Wave IV PGS Risk-Tolerance

Data Layouts

Wave IV PGS Risk-Tolerance

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SPGS_RISK - PGS_RISK_PLINK_GWAS	
Туре	Numeric (Double)
Description	Polygenic score for general risk tolerance, obtained using classic PLINK method and standard GWAS results. Note these are reverse-coded.

SPGS_RIS2 - PGS_RISK_LDPRED_GWAS	
Туре	Numeric (Double)
Description	Polygenic score for general risk tolerance, obtained using LDpred method and standard GWAS results

SPGS_RIS3 - PGS_RISK_LDPRED_MTAG	
Туре	Numeric (Double)
Description	Polygenic score for general risk tolerance, obtained using LDpred method and results from multivariate analysis of adventurousness, automobile speeding propensity, drinks per week, ever smoker, number of sexual partners, and lifetime cannabis use

% PC1	
Туре	Numeric (Double)
Description	1st principal component (PC) of the covariance matrix of the individuals genotypic data

№ PC2	
Туре	Numeric (Double)
Description	2nd principal component (PC) of the covariance matrix of the individuals genotypic data

№ PC3	
Туре	Numeric (Double)
Description	3rd principal component (PC) of the covariance matrix of the individuals genotypic data

∞	
PC4	
Туре	Numeric (Double)
Description	4th principal component (PC) of the covariance matrix of the individuals genotypic data
	genotypic data
№ PC5	
Туре	Numeric (Double)
Description	5th principal component (PC) of the covariance matrix of the individuals genotypic data
⋘ PC6	
Туре	Numeric (Double)
Description	6th principal component (PC) of the covariance matrix of the individuals
	genotypic data
№ PC7	
Туре	Numeric (Double)
Description	7th principal component (PC) of the covariance matrix of the individuals genotypic data
	Seriot/pio data
% PC8	
Туре	Numeric (Double)
Description	8th principal component (PC) of the covariance matrix of the individuals genotypic data
% PC9	
Туре	Numeric (Double)
Description	9th principal component (PC) of the covariance matrix of the individuals2
	genotypic data
⋘ PC10	
Туре	Numeric (Double)
Description	10th principal component (PC) of the covariance matrix of the individuals 2
	genotypic data