

## Wave IV PGS Risk-Tolerance

### Title

Wave IV PGS Risk-Tolerance

### Abstract

Contains polygenic scores for general risk tolerance, adventurousness and risky behaviors in the driving, drinking, smoking and sexual domains. Polygenic scores were created for unrelated, Add Health participants of European Ancestry. Score metrics were generated using Plink, LDpred or MTAG software using the UK Biobank GWAS study.

Variables: 13

Observations: 4,755

### Variables

#### Wave IV PGS Risk-Tolerance

- [PGS\\_RISK - PGS\\_RISK\\_PLINK\\_GWAS](#)
- [PGS\\_RIS2 - PGS\\_RISK\\_LDPRD\\_GWAS](#)
- [PGS\\_RIS3 - PGS\\_RISK\\_LDPRD\\_MTG](#)
- [PC1](#)
- [PC2](#)
- [PC3](#)
- [PC4](#)
- [PC5](#)
- [PC6](#)
- [PC7](#)
- [PC8](#)
- [PC9](#)
- [PC10](#)

#### PGS\_RISK - PGS\_RISK\_PLINK\_GWAS

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

#### PGS\_RIS2 - PGS\_RISK\_LDPRD\_GWAS

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

 PGS_RIS3 - PGS_RISK_LDPRED_MTAG	
Type	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	
Type	Numeric (Double)
Description	1st principal component (PC) of the covariance matrix of the individuals' genotypic data

 PC2	
Type	Numeric (Double)
Description	2nd principal component (PC) of the covariance matrix of the individuals' genotypic data

 PC3	
Type	Numeric (Double)
Description	3rd principal component (PC) of the covariance matrix of the individuals' genotypic data

 PC4	
Type	Numeric (Double)
Description	4th principal component (PC) of the covariance matrix of the individuals' genotypic data

 PC5	
Type	Numeric (Double)
Description	5th principal component (PC) of the covariance matrix of the individuals' genotypic data

 PC6	
Type	Numeric (Double)
Description	6th principal component (PC) of the covariance matrix of the individuals' genotypic data

 PC7	
Type	Numeric (Double)

Description	7th principal component (PC) of the covariance matrix of the individuals¶ genotypic data
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PC8	
Type	Numeric (Double)
Description	8th principal component (PC) of the covariance matrix of the individuals¶ genotypic data

PC9	
Type	Numeric (Double)
Description	9th principal component (PC) of the covariance matrix of the individuals¶ genotypic data

PC10	
Type	Numeric (Double)
Description	10th principal component (PC) of the covariance matrix of the individuals¶ genotypic data

## Logical Products

Wave IV PGS Risk-Tolerance	
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### Data Layouts

Wave IV PGS Risk-Tolerance	
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### Wave IV PGS Risk-Tolerance

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

PGS_RIS2 - PGS_RISK_LDPRED_GWAS	
Type	Numeric (Double)
Description	Polygenic score for general risk tolerance, obtained using LDpred method and standard GWAS results

PGS_RIS3 - PGS_RISK_LDPRED_MTAG	
Type	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	
Type	Numeric (Double)
Description	1st principal component (PC) of the covariance matrix of the individuals' genotypic data
 PC2	
Type	Numeric (Double)
Description	2nd principal component (PC) of the covariance matrix of the individuals' genotypic data
 PC3	
Type	Numeric (Double)
Description	3rd principal component (PC) of the covariance matrix of the individuals' genotypic data
 PC4	
Type	Numeric (Double)
Description	4th principal component (PC) of the covariance matrix of the individuals' genotypic data
 PC5	
Type	Numeric (Double)
Description	5th principal component (PC) of the covariance matrix of the individuals' genotypic data
 PC6	
Type	Numeric (Double)
Description	6th principal component (PC) of the covariance matrix of the individuals' genotypic data
 PC7	
Type	Numeric (Double)
Description	7th principal component (PC) of the covariance matrix of the individuals' genotypic data

 PC8		
Type	Numeric (Double)	
Description	8th principal component (PC) of the covariance matrix of the individuals' genotypic data	

 PC9		
Type	Numeric (Double)	
Description	9th principal component (PC) of the covariance matrix of the individuals' genotypic data	

 PC10		
Type	Numeric (Double)	
Description	10th principal component (PC) of the covariance matrix of the individuals' genotypic data	

## Physical Instances

 Wave IV PGS Risk-Tolerance		
Title	Wave IV PGS Risk-Tolerance	
File Name	riskpgs.sas7bdat	
Case Quantity	4755	
Variable Count	13	

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

Valid	Invalid	Minimum	Maximum	Mean	StdDev
4755	0	-3.276866...	3.6309792...	-7.32E-11	1E+0

 PGS_RIS2 - PGS_RISK_LDPRED_GWAS		
Type	Numeric (Double)	
Description	Polygenic score for general risk tolerance, obtained using LDpred method and standard GWAS results	

Valid	Invalid	Minimum	Maximum	Mean	StdDev
4755	0	-3.295509...	3.3845553...	1.21E-10	1E+0

 PGS_RIS3 - PGS_RISK_LDPRED_MTAG		
Type	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				
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Valid	Invalid	Minimum	Maximum	Mean	StdDev
4755	0	-3.218097...	3.6649067...	-2.93E-10	0.9999999...

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

Valid	Invalid	Minimum	Maximum	Mean	StdDev
4755	0	-0.069079...	0.0231721...	2.18E-7	0.0144819...

 PC2					
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Type	Numeric (Double)				
Description	2nd principal component (PC) of the covariance matrix of the individuals' genotypic data				

Valid	Invalid	Minimum	Maximum	Mean	StdDev
4755	0	-0.029585...	0.0783639...	-6.87E-6	0.0144841...

 PC3					
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Type	Numeric (Double)				
Description	3rd principal component (PC) of the covariance matrix of the individuals' genotypic data				

Valid	Invalid	Minimum	Maximum	Mean	StdDev
4755	0	-0.039884...	0.0325929...	4.27E-6	0.0144807...

 PC4					
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Type	Numeric (Double)				
Description	4th principal component (PC) of the covariance matrix of the individuals' genotypic data				

Valid	Invalid	Minimum	Maximum	Mean	StdDev
4755	0	-0.040301...	0.0676110...	-1.01E-6	0.0144760...

 PC5					
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Type	Numeric (Double)				
Description	5th principal component (PC) of the covariance matrix of the individuals' genotypic data				

Valid	Invalid	Minimum	Maximum	Mean	StdDev
4755	0	-0.040139...	0.0653000...	4.13E-6	0.0144626...

 PC6

Type	Numeric (Double)
Description	6th principal component (PC) of the covariance matrix of the individuals¶ genotypic data

Valid	Invalid	Minimum	Maximum	Mean	StdDev
4755	0	-0.043688...	0.0640709...	2.15E-6	0.0144829...

 PC7

Type	Numeric (Double)
Description	7th principal component (PC) of the covariance matrix of the individuals¶ genotypic data

Valid	Invalid	Minimum	Maximum	Mean	StdDev
4755	0	-0.058740...	0.0648081...	-2.31E-5	0.0144633...

 PC8

Type	Numeric (Double)
Description	8th principal component (PC) of the covariance matrix of the individuals¶ genotypic data

Valid	Invalid	Minimum	Maximum	Mean	StdDev
4755	0	-0.054538...	0.0590273...	2.19E-5	0.0144884...

 PC9

Type	Numeric (Double)
Description	9th principal component (PC) of the covariance matrix of the individuals¶ genotypic data

Valid	Invalid	Minimum	Maximum	Mean	StdDev
4755	0	-0.060507...	0.0531201...	1.13E-5	0.0144751...

 PC10

Type	Numeric (Double)
Description	10th principal component (PC) of the covariance matrix of the individuals¶ genotypic data

Valid	Invalid	Minimum	Maximum	Mean	StdDev
4755	0	-0.051092...	0.0544148...	1.05E-5	0.0144841...