

# Deciphering the genetics of speech & language disorders



Simon E. Fisher

Language and Genetics  
Max Planck Institute for Psycholinguistics  
Nijmegen, the Netherlands



---

---

---

---

---

---

---

---

# An instinct to acquire an art



Hmmm...  
Little sisters are  
strange....



Gold was my  
favourite colour but  
now I like purple!

Simon E. Fisher

Max Planck Institute, Nijmegen

---

---

---

---

---

---

---

---

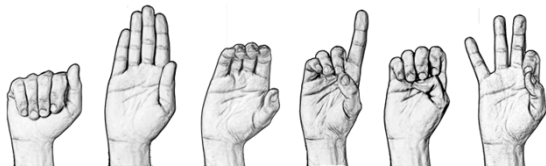


Image © William Vicars, at Lifeprint.com

Simon E. Fisher

Max Planck Institute, Nijmegen

---

---

---


---

---

---

---

---



**Simon E. Fisher** **Max Planck Institute, Nijmegen**

---

---

---

---

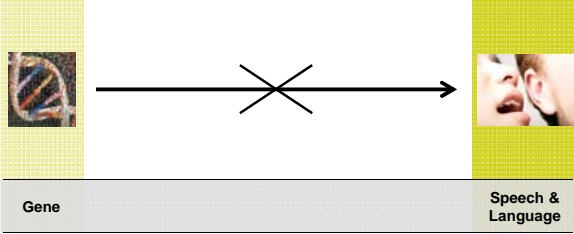
---

---

---

---

### The myth of the abstract gene



Genes should not be viewed as abstract entities with a mysterious ability to directly control a person's behaviour or cognitive skills

---

---

---

---

---

---

---

---

### Beyond the abstract gene....

TCGATTAGCTAGAGGAGAGGAGAGTTCGATTCCTGGGGCTCTCTCGGATAGTAGATAGCAGTACAGCATCGG  
 GATGCATGCTATTAGCTAGCTAGCAGTAGTACGTAGCTCTGATCGATCGGTAGCTAGCTATGCTAGCT  
 AGTCGAGGCGAAGGGGGCTCAGAGATCTCTCTCTCTCTCTCGATAGCTAGCTAGCTAGCTAGCTAGC  
 GAGGAGTAGCTAGTTCGATCGTCTCGAGAGGACTCTCTGGGGCTCTCGCTAGGGAGGAGAGGAGAGT  
 CTGTATGCTAGCTGCATGCTACGGAGGGATAGTAGATAGCAGTACCGTTAAATCGTACTGGAGCTAGC  
 TAGGGCGGATAGCTAGCGATCGATCGAGGGGGCTCTCTCGATCGATCGATCGAGCATCGGATCGAGGAG  
 CGAGGAGCTAGCGTAGTTCGACTATGCGAGCGATCTATGCTATCGAGGGGGCTCTCTCGAATTGTAG  
 CTAGTGATCGTAGCTAGCTAGAGAGGAGAGATCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT  
 AGCTAGTACGATGCTAGCTAGCTAGCTAGCTAGCTAGCTAGCTAGCTAGCTAGCTAGCTAGCTAGCTA  
 CGTAGCTATATTATTCTAGCATCGTAGGGATTTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT  
 GTGATCGTAGCTAGCTAGAGAGTATGCTAGCTAGGGAGGAGGAGAGCGACTGATCGATCGGTAGCTAG  
 CGATCGACTATGCGAGCGATGCTAGCTAGCTAGCTAGCTAGCTAGCTAGCTAGCTAGCTAGCTAGCTA  
 TTACGTAGCTAGCTAGCTAGCTAGTCACTGCTAGCTAGCTAGCTAGCTAGCTAGCTAGCTAGCTAGCTA  
 TTTACAGTGCATGCTAGCTAGCTAGCTAGCTAGCTAGCTAGCTAGCTAGCTAGCTAGCTAGCTAGCTA  
 CTCTTACTATGCGAGCGATGCTAGCTAGCTAGCTAGCTAGCTAGCTAGCTAGCTAGCTAGCTAGCTA  
 AGCTAGGGCGGATAGCTAGCGACGGCTAGCTAGCTAGCTAGCTAGCTAGCTAGCTAGCTAGCTAGCTA  
 TAGCTAGAGGAGAGGAGAGTTCGACTGATCGATCGGTAGCTAGCAAGCGCTAGCTAACTCTCTGGGGCT  
 CTCTCGGATAGTAGATAGCAGTACAGTTCGGGATGCATGCTATTTTACGTAGCTAGCTAGCTAGCTAGCT  
 AGTGCATGCTACGTTAGCTAGCTAGCTAGCTAGCTAGCTAGCTAGCTAGCTAGCTAGCTAGCTAGCTAG  
 CTAGCTCTCGGATAGTAGATAGCAGTACAGCATCGGGATGCATGCTATTACTATGCGCGAGCGATCGAGC  
 ATCGACAGCGATCGACTAGAGGAGAGGAGATCTCTCTTACTATGCGAGCTTACGGAGAGGAGAGGAGG  
 ACTGATCGATCGGTAGCTAGCTATGCTATCGAGGGGGCTCTCTCGAGAGGACTCTCTGGGGCTCTCTC

**Simon E. Fisher** **Max Planck Institute, Nijmegen**

---

---

---

---


---

---

---

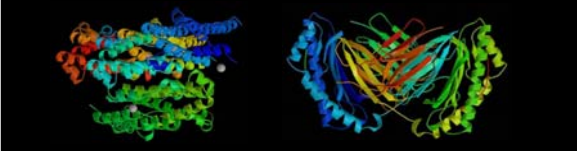
---

### Cracking the code



AUGAUGCAGGAAUCUGCGACAGAGACAAUAAGCAACAGUCAAUG

MetMetGlnGluSerAlaThrGluThrIleSerAsnSerSerMet



Simon E. Fisher Max Planck Institute, Nijmegen

---

---

---

---


---

---

---

---

### Genome, Interrupted



Promoter


transcription

Exon Intron Exon Intron Exon Intron Exon AAAAAAAAAA

splicing

AUG UGA AAAAAAAAAA

translation



Simon E. Fisher Max Planck Institute, Nijmegen

---

---

---

---



---

---

---

---

### Variety is the spice of life



Simon E. Fisher Max Planck Institute, Nijmegen

---

---

---

---


---

---

---

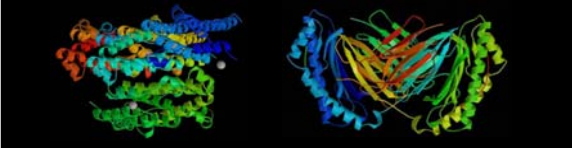
---

### Cracking the code



AUGAUGCAGGAAUCUGCGACA**A**AGACAAUAAGCAACAGUCAAUG

MetMetGlnGluSerAlaThr**Lys**ThrIleSerAsnSerSerMet



Simon E. Fisher Max Planck Institute, Nijmegen

---

---

---

---


---

---

---

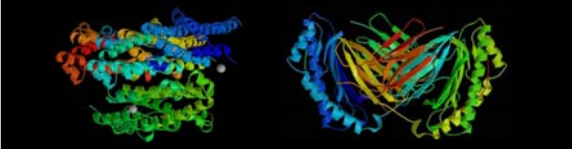
---

### Cracking the code



AUGAUGCAGGAAUCUGCGACA**U**AGACAAUAAGCAACAGUCAAUG

MetMetGlnGluSerAlaThr**Stop**



Simon E. Fisher Max Planck Institute, Nijmegen

---

---

---

---


---

---

---

---

### Genome, Interrupted



Promoter


transcription

Exon Intron Exon Intron Exon Intron Exon AAAAAAAAAA

splicing

AUG UGA AAAAAAAAAA

translation



Simon E. Fisher Max Planck Institute, Nijmegen

---

---

---

---

---

---

---

---

## Lost for words



- A subset of children grow up in language-rich environments, but do not develop normal skills
- Sometimes there is no overt cause e.g. cerebral palsy, hearing loss, mental retardation, cleft palate
- Developmental language disorders are often found to cluster in families
- Twin studies consistently indicate high heritability i.e. role for genetic factors



Simon E. Fisher

Max Planck Institute, Nijmegen

---

---

---

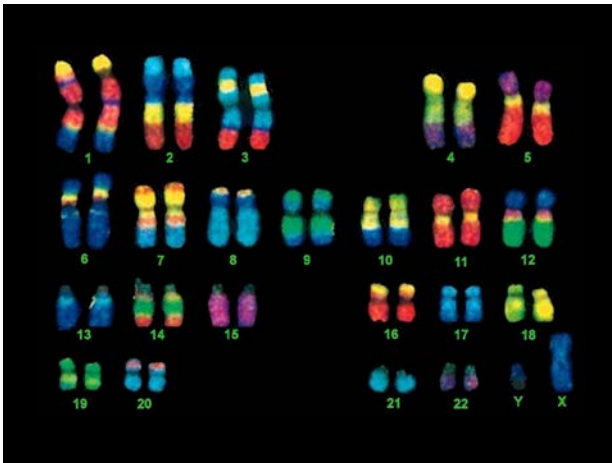
---

---

---

---

---



---

---

---

---

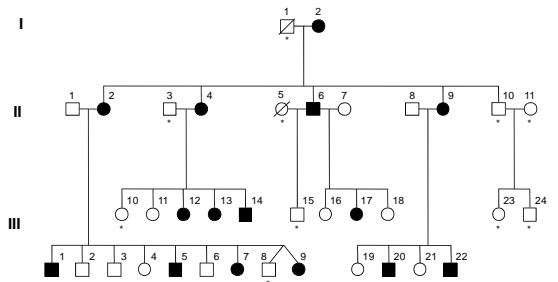
---

---

---

---

## The KE family



Fisher et al. (1998) *Nature Genet* 18: 168-70

Simon E. Fisher

Max Planck Institute, Nijmegen

---

---

---

---

---

---

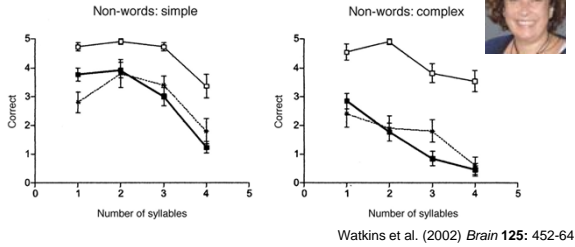
---

---

## Childhood apraxia of speech



- Difficulties in the learning and production of sequences of mouth movements, impairs speech development



Watkins et al. (2002) *Brain* 125: 452-64

Simon E. Fisher

Max Planck Institute, Nijmegen

---

---

---

---

---

---

---

---

---

---

## Simple inheritance, complex phenotype



- Difficulty learning/producing sequences of complex orofacial movements underlying speech
- Deficits in wide range of language-related skills:
  - Spoken AND written language
  - Impairment extends to receptive domain
  - Comprehension and production of grammar affected
- Non-verbal deficits are not central to disorder:
  - Unaffected individuals with low non-verbal IQ
  - Affected individuals with normal non-verbal IQ
  - Deficits in verbal cognition more severe/wide-ranging

Simon E. Fisher

Max Planck Institute, Nijmegen

---

---

---

---

---

---

---

---

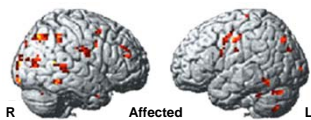
---

---

## Imaging the brain



Functional MRI – silent verb generation task



Liegeois et al. (2003) *Nature Neurosci* 6: 1230-7

Simon E. Fisher

Max Planck Institute, Nijmegen

---

---

---

---

---

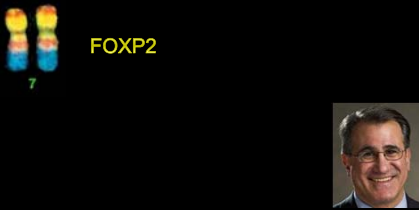
---

---

---

---

---



**FOXP2**

Fisher et al. (1998) *Nature Genet* 18: 168-70; Lai et al. (2001) *Nature* 413: 519-23

---

---

---

---

---

---

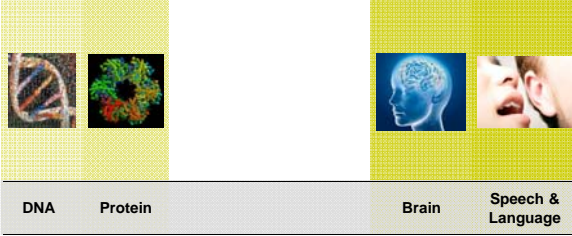
---

---

---

---

### Mind the gap!



**DNA**      **Protein**      **Brain**      **Speech & Language**

Simon E. Fisher      Max Planck Institute, Nijmegen

---

---

---

---

---

---

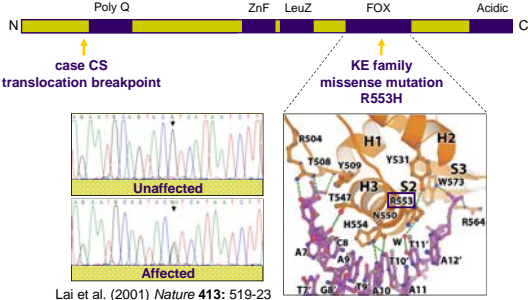
---

---

---

---

### The FOXP2 protein



**case CS translocation breakpoint**

**KE family missense mutation R553H**

Lai et al. (2001) *Nature* 413: 519-23

Simon E. Fisher      Max Planck Institute, Nijmegen

---

---

---

---

---

---

---

---

---

---

### The FOXP2 protein

MacDermot et al. (2005) *Am J Hum Genet* 76: 1074-80

Simon E. Fisher Max Planck Institute, Nijmegen

---

---

---

---

---

---

---

---

### FOXP2 is a transcription factor

Simon E. Fisher Max Planck Institute, Nijmegen

---

---

---

---

---

---

---

---

### Mind the gap!

DNA Protein Cell/Neuron Brain Speech & Language

Simon E. Fisher Max Planck Institute, Nijmegen

---

---

---

---

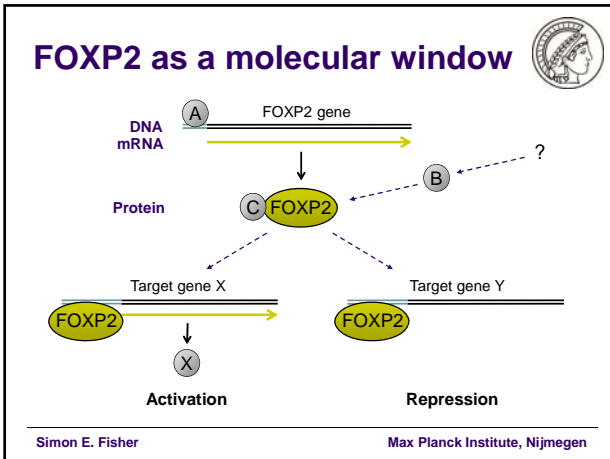
---

---

---

---






---

---

---

---

---

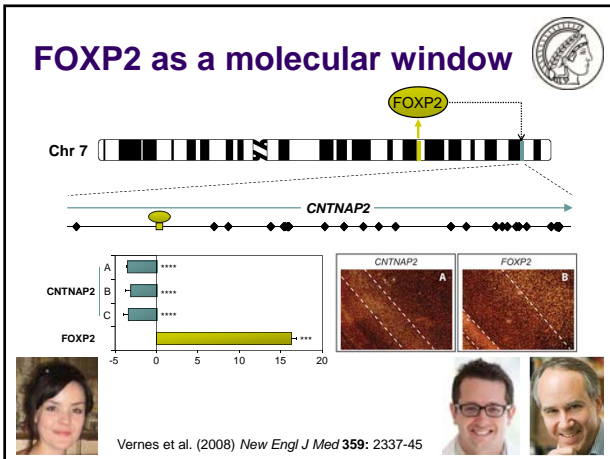
---

---

---

---

---




---

---

---

---

---

---

---

---

---

---

### FOXP2 as a molecular window

Chr 7: FOXP2 gene location

CNTNAP2 gene structure

- 184 families with common forms of language impairment; CNTNAP2 “risk” variants correlated with reduced abilities
- These variants are independently associated with delayed language (“age-at-first-word”) in autistic children
- Alleles are correlated with early language in general population

Vernes et al. (2008) *NEJM* 359: 2337-45;  
 Alarcón et al. (2008) *Am J Hum Genet* 82:150-9;  
 Whitehouse et al. (2011) *G2B* 10: 451-6

---

---

---

---

---

---

---

---

---

---

## A tale of two genes



- Distinct types of genetic effects:
- **FOXP2** = rare coding mutations change the sequence of the encoded protein, causing a monogenic speech/language disorder
- **CNTNAP2** = common variants outside the protein coding regions are associated with increased risk of language problems
- Converge on a shared functional pathway

Simon E. Fisher

Max Planck Institute, Nijmegen

---

---

---

---

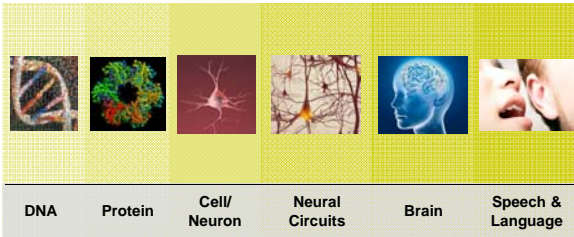
---

---

---

---

## Mind the gap!



Simon E. Fisher

Max Planck Institute, Nijmegen

---

---

---

---

---

---

---

---

## A tricky question.....



Does this mouse have a speech and language disorder?

Simon E. Fisher

Max Planck Institute, Nijmegen

---

---

---

---

---

---

---

---

## What is Foxp2 doing in a mouse?

Human 715 aa: N [D] [2] [Q] [N] [S] [FOX] [R553H] [R552H] [Acidic] C

Mouse 714 aa: N [E] [-] [T] [N] [FOX] [R553H] [R552H] C

- Mouse Foxp2 differs from human protein by only 3 substitutions, and slightly shorter stretch of Q residues
- Switched on in corresponding regions in rodent and human brain => includes neuronal subpopulations in the cortex, basal ganglia, thalamus & cerebellum
- Circuits implicated in multimodal sensory processing, sensorimotor integration and motor-skill learning

Simon E. Fisher Max Planck Institute, Nijmegen

---

---

---

---

---

---

---

---

---

## Foxp2 and neural connectivity

number branches } Process length  
number processes }

■ Normal  
□ R552H

Vernes et al. (2011) *PLoS Genet* 7(7): e1002145

Simon E. Fisher Max Planck Institute, Nijmegen

---

---

---

---

---

---

---

---

---

## Voluntary motor-skill learning

De Bono et al. (2006) *Am J Physiol Regul Integr Comp Physiol* 290: R926-34

Disruption of one Foxp2 copy leads to normal base-line motor behaviour, but significant deficits in motor-skill learning Groszer et al. (2008) *Curr Biol* 18: 354-62

Simon E. Fisher Max Planck Institute, Nijmegen

---

---

---

---

---

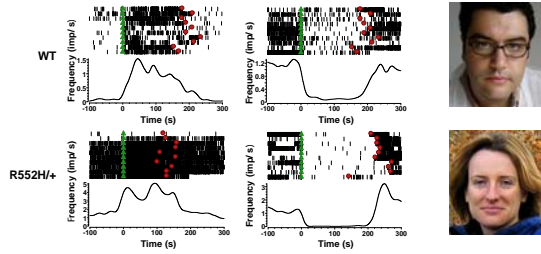
---

---

---

---

## Foxp2 and striatal plasticity



French et al. (2011)  
*Mol Psychiatry* doi: 10.1038/mp.2011.105

Simon E. Fisher

Max Planck Institute, Nijmegen

---

---

---

---

---

---

---

---

---

---

## Not the “gene for language”



- Versions of FOXP2 found in highly similar form in vertebrate species that lack speech and language
- Ancient role(s) in brain of common ancestor?
- FOXP2 likes moonlighting – it is also switched on in parts of developing lung, intestines and heart
- Like other regulatory genes, FOXP2 does several different jobs, interacting with other factors

Simon E. Fisher

Max Planck Institute, Nijmegen

---

---

---

---

---

---

---

---

---

---



---

---

---

---

---

---

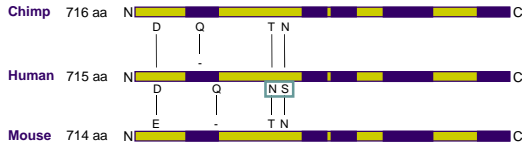
---

---

---

---

# FOXP2 in primate evolution



When it comes to FOXP2 protein, a chimpanzee is more similar to a mouse than it is to a human....



Simon E. Fisher

Max Planck Institute, Nijmegen

---

---

---

---

---

---

---

---

---

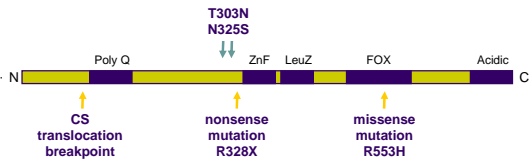
---

---

---

Enard et al. (2002) *Nature* 418: 869-72

### SUBSTITUTIONS ON HUMAN LINEAGE



### MUTATIONS IN DISORDERS

Lai et al. (2001) *Nature* 413: 519-23; MacDermot et al. (2005) *AJHG* 76: 1074-80

---

---

---

---

---

---

---

---

---

---

---

---



Carroll SB (2003) *Nature* 422: 849-857

Simon E. Fisher

Max Planck Institute, Nijmegen

---

---

---

---

---

---

---

---

---

---

---

---

## Molecular evolution of FOXP2



- Accelerated change of FOXP2 on human lineage after split from chimpanzee
- This does not mean that FOXP2 was *the* driving force behind evolution of spoken language
- Modifications of ancient role(s) in neural circuits of the vertebrate brain
- Just one piece of a complex jigsaw
- **Can we empirically test whether evolutionary changes have any functional significance?**

Simon E. Fisher

Max Planck Institute, Nijmegen

---

---

---

---

---

---

---

---



---

---

---

---

---

---

---

---



---

---

---

---

---

---

---

---

Enard et al. (2009) *Cell* **137**: 961-71  
 Reimers-Kipping et al. (2011) *Neuroscience* **175**: 75-84

Konopka et al. (2009) *Nature* **462**: 213-7

**SUBSTITUTIONS ON HUMAN LINEAGE**

**MUTATIONS IN DISORDERS**

Groszer et al. (2008) *Curr Biol* **18**: 354-62  
 Gaub et al. (2010) *Genes Brain Behav* **9**: 390-401  
 French et al. (2011) *Mol Psychiatry* doi: 10.1038/mp.2011.105  
 Kurt et al. (2012) *PLoS One* **7**: e33130

---

---

---

---

---

---


---

---

---

---

### The FOXP2 story



- Missense & nonsense mutations in FOXP2 cause a severe speech & language disorder in humans
- FOXP2 is a regulatory gene – its downstream targets (e.g. CNTNAP2) offer entrypoints into neural pathways involved in speech & language
- Not a “gene for speech” - versions of FOXP2 in many species e.g. affects motor learning in mice
- Helps modulate neural plasticity of brain networks, including cortico-basal ganglia circuitry
- Our unique capacity for complex spoken language is built on evolutionarily ancient systems

---

---

---

---

---

---


---

---



---

---

### The next generation.....



- Initial human genome sequence: 10 yrs, \$3 billion
- By 2010, a human genome could be sequenced in less than a month, for <\$30,000
- Now: Genome in a few days for <\$4,000
- The era of personal genomics

Simon E. Fisher                      Max Planck Institute, Nijmegen

---

---

---

---

---

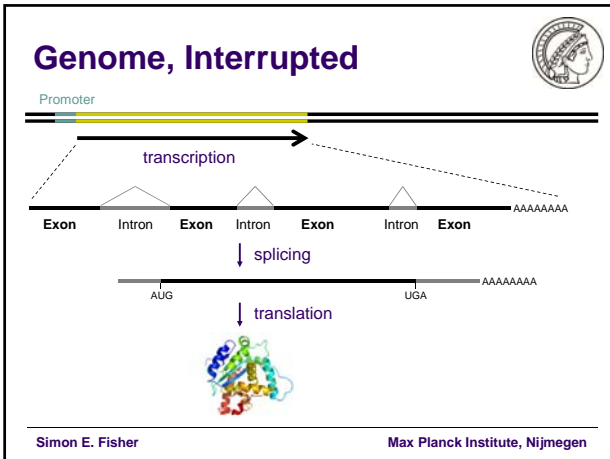
---

---

---

---

---




---

---

---

---

---

---

---

---

### Exome sequencing in autism

- Autism = pervasive impairment in language, communication and social reciprocity, restricted interests or stereotyped behaviours
- Hypothesis: de novo (i.e. new non-inherited) protein-coding mutations may contribute
- Study families where a single child has autism:

Simon E. Fisher  
Max Planck Institute, Nijmegen

---

---

---

---

---

---

---

---

### Exome sequencing in autism

Simon E. Fisher  
Max Planck Institute, Nijmegen

---

---

---

---

---

---

---

---



## FOXP1? Sounds familiar....



- The most similar gene in humans to FOXP2
- Like FOXP2, it encodes a protein which regulates the transcription of other genes
- In some parts of the brain, FOXP1 and FOXP2 are switched on together (co-expressed)
- The proteins can directly interact with each other and may work together to regulate target genes



Simon E. Fisher

Max Planck Institute, Nijmegen

---

---

---

---

---

---

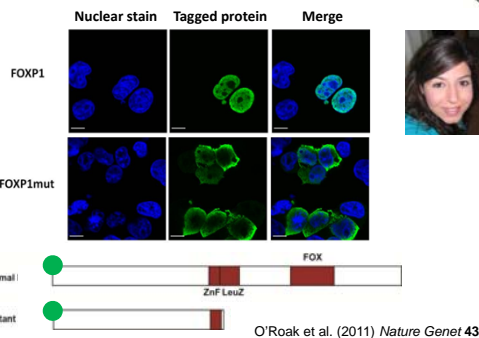
---

---

---

---

## Testing the FOXP1 mutation



Simon E. Fisher

Max Planck Institute, Nijmegen

---

---

---

---

---

---

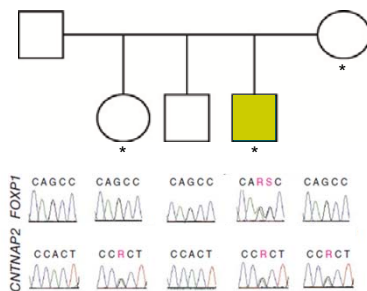
---

---

---

---

## Same child, second mutation...



O'Roak et al. (2011) Nature Genet 43: 585-9

Simon E. Fisher

Max Planck Institute, Nijmegen

---

---

---

---

---

---

---

---

---

---

### FOXP1 regulates CNTNAP2 ?

O'Roak et al. (2011) *Nature Genet* 43: 585-9

Simon E. Fisher Max Planck Institute, Nijmegen

---

---

---

---

---

---

---

---

---

---

### FOXP1-CNTNAP2 dual-hit model

- FOXP1: CAGCC, CAGCC, CAGCC, CAGCC, CAGCC
- CNTNAP2: CCACT, CCRCT, CCACT, CCRCT, CCRCT

- FOXP1 mutation yields elevated CNTNAP2 levels => amplifies deleterious effects of damaged CNTNAP2?

---

---

---

---

---

---

---

---

---

---

### NGS in language impairments

- Next generation sequencing coupled to cell based assays is a powerful tool
- We recently completed exome sequencing in 50 individuals with language disorder
- Knowledge of functional neurogenetic pathways will help to identify the critical variants
- Insights from language disorders promise to shed light on the contribution of genes to normal speech and language processing

Simon E. Fisher Max Planck Institute, Nijmegen

---

---

---

---

---

---

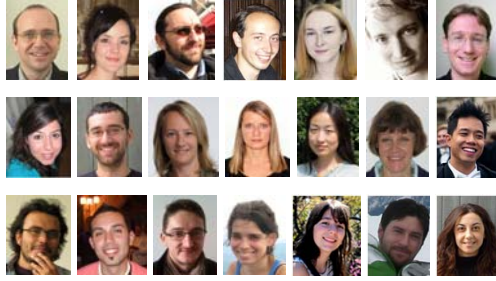
---

---

---

---

## Language and Genetics, MPI



Simon E. Fisher

Max Planck Institute, Nijmegen

---

---

---

---

---

---

---

---

## Special thanks to.....



MPI Nijmegen, Netherlands  
 Sonja Vernes, Pelagia Derizioti  
 Oxford University, UK  
 Tony Monaco, Cecilia Lai  
 Kay MacDermot, Dianne Newbury  
 Jérôme Nicod, Rathi Pulyadi  
 Joe De Bono, David Keays  
 Pete Oliver, Kay Davies  
 Nick Rawlins, Dorothy Bishop  
 Oxford Radcliffe Hospital, UK  
 Jane Hurst  
 ICH, London, UK  
 Faraneh Vargha-Khadem  
 The SLI Consortium, UK

MPI Leipzig, Germany  
 Wolfgang Enard, Svante Pääbo  
 INSERM, Paris, France  
 Matthias Groszer  
 CNP, Lisbon, Portugal  
 Catherine French, Rui Costa  
 UCLA, Los Angeles, USA  
 Brett Abrahams, Liz Spiteri,  
 Dan Geschwind  
 Wash U, Seattle, USA  
 Brian O'Roak, Evan Eichler  
 UWA, Perth, Australia  
 Andrew Whitehouse, Craig Pennell

**Funding:** Royal Society, Wellcome Trust, Autism Speaks,  
 Simons Foundation Autism Research Initiative, Max Planck Society

---

---

---

---

---

---

---

---