

Genetic Relationship between Schizophrenia and Nicotine Dependence

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Supplementary Table S1. Data sets used in FTND and cotinine genome-wide meta-analyses

Samples			SCZ risk	Reference
	N (phenotype)	Phenotype	score	
ALSPAC	258/161/193	COTININE/FTND/TFC	No	[1]
CARDIA	387	COTININE	No	[2]
FinnTwin	145	COTININE	No	[3-5]
FINRISK	218	COTININE	No	[6, 32]
Framingham	93	COTININE	No	[7]
GenMets	485	COTININE	No	[8]
MESA	189	COTININE	No	[9]
NESDA	808	COTININE	No	[10]
NTR	897/2436/2484	COTININE/FTND/TFC	No	[11-16]
TwinsUK	676	COTININE	No	[17]
YFS	392	COTININE	No	[18]
SAGE	2042/2042	FTND/TFC	Yes	[19-21]
SC	2479/2479	FTND/TFC	Yes	[22,23]
MGS	2496/2496	FTND/TFC	No	[24]
COPD	320/320	FTND/TFC	Yes	[25]
ISIB	299/265	FTND/TFC	Yes	[26]
CIDR370v1	166/574	FTND/TFC	Yes	[27]
CIDR370v3	251/598	FTND/TFC	Yes	[27]
CIDRc3	1710	TFC	Yes	[28]
CIDRc4	477	TFC	Yes	[29]
EAGLE	3051	FTND	Yes	[30]
AUTW	190/190	FTND/TFC	No	-
CEDAR	294/294	FTND/TFC	No	-
FIN1	1563/1573	FTND/TFC	No	[3-5]
FIN2	811/811	FTND/TFC	No	[3-5]
Yale_Penn	2114/2117	FTND/TFC	No	[31]

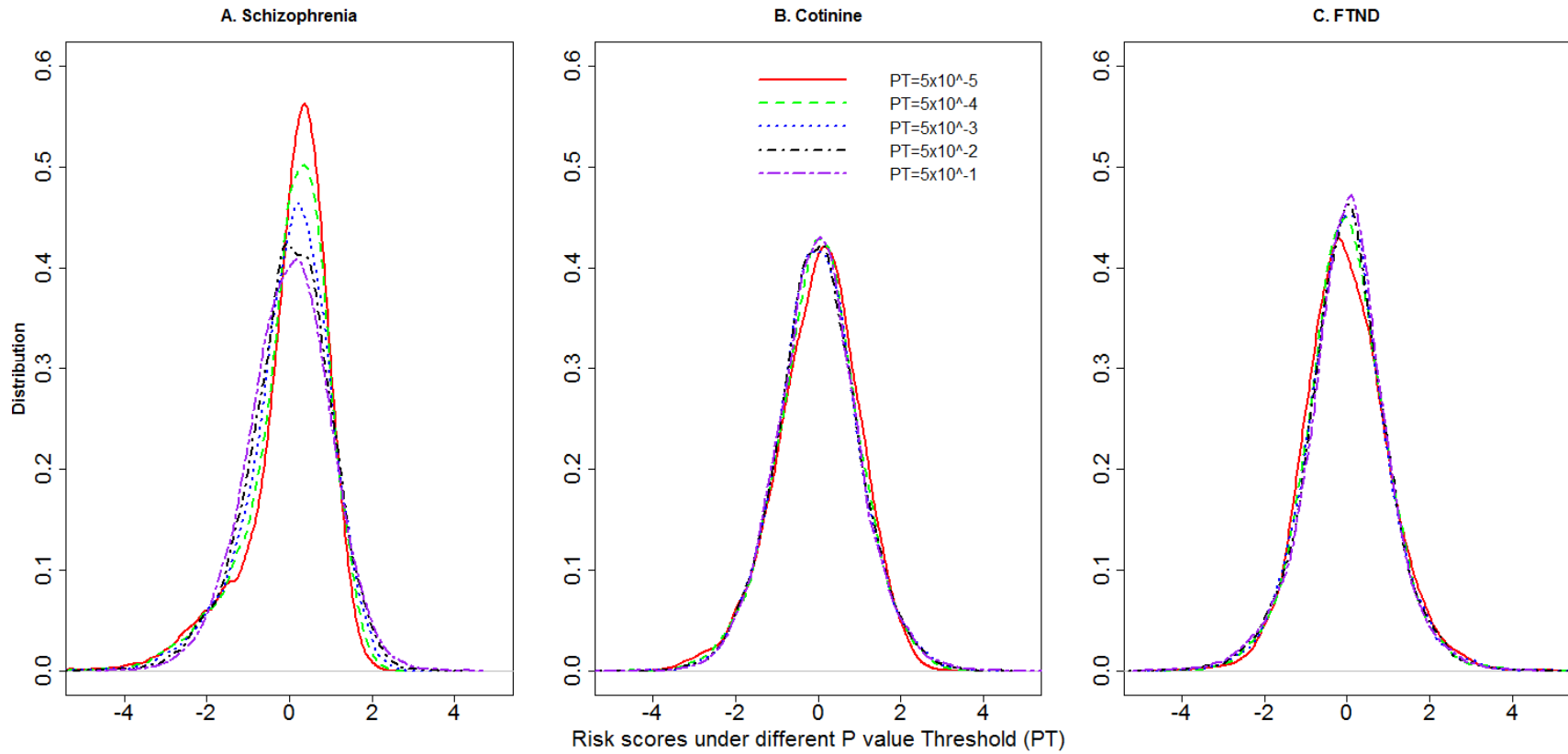
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Distribution of Risk Scores



Supplementary Figure S1. Distributions of genetic risk scores used in this study.