Prevalence and correlates of attention-deficit/hyperactivity disorder among youth with 16p11.2 copy number variation

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INTRODUCTION

- Attention-deficit/hyperactivity disorder (ADHD) is a neurodevelopmental syndrome affecting about 5% of children worldwide[1]. Like many behavioral disorders, ADHD is a phenotypically heterogeneous construct. The type and severity of symptoms present[2, 3] can vary across cases, as can age of onset[4]. Gender[5], intelligence[6], and co-occurring autism spectrum disorder (ASD)[7] may also influence the disorder's presentation. These factors all tend to complicate diagnosis and treatment.
- One way to constrain heterogeneity is by focusing on subgroups of individuals with ADHD associated with specific genetic risk factors. 16p11.2 copy number variation (CNV) is one such factor. 16p11.2 deletion (16p11.2del) and 16p11.2 duplication (16p11.2dup) are each associated with a neurodevelopmental syndrome that includes ADHD[8].
- The present study analyzed data from the Simons Variations in Individuals Project (VIP) cohort to compare the prevalence and correlates of ADHD among youth carrying 16p11.2del or 16p11.2dup and their noncarrier biological siblings.with 16p11.2 duplications (16p11.2dup), youth with 16p11.2 deletions (16p11.2del), and their noncarrier biological siblings.

METHODS

- As shown in Figure 1, the Simons VIP cohort was progressively narrowed down to youth only; to youth who underwent structured assessment for ADHD using the NIMH Diagnostic Interview Schedule for Children (DISC-Youth) or Diagnostic Interview Schedule for Young Children (DISC-YC); and finally to youth with symptom-count-level data available from those measures.
- For each member of the sample, DISC data were used to calculate variables representing presence of ADHD diagnosis and number of ADHD symptoms endorsed in the past year.
- 16p11.2dup carriers, 16p11.2del carriers, and noncarriers were compared in terms of age, IQ, number of ADHD symptoms, ADHD diagnosis, female gender, and ASD diagnosis. Pairwise post-hoc comparisons were carried out using Tukey's procedure (for continuous variables) or Bonferroni-corrected chi-square (for categorical variables).
- For the primary analysis, ADHD diagnosis was used as the outcome variable in a logistic regression model that included 16p11.2del carrier status, 16p11.2dup carrier status, age, full-scale IQ, gender, and co-occurring ASD diagnosis as predictors.
- Subsequent regressions were conducted among 16p11.2del carrier, 16p11.2dup carrier, and noncarrier family member subgroups.
- An exploratory linear regression analysis was also conducted using number of ADHD symptoms (rather than categorical diagnosis) as the outcome. Generalized estimating equations (GEEs) were used to estimate parameters in all regressions to control for correlations between family members.

Simons VIP Cohort			
* 665 subjects - 137 16p11.2del			Youth with
- 119 youth	Youth	Youth with DISC	symptom-count data
- 18 adults	* 324 subjects	* 225 subjects	* 179 subjects
- 130 16p11.2dup	- 119 16p11.2del	- 101 16p11.2del	- 79 16p11.2del
- 69 youth	- 69 16p11.2dup	- 55 16p11.2dup	- 47 16p11.2dup
- 61 adults	- 136 noncarrier	- 69 noncarrier	- 53 noncarrier
- 398 noncarrier			
- 136 youth			
- 262 adults	Figure 1: Sample select	ion process	

able 1: Characteristics of 16p11.2 CNV carriers and their non-carrier siblings

2.523

1.178

ASD Dx

Table 1: Chara	cteristi	cs of 16p	511.2 C	NV carr	iers and	their i	non-carr	ier siblii	ngs.			
Characteristic Total (n =		tal 16p11 = 179) (n = 47		p11.2dup 16p1		.2del	Noncar	rier	Main effect	Post-hoc comparisons		
				7)	(n = 79)		(n = 53))			1100	
	M	SD	M	SD	M	SD	M	SD	ANOVA p	Tukey's		b
Age	7.83	4.20	6.65	4.15	7.45	3.92	9.45	4.25	0.002	Comparison dup-del dup-noncarrier del-noncarrier dup-del dup-noncarrier del-noncarrier dup-del dup-noncarrier		0.54 0.002 0.02 0.31 < 0.001 < 0.001
IQ	87.87	19.98	77.87	20.47	82.33	14.99	104.98	14.95	< 0.001			
ADHD Sx	6.78	6.26	8.79	6.62	7.48	5.85	3.94	5.59	< 0.001			0.46 < 0.001 0.003
	#	%	#	%	#	%	#	%	$\chi^2 p$	Pairwise Company	7.0	p (corrected)
Female gender	80	45	25	53	36	46	19	36	0.22	dup-del dup-noncarrier del-noncarrier		1.00 0.37 1,00
ASD Dx	25	14	9	19	14	18	2	4	0.04	dup-del dup-noncarrier del-noncarrier		1.00 0.10 0.10
ADHD Dx	78	44	24	52	42	53	12	23	0.001	dup-del dup-no del-non	ncarrier	1.00 0.003 0.02
Table 2: Predict	tors of	ADHD	diagnos	sis amon	g 16p1	1.2 CN	IV carrie	ers and t	heir non-carr			
Predictor		β		SE			OR (9	95% CI			p	
16p11.2del car	rier	0.910	.910				· ·	2.49 (0.966 - 6.390)			0.059	
16p11.2dup ca		0.727		0.532	0.482			2.07 (0.729 — 5.874)			0.172	
Age		0.006		0.003			1.01 (0.999 — 1.012)			0.099		
IQ		-0.025		0.010					<u> </u>		0.016	
Female gender		0.517			0.324			0.97 (0.956 — 0.995) 1.68 (0.888 — 3.166)			0.010	
ASD Dx		0.300		0.509			1,35 (0.498 — 3.662)			0.555		
Table 3: Predict			diagnos		g dupli	cation o		<u> </u>	,			
Predictor		β		SE	OT			95% CI			p	
Age	1	0.014		0.008		- 	1.01 (0.997 — 1.030)			0.098		
IQ	-0.018			0.015			0.98 (0.952 — 1.010)			0.231		
Female gender		$\frac{0.634}{0.271}$		0.697				1.89 (0.481 — 7.390) 0.69 (0.119 — 4.010)			0.363	
ASD Dx Table 4. Predict		-0.371	diagna	0.898	a dalaci	0.5		`	— 4.010)		0.680	
Table 4: Predict	tors of	R	diagnos		g deleti	on carr					6	
Predictor	ı	β			SE 0.005			OR (95% CI)		<i>p</i>		
Age			0.005			1.0 (0.997 - 1.020) 0.98 (0.946 - 1.010)		0.150				
IQ Female gender				0.017			0.98 (0.946 - 1.010) 1.5 (0.673 - 3.650)		0.190			
ASD Dx			0.601		<u> </u>	1.8 (0.547 - 5.760)		0.430				
Table 5: Predict			diagnos		g non-c	carrier s	<u> </u>		,			
Predictor		β		SE				95% CI			p	
Age		-0.013		0.007			 		- 1.00)		0.073	
IQ		-0.110		0.040		0.90	0.90 (0.829 — 0.968)			0.006		
Female gender		0.976		0.860				`	<u> </u>		0.256	
ASD Dx		-0.712		1.088		11-2-6			<u>4.110)</u>	•1_1_	0.508	
Table 6: Predict	tors of	ADHD	sympto	_	ng 16p	11.2 C			their non-car	rier siblii	ngs.	
Predictor 16p11.2dup ca	rrier	<u>В</u> 11.057		SE 2.608				% CI)	<u> </u>		0.108	
16p11.2del car		2.148		1.335			0.11 (-0.057 — 0.272)			0.201		
Age		-0.004		0.007					<u>8 — 0.081)</u>		0.545	
IQ		<u>-0.069</u>		0.0229	9				$\frac{50.078)}{0.271}$		0.003	
Female gender		1.606		0.915				`	$\frac{-0.271)}{0.260}$		0.079	



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RESULTS

- Relative to noncarriers, 16p11.2dup and 16p11.2del carriers were younger, had lower IQs, and had more ADHD symptoms. They were also more likely to have a diagnosis of ADHD or ASD (Table 1). Among the entire sample, ADHD diagnosis was weakly negatively predicted by IQ (OR 0.97 for every point decrease, 95% CI 0.956 0.995, p = 0.016) (Table 2). This relationship was not identified in regressions conducted among duplication carriers only (Table 3) or deletion carriers only (Table 4), but remained in a regression carried out among noncarriers only (OR 0.90 for every point decrease, 95% CI 0.829 0.968, p = 0.006) (Table 5).
- In the exploratory regression against number of ADHD symptoms endorsed in the past year, IQ negatively predicted ADHD symptoms, with 0.07 fewer symptoms for every point decrease (standardized effect -0.22, 95% CI -0.365 -0.078, p 0.003). ASD was a positive predictor: subjects with ASD had 2.5 more ADHD symptoms than those without (standardized effect 0.14, 95% CI 0.012 0.269, p = 0.032).

CONCLUSIONS

• The finding that IQ negatively predicts ADHD diagnosis in our entire analysis sample and among noncarriers is consistent with evidence that children with ADHD tend to have lower IQ than their unaffected peers[9]. The lack of an association between IQ and ADHD within 16p11.2dup or 16p11.2del carriers is interesting given that ADHD is more prevalent, and IQ is on average lower, in those groups. It may be that IQ differences are a better predictor of ADHD among neurotypical youth than those who may already have some level of cognitive compromise. The finding that ASD diagnosis predicts number of ADHD symptoms but not ADHD diagnosis per se is consistent with emerging evidence that phenotypic traits of ASD and ADHD can tend to overlap[10], which may be particularly true in those affected by the 16p11.2 CNV. In the future, with the recruitment of more 16p11.2 CNV carriers, it may be possible to characterize their overlapping symptoms in more detail.

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Contact

0.032

|0.14 (0.012 - 0.269)|

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