

Influence of dopamine-related genes on craving, impulsivity, and aggressiveness in Korean males with alcohol use disorder

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Background

The development of alcohol use disorder (AUD) progresses from impulsive drinking to compulsive alcohol intake via repeated binge drinking, withdrawal, and craving. The dopamine system, which plays a crucial role in various cognitive and reward processes, as well as risk-taking and impulsive behaviors, is a promising target for understanding the underlying molecular mechanism and treatment of AUD.

Recent evidence has suggested that abnormality of dopaminergic neurotransmission is implicated in the pathophysiology of AUD. Several studies have shown associations between dopamine-related gene variants and alcohol-related severity and cravings in patients with AUD. The aim of the study was to examine the genetic influence of dopamine system on the problematic drinking, impulsivity and aggression in a Korean male population with alcohol use disorder (AUD).

Methods

Participants: 295 males with AUD (Korean)

Measurements: Alcohol Use Disorders Identification Test (AUDIT), Obsessive-Compulsive Drinking Scale (OCDS), UPPS-P Impulsive Behavior Scale (UPPS-P), Buss-Perry Aggression Questionnaire (BPAQ) and Beck Depression Inventory (BDI)

Genetic polymorphism selection and genotyping

- 6 polymorphisms of dopamine-related genes, 1) 5 SNPs - rs4532 in *DRD1*, rs2283265 in *DRD2*, rs6280 in *DRD3*, rs1800497 in *ANKK1* and rs4680 in *COMT* genes

2) VNTRs of the *DAT1* gene

The analyses were conducted with the R package SNPAssoc, statistical significance was set at $p < 0.0083$ after Bonferroni correction.

Results

A significant association was detected between *DRD3* SNP rs6280 and OCDS score ($p = 0.0061$). With respect to impulsivity and aggression, rs4532 of *DRD1* was significantly related to UPPS-P score ($p = 0.0037$). Although it did not reach statistical significance after correction for multiple comparisons, rs4532 also showed nominally significant association ($p = 0.0261$) with BPAQ score.

Table 1. Socio-demographic and clinical characteristics of the study sample

	^a AUD (n=295)
Age, years	48.40 ± 7.87
Education, years	13.62 ± 3.91
Duration of AUD, years	17.62 ± 10.46
AUDIT	26.70 ± 7.69
OCDS	19.07 ± 7.29
UPPS-P	140.47 ± 18.05
BPAQ	71.54 ± 18.11
BDI	18.98 ± 12.58

Table 2. Characteristics of dopamine-related gene variants in the study

Gene	rs number	Chr	Position ^a	Geno ^b	P _{HWE} ^c	MAF	D/d ^d	Function
<i>DRD1</i>	rs4532	5	174870150	99.3	0.3990	0.119	A/G	5 prime UTR variant
<i>DRD2</i>	rs2283265	11	113285536	99.0	0.9061	0.457	G/T	Intron variant
<i>DRD3</i>	rs6280	3	113890815	98.3	0.4604	0.272	A/G	Missense variant
<i>ANKK1</i>	rs1800497	11	113270828	98.3	0.7210	0.434	G/A	Missense variant
<i>COMT</i>	rs4680	22	19951271	99.7	0.5607	0.276	G/A	Missense variant
<i>DAT1</i>	VNTR	5	1393746-1393824	99.7	0.2687	0.06	10R/≤9R	3 prime UTR variant

Table 3. The effects of dopamine-related gene variants on the severity of AUD

rs number	D/d ^a	DD/Dd/d ^b	AUDIT				OCDS					
			DD ^c	Dd ^c	dd ^c	Mean difference (95% CI)	p ^d	DD ^c	Dd ^c	dd ^c	Mean difference (95% CI)	p ^d
rs4532	A/G	225/66/2	27.20 ± 0.50	25.06 ± 0.97		-1.724 (-3.66, 0.22)	0.08	19.61 ± 0.49	17.13 ± 0.81		-2.128 (-3.82, -0.44)	0.01
rs2283265	G/T	85/147/60	26.87 ± 0.83	26.82 ± 0.63	26.35 ± 1.04	0.0309 (-1.14, 1.20)	0.95	19.52 ± 0.84	19.35 ± 0.61	17.78 ± 0.85	-0.4543 (-1.48, 0.57)	0.38
rs6280	A/G	156/110/24	26.14 ± 0.61	27.65 ± 0.75	26.88 ± 1.48	1.1839 (-0.10, 2.46)	0.07	18.53 ± 0.56	19.29 ± 0.70	21.67 ± 1.74	1.5753 (0.46, 2.69)	0.0061
rs1800497	G/A	91/146/53	26.86 ± 0.81	26.64 ± 0.62	26.30 ± 1.16	0.0168 (-1.18, 1.21)	0.97	19.78 ± 0.81	19.18 ± 0.60	17.53 ± 0.91	-0.6972 (-1.75, 0.35)	0.19
rs4680	G/A	152/122/20	26.74 ± 0.63	26.99 ± 0.69	24.55 ± 1.69	-0.4360 (-1.76, 0.88)	0.51	19.24 ± 0.57	18.80 ± 0.68	19.30 ± 1.80	-0.1453 (-1.31, 1.01)	0.80
<i>DAT1</i> VNTR	≥10R/≤9R	261/31/2	26.55 ± 0.48	27.52 ± 1.30		1.538 (-1.05, 4.12)	0.24	19.10 ± 0.46		18.70 ± 1.05	0.4090 (-1.88, 2.70)	0.72

Table 4. The effects of dopamine-related gene variants on impulsivity and aggression in AUD

rs number	D/d ^a	DD/Dd/d ^b	UPPS-P				BPAQ					
			DD ^c	Dd ^c	dd ^c	Mean difference (95% CI)	p ^d	DD ^c	Dd ^c	dd ^c	Mean difference (95% CI)	p ^d
rs4532	A/G	225/66/2	142.0 ± 1.15	134.6 ± 2.32		-6.367 (-10.63, -2.11)	0.0037	72.92 ± 1.19	66.99 ± 2.26		-5.3038 (-9.95, -0.65)	0.02
rs2283265	G/T	85/147/60	140.8 ± 2.06	142.0 ± 1.46	136.1 ± 2.24	-1.242 (-3.86, 1.38)	0.35	69.59 ± 2.03	72.93 ± 1.54	70.62 ± 2.09	1.3059 (-1.53, 4.14)	0.36
rs6280	A/G	156/110/24	139.7 ± 1.42	141.8 ± 1.79	139.4 ± 3.81	1.7072 (-1.18, 4.59)	0.24	72.62 ± 1.37	70.55 ± 1.83	68.54 ± 3.54	-1.4920 (-4.55, 1.56)	0.33
rs1800497	G/A	91/146/53	141.5 ± 1.96	141.9 ± 1.50	134.7 ± 2.22	-2.0831 (-4.75, 0.58)	0.12	70.26 ± 1.95	72.93 ± 1.53	70.11 ± 2.31	0.9170 (-1.96, 3.80)	0.53
rs4680	G/A	152/122/20	141.3 ± 1.43	139.2 ± 1.69	140.5 ± 3.89	-1.2109 (-4.14, 1.72)	0.41	72.11 ± 1.49	69.98 ± 1.49	74.05 ± 4.88	-0.4783 (-3.59, 2.64)	0.76
<i>DAT1</i> VNTR	≥10R/≤9R	261/31/2	140.6 ± 1.12	139.1 ± 3.22		0.0316 (-5.79, 5.86)	0.99	71.96 ± 1.13		67.73 ± 2.93	-2.715 (-8.98, 3.54)	0.39

Conclusion

Our results support that genetic variations of dopamine system may contribute to alcohol cravings and impulsivity in patients with AUD.