SFARI

Simons Foundation Autism Research Initiative

Annual Science Meeting Washington, DC September 12 – 14, 2010

MEETING AGENDA

SUNDAY, SEPTEMBER 12

2:00-5:00	Registration				
5:00-7:00	introductions; All SFARI Investigators give a brief (one-rr				
	of their SFARI-supported research.				
7:00-7:45	Hors d'oeuvres and cocktails				
7:46-9:46	Dinner & Evening Program				
	Welcome: JIM Simons				
	Keinarks: Genry Fischaddii Keynete Address: Dichard Litten, Human Hynertension:				
	General-Dopulation Therapeutics				
	General-Population merapeutics				
MONDAY, SEPTEN	IBER 13				
7:30-8:30 Brea	lkfast				
	Session #1: Cognition and Behavior				
	Session Moderator. Revin Feiphiey				
8:30-9:00	Ami Klin / Warren Jones				
9:00-9:30	Joseph Piven				
9:30-10:00	Randy Buckner				
10:00-10:30	Break				
10:30-11:00	Rebecca Saxe				
11:00-11:30	Marlene Behrmann				
11:30-12:00	Open Discussion				
12:00-1:10	Lunch				

Session #2: Gene Discovery and Expression Part I Session Moderator: Lauren Weiss

1:10-1:40	Mike Wigler
1:40-2:10	Matt State
2:10-2:40	Evan Eichler
2:40-3:10	Break

Session #3: Gene Discovery and Expression Part II Session Moderator: Huda Zoghbi

3:10-3:30	Stephen Warren
3:30-3:50	Isaac Kohane

3:50-4:10	Barbara Wold			
4:10-4:30	Break			

4:30-4:50Eric Courchesne4:50-5:10Dan Arking5:10-5:30Dan Geschwind5:30-6:00Open Discussion7:30-8:00Hors d'oeuvres and cocktails8:00-9:30Dinner & Evening ProgramIntroduction: Marilyn Simons
"Bye" A movie by Anthony Morrison
Remarks: Catherine Lord

TUESDAY, SEPTEMBER 14TH

7:30-8:30 Breakfast

Session # 4: Molecular Mechanisms / Synaptic Biology Part I Session Moderator: Bernardo Sabatini

- 8:0-8:50 Guoping Feng
- 8:60-9:10 Ricardo Dolmetsch
- 9:10-9:30 Hazel Sive
- 9:30-9:50 Break
- 9:50-10:10 Maria Karayiorgou
- 10:10-10:30 Ben Philpot
- 10:30-11:00 Open Discussion and Break

Session # 5: Molecular Mechanisms / Synaptic Biology Part II Session Moderator: Gerry Fischbach

- 11:00-11:30 Luis:Parada
 - 11:30-12:00 Thomas Südhof
 - 12:00-12:45 Eric Kandel, The Long and Short of Autism Research 2010
 - 1:00 Lunch & Departure

Copy Number Variant – Deletion (M. Wigler)



Copy Number Variant - Duplication



Gene Discovery – Copy Number Variants (CNVs) E. Eichler





Markers of Functional Specialization: Asymmetry (R. Buckner)





Research Report

Atypical functional lateralization of language in autism spectrum disorders

Natalia M. Kleinhans^{a,*}, Ralph-Axel Müller^{b,d}, David N. Cohen^c, Eric Courchesne^{e,f}

^aDepartment of Radiology, University of Washington, Seattle, WA 98195, USA ^bDepartment of Psychology, San Diego State University, San Diego, CA 92120, USA ^cGeorge Washington University School of Medicine, Washington, DC 20037, USA ^dDepartment of Cognitive Science, University of California, San Diego, CA 92093, USA ^cDepartment of Neuroscience, University of California, San Diego, CA 92093, USA ^cDildren's Hospital Research Center, San Diego, CA 92123, USA

ARTICLE INFO

ABSTRACT

Article history: Accepted 29 April 2008 Available online 14 May 2008

Keywords: fMRI Asymmetry Letter fluency Category fluency Frontal lobe Impaired language is a prominent behavioral marker of autism spectrum disorders (ASD), but its neurobiological underpinnings are incompletely understood. We studied letter and category fluency in 14 high functioning ASD individuals and 14 age-matched controls. Each fluency condition was compared to self-paced repetition of the word "nothing." Responses were recorded to monitor performance. In letter fluency, the ASD group had significantly greater activation than controls in the right frontal and right superior temporal lobes. Betweengroup differences were not observed in left prefrontal cortex. By examining functional asymmetry in frontal cortex, we found that the ASD group had significantly reduced lateralization of activation patterns in letter fluency compared to the controls. In category fluency, no between-group differences in lateralization were found, in light of greater bilateral activation in controls. These findings indicate reduced hemispheric differentiation for certain verbal fluency tasks in ASD, consistent with some previous evidence of atypical functional and structural asymmetries in autism. Abnormal functional organization may contribute to the language impairment seen in ASD.

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1. Introduction

Atypical language development is a prominent behavioral marker of autism spectrum disorders (ASD). In young autistic children, language deficits are among the most salient overt symptoms. Lack of spoken language by 2 years of age is often the first indicator to impel parents to seek professional advice (De Giacomo and Fombonne, 1998) or to be recognized as a significant risk factor by pediatricians. The timing of language

acquisition is a key predictor of functional outcome; acquisition of useful speech by 5-6 years of age has been associated with better educational and functional attainment in adulthood (Howlin et al., 2000).

The severity of language deficits in individuals with ASD varies markedly. Approximately half of all individuals with autistic disorder remain nonverbal throughout life, while other individuals may develop fluent language and extensive vocabularies (Volkmar et al., 2000). However, even in high

Challenges

Very little is known about diversity in systems-level brain architecture.

What is normal and what is atypical is unclear.

Genetic influences emerge from rare variants or may exert small effects necessitating large samples for many forms of discovery. Need an approach to explore diversity across normal individuals.

- Robust within individuals.
- Amenable to collection of large samples.
- Provides a foundation for genetic explorations.
- Can be connected to behavior.
- Extendable to ASD and related groups.

Paternal imprinting of Ube3a (B. Philpot)





Autism



Therapeutic Strategy



Unsilencing of paternal Ube3a throughout brain



Unsilencing of paternal Ube3a throughout brain





Function of Ube3a



modified from Dan, 2008

Ube3a in Autism Spectrum Disorders

• Deletions or mutations of *UBE3A* underlie Angelman syndrome, a severe mental retardation and autism spectrum disorder

(Wagstaff and Beaudet labs, 1997; Peters et al., 2004)

• Maternal duplication of the 15q11-13 region, spanning the UBE3A gene, is one of the most identifiable genetic causes of autism

(Cook et al., 1997; Schroer et al., 1998; Glessner et al., 2009)

• UBE3A is required for experience-driven plasticity during critical periods

(Yashiro et al., 2009; Sato and Stryker, 2010)

Lesson #3: Two-Hit CNV Model of Disease

Disorder	Cases (n)	Total	Incidence	de novo	Second	Second	<i>p</i> -value
		cases	(%)	(%)	hits	hit (%)	
Smith-Magenis syndrome	25	20,647	0.12	100%	0	0%	
17q21.31 syndrome	29	20,647	0.14	100%	0	0%	
Williams syndrome	60	20,647	0.29	100%	3	5%	0.52
DiGeorge syndrome	113	20,647	0.54	76.4%	8	7%	0.17
15q13.3 deletion	66	20,647	0.32	24%	6	9.1%	0.1
16p11.2 deletion	91	20,647	0.44	76.4%	9	9.9%	0.03
22q11.2 duplication		26,176	0.37	22.5%	11	11.3%	0.011
1q21.1 deletion		25,866	0.37	33%	11	11.2%	0.01
16p12.1 deletion		21,127	0.19	4.76%	9	21.9%	0.0002
Controls ¹	471	5285			21	4.4%	
Controls		5285			21	0.39%	

(unconditioned)2

1For comparison, controls were conditioned to have at least one large CNV (>500 kbp) and then the number of second hits in these cases were counted. 2controls were not (conditioned)2 for first hit. This gives an estimate of two hits in the general population

compared to affected individuals

Summary (E. Eichler 9/2010)

- Large CNVs (~500 kbp) account for ~11% of SSC– inheritance assessment suggests that 1/3 rd are *de novo*.
- Refinement of 17 targeted regions identifies candidate smaller CNVs in 13.8% of samples eg. CDH8 and ACACA
- New hotspots emerging eg. 15q25 highlighting specific genes IMMP2L, ATXBP1, CTNNA3 (1-2% of patients)
- Exome sequencing on 17 trios—reveals de novo and inherited mutations in strong candidates (FOXP1, GRIN2B, CNTNAP2, SHANK2).

Copy Number Variants



General Strategy

