

**BIOGRAPHICAL SKETCH**

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NAME <b>James S. Sutcliffe</b>	POSITION TITLE <b>Associate Professor</b>		
eRA COMMONS USER NAME sutclijs			
EDUCATION/TRAINING (Begin with baccalaureate or other initial professional education, such as nursing, and include postdoctoral training.)			
INSTITUTION AND LOCATION	DEGREE (if applicable)	YEAR(s)	FIELD OF STUDY
Auburn University, Auburn, AL	B.S.	1986	Biochemistry
Emory University, Atlanta GA	Ph.D.	1992	Human Genetics
Baylor College of Medicine	Post-Doc	1992-1997	Human Genetics

**A. Positions:**

1985-1986 Research Assistant, Southeast Experimental Station, U.S. Forestry Service, Auburn, AL  
 1986-1992 Graduate Assistant, Department of Biochemistry, Emory University, Atlanta, GA  
 1992-1997 Res. Assoc., Department of Molecular and Human Genetics, Baylor College of Medicine  
 Houston, TX  
 1997-2005 Assistant Professor, Department of Molecular Physiology and Biophysics  
 1997- Investigator, Center for Molecular Neuroscience, Vanderbilt Kennedy Center for Research on  
 Human Development and Center for Human Genetics Research; Vanderbilt University,  
 Nashville, TN  
 2005- Associate Professor, Department of Molecular Physiology & Biophysics  
 2006- Associate Professor, Department of Psychiatry (secondary appointment)

**B. Selected peer-reviewed publications (in chronological order):**

Verkerk AJMH, Pieretti M, Sutcliffe JS, Fu YH, Kuhl DPA, Pizzuti A, Reiner O, Richards S, Victoria MF, Zhang F, Eussen BE, Van Ommen GJB, Blonden LAJ, Riggins GJ, Chastain JL, Kunst CB, Caskey CT, Nelson DL, Oostra BA and Warren ST: Identification of a gene (*FMR-1*) containing a CGG repeat coincident with a fragile X breakpoint cluster region which exhibits length variation in fragile X syndrome. *Cell*, 65: 905-914 (1991).

Sutcliffe JS, Zhang F, Nelson DL, Caskey CT and Warren ST: PCR amplification and analysis of yeast artificial chromosomes. *Genomics*, 13: 1303-1306 (1992).

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- Matsuura T, Sutcliffe JS, Fang P, Galjaard R-J, Jiang Y-h, Benton CS, Rommens JM and Beaudet AL: *De novo* truncating mutations in E6-AP ubiquitin-protein ligase (*UBE3A*) in Angelman syndrome. *Nature Genet*, 15: 74-77 (1997).
- Albrecht A, Sutcliffe JS, Cattanach BM, Beechey CV, Armstrong D, Eichele G and Beaudet AL: Imprinted expression of the murine Angelman syndrome gene *Ube3a* in hippocampal and Purkinje neurons. *Nature Genet*, 17: 75-78 (1997).
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**C. Research Support**

**Ongoing Support:**

2 R01 MH061009 (Sutcliffe) NIH/NIMH Genetic Analysis of 15q11-q13 in Autism	5/7/04-4/30/09
1 R01 NS049261 (Sutcliffe) NIH/NINDS Unraveling the genetic etiology of autism	4/04/06-3/31/11
Simons Simplex Autism Family Collection (Sutcliffe) Simons Foundation Simons Simplex Collection Consortium	12/1/06-3/31/10
Autism Genome Project Consortium (Sutcliffe) Autism Speaks	1/1/07-12/31/09
1 P50 HD055751 (Cook) NIH/NICHD ACE: Translational studies of insistence of sameness in autism spectrum disorders	6/1/07-5/31/12
Simons Simplex Collection Genetics Consortium (Sutcliffe) Simons Foundation	9/1/09-8/31/10
1 R01 MH089482 (Sutcliffe) NIH/NIMH Elucidating the Genetic Architecture of Autism by Deep Genomic Sequencing	9/20/09-9/29/11

**Past Support:**

Autism Speaks Augmentation Award (Sutcliffe) Autism Speaks Altered regulation of SERT is mediated by autism associated functional variation in $\beta$ 3 integrin	4/1/07-3/30/08
NIH R01 HD35684 (Dykens) NIH/NICHD Prader-Willi syndrome: correlates of compulsivity	7/1/03-3/31/08