-Density Lipoproteins - LDL)

Type	Numeric (Double)
Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Notes	Computed Variable, for details see documentation M2MR_P6_PolygenicRiskScore_Documentation_20211129 for PRSice method.

Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	-3.480962...	2.7824664...	-0.116445...	0.9596791...

BRA6DPRSMDD - Polygenic Risk Score (Major Depressive Disorder)

Type	Numeric (Double)
Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Notes	Computed Variable, for details see documentation M2MR_P6_PolygenicRiskScore_Documentation_20211129 for PRSice method.
Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	-1.099910...	-0.244777...	-0.685492...	0.1317521...

BRA6DPRSNEURO - Polygenic Risk Score (Neuroticism)

Type	Numeric (Double)
Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Notes	Computed Variable, for details see documentation M2MR_P6_PolygenicRiskScore_Documentation_20211129 for PRSice method.
Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	-3.454321...	3.4953745...	-0.023058...	1.0899349...

BRA6DPRNSP - Polygenic Risk Score (Number of Sexual Partners)

Type	Numeric (Double)
Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Notes	Computed Variable, for details see documentation M2MR_P6_PolygenicRiskScore_Documentation_20211129 for PRSice method.
Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	-2.863686...	1.5027934...	-0.581720...	0.6114793...

BRA6DPRSOCD - Polygenic Risk Score (Obsessive Compulsive Disorder)

Type	Numeric (Double)
Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Notes	Computed Variable, for details see documentation M2MR_P6_PolygenicRiskScore_Documentation_20211129 for PRSice method.
Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	-0.445111...	1.8550141...	0.6766934...	0.3511834...

BRA6DPRSPTSD - Polygenic Risk Score (Post Traumatic Stress Disorder - PTSD)

Type	Numeric (Double)
Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Notes	Computed Variable, for details see documentation M2MR_P6_PolygenicRiskScore_Documentation_20211129 for PRSice method.
Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	-1.121618...	-0.203248...	-0.680020...	0.1517152...

BRA6DPRSRA - Polygenic Risk Score (Rheumatoid Arthritis)

Type	Numeric (Double)
Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Notes	Computed Variable, for details see documentation M2MR_P6_PolygenicRiskScore_Documentation_20211129 for PRSice method.
Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	-1.526980...	0.2880717...	-0.680569...	0.2636633...

BRA6DPRSRT - Polygenic Risk Score (Risk Tolerance)

Type	Numeric (Double)
------	------------------

Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Notes	Computed Variable, for details see documentation M2MR_P6_PolygenicRiskScore_Documentation_20211129 for PRSice method.
Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	-3.434965...	3.4464570...	0.0452065...	1.1067631...

BRA6DPRSSCHIZ - Polygenic Risk Score (Schizophrenia)

Type	Numeric (Double)
Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Notes	Computed Variable, for details see documentation M2MR_P6_PolygenicRiskScore_Documentation_20211129 for PRSice method.
Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	-1.293660...	-0.015740...	-0.652026...	0.2079331...

BRA6DPRSSWB - Polygenic Risk Score (Subjective Well-Being)

Type	Numeric (Double)
Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Notes	Computed Variable, for details see documentation M2MR_P6_PolygenicRiskScore_Documentation_20211129 for PRSice method.

Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	-2.134264...	2.5158390...	0.4224191...	0.6706862...

BRA6DPRSSD - Polygenic Risk Score (Suicide Death)

Type	Numeric (Double)
Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Notes	Computed Variable, for details see documentation M2MR_P6_PolygenicRiskScore_Documentation_20211129 for PRSice method.
Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	-1.743362...	0.5604983...	-0.631794...	0.3087871...

BRA6DPRSCHOL - Polygenic Risk Score (Total Cholesterol)

Type	Numeric (Double)
Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Notes	Computed Variable, for details see documentation M2MR_P6_PolygenicRiskScore_Documentation_20211129 for PRSice method.
Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	-3.125151...	2.3057836...	-0.333405...	0.9425048...

BRA6DPRSTRI - Polygenic Risk Score (Triglycerides)

Type	Numeric (Double)
Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Notes	Computed Variable, for details see documentation M2MR_P6_PolygenicRiskScore_Documentation_20211129 for PRSice method.
Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	-3.381444...	2.2028101...	-0.524849...	0.7644538...

BRA6DPRST2D - Polygenic Risk Score (Type 2 Diabetes)

Type	Numeric (Double)
Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Notes	Computed Variable, for details see documentation M2MR_P6_PolygenicRiskScore_Documentation_20211129 for PRSice method.
Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	-1.562409...	0.3437915...	-0.676932...	0.3343980...

BRA6DPRSUC - Polygenic Risk Score (Ulcerative Colitis)

Type	Numeric (Double)
------	------------------

Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Notes	Computed Variable, for details see documentation M2MR_P6_PolygenicRiskScore_Documentation_20211129 for PRSice method.
Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	-0.052816...	1.4750011...	0.5881649...	0.2169020...

BRA6DPRSWHR - Polygenic Risk Score (Waist to Hip Ratio)

Type	Numeric (Double)
Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Notes	Computed Variable, for details see documentation M2MR_P6_PolygenicRiskScore_Documentation_20211129 for PRSice method.
Display Values	2
Stats	2

			Frequency	% of total	% of valid
Missing	9.00	INAPP	825	38.95%	
		Total	825	38.95%	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
1293	825	-1.907280...	0.7158829...	-0.601886...	0.3523350...

BRA6DPRSCSALZ - Polygenic Risk Score-CS Method (Alzheimers Disease)

Type	Numeric (Double)
Numeric Details	Decimals: 2
BackwardSkip	(BRA6DPAVAIL = 2)
Notes	Computed Variable, for details see documentation M2MR_P6_PolygenicRiskScore_Documentation_20211129 for PRS-CS method.

Display Values	2
Stats	2