

MILAN, Italy

7<sup>th</sup>-8<sup>th</sup> october  
2016



# 1<sup>st</sup> European Conference on GLUT1 Deficiency

## GLUT1 DS: The Genetic Approach

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AUO

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Firenze – Italy



1<sup>st</sup> European Conference on Glut1 Deficiency

Milan, Italy

7<sup>th</sup>-8<sup>th</sup> October 2016

# Outline

- Treatable, highly *epileptogenic*, encephalopathy
- SLC2A1 or GLUT1 gene
- Inheritance
- Genotype
- Phenotype
- Genotype-phenotype correlations



# *SLC2A1 gene*

SOLUTE CARRIER FAMILY 2 (FACILITATED GLUCOSE TRANSPORTER), MEMBER 1; SLC2A1

*Alternative titles; symbols*

GLUCOSE TRANSPORTER 1; GLUT; GLUT1  
ERYTHROCYTE/HEPATOMA GLUCOSE TRANSPORTER

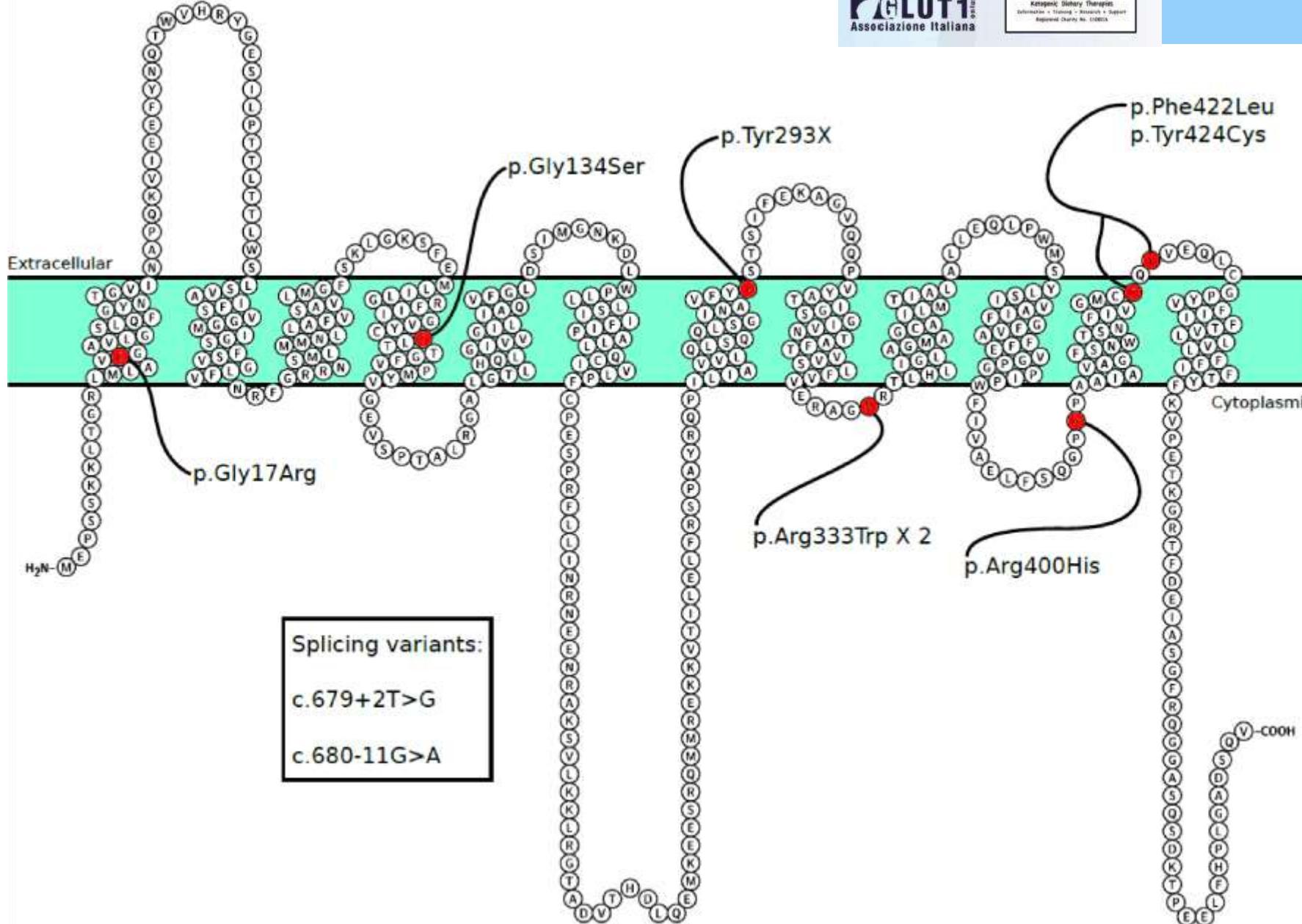
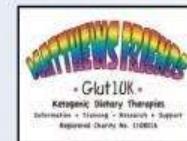
Humans have 2 copies of the gene  
Maternal copy + paternal copy

*HGNC Approved Gene Symbol: SLC2A1*

*Cytogenetic location: 1p34.2      Genomic coordinates (GRCh38): 1:42,925,374-42,959,175* (from NCBI)

## Gene-Phenotype Relationships

Location	Phenotype	Phenotype MIM number	Inheritance <small>(in progress)</small>	Phenotype mapping key
1p34.2	Dystonia 9	601042	AD	3
	GLUT1 deficiency syndrome 1, infantile onset, severe	606777	AD, AR	3
	GLUT1 deficiency syndrome 2, childhood onset	612126	AD	3
	Stomatin-deficient cryohydrocytosis with neurologic defects	608885	AD	3
	{Epilepsy, idiopathic generalized, susceptibility to, 12}	614847	AD	3



Mutations might be:

- Missense
- Frameshift
- Non sense
- Truncating
- deletions



## How should the gene be analyzed?

- Sanger-based automated sequencing → to uncover missense or truncating mutations
- Sequencing must include regulatory sequences of the SLC2A1 gene, e.g. promoter sequences and/or sequences deep in introns
- About 10% of GLUT1-DS might also be caused by (partial) deletion of the SLC2A1 gene.
- Multiplex Ligation-dependent Probe Amplification (MPLA) must be performed in patients with suspected GLUT1 deficit

# Inheritance



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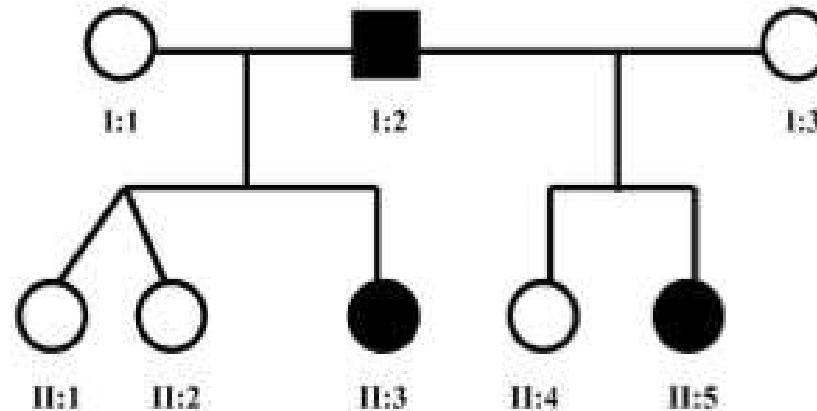
© 2001 Oxford University Press

*Human Molecular Genetics*, 2001, Vol. 10, No. 1 63–68

## Autosomal dominant transmission of GLUT1 deficiency

Jörg Klepper<sup>1,+,§</sup>, Michèl Willemsen<sup>2,+</sup>, Aad Verrrips<sup>2</sup>, Elena Guertsen<sup>1</sup>, Ralf Herrmann<sup>1</sup>, Christian Kutzick<sup>1</sup>, Anne Flörcken<sup>1</sup> and Thomas Voit<sup>1</sup>

- Several affected family members over different generations
- Mutations are transmitted by mildly affected parents
- Mosaic mutations in the transmitting parent
- Non penetrant transmitting parent



## Autosomal dominant Glut-1 deficiency syndrome and familial epilepsy

Brockmann K, Wang D, Korenke CG, von Moers A, Ho YY, Pascual JM, Kuang K, Yang H, Ma L, Kranz-Eble P, Fischbarg J, Hanefeld F, De Vivo DC.

*Ann Neurol* 2001;50(4):476-85.



# Inheritance

Milan, Italy  
7<sup>th</sup>-8<sup>th</sup> October 2016

*Neuropediatrics*. 2009 Oct;40(5):207-10. doi: 10.1055/s-0030-1248264. Epub 2010 Mar 10.

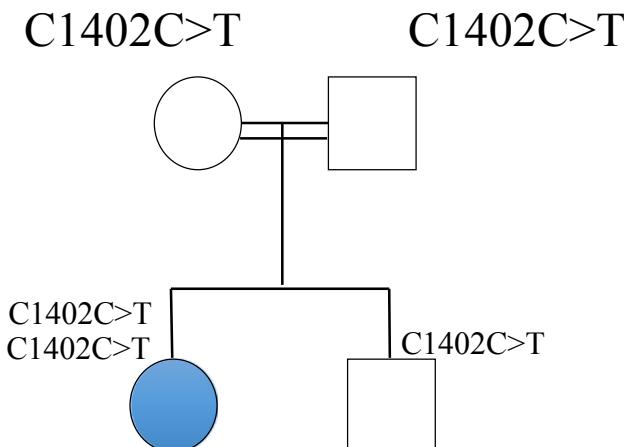
## Autosomal recessive inheritance of GLUT1 deficiency syndrome.

Klepper J<sup>1</sup>, Scheffer H, Elsaïd MF, Kamsteeg EJ, Leferink M, Ben-Omran T.

### Author information

#### Abstract

GLUT1 deficiency syndrome (GLUT1DS) is understood as a monogenetic disease caused by heterozygous SLC2A1 gene mutations with autosomaldominant and sporadic transmission. We report on a six-year-old girl from an inbred Arab family with moderate global developmental delay, epilepsy, ataxia, hypotonia, and hypoglycorrachia (CSF glucose 36 mg/dL; CSF lactate 1.09 mmol/L; CSF/blood glucose ratio 0.44). Molecular analysis of the SLC2A1 gene identified a novel homozygous c1402C>T (p. Arg468Trp) mutation in exon 10 in the index patient and her asymptomatic younger sister. The mutation was absent in 120 control alleles of healthy individuals as well as in 400 alleles of other GLUT1DS patients. Arg468 represents a highly conserved, functionally important amino acid residue in the GLUT1 carboxy-terminus essential for substrate recognition and transport. Both unaffected parents were heterozygous for the mutation. A younger brother and two family members were healthy and carried the GLUT1 wild type. A ketogenic diet effectively controlled seizures in the index patient. We conclude that GLUT1DS can be transmitted as an autosomal recessive disease and provide new insights into genetic counselling for this treatable disorder.



# Inheritance

## Glut1 Deficiency: Inheritance Pattern Determined by Haploinsufficiency

Michael Rotstein, MD<sup>1</sup>, Kristin Engelstad, BS<sup>1</sup>, Hong Yang, MD<sup>1</sup>, Dong Wang, MD<sup>1</sup>, Brynn Levy, M.Sc.(Med), Ph.D<sup>2</sup>, Wendy K. Chung, MD<sup>3</sup>, and Darryl C. De Vivo, MD.<sup>1</sup>

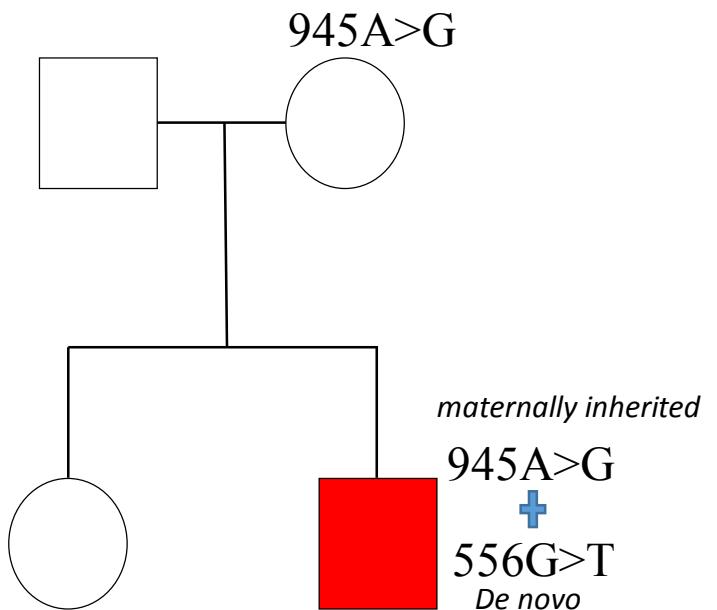
<sup>1</sup> Colleen Giblin Laboratories for Pediatric Neurology Research, Columbia University, New York, NY USA

<sup>2</sup> Clinical Cytogenetics Laboratory, Columbia University, New York, NY USA

<sup>3</sup> Department of Pediatrics, Columbia University, New York, NY USA

### Abstract

Two families manifesting Glut1 deficiency syndrome (Glut1 DS) as an autosomal recessive trait are described. In one family, a severely affected boy inherited a mutated allele from his asymptomatic heterozygous mother. A *de novo* mutation developed in the paternal allele producing compound heterozygosity. In another family, two mildly affected sisters inherited mutations from their asymptomatic heterozygous consanguineous parents. RBC glucose uptake residual activity, a surrogate of haploinsufficiency, correlated with the clinical severity. These cases demonstrate that Glut1 DS may present as an autosomal recessive trait. The clinical pattern of inheritance is determined by the relative pathogenicity of the mutation and the resulting degree of haploinsufficiency.





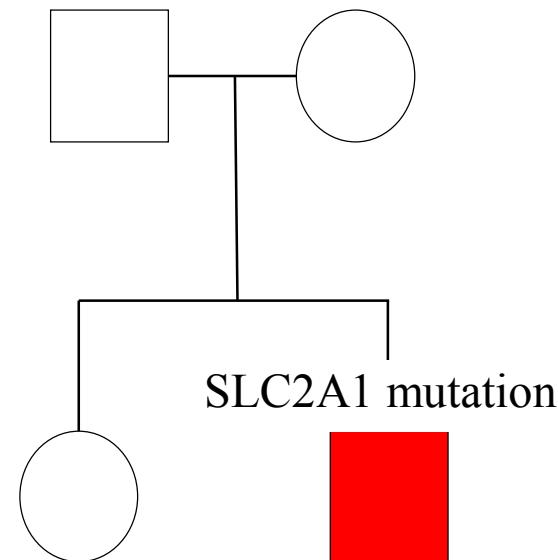
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## Inheritance

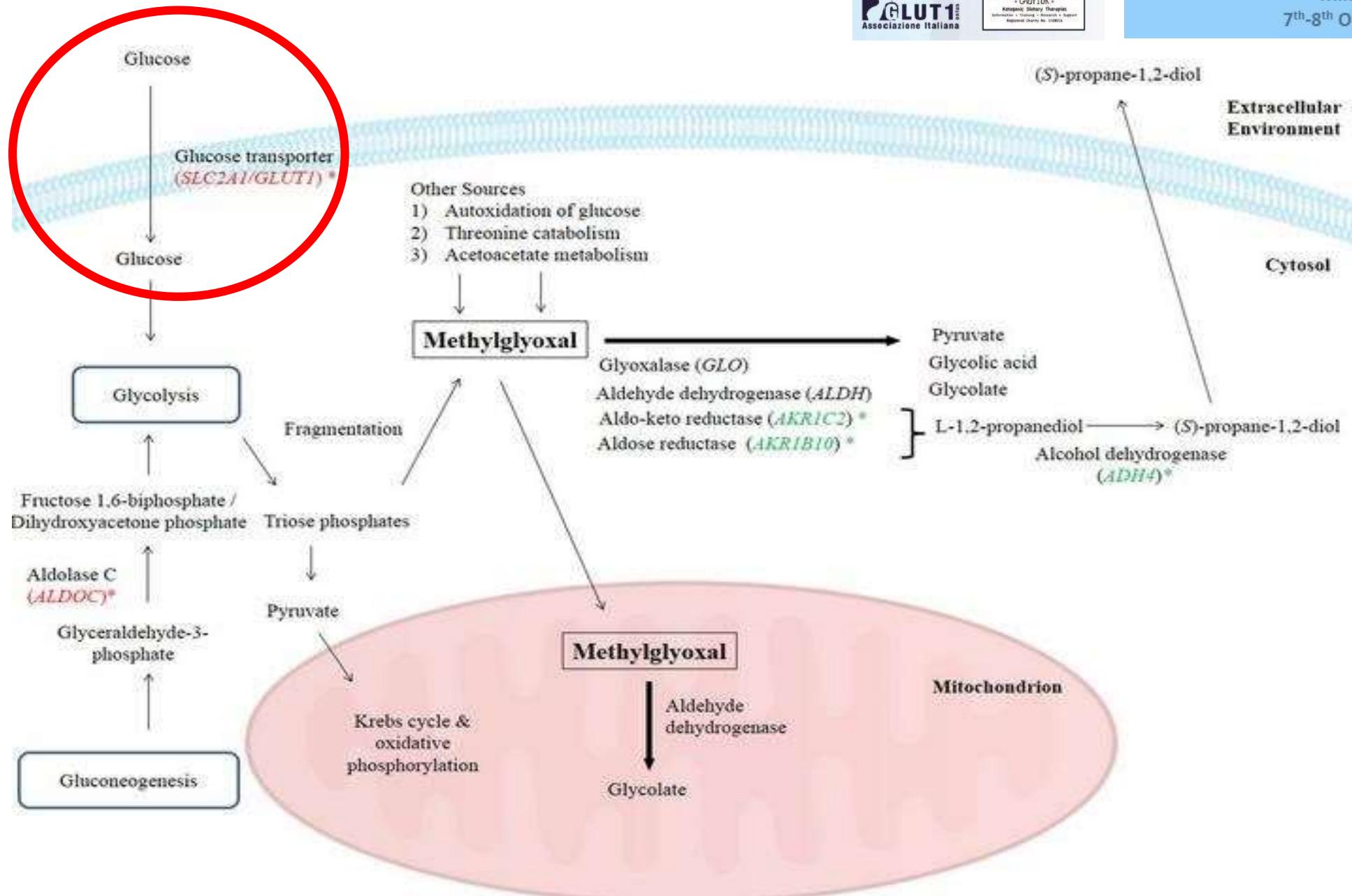
GLUT1-DS often presents as a sporadic disease with ‘de novo’ mutations producing haploinsufficiency and conferring symptomatic heterozygosity



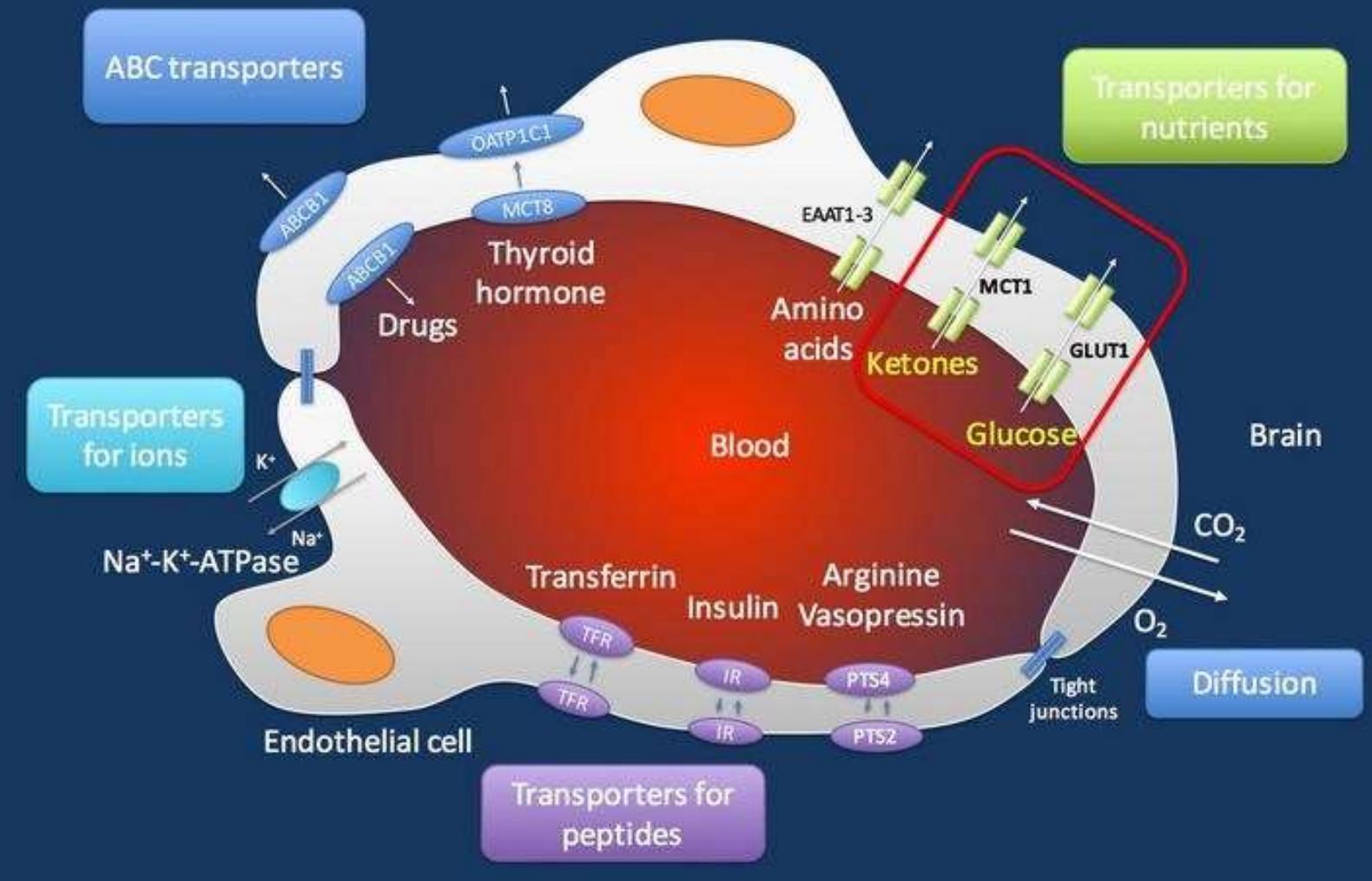


## Genotype

- Several hot spots for recurrent mutations have been identified
  - Asn34
  - Gly91
  - Ser113
  - Arg126
  - Arg153
  - Arg264
  - Thr295
  - Arg333
- All mutations either lead to absence or loss of function of one of the SLC2A1 alleles



## SLC2A1 (Glut1) and the blood-brain-barrier (BBB)





# Rare disorder?

2011: close to 200 patients identified worldwide  
2016: thousand/s patients..

2004 to 2008

132 requests to analyse SLC2A1 gene

54 mutations<sup>↓</sup> (41%) (French study)

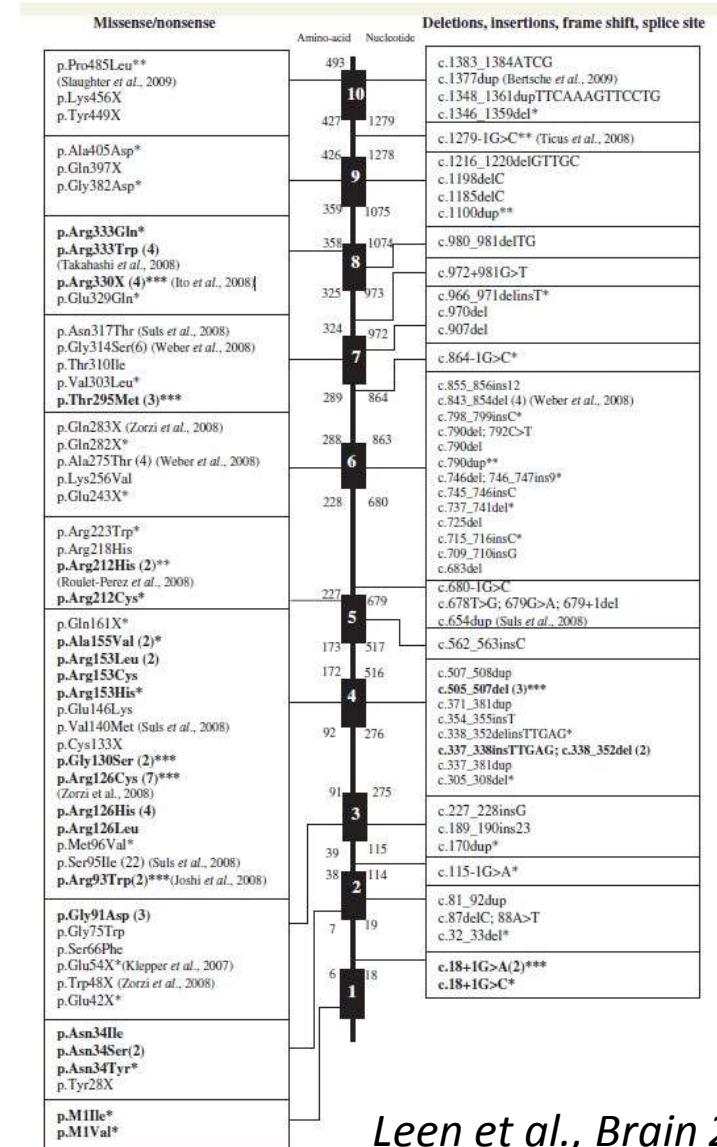
6/207= 2.8% - Danish study

Estimated frequency 1:83,000

Larsen et al., Epilepsia 2015

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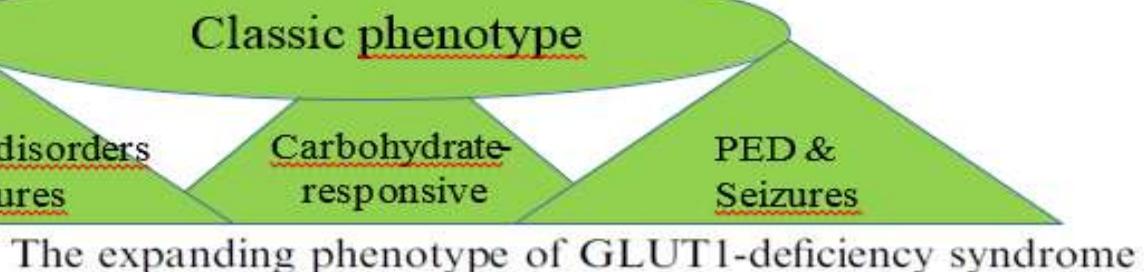
Leen et al., *Brain* 2010

*N Engl J Med*  
1991;325(10):703-9.

**DEFECTIVE GLUCOSE TRANSPORT  
ACROSS THE BLOOD-BRAIN BARRIER AS  
A CAUSE OF PERSISTENT  
HYPOGLYCORRHACHIA, SEIZURES, AND  
DEVELOPMENTAL DELAY**

DARRYL C. DE VIVO, M.D.,  
ROSARIO R. TRIFILETTI, M.D., PH.D.,  
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GABRIEL M. RONEN, M.D.,  
RAMIN A. BEHMAND, B.S.,  
AND SAMI I. HARIK, M.D.

1991



Knut Brockmann Brain & Development 31 (2009) 545–552

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N Engl J Med 1991;325:703–9.

**Table 1. Blood and Cerebrospinal Fluid Values for Glucose and Lactate in Two Children with Defective Glucose Transport across the Blood–Brain Barrier.\***

AGE	BLOOD GLUCOSE	BLOOD LACTATE	CSF GLUCOSE	CSF LACTATE†	CSF GLUCOSE:BLOOD GLUCOSE‡
<i>mo</i> <i>millimoles per liter</i>					
<b>Patient 1</b>					
2.5	5.5	—	1.6	—	0.29
4.5	4.7	—	0.88	0.41	0.19
6.0	4.8	—	1.06	—	0.22
7.5	5.8	1.3	1.89	0.31	0.33
<b>Patient 2</b>					
5.5	—	—	1.4	1.0	—
5.8	6.7	—	1.6	1.3	0.24
6.0	4.6	1.0	1.5	0.9	0.33
9.0	9.4§	2.4	1.8	1.5	0.19
17.5	5.2	1.3	1.8	1.2	0.35
27.8	5.5	1.4	1.7	1.3	0.31



## Classic – De Vivo – GLUT1-DS phenotype

- Infantile onset seizures
- Delayed development
- Deceleration of head growth
- Paroxysmal events of non-epileptic nature
- Hypoglycorrachia and reduced CNS/ blood glucose ratio
- Dramatic response to KD

# SLC2A1 mutations and epilepsy type/syndromes

## Early-Onset Absence Epilepsy Caused by Mutations in the Glucose Transporter GLUT1

Arvid Suls, MSc,<sup>1-3</sup> Saul A. Mullen, MBBS,<sup>4</sup>

Yvonne G. Weber, MD,<sup>5</sup> Kristien Verhaert, MD,<sup>5</sup>

Berten Ceulemans, PhD, MD,<sup>3,6,7</sup> Renzo Guerrini, MD,<sup>8</sup>

Thomas V. Wuttke, MD,<sup>3,9</sup> Alberto Salvo-Vargas,<sup>3,9</sup>

Liesbet Deprez, PhD,<sup>1-3</sup> Lieven R. F. Claes, PhD,<sup>1-3</sup>

Albena Jordanova, PhD,<sup>1-3</sup> Samuel F. Berkovic, MD, FRS,<sup>4</sup>

Holger Lerche, MD,<sup>3,9</sup> Peter De Jonghe, PhD, MD,<sup>1-3,10</sup>

and Ingrid E. Scheffer, PhD, MBBS<sup>4,10</sup>

*Ann Neurol.* 2009 Sep;66(3):415-9

12% (4/34) early-onset absence  
epilepsy of patients

- 2 mutations de novo
- 2 mutations familial



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## Glucose Transporter 1 Deficiency as a Treatable Cause of Myoclonic Astatic Epilepsy

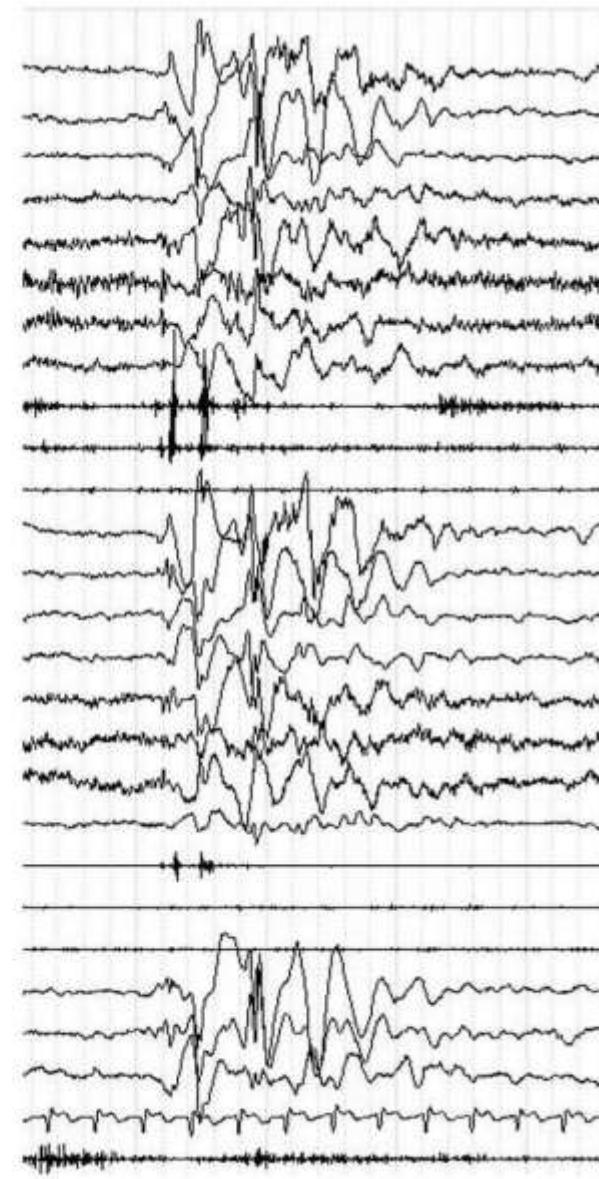
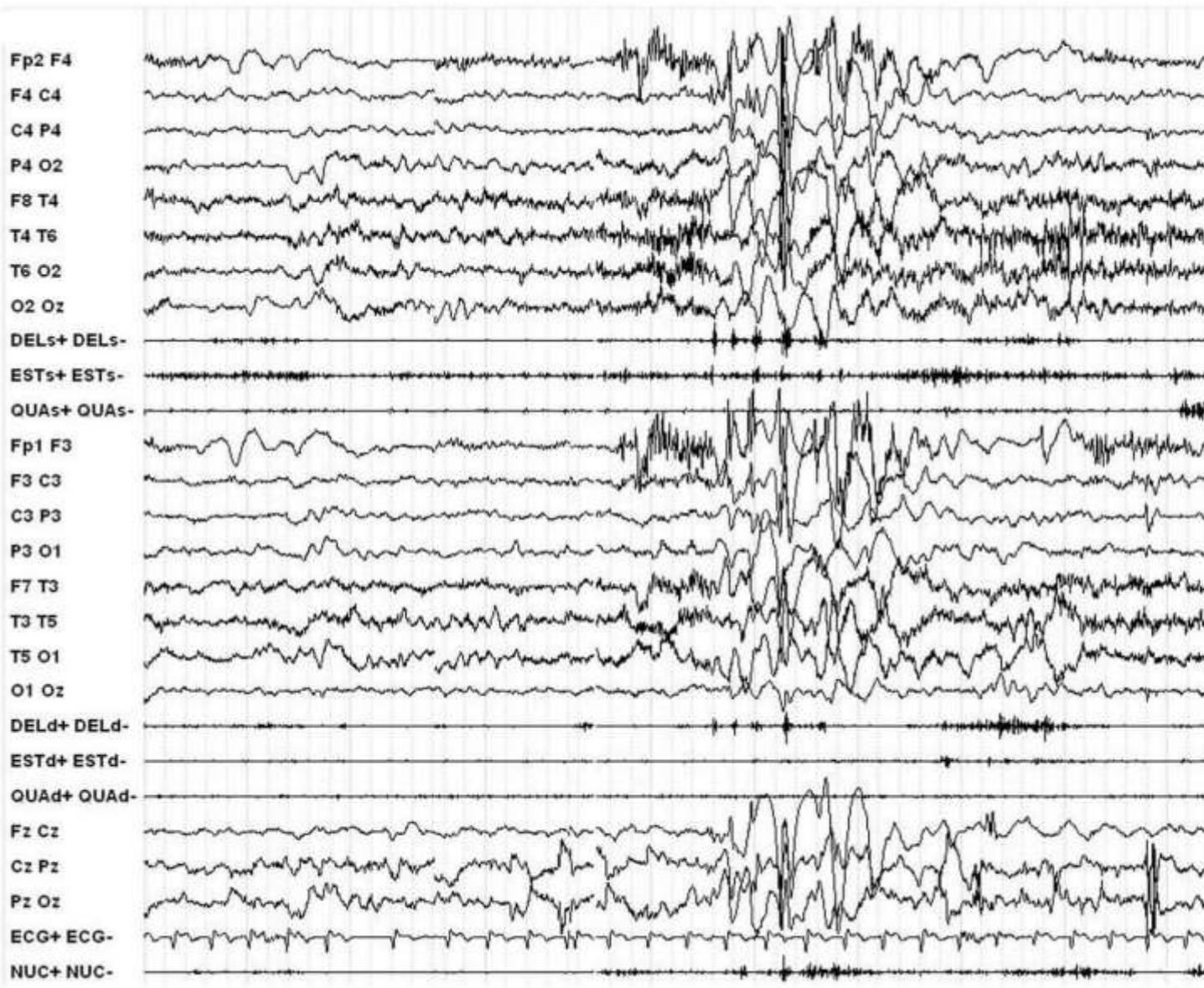
ARCH NEUROL/VOL 68 (NO. 9), SEP 2011

Saul A. Mullen, MBBS; Carla Marini, MD, PhD; Arvid Suls, PhD; Davide Mei, BSc; Elvio Della Giustina, MD; Daniela Buti, MD; Todor Arsov, MD, PhD; John Damiano, BSc; Kate Lawrence, BSc; Peter De Jonghe, MD, PhD; Samuel F. Berkovic, MBBS, MD; Ingrid E. Scheffer, MBBS, PhD; Renzo Guerrini, MD

Table. Clinico-molecular Features of the Patients With MAE and GLUT1 Deficiency

Patient/ Sex/ Age, y <sup>a</sup>	Sz Onset Age/Type	EEG Findings	AEDs/Sz Outcome	KD/Age at Onset	Cognitive Features	Neurological Examination	Movement Disorder Onset Age/Type	CSF Glucose Level/ CSF to Blood Glucose Level Ratio	SLC2A1 Mutation
1/M/15	<36 mo/ My, My-At	GSW, GPSW	VPA + ESM; 5 y sz-free	Yes/ 13.5 y	Normal early development, progressive regression; at 5 y WPSI FSIQ score, 78; 6 y WPSI FSIQ score, 63; 7 y WPSI FSIQ score, 59; 13 y WISC-R FSIQ score <40	Tremor, mild dysarthric speech	6 y/PED > lower limbs (>R); ↑ exercise; facial grimaces; duration 30-180 min; monthly frequency	35 mg/dL/NA	c.997C>T; p.Arg333Trp; parents, NA
2/M/4	8 mo/ My; 12 mo/ My-At	2.5- to 3-Hz GSW	VPA 1 y; sz-free	Yes/ 2.5 y	Mild motor and speech delay at 2 y and 2 mo (Griffiths score at 2 y, 62 and at 8 mo, 66)	Ataxia, dysarthric speech, poor motor performances, deceleration of head growth	None	32 mg/dL/0.42	c.1199G>A; p.Arg400His; de novo
3/M/12	24 mo/ My-At, My	GSW	VPA + TPM; 4 y sz-free	No	Mild motor and speech delay, progressive regression to severe ID	Dysarthric speech, ataxia, poor motor performances	6-7 y/PED > lower limbs (>L); ↑ exercise	37 mg/dL/0.42	c.997C>T; p.Arg333Trp; de novo
4/M/28	48 mo/ At, Ab, GTCS, non-CSE	GSW	VPA + LTG; 4 y sz-free	No	Mild ID	Normal	None	NA	c.971C>T; p.Ser324Leu; maternal





# The role of *SLC2A1* mutations in myoclonic astatic epilepsy and absence epilepsy, and the estimated frequency of GLUT1 deficiency syndrome



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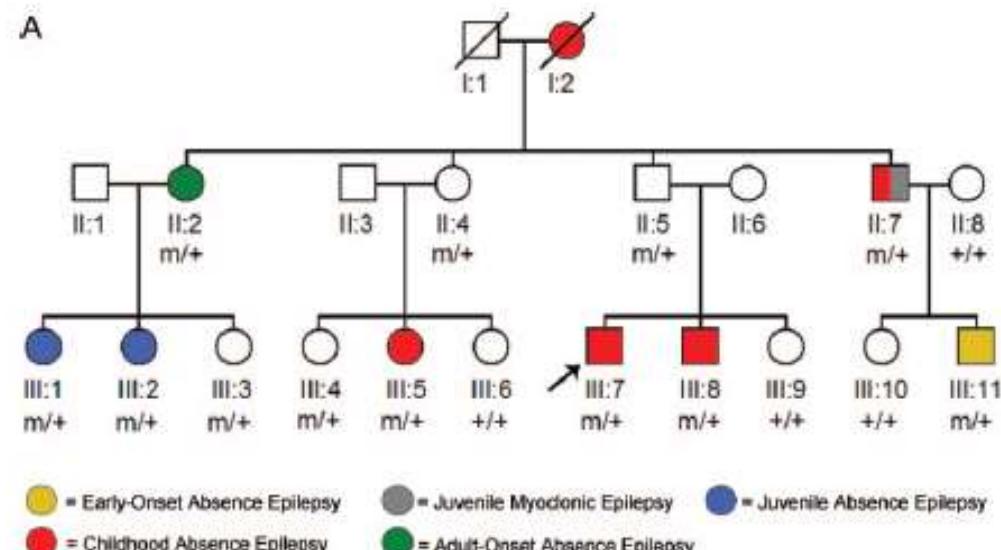
Larsen J, Johannessen KM, Ek J, Tang S, Marini C, Blichfeldt S, Kibaek M, von Spiczak S, Weckhuysen S, Frangu M, Neubauer BA, Uldall P, Striano P, Zara F; MAE working group of EuroEPINOMICS RES Consortium, Kleiss R, Simpson M, Muhle H, Nikanorova M, Jepsen B, Tommerup N, Stephani U, Guerrini R, Duno M, Hjalgrim H, Pal D, Helbig I, Møller RS. Epilepsia. 2015;56(12):e203-8.

Mutations in SCL2A1 were detected in 5 (10%) of 50 patients with early onset absence epilepsy and in 1 (2.7%) of 37 patients with unselected epilepsies, ID and movement disorders

GLUT1 mutations are a rare cause of familial idiopathic generalized epilepsy

Striano P, Weber YG, Toliat MR, Schubert J, Leu C, Chaimana R, Baulac S, Guerrero R, LeGuern E, Lehesjoki AE, Polvi A, Robbiano A, Serratosa JM, Guerrini R, Nürnberg P, Sander T, Zara F, Lerche H, Marini C; EPICURE Consortium.

Neurology. 2012;78(8):557-62.



From splitting GLUT1 deficiency syndromes to overlapping phenotypes

Marie Hully <sup>a,\*</sup>, Sandrine Vuillaumier-Barrot <sup>b</sup>, Christiane Le Bizec <sup>b</sup>, Nathalie Boddaert <sup>c</sup>, Anna Kaminska <sup>d</sup>, Karine Lascelles <sup>e</sup>, Pascale de Lonlay <sup>f</sup>, Claude Cances <sup>g</sup>, Vincent des Portes <sup>h</sup>, Agathe Roubertie <sup>i</sup>, Diane Doummar <sup>j</sup>, Anne LeBihanic <sup>k</sup>, Bertrand Degos <sup>l</sup>, Anne de Saint Martin <sup>m</sup>, Elisabeth Flori <sup>n</sup>, Jean Michel Pedespan <sup>o</sup>, Alice Goldenberg <sup>p</sup>, Catherine Vanhulle <sup>q</sup>, Soumeya Bekri <sup>r</sup>, Anne Roubergue <sup>s</sup>, Bénédicte Heron <sup>j</sup>, Marie-Anne Cournelle <sup>s</sup>, Alice Kuster <sup>t</sup>, Alexis Chenouard <sup>t</sup>, Marie-Noelle Loiseau <sup>u</sup>, Vassili Valayannopoulos <sup>f</sup>, Nicole Chemaly <sup>a</sup>, Cyril Gitiaux <sup>a</sup>, Nathalie Seta <sup>b</sup>, Nadia Bahi-Buisson <sup>a</sup>

**Epilepsy presentation at onset potentially predicts the outcome of patients with GLUT1-DS, since those with myoclonic seizures had a more ID, movement disorders and pyramidal signs than those with EOAE**

53 point mutations and 5 deletions

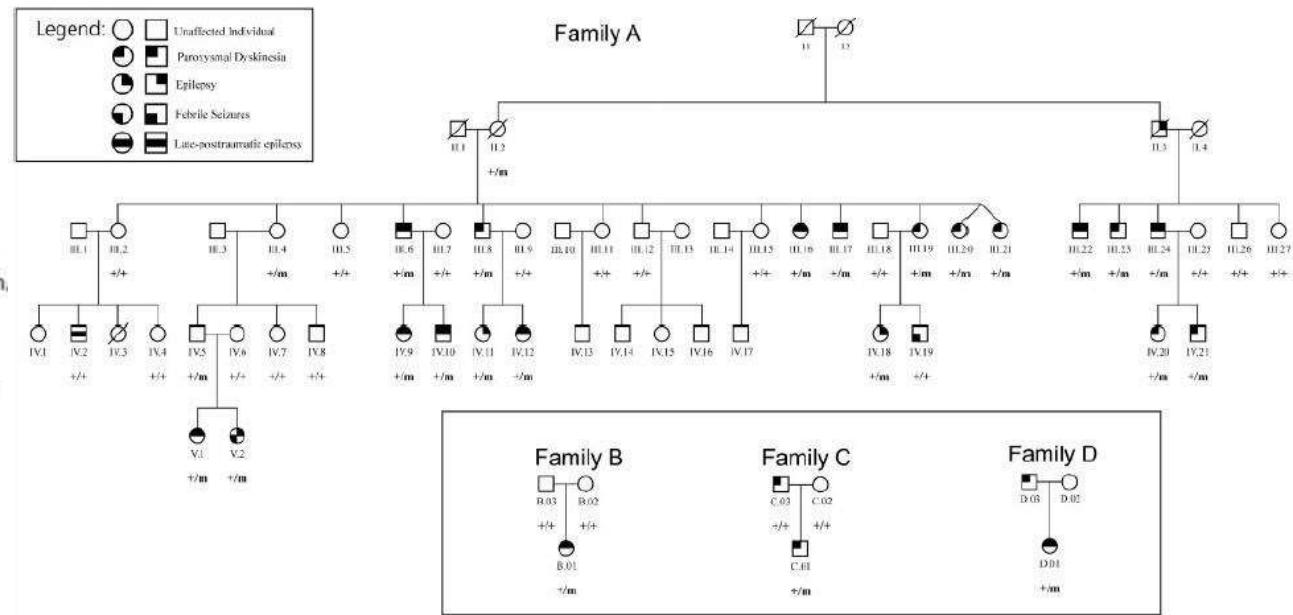
87.5% = classic phenotype with

- intellectual deficiency (41.7%)
- epilepsy (75%)
- movement disorder (29%)
- initial symptoms at a medium age of 7.5 months
- 52%: myoclonic seizures
- 38%: Early Onset Absence Epilepsy
- Three patients evolved from a classic phenotype during early childhood to a movement disorder predominant phenotype at a late childhood/adulthood

# Paroxysmal exercise-induced dyskinesia and epilepsy is due to mutations in *SLC2A1*, encoding the glucose transporter GLUT1

*Brain* (2008), 131, 1831–1844

Arvid Suls,<sup>1,2,3</sup> Peter Dedeken,<sup>4</sup> Karolien Goffin,<sup>5</sup> Hilde Van Esch,<sup>6</sup> Patrick Dupont,<sup>5</sup> David Cassiman,<sup>7</sup> Judith Kempfle,<sup>8,9</sup> Thomas V. Wuttke,<sup>8,9</sup> Yvonne Weber,<sup>8</sup> Holger Lerche,<sup>8,9</sup> Zaid Afawi,<sup>10</sup> Wim Vandenberghe,<sup>4</sup> Amos D. Korczyn,<sup>11</sup> Samuel F. Berkovic,<sup>12</sup> Dana Ekstein,<sup>13</sup> Sara Kivity,<sup>14</sup> Philippe Ryvlin,<sup>15</sup> Lieve R. F. Claes,<sup>12,3</sup> Liesbet Deprez,<sup>12,3</sup> Snezana Maljevic,<sup>8,9</sup> Alberto Vargas,<sup>8,9</sup> Tine Van Dyck,<sup>1,2,3</sup> Dirk Goossens,<sup>3,16</sup> Jurgen Del-Favero,<sup>3,16</sup> Koen Van Laere,<sup>5</sup> Peter De Jonghe<sup>1,2,3,7</sup> and Wim Van Paesschen<sup>4</sup>



With intellectual disability and ataxia

*J Inher Metab Dis.* 2003;26(6):559-63.

## GLUT-1 deficiency without epilepsy--an exceptional case.

Overweg-Plandsoen WC<sup>1</sup>, Groener JE, Wang D, Onkenhout W, Brouwer OF, Bakker HD, De Vivo DC.

*Arch Neurol.* 2009 Nov;66(11):1410-4. doi: 10.1001/archneurol.2009.236.

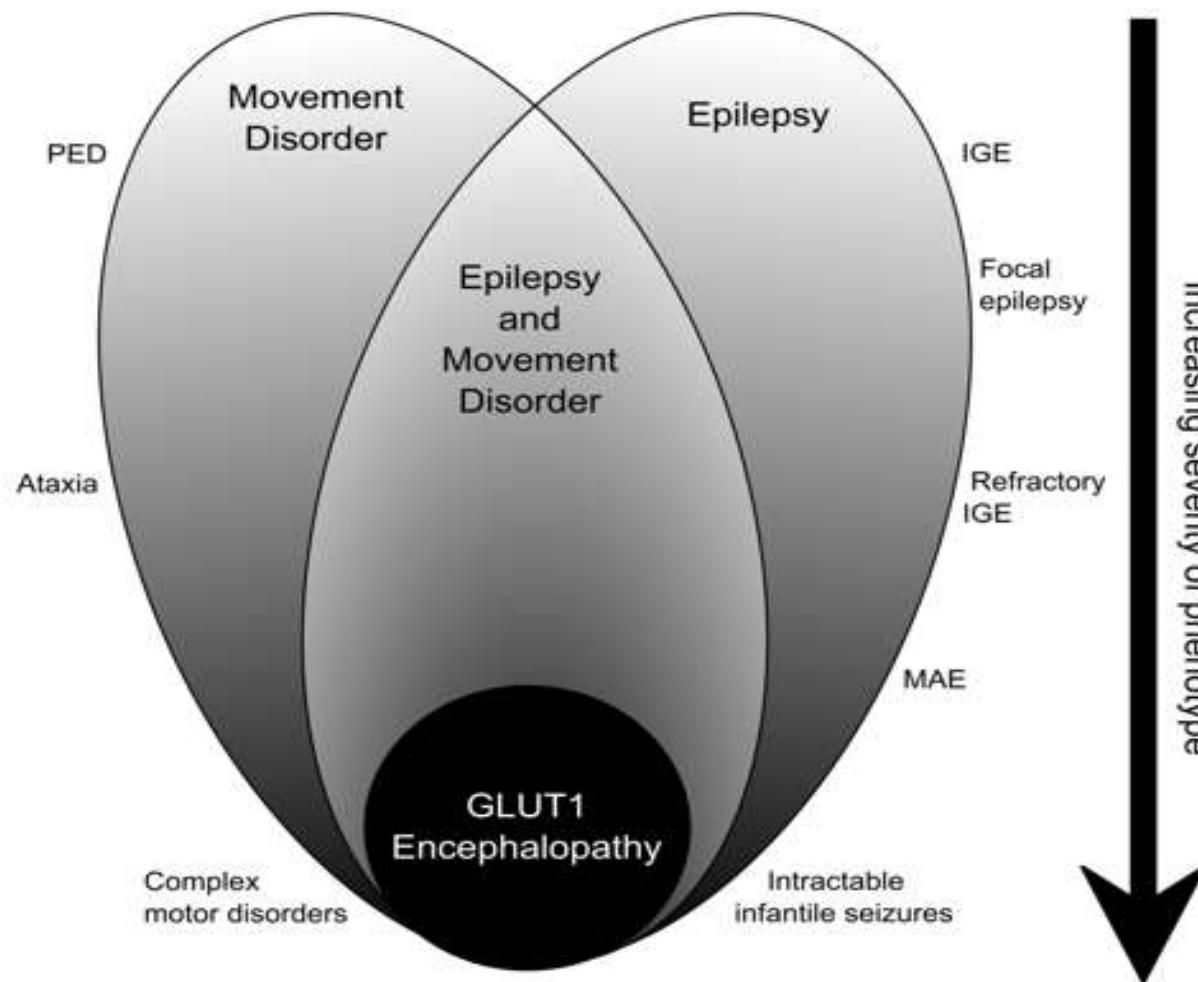
## Childhood chorea with cerebral hypotrophy: a treatable GLUT1 energy failure syndrome.

Pérez-Dueñas B<sup>1</sup>, Prior C, Ma Q, Fernández-Alvarez E, Setoain X, Artuch R, Pascual JM.

*J Inher Metab Dis.* 2011 Apr;34(2):483-8. doi: 10.1007/s10545-010-9264-6. Epub 2011 Jan 13.

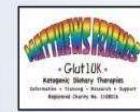
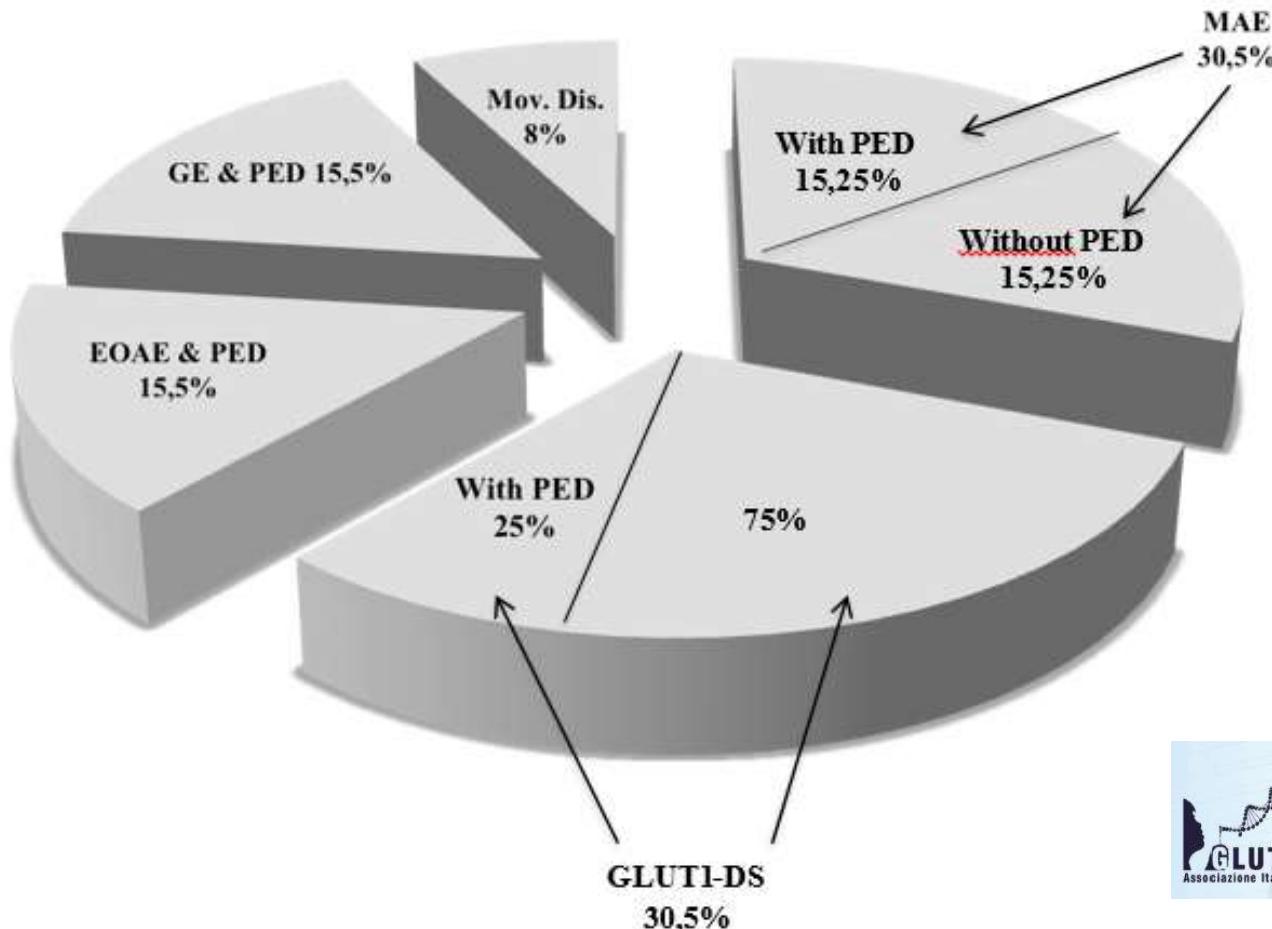
## Dystonic tremor caused by mutation of the glucose transporter gene GLUT1.

Rouberque A<sup>1</sup>, Apartis E, Mesnage V, Doummar D, Trocello JM, Roze E, Taieb G, De Villemeur TB, Vuillaumier-Barrot S, Vidailhet M, Levy R.



Mullen et al 2011

Graphic representation of the spectrum of disorders associated with *SLC2A1* gene mutations in our cohort (13 patients) (*unpublished data*)

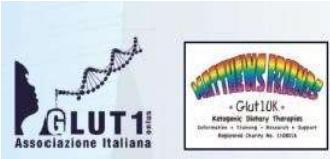


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**Legend**

EOAE: early onset absence epilepsy; GE: generalized epilepsy; GLUT1-DS: GLUT1 deficiency syndrome; Mov. Dis: movement disorder; PED: paroxysmal exercise induced dyskinesia

# Genotype and phenotype correlations



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- Type of mutation with the severity of intellectual disability (ID)

Mild ID in

- 79% of patients with missense mutations
- 26% of patients with nonsense, frame shift, splice site or translation initiation mutation or multiple exon deletion

- Presence or movement disorders

- 63% in patients with missense mutations
- 88% with other type of mutations

- **100% multiple exon deletion have: early onset classical phenotyp**

## Correlations

### genotype

mutations with truncating effect

missense mutations

### CFS/blood glucose ratio

mean 0.30

mean 0.40

### phenotype

early onset GLUT1DS

late onset GLUT1-DS or non classical phenotype

# From our cohort of patients carrying SLC2A1 mutations



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MUTATION	SYNDROME	GLYCORRHACHIA	
MISSENSE	4 MAE	<ul style="list-style-type: none"> <li>mean: 36.5 mg/dl</li> </ul>	<ul style="list-style-type: none"> <li>No Genotype-phenotype correlation</li> <li>Missense mutations: wide range of manifestations from early onset absence epilepsy to GLUT1-DS</li> </ul>
	1 IGE	<ul style="list-style-type: none"> <li>median: 37 mg/dl</li> </ul>	
	1 PED	<ul style="list-style-type: none"> <li>range: 31 mg/dl- 43 mg/dl</li> </ul>	
	1 DE VIVO		
SPLICING	1 EOAE 1 DE VIVO	<ul style="list-style-type: none"> <li>mean: 35 mg/dl</li> <li>median: 35 mg/dl</li> <li>range: 32 mg/dl-38 mg/dl</li> </ul>	<ul style="list-style-type: none"> <li>Mutations with truncating effect: GLUT1-DS</li> <li>2 patients with same missense mutation and identical phenotype: myoclonic astatic epilepsy with childhood onset PED and adolescent onset behavioral disturbance</li> </ul>
TRUNCATING	1 DE VIVO	31 mg/dl	
FRAME SHIFT	1 DE VIVO	29 mg/dl	(unpublished data)



## Conclusion

If GLUT1DS is suspected:

A Lumbar puncture first

B Genetic testing first

- ✓ Individual cases with normal glyccorrachia
- ✓ Relying on hypoglycorrachia alone might be unhelpful for a definite diagnosis in individual cases- GLUT1 deficit might remain undiagnosed and untreated



# Conclusion

- Genetic testing to confirm the clinical diagnosis
  - *SLC2A1* sequencing
  - *SLC2A1* MLPA (to detect deletions)
  - Parental analysis to explore inheritance
  - Genotype-phenotype correlation
- Mutations with truncating functional effect have a more severe phenotype