Wave IV PGS Risk-Tolerance		
Number of observations: 4,755		

			PGS_RISK	num 21
4,755		range -3.2768 to 3.6309		
*Polygenic score for general risk tolerance, obtained using classic PLINK method and standard GWAS results. Note these are reverse-coded.				

			PGS_RIS2	num 21
4,755		range -3.2955 to 3.3845		
*Polygenic score for general risk tolerance, obtained using LDpred method and standard GWAS results				

			PGS_RIS3	num 21
4,755		range -3.2181 to 3.6649		
*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	num 21
4,755		range -0.0690 to 0.0231		
*1st principal component (PC) of the covariance matrix of the individuals genotypic data				

			PC2	num 21
4,755		range -0.0295 to 0.0783		
*2nd principal component (PC) of the covariance matrix of the individuals genotypic data				

			PC3	num 21
4,755		range -0.0398 to 0.0325		
*3rd principal component (PC) of the covariance matrix of the individuals genotypic data				

			PC4	num 21
4,755		range -0.0403 to 0.0676		
*4th principal component (PC) of the covariance matrix of the individuals genotypic data				

			PC5	num 21
4,755		range -0.0401 to 0.0653		
*5th principal component (PC) of the covariance matrix of the individuals genotypic data				

			PC6	num 21
4,755		range -0.0436 to 0.0640		
*6th principal component (PC) of the covariance matrix of the individuals genotypic data				

			PC7	num 21
4,755		range -0.0587 to 0.0648		
*7th principal component (PC) of the covariance matrix of the individuals genotypic data				

			PC8	num 21
4,755		range -0.0545 to 0.0590		
*8th principal component (PC) of the covariance matrix of the individuals genotypic data				

			PC9	num 21
4,755		range -0.0605 to 0.0531		
*9th principal component (PC) of the covariance matrix of the individuals genotypic data				

			PC10	num 21
4,755		range -0.0510 to 0.0544		
*10th principal component (PC) of the covariance matrix of the individuals genotypic data				