

6th Symposium on *ATP1A3* in Disease (*ATP1A3* 2017)

September 21-22, 2017. Tokyo, Japan

PROGRAM & ABSTRACTS



by Miya

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Main Host: National Center of Neurology and Psychiatry (NCNP)
Co-Host: Japan AHC Family Association (JAFA)
Support: Japanese Society of Child Neurology

6th Symposium on *ATP1A3* in Disease (*ATP1A3* 2017)

September 21-22, 2017, Tokyo, Japan

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Venue: Palace Hotel Tachikawa

4F Rose room

2-4-15 Akebono-cho, Tachikawa city, Tokyo, Japan 190-0012

Organizer: Masayuki Sasaki, M.D.
Department of Child Neurology,
National Center of Neurology and Psychiatry

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Greetings from the Organizer of *ATP1A3* 2017

Dear Colleagues and Friends,

It is our great pleasure to host the **6th Symposium on *ATP1A3* in Disease** in Tokyo, Japan.



The 1st Symposium on *ATP1A3* in Disease was held in Brussels, Belgium in December 2012, when it was first discovered that alternating hemiplegia of childhood (AHC) was caused by *ATP1A3* mutations. Since then the symposiums have been held in Europe and the US. This is the first time that the symposium is held in Asia.

ATP1A3 encodes the Na⁺/K⁺-ATPase α 3 subunit which is expressed mainly in neurons in the central nervous system. Na⁺/K⁺-ATPase is an essential transmembrane protein, which is ubiquitously expressed in all animal cell membranes. The Na⁺/K⁺-ATPase transports and exchanges 3Na⁺ and 2K⁺ through the plasma membrane using energy from hydrolysis of an ATP. This pump function was discovered 60 years ago (1957), and it is said that 30% of cellular ATPs are used by Na⁺/K⁺-ATPase in living creatures.

Thirteen years ago (2004), the first human disease caused by *ATP1A3* mutations was discovered. It was rapid-onset dystonia–parkinsonism (RDP), which mainly occurs in adulthood. Eight years later (2012), three independent research groups (Heinzen et al, Rosewich et al, and Ishii et al) reported that AHC was caused by *ATP1A3* mutations. AHC mainly occurs in early childhood. Since then several rare conditions other than RDP or AHC have been demonstrated to be caused by *ATP1A3* mutations. However, effective treatments for *ATP1A3*-related diseases have not yet been established. In this symposium, we would like to discuss the characteristics of *ATP1A3*-related diseases, the physiological functions of the Na⁺/K⁺-ATPase, and the possible new treatment methods for *ATP1A3*-related diseases.

We have invited experts on clinical research of *ATP1A3*-related diseases and basic research of the Na⁺/K⁺-ATPase from all over the world. In addition, some Japanese experts on the relevant and very exciting fields will give us great talk. We welcome many attendants who are interested in Na⁺/K⁺-ATPase and *ATP1A3*-related diseases. We hope that all attendants will enjoy this symposium and the pleasant autumn season of Japan.

Masayuki Sasaki, M.D.,
Organizer of the **6th Symposium on *ATP1A3* in Disease**

Organizing committee

Organizer: Masayuki Sasaki, M.D. Department of Child Neurology,
National Center of Neurology and Psychiatry (NCNP)

Organizing Committee:

Takao Takahashi, M.D., Professor, Department of Pediatrics, Keio University

Atsushi Araki, M.D., Director, Department of Pediatrics, Noe Hospital

Hitoshi Osaka, M.D., Professor, Department of Pediatrics, Jichi Medical University

Yoshiaki Saito, M.D. Associate Professor, Department of Child Neurology,
Tottori University

Shin-ichi Hirose, M.D. Professor, Department of Pediatrics, Fukuoka University

Takashi Shiihara M.D. Director, Department of Child Neurology,
Gunma Children's Hospital

Secretariats

Yuko Motohashi, M.D., Department of Child Neurology, NCNP

Eri Takeshita, M.D., Department of Child Neurology, NCNP

Noriko Nakamura, Department of Child Neurology, NCNP

Registration

Registration Fee

The registration fee covers the Symposium brochure (Program and abstracts), the coffee breaks and the lunches on 21 and 22 September 2017, and the social events on 21 September 2017.

| Category | Standard Fee | Reduced Fee * (JPY 6,000 reduction) |
|---|-------------------------|-------------------------------------|
| First early bird April 1 – May 31 | JPY 36,000 | JPY 30,000 |
| Second early bird June 1 – July 21 | JPY 40,000 | JPY 34,000 |
| Standard registration July 22 – Sep 22 | JPY 44,000 | JPY 38,000 |
| Special parents discount | If both parents attend, | 1/2 discount for the Second person |

* For those who register The 15th International Conference on Na,K-ATPase and Related Transport ATPases (Otsu, September 24-30, 2017), or The 23rd World Congress of Neurology (Kyoto, September 16-21, 2017)

On-site Registration

Registration desk will open at

Day 1: Thursday, Sep 21 8:30-18:00

Day 2: Friday, Sep 22 8:30-12:00

Social event

Conference Dinner

Date: Thursday, Sep 21 18:30-20:30

Venue: Meeting room, 4th floor, Palace Hotel Tachikawa

Instructions for presentations

(1) To all oral presenters

We prepare WINDOWS7 PC and one projector.

A single projection screen without sound is available for presentation.

Please use your own PC for the presentation.

Don't forget to the following items; AC Adaptor & Connection cables.

Please ensure that your computer is equipped with the monitor connector of mini D-sub 15 pins.



If your computer does not have this connection, please bring an appropriate converter with you.

Please keep your presentation time.

For example, (20+5) means 20 minutes presentation and 5 minutes discussion time.

Please complete your presentation data in English.

(2) To all poster presenter

All poster presenters are requested to make a registration at this Symposium.

Poster area is prepared in front of the Symposium hall.

Poster discussion time is scheduled twice in the last 30 minutes in both lunch times.

Poster presentation is required to stand by your own poster at the presentation time.

No chairperson will stand by.

Installation: 8:30-9:00 Sep 21

Presentation 13:30-14:00 Sep 21, 12:30-13:00 Sep 22

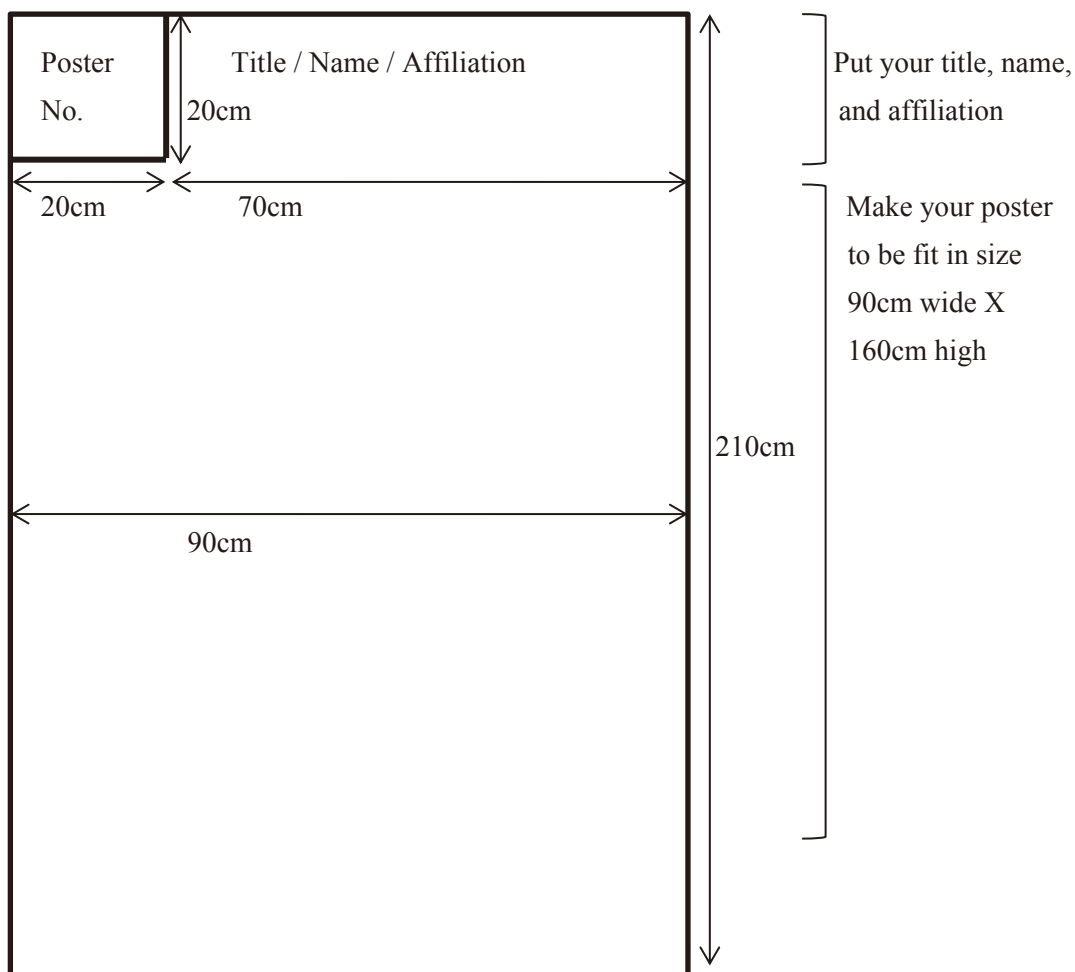
Removal 15:10-15:30 Sep 22 (All posters will be removed)

Poster panel size is 90cm wide X 210cm high.

For poster itself, recommendable size is 90cm X 160cm.

Poster number will be prepared with push pins for your poster in size 20cm X 20cm.

Please prepare a banner to show your Title, Name, and Affiliation in size 70cm X 20cm.



Program (invited speakers and titles)

September 21

8:30- Registration

9:00-9:20 Opening address

Masayuki Sasaki, National Center of Neurology and Psychiatry, Japan
Overview of Clinical Studies on Alternating Hemiplegia of Childhood

9:20-11:00 Session 1. Clinical Aspects and Genotype/Phenotype Correlations in
ATP1A3-related Disease (Chair Mohamad Mikati and Allison Brashear)

1. **Hendrik Rosewich**, Georg August Medical University Göttingen, Germany. (20+5)
Clinical and genetic spectrum of *ATP1A3*-related disorders
2. **Allison Brashear**, Wake Forest University School of Medicine, USA. (20+5)
Rapid-onset dystonia-parkinsonism (RDP): Clinical and Genetic Updates
3. **Yasunari Sakai**, Kyushu University, Japan (10+5)
Relapsing encephalopathy with cerebellar ataxia (RECA)
4. **Naoko Ishihara**, Fujita Health University, Japan (10+5)
Catastrophic early life epilepsy with *ATP1A3* mutation
5. **Kathleen J. Sweadner**, Massachusetts General Hospital, USA (10+5)
Genotype-Structure-Phenotype: what makes *ATP1A3* mutations different?

11:00-11:20 Coffee break

11:20-12:40 Session 2. Clinical Diversity in AHC and other *ATP1A3*-related Disease
(Chair Kathy Swoboda, Hendrik Rosewich)

1. **Eleni Panagiotakaki**, University Hospitals of Lyon, France (20+5)
Clinical profile of patients with *ATP1A3* mutations in AHC in an International Cohort
2. **Kathryn J. Swoboda**, Massachusetts General Hospital, USA (20+5)
Transcriptome analysis and clinical database/biorepository of Alternating Hemiplegia of Childhood patients with *ATP1A3* mutations for potential therapeutic discovery
3. **Simona Balestrini**, UCL Institute of Neurology, UK (20+5)
Cardiac phenotype in *ATP1A3* related-syndromes

12:40-14:00 Lunch

Restaurant (1F, iL PePe, Palace Hotel Tachikawa)

13:30-14:00 Poster presentation (1) 15 Posters for 2 days at the 4th floor

14:00-15:00 Special lecture 1

(Chair Takao Takahashi: Keio University School of Medicine, the President of Japan Pediatric Society and the President of Japanese Society of Child Neurology)

Hideyuki Okano, Keio University School of Medicine, Japan (50+ α)

Modeling Human Diseases with iPS cells and Genetically Modified Non-Human Primates

15:00-16:00 Special lecture 2

(Chair Bente Vilsen)

Chikashi Toyoshima, The University of Tokyo, Japan (50+ α)

Structural biology of Na⁺,K⁺-ATPase: towards understanding of mutations in *ATP1A3*

16:00-16:20 Coffee break

16:20-18:00 Session 3. Basic research topics on *ATP1A3* (Chair Jan Koenderink)

1. **Arn M.J.M. van den Maagdenberg**, Leiden University, The Netherlands (20+5)
New causative gene(s) in AHC?
2. **Jan B. Koenderink**, Radboud University, the Netherland (20+5)
Biochemical consequences of *ATP1A3* mutations
3. **Bente Vilsen**, Aarhus University, Denmark (20+5)
***ATP1A3* neurological disease mutations affecting Na⁺ binding: Structural and functional perspectives and rescue of compromised function continued**

18:30- Conference Dinner (the same floor as the meeting)

Family talk

1. DVD. Human Timebomb
2. Talk from the Japanese AHC Family Association
3. Europe and the US Family

September 22

9:00-10:20 Session 4. Model animal studies in *ATP1A3*-related disease

(Chair Mohamad Mikati)

1. **Steven Clapcote**, University of Leeds, UK (20+5)
Development of new treatments in the *Myshkin* mouse model of AHC
2. **Karin Lykke-Hartmann**, Aarhus University, Denmark (20+5)
Hypothermia-induced dystonia and abnormal cerebellar activity in a mouse model with a single disease-mutation in the sodium-potassium pump
3. **Mohamad A. Mikati**, Duke University, USA (20+5)
Study of Knock-in model mouse; Development of new treatments

10:20-10:40 Coffee break

10:40-11:10 Special lecture 3 (Chair Hitoshi Osaka: Jichi Medical school)

Minako Hoshi, Institute of Biomedical Research and Innovation, Japan (25+5)

Na⁺, K⁺-ATPase α 3 is a Death Target of Alzheimer Amyloid- β Assembly

What Shall We Do Next towards A Better Understanding of Na⁺, K⁺-ATPase α 3's Role in Health and Disease?

11:10-11:40 Special lecture 4 (Chair Hitoshi Osaka)

Noriyuki Matsuda, Tokyo Metropolitan Institute of Medical Science, Japan (25+5)

Mitochondrial autophagy and familial Parkinson's disease

11:40-13:00 Lunch

Restaurant (1F iL PePe, Palace Hotel Tachikawa)

12:30-13:00 Poster Presentation (2) the same posters as the day 1

13:00-14:40 Session 5. Treatment trials (Chair Hendrik Rosewich, Kathy Swoboda)

1. **Atsushi Ishii**, Fukuoka University, Japan (20+5)

Treatment with adenosine- 5'- triphosphate for AHC

2. **Steven Petrou**, University of Melbourne, Australia (20+5)

Loss of function *ATP1A3* mutations differentiated by pre-steady-state analysis

3. **Kevin C. Ess**, Vanderbilt University, USA (20+5)

Modeling Alternating Hemiplegia of Childhood using Patient Derived Stem Cells

4. **Alfred L. George, Jr**, Northwestern University, USA (20+5)

Modeling Alternating Hemiplegia of Childhood in Induced Pluripotent Stem Cells

14:40-15:00 General discussion (Chair Allison, Mohamad, Tsveta)

The function of the Na⁺/K⁺-ATPase α 3 isoform and the new treatment methods

15:00-15:10 Conclusions and future priorities (Hendrik Rosewich)

[Symposium closed]

15:30-17:30

Optional event: A stroll in the Showa Memorial Park (about 2 hour course)

Organizer Speech

Overview of Clinical Studies on Alternating Hemiplegia of Childhood

Masayuki Sasaki, M.D.

Department of Child Neurology, National Center of Neurology and Psychiatry

Alternating hemiplegia of childhood (AHC) was first reported as a complicated migraine beginning in infancy (Verret S and Steel JC, *Pediatrics* 1971). Diagnostic criteria were proposed (Aicardi J, *Dev Med Child Neurol* 1980) and have long been used. Flunarizine was first reported to be effective in one case (Casaer P, *Lancet* 1984); subsequently, several groups have observed that flunarizine is effective in reducing the duration and severity of hemiplegic attacks (Casaer P, *Neuropediatrics* 1987, Silver K, *Neurology* 1993, Bourgeois M, *J Pediatr* 1993, Sasaki M, *Brain Dev* 2001) but not in curing the disease.

Large patient studies have revealed clinical courses and no diagnostic clue in patients with AHC (Sakuragawa N, *Brain Dev* 1993, Mikati MA, *Pediatr Neurol* 2000, Sweney MT, *Pediatrics* 2009, Panagiotakaki E, *Brain* 2010). All studies showed that brain magnetic resonance imaging and electroencephalogram failed to show specific abnormalities in patients with AHC.

The pathophysiology of AHC in terms of mitochondrial function (Arnold DL, *Ann Neurol* 1993, Kemp GJ, *Ann Neurol* 1995), brain perfusion (Aminian A, *Ann Neurol* 1993, Siemes H, *Dev Med Child Neurol* 1993), brain glucose metabolism (de Silva EA, *Ann Neurol* 1996, Sasaki M, *Brain Dev* 2009), and small-vessel studies (Auvin S, *Neurology* 2006, Sasaki M, *Brain Dev* 2011) has been investigated by many groups. However, these studies could not reveal the pathophysiology of AHC.

A point mutation in *ATPIA2*, which encodes the Na⁺/K⁺-ATPase α 2 subunit, was noted in only one autosomal dominant AHC family (Swoboda KJ, *Ann Neurol* 2004) but no follow-up studies have confirmed these results (Boileau S, *Dev Med Child Neurol* 2008).

Calcium and other ion channels have been studied, but only negative results have been reported (Haan J, *Cephalalgia* 2000, de Vries B, *Cephalalgia* 2008, Hirose S, Sasaki M, unpublished data).

In 2012, three study groups independently discovered the causative gene *ATPIA3* (Heinzen EL, *Nat Genet* 2012, Rosewich H, *Lancet Neurol* 2012, Ishii A, *PLoS One* 2013), which encodes the Na⁺/K⁺-ATPase α 3 subunit. Rapid-onset dystonia–parkinsonism (RDP) had been known to be caused by the same gene abnormalities (de Carvalho AP, Brashear A, *Neuron* 2004). Therefore, AHC could be an allelic disorder of RDP. A genotype–phenotype correlation was seen in *ATPIA3*-related disorders (Heinzen EL, *Lancet Neurol* 2014, Rosewich H, *Neurology* 2014).

Before the causative gene of AHC was uncovered, some patients presented with sudden death or severe deterioration (Neville BG, *Dev Med Child Neurol* 2007, Saito Y, *Epilepsy Res* 2010), but the etiology of irreversible deterioration was not known. Our observation on the relation between clinical course and gene mutation type revealed that there could be genotype–phenotype correlations even in AHC (Sasaki M, *Neurology* 2014). Larger observational studies have reported the same tendencies (Viollet L, *PLoS One* 2015, Panagiotakaki E, *Orphanet J Rare Dis* 2015).

New phenotypes have been reported, and the concepts of *ATPIA3*-related neurological disorders are expanding (Demos MK, *Orphanet J Rare Dis* 2014, Dard R, *Dev Med Child Neurol* 2015, Paciorkowski AR, *Epilepsia* 2015).

Precise mechanisms underlying the symptoms and gene abnormalities have been studied using animal models (Clapcote SJ, *Proc Natl Acad Sci USA* 2009, Ikeda K, *J Physiol* 2013, Hunanyan AS, *Epilepsia* 2015) or cell models (Li M, *Neurobiol Dis* 2015) but have not yet been completely elucidated. Curable treatment methods are currently being investigated using new technologies.

Masayuki SASAKI

Present Position

Director, Department of Child Neurology,
National Center of Neurology and Psychiatry, Kodaira, Tokyo, Japan

Education

- 1983 MD, Niigata University School of Medicine, Niigata, Japan
1983-1988 Residency of Pediatrics, Niigata University Hospital
1988-1990 Residency of Child Neurology, Department of Child Neurology, National Center of Neurology and Psychiatry, Kodaira, Tokyo, Japan
1990-1992 Fellow, Division of Inherited Metabolic Disorders, National Institute of Neuroscience, National Center of Neurology and Psychiatry
1992 PhD (Dr. of Medical Science), Toho University
1992-1994 Visiting fellow, Myelin Section, Laboratory of Molecular and Cellular Neurosciences, National Institute of Neurological Disorders and Stroke (NINDS), National Institute of Health (NIH), Bethesda, Maryland, USA



Academic Appointments

- 1994-1996 Department of Child Neurology, National Center of Neurology and Psychiatry
1996-2002 Section Chief, Department of Child Neurology, National Center of Neurology and Psychiatry
2002-Present Director, Department of Child Neurology, National Center of Neurology and Psychiatry

Publications

On alternating hemiplegia of childhood (AHC)

1. Saito Y, Sakuragawa N, **Sasaki M**, Sugai K, Hashimoto T: A case of alternating hemiplegia of childhood with cerebellar atrophy. *Pediatr Neurol* 1998;19:65-68.
2. **Sasaki M**, Sakuragawa N, Osawa M: Long-term effect of flunarizine on patients with alternating hemiplegia of childhood. *Brain Dev* 2001;23:303-305.
3. **Sasaki M**, Sakuma H, Fukushima A, Yamada KI, Ohnishi T, Matsuda H. Abnormal cerebral glucose metabolism in alternating hemiplegia of childhood. *Brain Dev* 2009;31:20-26.
4. Saito Y, Inui T, Sakakibara T, Sugai K, Sakuma H, **Sasaki M**. Evolution of hemiplegic attacks and epileptic seizures in alternating hemiplegia of childhood. *Epilepsy Res* 2010;90:248-258.
5. **Sasaki M**, Matsufuji H, Inui T, Arima K. Absence of small-vessel abnormalities in alternating hemiplegia of childhood. *Brain Dev* 2011;33:390-393.
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7. Ishii A, Saito Y, Mitsui J, Ishiura H, Yoshimura J, Arai H, Yamashita S, Kimura S, Oguni H, Morishima S, Tsuji S, **Sasaki M**, Hirose S. Identification of *ATPIA3* mutations by exome sequencing as the cause of alternating hemiplegia of childhood in Japanese patients. *PLoS One* 2013;8:e56120.
8. **Sasaki M**, Ishii A, Saito Y, Hirose S. Intermediate form between alternating hemiplegia of childhood and rapid-onset dystonia-parkinsonism. *Mov Disord* 2014;29:153-154.
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 11. Rosewich H, Sweney MT, DeBrosse S, Ess K, Ozelius L, Andermann E, Andermann F, Andrasco G, Belgrade A, Brashear A, Ciccodicola S, Egan L, George AL Jr, Lewelt A, Magelby J, Merida M, Newcomb T, Platt V, Poncelin D, Reyna S, **Sasaki M**, Sotero de Menezes M, Sweadner K, Viollet L, Zupanc M, Silver K, Swoboda K. Research conference summary from the 2014 International Task Force on *ATPIA3*-Related Disorders. *Neurol Genet* 2017;3: e139.
 12. **Sasaki M**, Ishii A, Saito Y, Hirose S. Progressive brain atrophy in alternating hemiplegia of childhood. *Mov Disord Clin Prac* 2017 May/June;4(3):406-411.

Select publications other than AHC

1. Kurachi Y, Oka A, Muzuguchi M, Ohkoshi Y, **Sasaki M**, Itoh M, Hayashi M, Goto Y, Takashima S: Rapid immunologic diagnosis of classic late infantile neuronal ceroid-lipofuscinosis. *Neurology* 2000;25:1676-1680.
2. Ohnishi T, Matsuda H, Hashimoto T, Kunihiro T, Nishikawa M, Uema T, **Sasaki M**: Abnormal regional cerebral blood flow in childhood autism. *Brain* 2000;123:1838-1844.
3. Shimojo Y, Osawa Y, Fukumizu M, Hanaoka S, Tanaka H, Ogata F, **Sasaki M**, Sugai K: Severe infantile dentatorubral pallidolusian atrophy with extreme expansion of CAG repeats. *Neurology* 2001;56:277-278.
4. Sudo A, **Sasaki M**, Sugai K, Matsuda H. Therapeutic effect and [123I]IMP SPECT findings of sodium dichloroacetate in a patient with MELAS. *Neurology* 2004;62:338-339.
5. **Sasaki M**, Takanashi J, Tada H, Sakuma H, Furushima W, Sato N. Diffuse cerebral hypomyelination with cerebellar atrophy and hypoplasia of the corpus callosum. *Brain Dev* 2009;31:582-587.
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10. **Sasaki M**, Ohba C, Iai M, Hirabayashi S, Osaka H, Hiraide T, Saitsu H, Matsumoto N. Sporadic infantile-onset spinocerebellar ataxia caused by missense mutations of the inositol 1,4,5-triphosphate receptor type 1 gene. *J Neurol* 2015;262:1278-1284.
11. Kadera H, Ohba C, Kato M, Maeda T, Araki K, Tajima D, Matsuo M, Hino-Fukuyo N, Kohashi K, Ishiyama A, Takeshita S, Motoi H, Kitamura T, Kikuchi A, Tsurusaki Y, Nakashima M, Miyake N, **Sasaki M**, Kure S, Haginoya K, Saitsu H, Matsumoto N. De novo *GABRA1* mutations in Ohtahara and West syndromes. *Epilepsia* 2016;57:566-73.

Special Invited Speakers

Special lecture 1

Modeling Human Diseases with iPS cells and Genetically Modified Non-Human Primates

Hideyuki Okano

Keio University School of Medicine, Japan

For effective modelling of human psychiatric/psychiatric disorders, we took advantage of iPS cell technologies and transgenic non-human primates. So far, we have established iPS cells from the patients of about 40 human psychiatric/psychiatric disorders, including Alzheimer disease^{1,2}, Parkinson disease^{3, 4, 5}, ALS⁶, Rett syndrome^{7,8}, Pelizaeus-Merzbacher disease⁹ (Kuroiwa-Numasawa et al., Stem Cell Reports, 2014), Lissencephaly¹⁰, retinitis pigmentosa¹¹ and Pendred Syndrome¹². Furthermore, for faithfully modeling the human disorders *in vivo*, we developed transgenic non-human primates (common marmosets) with germline transmission¹³. In the present talk, we also wish to mention our recent data of generation of common marmoset transgenic models of neurodegenerative diseases, including Parkinson disease, Alzheimer disease and ALS. Furthermore, we could generate knock-out technologies of common marmoset using genome editing technologies¹⁴, which can be applied for generation of models of autism and psychiatric disorders. At the end, I will mention about Brain Mapping Projects in Japan, in which investigation of common marmoset brains plays key roles^{15, 16, 17}.

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3. Imaizumi Y *et al.*: Mitochondrial dysfunction associated with increased oxidative stress and alpha-synuclein accumulation in PARK2 iPSC-derived neurons and postmortem brain tissue. **Mol Brain.** 5(1): 35, 2012.
4. Ohta E *et al.*: I2020T mutant LRRK2 iPSC-derived neurons in the Sagami-hara family exhibit increased Tau phosphorylation through the AKT/GSK-3 **Hum Mol**

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 10. Bamba Y *et al.*: *In vitro* characterization of neurite extension using induced pluripotent stem cells derived from lissencephaly patients with TUBA1A missense mutations. **Mol Brain** 9(1):70, 2016.
 11. Yoshida T *et al.*: The use of induced pluripotent stem cells to reveal pathogenic gene mutations and explore treatments for retinitis pigmentosa. **Mol Brain**. 7 (1): 45, 2014.
 12. Hosoya M *et al.*: Cochlear cell modeling using disease-specific iPSCs unveils a degenerative phenotype and suggests treatments for congenital progressive hearing loss. **Cell Reports**, 18(1):68-81, 2017.
 13. Sasaki E *et al.*: Generation of transgenic non-human primates with germ line transmission. **Nature**, 459(7246): 523-527, 2009.
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 15. Okano H, Miyawaki A, Kasai K.: Brain/MINDS: Brain-Mapping Project in Japan. **Philos Trans R Soc Lond B Biol Sci**. 370: 20140310, 2015.
 16. Okano H, Yamamori T: How can brain mapping initiatives cooperate to achieve the same goal? **Nat Rev Neurosci** 17, 733-734, 2016.
 17. Okano H, Sasaki E, Yamamori T, Iriki A, Shimogori T, Yamaguchi Y, Kasai K and Miyawaki A.: Brain/MINDS: a Japanese National Brain Project for Marmoset Neuroscience, **Neuron**, 92 (3): 582-590, 2016.

Hideyuki Okano, M.D., Ph.D.

Position:

- Dean, Keio University School of Medicine, Tokyo, Japan.
- Professor, Department of Physiology, Keio University School of Medicine, Tokyo, Japan
- Team Leader, Laboratory for Marmoset Neural Architecture, Brain Science Institute RIKEN

Education:

- 1988 Ph.D. (Dr. of Medical Science), Keio University
- 1983 M.D. Keio University School of Medicine

Employment:

- 2015~Present** **Dean, Keio University School of Medicine**
- 2014~Present: Team Leader, Laboratory for Marmoset Neural Architecture, Brain Science Institute RIKEN
- 2009~Present University of New South Wales Visiting Professor, Australia
- 2008~Present University of Queensland Honorary Professor in the Queensland Brain Institute, Australia
- 2001~Present** **Professor, Department of Physiology, Keio University School of Medicine.**
- 2007~ 2015 Dean, Graduate School of Medicine, Keio University.
- 1997-2001: Professor, Division of Neuroanatomy, Department of Neuroscience, Osaka University Graduate School of Medicine
- 1994-1997: Professor, Department of Molecular Neurobiology, Institute of Basic Medical Sciences, University of Tsukuba.
- 1992-1994: Instructor, Department of Molecular Neurobiology, Institute of Medical Science, University of Tokyo.
- 1989-1993 Postdoctoral Research Fellow, Department of Biological Chemistry, The Johns Hopkins University School of Medicine.
- 1985-1989: Instructor, Institute for Protein Research, Osaka University.
- 1983-1985: Instructor, Department of Physiology, Keio University School of Medicine.



Board of Director

- The International Society for Stem Cell Research (ISSCR)*
- The Japan Neuroscience Society (JNS)* (Vice President)
- The Japanese Society of Inflammation and Regeneration (JSIR)*
- The Japanese Society for Neurochemistry (JSN)*
- The Japanese Society for Regenerative Medicine (JSRM)*

Editor and Editorial Boards

Inflammation and Regeneration (Editor-in-Chief)

Stem Cell Reports (Associate Editor)

eLife (Board of Editor)

Scientific Reports (Editorial Board)

Cell Stem Cell (Editorial Board)

Stem Cells (Editorial Board)

Genes to Cells (Associate Editor)

Journal of Neuroscience Research (Associate Editor)

Neuroscience Research (Associate Editor)

Stem Cells and Development (Editorial Board)

Development, Growth & Differentiation (Editor)

Developmental Neuroscience (Editorial Board (past))

Major Grants:

Project Leader, the Brain Mapping by Integrated Neurotechnologies for Disease Studies (Brain/MINDS) supported by MEXT

Principal Investigator, Program for Intractable Disease Research Utilizing Disease-specific iPS Cells funded by the Japan Science and Technology Agency (JST)

Principal Investigator, Research Center Network for Realization of Regenerative Medicine supported by JST

Awards:

2017 DGD Editor-in-Chief Prize

2016 Faculty Award for Internationalization: Impact Factor Most Outstanding Lab Award (from Keio University)

2016 Molecular Brain Award (from The Association for the Study of Neurons and Diseases (A.N.D.))

2014 Erwin von Bälz Award (from Boehringer Ingelheim GmbH)

2013 Stem Cell Innovator Award (from GeneExpression Systems & Apasani Research Conference USA)

2011 The Johnson & Johnson Innovation Award

2009 A Medal of Honor with Purple Ribbon (from Japanese Emperor)

2008 Inoue Prize for Science (from Inoue Foundation for Science)

2007 Lead Reviewer Award (from *Stem Cells*)

2006 Minister Award of Ministry of Education, Culture, Sports, Science and Technology

2004 Medical Award of The Japan Medical Association (from The Japan Medical Association)

2004 Distinguished Scientist Award (from University of Catania School of Pharmacy)

2004 Gold Medal, Tokyo Techno-Forum 21 Award (from Tokyo Techno Forum 21)

2001 Naka-akira Tsukahara Award (from Brain Science Foundation)

1998 Kitasato Award (from Sanshi-Kai, Keio University School of Medicine)

1995 Yoshihiro Kato Memorial Award (from Yoshihiro Kato Memorial Foundation)

1988 Sanshikai Award (from Sanshi-Kai, Keio University School of Medicine)

Special lecture 2

Structural biology of Na⁺,K⁺-ATPase: towards understanding of mutations in *ATP1A3*

Chikashi Toyoshima

Institute of Molecular and Cellular Biosciences

The University of Tokyo

We have been working on crystal structure studies of two P-type ion transporting ATPases, namely, Ca²⁺-ATPase (SERCA) and Na⁺,K⁺-ATPase. As crystallization of Na⁺,K⁺-ATPase is much more difficult than that of SERCAs, structural and biochemical studies of Na⁺,K⁺-ATPase are far behind those of SERCA. Furthermore, Na⁺,K⁺-ATPase is a much more complex system than SERCA and finely regulated. As locations of regulatory proteins (i.e. FXYD with Na⁺,K⁺-ATPase and phospholamban / sarcolipin with SERCA) in their 3D structures are different, the mode of regulation is likely to be different. Yet similarity between the two ATPases is strong enough to provide us with insight into the structural and functional consequences of mutations identified in *ATP1A3*. In this talk, I will try to explain, in structure terms, the functional consequences of some of the disease-related mutations. It might be of particular interest that several compounds are now found with SERCA that increase the maximum ATPase activity, as it might be possible to find similar compounds that amend impaired Na⁺,K⁺-ATPase activity.

Chikashi TOYOSHIMA

▪ Present Position

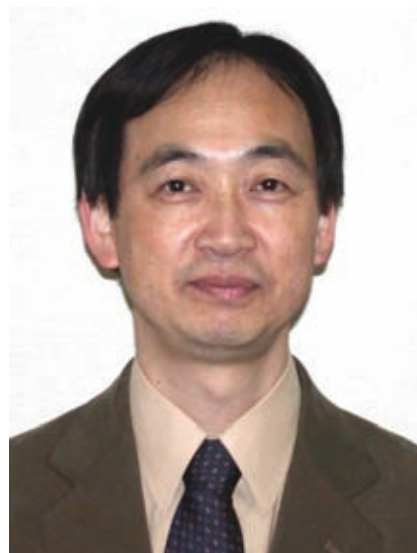
Professor, Institute of Molecular Cellular Biosciences, the University of Tokyo, Tokyo, Japan

▪ Education

- 1978 BSc, Department of Physics, Faculty of Science, the University of Tokyo
1978-1983 DSc, Department of Physics, Division of Science, the University of Tokyo

▪ Academic Appointments

- 1984-1986 Research Associate at the Department of physics, the University of Tokyo,
1986-1988 Postdoctoral fellow, Department of Cell Biology, Stanford University
1988-1989 Scientific staff, Medical Research Council Laboratory of Molecular Biology, Cambridge, UK
1989-1989 Research scientist, Frontier Research Program (RIKEN).
1990-1994 Associate professor, Department of Biological Sciences, Faculty of Bioscience and Biotechnology, Tokyo Institute of Technology
1994-present Professor, Institute of Molecular and Cellular Biosciences, the University of Tokyo
2005-present Foreign Associate of the National Academy of Science, USA.
2008 Hitchcock Professor, University of California, Berkeley, USA
2010-present Director of the Center for Challenging Proteins, Institute of Molecular and Cellular Biosciences, the University of Tokyo



▪ Publication

1. Y. Norimatsu and C. Toyoshima, Protein-lipid interplay revealed with crystals of a calcium pump. *Nature* In press
2. C. Toyoshima, The road to understanding an ion pump, *Physica Scripta*, 91, 042501 (2016)
3. F. Cornelius, M. Habeck, R. Kanai, C. Toyoshima and S.J.D. Karlish: General and specific lipid-protein interactions in Na,K-ATPase. *Biochim. Biophys. Acta.* 1848, 1729-1743 (2015)
4. H. Ogawa, F. Cornelius, A. Hirata and C. Toyoshima: Sequential substitution of K⁺ bound to Na⁺,K⁺-ATPase visualised by X-ray crystallography. *Nature Commun.* 6, 8004 (2015)
5. M. Habeck, H. Haviv, A. Katz, E. Kapri-Pardes, S. Ayciriex, A. Shevchenko, H. Ogawa, C. Toyoshima, and S.J.D. Karlish: Stimulation, inhibition or stabilization of Na,K-ATPase caused by specific lipid interactions at distinct sites. *J. Biol. Chem.* 290 4829-4842 (2015)
6. R. Kanai, H. Ogawa, B. Vilsen, F. Cornelius and C. Toyoshima: Crystal structure of a Na⁺-bound Na⁺,K⁺-ATPase preceding the E1P state. *Nature* 502, 201-206 (2013)

7. C. Toyoshima, S. Iwasawa, H. Ogawa, A. Hirata, J. Tsueda and G. Inesi: Crystal structures of the calcium pump and sarcolipin in the Mg^{2+} -bound E1 state. *Nature* **495**, 260-264 (2013)
8. F. Cornelius, R. Kanai, and C. Toyoshima: A structural view on the functional importance of the sugar moiety and steroid hydroxyls of cardiotonic steroids in binding to Na,K-ATPase. *J. Biol. Chem.* **288**, 6602-6616 (2013)
9. H. Ogawa, T. Shinoda, F. Cornelius and C. Toyoshima: Crystal structure of the sodium-potassium pump (Na^+ , K^+ -ATPase) with bound potassium and ouabain. *Proc. Nat. Acad. Sci. USA* **106**, 13742-13747 (2009)
10. T. Shinoda, H. Ogawa, F. Cornelius and C. Toyoshima: Crystal structure of the sodium-potassium pump at 2.4 Å resolution. *Nature* **459**, 446-450 (2009)
11. C. Toyoshima, Y. Norimatsu, S. Iwasawa, T. Tsuda and H. Ogawa: How processing of aspartylphosphate is coupled to lumenal gating of the ion pathway in the calcium pump. *Proc. Nat. Acad. Sci. USA*. **104**, 19831-19836 (2007)
12. C. Toyoshima, H. Nomura and T. Tsuda: Lumenal gating mechanism revealed in calcium pump crystal structures with phosphate analogues. *Nature* **432**, 361-368 (2004)
13. C. Toyoshima and T. Mizutani: Crystal structure of the calcium pump with a bound ATP analogue. *Nature* **430**, 529-535 (2004)
14. C. Toyoshima and H. Nomura: Structural changes in the calcium pump accompanying the dissociation of calcium. *Nature* **418**, 605-611 (2002)
15. C. Toyoshima, M. Nakasako, H. Nomura and H. Ogawa: Crystal structure of the calcium pump of sarcoplasmic reticulum at 2.6 Å resolution. *Nature* **405**, 647-655 (2000)

Special lecture 3

Na⁺, K⁺-ATPase α 3 Is a Death Target of Alzheimer Amyloid- β Assembly What Shall We Do Next Towards A Better Understanding of Na⁺, K⁺-ATPase α 3's Role In Health And Disease?

Minako Hoshi, Ph.D.

Director, Center for Brain and Neurodegenerative Disease Research, Institute of Biomedical Research and Innovation, Foundation for Biomedical Research and Innovation

Alzheimer's disease (AD) impairs a person's neural network involved in cognitive function by affecting neuronal synaptic connections and degenerating neurons themselves. Amyloid β -protein (A β), a small protein produced by proteolytic cleavages of amyloid precursor protein (APP) in a physiological pathway, has been considered to play a primary role in such loss of synapses and neurons. Recent studies have shown that A β has an ability to form structurally distinct assemblies, which exert different toxic functions through different targets. Our laboratory has long been focusing on understanding mechanisms of neurodegeneration in AD and has identified A β assemblies from AD patient brains, termed amylospheroids (ASPD), as responsible for neurodegeneration (Hoshi et al. PNAS2003, Noguchi et al. JBC2009). Recently, we discovered that the neuron-specific α 3 subunit of the Na⁺, K⁺-ATPase pump (NAK α 3), the catalytic subunit that is essential for neuronal excitability, is a toxic target for ASPD (Ohnishi et al. PNAS2015). This is a new system that involves pre-synaptic calcium hyperactivation, which is triggered by impairing NAK α 3-derived NAK pump activity, leading to neurodegeneration (Figure 1). We also discovered that ASPD-binding tetrapeptides blocked the ASPD:NAK α 3 interaction and protected mature neurons from ASPD neurotoxicity. Interestingly, a French group who have long worked on neurotoxicity of α -synuclein assemblies independently found that NAK α 3 is a toxic target for α -synuclein assemblies and the α -synuclein: NAK α 3 interaction is related to neurodegeneration in Parkinson's disease (Shrivastava et al. EMBO J 34, 2408-2423, 2015). Surprisingly, ASPD and α -synuclein share the essential binding region in the fourth extracellular loop of NAK α 3. At the seminar, I would like to present how we identified first ASPD, and then NAK α 3 as a toxic target of ASPD. I would be happy to discuss about what we shall do as a next to uncover distribution and function of NAK α 3 in health and disease.

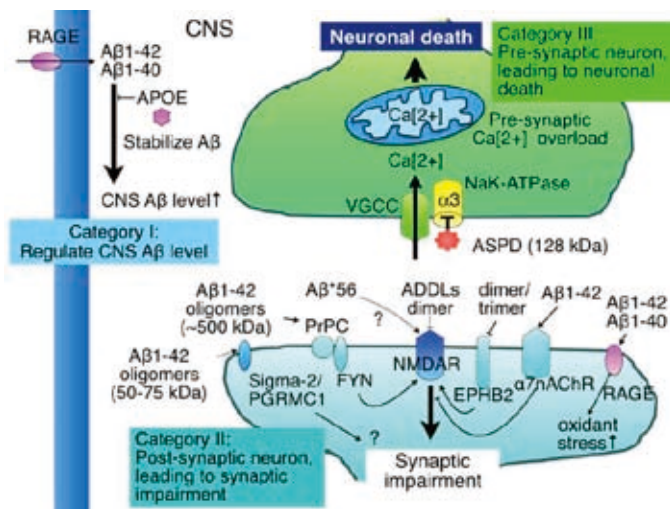


Figure 1. A β receptor/ligand systems.

MINAKO HOSHI

Degrees: Ph. D. (Biochemistry), The University of Tokyo (1991)
M. Sci. (Biochemistry), The University of Tokyo (1988)
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Previous posts: Associate Professor, Department of Anatomy and Developmental Biology,
Graduate School of Medicine, Kyoto University (2009-2017)
Principal Researcher, Mitsubishi Kagaku Institute of Life Science (2001-2009)
Associate Professor, Graduate School of Bioscience and Biotechnology, Tokyo
Institute of Technology (2004-2008)
Principal Investigator, a Proposal-Oriented Research Promotion Program from
Japanese Science and Technology Agency (2000-2003)
Researcher (Project Head), Mitsubishi Kagaku Institute of Life Sciences
(1997-2001)
Associate Scientist, Mitsubishi Kagaku Institute of Life Sciences (1996-1997)
Researcher, Neuropathology, Tokyo Metropolitan Institute of Gerontology
(1996-1997)
Postdoctoral Fellow, Mitsubishi Kagaku Institute of Life Sciences (1994-1996)
National Institute Postdoctoral Fellow from Japanese Science and Technology
Agency, National Institute of Neuroscience, National Center of Neurology
and Psychiatry (1991-1994)
Research Fellow of Japanese Society for the Promotion of Science (1991)

Recent Selected Publications

1. Hoshi M, Sato M, Matsumoto S, Noguchi A, Yasutake K, Yoshida N, Sato K. Spherical aggregates of beta-amyloid (amylospheroid) show high neurotoxicity and activate tau protein kinase I/glycogen synthase kinase-3beta. *Proc Natl Acad Sci USA*. 2003 May 27;100(11):6370-6375.
2. Roychoudhuri R, Yang M, Hoshi MM, Teplow DB. Amyloid beta-protein assembly and Alzheimer disease. *J Biol Chem*. 2009 Feb 20;284(8):4749-4753.
3. Noguchi A, Matsumura S, Dezawa M, Tada M, Yanazawa M, Ito A, Akioka M, Kikuchi S, Sato M, Ideno S, Noda M, Fukunari A, Muramatsu S, Itokazu Y, Sato K, Takahashi H, Teplow DB, Nabeshima Y, Kakita A, Imahori K, Hoshi M. Isolation and characterization of patient-derived, toxic, high mass amyloid beta-protein (A β) assembly from Alzheimer disease brains. *J Biol Chem*. 2009 Nov 20;284(47):32895-32905.
4. Matsumura S, Shinoda K, Yamada M, Yokojima S, Inoue M, Ohnishi T, Shimada T, Kikuchi K, Masui D, Hashimoto S, Sato M, Ito A, Akioka M, Takagi S, Nakamura Y, Nemoto K, Hasegawa Y, Takamoto H, Inoue H, Nakamura S, Nabeshima Y, Teplow DB, Kinjo M, Hoshi M. Two distinct

- amyloid beta-protein (A β) assembly pathways leading to oligomers and fibrils identified by combined fluorescence correlation spectroscopy, morphology, and toxicity analyses. *J Biol Chem*. 2011 Apr 1;286(13):11555-11562.
5. Xiao Y, Ma B, McElheny D, Parthasarathy S, Long F, Hoshi M, Nussinov R, Ishii Y. A β (1-42) fibril structure illuminates self-recognition and replication of amyloid in Alzheimer's disease. *Nat Struct Mol Biol*. 2015 Jun;22(6):499-505.
 6. Parthasarathy S, Inoue M, Xiao Y, Matsumura Y, Nabeshima Y, Hoshi M, Ishii Y. Structural Insight into an Alzheimer's Brain-Derived Spherical Assembly of Amyloid β by Solid-State NMR. *J Am Chem Soc*. 2015 May 27;137(20):6480-6483.
 7. Ohnishi T, Yanazawa M, Sasahara T, Kitamura Y, Hiroaki H, Fukazawa Y, Kii I, Nishiyama T, Kakita A, Takeda H, Takeuchi A, Arai Y, Ito A, Komura H, Hirao H, Satomura K, Inoue M, Muramatsu S, Matsui K, Tada M, Sato M, Saijo E, Shigemitsu Y, Sakai S, Umetsu Y, Goda N, Takino N, Takahashi H, Hagiwara M, Sawasaki T, Iwasaki G, Nakamura Y, Nabeshima Y, Teplow DB, Hoshi M. Na, K-ATPase $\alpha 3$ is a death target of Alzheimer patient amyloid- β assembly. *Proc Natl Acad Sci USA*. 2015 Aug 11;112(32):E4465-4474.

Mitochondrial autophagy and familial Parkinson's disease

Noriyuki Matsuda,

Tokyo Metropolitan Institute of Medical Science

Parkinson's disease (PD) is a common movement disorder characterized by dopaminergic neuronal loss. The majority of PD cases are sporadic, however, the discovery of the genes linked to hereditary forms of PD has provided important insights into the molecular mechanisms. For example, functional analysis of the recessive familial PD-related genes has revealed that the disease is relevant to mitochondrial quality control. This is consistent with the prior idea that several cases of sporadic and chemical-induced PDs have been associated with mitochondrial dysfunction.

We now focus on *PINK1* and *PARKIN*, responsible genes for hereditary recessive PD. *PINK1* and *PARKIN* encode Ser/Thr kinase and ubiquitin ligase (E3), respectively. We revealed that when the mitochondrial membrane potential decreased, PINK1 phosphorylates ubiquitin at Ser65, and the phosphorylated ubiquitin functions as an activator for E3 function of Parkin (Koyano, Nature 2014). Moreover, phosphorylated poly-ubiquitin chain on damaged mitochondria recruits Parkin to damaged mitochondria by functioning as a Parkin receptor (Okatsu, JCB 2015). Consequently, trio of PINK1, Parkin, and phospho-ubiquitin tag outer membrane proteins on depolarized mitochondria with ubiquitin. Damaged mitochondria are then targeted for selectively degradation, because ubiquitin functions as a signal for degradation by the proteasome and autophagy. Impairment of this process predisposes to familial PD. Summary of the latest knowledge for relationship between mitochondrial quality control, autophagy, and Parkinson's disease will be discussed.

Noriyuki MATSUDA

[Present Position]

Project Leader, Ubiquitin Project,
Tokyo Metropolitan Institute of Medical Science, Tokyo, Japan



[Education]

1991 - 1995 Bachelor of Science, University of Tokyo, Japan
1995 - 1997 Master of Science, University of Tokyo, Japan
1997 - 2001 Ph.D. (Biological Sciences), University of Tokyo, Japan

[Career]

2001 - 2002 Special Postdoctoral Researcher, Molecular Membrane Biology Laboratory, RIKEN, Japan.
2002 - 2007 Postdoctoral Fellow, Department of Molecular Oncology, Tokyo Metropolitan Institute of Medical Science (TMIMS), Japan.
2007 - 2008 Senior Scientist, Systems and Structural Biology Center of RIKEN, Japan.
2008 - Present Senior Researcher (2008-2011), Chief Researcher (2011-2013), Associate Director-Researcher (2013 - 2015), and Project Leader (2015 -) of Ubiquitin Project, TMIMS, Japan.

[Selected Publications]

- 1) Yamano, K., Queliconi, B.B., Koyano, F., Saeki, Y., Hirokawa, T., Tanaka, K., and **Matsuda, N.** Site-specific interaction mapping of phosphorylated ubiquitin to uncover Parkin activation. **J. Biol. Chem.**, 290(42), 25199-25211 (2015)
- 2) Okatsu, K., Koyano, F., Kimura, M., Kosako, H., Saeki, Y., Tanaka, K., and **Matsuda, N.** Phosphorylated ubiquitin chain is the genuine Parkin receptor. **Journal of Cell Biology**, 209(1), 111-128 (2015)
- 3) Koyano, F., Okatsu, K., Kosako, H., Tamura, Y., Go, E., Kimura, M., Kimura, Y., Tsuchiya, H., Yoshihara, H., Hirokawa, T., Endo, T., Fon, E-A., Trempe, J-F., Saeki, Y., Tanaka, K., and **Matsuda, N.** Ubiquitin is phosphorylated by PINK1 to activate Parkin. **Nature**, 510(7503), 162-166 (2014)
- 4) Okatsu, K., Oka, T., Iguchi, M., Imamura, K., Kosako, H., Tani, N., Kimura, M., Go, E., Koyano, F., Funayama, M., Shiba-Fukushima, K., Sato, S., Shimizu, H., Fukunaga, Y., Taniguchi, H., Komatsu, M., Hattori, N., Mihara, K., Tanaka, K., and **Matsuda, N.** PINK1 autophosphorylation upon membrane potential dissipation is essential for Parkin recruitment to damaged mitochondria. **Nature Commun.** 3, e1016 (2012)
- 5) **Matsuda, N.**, Sato, S., Shiba, K., Okatsu, K., Saisho, K., Gautier, C., Sou, Y-S., Saiki, S., Kawajiri, S., Sato, F., Kimura, M., Komatsu, M., Hattori, N. and Tanaka, K. PINK1 stabilized by mitochondrial depolarization recruits Parkin to damaged mitochondria and activates latent Parkin for mitophagy. **Journal of Cell Biology**, 189(2), 211-221 (2010)

Invited Speakers

Session 1-1

Clinical and genetic spectrum of *ATP1A3*-related disorders

Hendrik Rosewich, MD, Assistant Professor

Department of Pediatrics and Adolescent Medicine, Division of Pediatric Neurology, University Medical Center Göttingen, Georg August University, Faculty of Medicine, Göttingen, Germany

P-type cation transport proteins establish and maintain electrochemical gradients for Na⁺ and K⁺ across the plasma membrane. Mutations in the $\alpha 3$ catalytic subunit, encoded by *ATP1A3*, almost exclusively expressed in neurons, are associated with at least 3 distinct, yet overlapping, neurologic syndromes: rapid-onset dystonia parkinsonism (RDP); alternating hemiplegia of childhood (AHC); and cerebellar ataxia, areflexia, pes cavus, optic atrophy, and sensorineural hearing loss (CAPOS). From the past descriptions of each entity, distinct phenotypes have emerged, demonstrating that classic AHC, classic RDP, and CAPOS syndrome constitute clinical prototypes in a continuous and still expanding phenotypic spectrum of *ATP1A3* related disorders.

This still expanding phenotypic spectrum is an excellent example how modern sequencing techniques can help to diagnose a condition not primarily thought to be associated with a certain gene (reverse genetics). Moreover, it causes a diagnostic challenge, even for experienced clinicians. Reviews about the currently known broad phenotypic spectrum are needed to provide useful diagnostic guidelines to physicians to select the right patients amongst suspected phenotypes. For this purpose, core clinical features, that in combination make an *ATP1A3* related disorder most likely, have to be extracted from the complex and wide-ranging known phenotypic spectrum.

Mutation updates of all patient mutations reported in the *ATP1A3* gene and clear assignability to a certain phenotype are needed for both, clinicians and basic scientists. Clinicians can discuss with the patients and parents the most likely clinical course, and researchers studying the cellular function of *ATP1A3* can interpret their data on the basis of the cellular and clinical phenotype. The understanding of the pathomechanism in combination with the clinical phenotype is mandatory to establish future therapeutic trials. For those, natural history studies at least for the clinically well-defined entities (AHC/RDP/CAPOS) are an indispensable prerequisite to define study endpoints in pharmacological trials.

Hendrik ROSEWICH

▪ Present Position

Assistant Professor, Department of Pediatric and Pediatric Neurology, University Medical Center Göttingen, Georg August University Göttingen, Germany



▪ Education and Academic Appointments

- 1996 - 2003 Medicine, Heinrich Heine University, Düsseldorf, Germany, 2003 Medical Licensure
- 2002 Predoctoral Fellowship Howard Hughes Medical Institute and Department of Pediatrics, Neurology and Molecular Biology, Johns Hopkins Medical Institutions Baltimore, U.S.A. (Director: Professor David Valle)
- 2003 - 2016 Clinical and Research Fellow in Pediatrics, Department of Pediatrics and Pediatric Neurology; University Medicine Göttingen, Germany (Director: Professor Dr. Jutta Gärtner)
- Since 2009 Clinical Board Certification for Pediatrics
- 2014 Habilitation and Venia legendi for Pediatrics, Georg August University, Göttingen, Germany
- Since 2016 Clinical Board Certification for Pediatric Neurology
- Since 2014 Assistant Professor, Senior Physician, Department of Pediatric and Pediatric Neurology, University Medical Center Göttingen, Georg August University Göttingen, Germany

▪ Publication

Rosewich H, Sweney MT, DeBrosse S, Ess K, Ozelius L, Andermann E, Andermann F, Andrasco G, Belgrade A, Brashear A, Ciccodicola S, Egan L, George AL Jr, Lewelt A, Magelby J, Merida M, Newcomb T, Platt V, Poncelin D, Reyna S, Sasaki M, Sotero de Menezes M, Sweadner K, Viollet L, Zupanc M, Silver K, Swoboda K. Research conference summary from the 2014 International Task Force on ATP1A3-Related Disorders. *Neurol Genet.* 2017 Mar 2;3(2):e139.

Rosewich H, Baethmann M, Ohlenbusch A, Gärtner J, Brockmann K.: A novel ATP1A3 mutation with unique clinical presentation. *J Neurol Sci.* 2014; 341(1-2):133-5.

Rosewich H, Ohlenbusch A, Huppke P, Schlotawa L, Baethmann M, Carrilho I, Fiori S, Lourenço CM, Sawyer S, Steinfeld R, Gärtner J, Brockmann K.: The expanding clinical and genetic spectrum of ATP1A3-related disorders. *Neurology.* 2014; 82(11):945-55.

Heinzen EL, Arzimanoglou A, Brashear A, Clapcote SJ, Gurrieri F, Goldstein DB, Jóhannesson SH, Mikati MA, Neville B, Nicole S, Ozelius LJ, Poulsen H, Schyns T, Sweadner KJ, van den Maagdenberg A, Vilsen B; ATP1A3 Working Group (**Rosewich H**): Distinct neurological disorders with ATP1A3 mutations. *Lancet Neurol.* 2014; 13(5):503-14.

Rosewich H, Thiele H, Ohlenbusch A, Maschke U, Altmüller J, Frommolt P, Zirn B, Ebinger F, Siemes H, Nürnberg P, Brockmann K, Gärtner J.: Heterozygous de-novo mutations in ATP1A3 in patients with alternating hemiplegia of childhood: a whole-exome sequencing gene-identification study. *Lancet Neurol.* 2012; 11(9):764-73.

Session 1-2

Rapid-onset dystonia-parkinsonism (RDP): Clinical and Genetic Updates

Allison Brashear, MD, MBA

Professor & Chair, Department of Neurology Walter C. Teagle Endowed Chair, Wake Forest School of Medicine, Winston Salem, NC USA

Christopher T. Whitlow, MD, PhD, MHA

Associate Professor of Radiology Division Head, Neuroradiology, Wake Forest School of Medicine Winston Salem, NC USA

Since 1993, with the original description of RDP, ATP1A3-related disorders have broadened to include not only RDP, but CAPOS, ataxia, and AHC. Multiple lines of evidence, including imaging, clinical examinations and pathology suggest that ATP1A3 mutations contribute to a complex cascade of signs and symptoms indicating wide spread impact on the brain. Imaging and human pathology suggest that ATP1A3 mutations lead to impairment of the cerebello-thalamo-cortical (CbTC) pathways. Our findings are supported by human brain pathology in patients who were symptomatic over a life time, demonstrating cell death in areas including the globus pallidus, subthalamic nucleus, red nucleus, inferior olivary nucleus, cerebellar purkinje and granule cell layers and dentate. When and how cell death occurs, how this may change over time and whether a reproducible biomarker can be followed for treatment or prevention is part of our current body of work.

In our preliminary investigation, data were analyzed from thirteen participants with ATP1A3 mutations (age range 14-65; mean \pm SD = 38 \pm 14) who were enrolled in a larger ongoing Institutional Review Board approved study of brain structure/function at Wake Forest School of Medicine. All subjects were diagnosed using a standard battery of movement disorder assessments, including Burke-Fahn-Marsden Dystonia Rating Scale (BFMDRS), Unified Parkinson's Disease Rating Scale motor subscore (UPDRS), and International Cooperative Ataxia Rating Scale (ICARS). Brain MRI, including structural anatomic and arterial spin labeling cerebral blood flow images were acquired using standard techniques. Our results suggest that there were more associations between measures of dystonia and cerebral blood flow in components of the CbTC compared to gray matter volume, suggesting that changes in brain function may play an important role in phenotypes associated with *ATPIA3* gene mutations. Whether MRI measures of brain structure and function can be used as biomarkers to track progression and/ or treatment over time is a larger goal of the study.

Overall heightened awareness of the broad spectrum of *ATPIA3* diseases and an understanding of how the pump may be altered by specific mutations opens the door for potential prevention and treatment. The overlap of clinical, imaging, genetic and bench work in studying *ATPIA3* disease is a unique opportunity to develop a new line of therapies for neurologic diseases.

Supported by NINDS grant 5R01NS058949-08 (AB)

Allison Brashear, MD, MBA

Allison Brashear, MD, MBA is Professor and Chair of the Department of Neurology at Wake Forest School of Medicine in Winston-Salem, NC (2005-present). She holds the Walter C. Teagle Chair of Neurology. Dr. Brashear also served as Interim Chair of the Department of Psychiatry.

Dr. Brashear is the principal clinician to describe a unique genetic form of dystonia-parkinsonism, Rapid-Onset Dystonia-Parkinsonism (RDP). Her group reported the genetic mechanism responsible for RDP (mutations in the Na/K ATPase alpha 3 subunit) in *Neuron* in July 2004. In 2008, NINDS funded her group to study these mutations in the ATP1A3 gene in more depth (R01NS058949; PI: Brashear). This work was refunded in April, 2015 for five years. The results are published in *Lancet Neurology*, *Brain*, *Neuron*, *Acta Neuropathologica*, *Neurology*, *Annals of Neurology* and *Movement Disorders*. In 2016 Dr. Brashear was appointed a permanent member of the NINDS NST-1 (K awards) study section.

Dr. Brashear is the lead principal investigator in many multi-center trials for the treatment of cervical dystonia and spasticity. She is the lead author on the pivotal paper, “Intramuscular Injection of Botulinum Toxin for the Treatment of Wrist and Finger Spasticity after a Stroke”, published in *The New England Journal of Medicine*. Her work with botulinum toxin has been published in *Lancet Neurology*, *Neurology*, *Movement Disorders* and *Archives of PMR*.

Dr. Brashear served on the American Academy of Neurology (AAN) Board of Directors from 2013-15. In 2015 she completed six year terms on the AAN Education Committee and the Medical Economic Management Committee. She served on the Board of Directors of the American Neurological Association (ANA), United Council for Neurological Subspecialties (UCNS, vice chair) and currently serves of council of the American Board of University Professors of Neurology (AUPN). In August, 2014 Dr. Brashear was appointed to a renewable four year term as a Director of the American Board of Psychiatry and Neurology (ABPN).

In 2008 the Wake Forest School of Medicine and North Carolina Baptist Hospital integrated to form the Wake Forest Baptist Medical Center. Dr. Brashear served as the first faculty member appointed to the Wake Forest Baptist Medical Center Board of Directors (2007-2013). She also served on the following WFBMC Board of Directors committees: finance, audit, clinical and academic affairs committees. She also serves as one of two faculty members of the WFBMC Capital Campaign Cabinet.

Dr. Brashear completed the Harvard School of Public Health Leadership program for physicians in 2004, and in 2007 she finished the year-long national program for women leaders, Executive Leadership Academic Medicine (ELAM). In 2014 she was selected as an AAMC Council of Dean’s Fellow. Dr. Brashear earned an MBA from Fuqua School of Business at Duke University in 2012, with additional certification in Health Sector Management.

Dr. Brashear graduated from DePauw University in 1983, Indiana University School of Medicine in 1987 and from Fuqua Business School at Duke University in 2012. She completed her training in Neurology at Indiana University School of Medicine in 1991.

Dr. Brashear and her husband, Clifford Ong, have two children, Richard (20) and Diane (17).



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ACADEMIC APPOINTMENTS

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Session 1-3

Relapsing encephalopathy with cerebellar ataxia (RECA)

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Abstract

Background: Alternating hemiplegia of childhood (AHC) is a rare neurological disorder characterized by recurrent hemiplegia, oculogyric and choreoathetotic involuntary movements. *De novo* mutations in *ATPIA3* cause AHC and its associated syndromes. It remains to be determined, however, whether rare variations in *ATPIA3* may lead to atypical neurological phenotypes.

Case: A 7-year-old Japanese boy started presenting with recurrent symptoms of generalized paralysis at 17 months of age. Hypotonia, rapidly bouncing or rotating ocular movements, dystonia and choreoathetosis persisted for more than a month after the onset. These symptoms recurred on febrile illness. Serial recordings of electroencephalogram and neuroimaging studies did not support the diagnosis of acute encephalopathy.

Results: Whole-exome sequencing identified a *de novo* mutation in *ATPIA3* (NM_152296.4: exon17: c.2266C>T: p.R756C). Previously reported cases carrying p.R756C or p.R756H mutations showed both overlapping and distinct phenotypes compared to those of our case. These symptoms were recently designated as relapsing encephalopathy with cerebellar ataxia (RECA). Molecular studies uncovering the pathogenic mechanisms of this disease are currently under progress.

Conclusion: This study confirmed that p.R756C mutation of *ATPIA3* caused atypical forms of AHC-associated disorders. The wide spectra of neurological phenotypes in AHC are linked to still unknown deficits in the molecular functions of *ATPIA3*.

Keywords:

Alternating hemiplegia of childhood (AHC), Whole-exome sequencing, relapsing encephalopathy with cerebellar ataxia (RECA), and *ATPIA3*

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• Publication

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Catastrophic early life epilepsy with *ATP1A3* mutation

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【Introduction】 Mutations of *ATP1A3* have been associated with various phenotypes such as Rapid Onset Dystonia-Parkinsonism (RDP), Alternating Hemiplegia of Childhood (AHC), and Cerebellar ataxia with Areflexia, Pes cavus, Optic atrophy, and Sensorineural hearing loss (CAPOS). Recently, two cases of catastrophic early life epilepsy with heterozygous mutations in *ATP1A3* were reported (Paciorkowski et al. *Epilepsia*. 2015), which indicates their mutations cause most severe phenotypes of *ATP1A3*-related disorder spectrum. However, gene functions and genotype-phenotype correlations were remained unclear. Here we report one child with catastrophic early life epilepsy, respiratory failure, postnatal microcephaly, and severe developmental disability, with a novel heterozygous mutation of *ATP1A3*.

【Subject】 A 2-year-old boy who was born to nonconsanguineous parents with normal delivery, and transferred to NICU because of respiratory failure at second day of birth. He also showed extreme hypotonia, episodic oculomotor abnormality and tachycardia, and frequent epileptic seizures. Mechanical ventilation was required to support his respiration. Epileptic seizures were intractable with treatment of multiple antiepileptic drugs, including extremely high dose Phenobarbital. In spite of many examinations including chromosomal analysis and muscle biopsy, etiology of his symptoms remained unknown.

【Methods and Results】 Whole exome sequencing analysis was performed for subject and his parents. First we filtered the data with the known gene list for severe brain abnormality but no candidate mutation was found. Then we filtered for de novo mutation and found one heterozygous mutation in *ATP1A3* (c.2736_2738CTTdel, p.Phe913del).

【Discussion】 Phe913 is a residue which has been highly conserved among vertebrates. There is a report that p.Val919del located in a same transmembrane domain reduces the activity of ATP1 α 3, suggesting that the current mutation also impairs the function. Two previously reported cases of catastrophic early life epilepsy had heterozygous mutations localized to the P domain of ATP1 α 3 (p.Gly358Val and p.Ile363Asn), and the observed mutations resulted in significant reduction of ATP1 α 3 activity in vitro. As allelic disorders of *ATP1A3*, e.g. RDP and AHC, appeared significant difference in phenotype, the cases with catastrophic early life epilepsy showed profound outcome and required much intensive care. Further functional studies are recommended to clarify the relationship between the mutation and such distinct phenotype.

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- **Education**

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2000-2004 PhD, Nagoya University Graduate School of Medicine
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- **Publication**

1. **Ishihara N**, Armsen W, Papadopoulos T, Betz H, Eulenburg V. Generation of a mouse line expressing Cre recombinase in glycinergic interneurons. *Genesis*. 2010 Jul;48(7):437-45.
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Session 1-5

Genotype-Structure-Phenotype: what makes *ATP1A3* mutations different?

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ATP1A3 is in the P2 family of ion transport ATPases with similar structures and mechanisms. There is strong sequence homology, and it is logical to expect that homologous DNA variants that are pathogenic in one gene would be pathogenic in all. Programs that predict the pathogenicity of VUS (variants of unknown significance) in patients are based on this. It is surprising, then, that the mutations actually found in patients differ very significantly between ATPases, comparing *ATP1A3* (AHC, RDP, etc.), *ATP1A2* (FHM2), and *ATP2A2* (a calcium ATPase, Darier disease, dominant inheritance). In Darier disease, mutations are distributed broadly in the protein structure, and can be truncations or frameshifts as well as missense. This is consistent with haploinsufficiency. For the Na,K-ATPases this is not the case. Almost all mutations are missense, indicating that the damaged protein must be present to cause the phenotypes.

The question is whether there is a genotype-phenotype relationship that underlies the specific symptoms that people exhibit. We analyzed the genetic changes (both random variants and human disease mutations) in all four of the Na,K-ATPase alpha subunit genes, and compared the locations of the disease mutations in crystal structures in E1-Na and E2-K conformations. The results are striking. The frequency of random variants differs among the genes, and shows the effects of negative selection. The specific patterns of disease mutations are equally different: highly restricted somatic mutations for *ATP1A1*, broadly distributed for *ATP1A2*, largely clustered for *ATP1A3*, and no mutations of *ATP1A4*. By examining the positions of mutations in the crystal structures, a genotype-phenotype relationship emerges. *ATP1A3* mutations that produce milder and more severe symptoms have different distributions in three-dimensional space. Finally, no mutations have been found in certain parts of the Na,K-ATPases, predicting that entirely new phenotypes will be discovered.

Supported by NIH grant NS058949 to A. Brashear.

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- **Present Position**

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- **Education**

1977 PhD, Harvard University, Cambridge, Massachusetts
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- **Academic Appointments**

1980-1981 Instructor, Department of Neurobiology, Harvard Medical School, Boston
1981-1987 Assistant Professor, Massachusetts General Hospital,
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- **Publications**

1. **Sweadner, KJ.** Enzymatic properties of separated isozymes of the Na,K-ATPase: substrate affinities, kinetic cooperativity, and ion transport stoichiometry. *J. Biol. Chem.* 1985; 260: 11508-11513.
2. McGrail, KM, Phillips, JM, and **Sweadner, KJ.** Immunofluorescent localization of three Na,K-ATPase isozymes in the rat central nervous system: Both neurons and glia can express more than one Na,K-ATPase. *J. Neurosci.* 1991; 11: 381-391.
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Session 2-1

Clinical profile of patients with *ATPIA3* mutations in AHC in an International Cohort

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In 2012, mutations in the *ATPIA3* gene (MIM 182350) were identified as the primary cause of AHC (AHC2, MIM 614820). Mutations in *ATPIA3* are found in approximately 75 % of cases and the disease is transmitted as an autosomal dominant trait. The mutations are usually de novo, but some have been found to be transmitted to offspring. Alternating hemiplegia of childhood (AHC) is a rare neurological disorder characterized by transient episodes of alternating hemiplegia/hemiparesis, dystonic attacks, paroxysmal abnormal ocular movements, epileptic seizures and episodes of autonomic dysfunction. The disease usually starts before 18 months of life and in the majority of patients before the age of 6 months. Plegic and tonic attacks disappear with sleep. Between attacks patients have an abnormal neurological examination often presenting ataxia, dystonia and other involuntary abnormal movements, and almost all present an intellectual disability. AHC has a prevalence of 1:100,000 children. There is a significant variability of the disease course between individuals and some genotype-phenotype correlations exist.

Clinical data from an international cohort of 155 AHC patients (84 females, 71 males; between 3 months and 52 years) were gathered using a specifically formulated questionnaire and analyzed relative to the mutational *ATPIA3* gene data for each patient.

RESULTS:

In total, 34 different *ATPIA3* mutations were detected in 85 % (132/155) patients. In general, mutations were found to cluster into five different regions. The most frequent mutations included: p.Asp801Asn (43 %; 57/132), p.Glu815Lys (16 %; 22/132), and p.Gly947Arg (11 %; 15/132). Of these, p.Glu815Lys was associated with a severe phenotype, with more severe intellectual and motor disability. p.Asp801Asn appeared to confer a milder phenotypic expression, and p.Gly947Arg appeared to correlate with the most favorable prognosis, compared to the other two frequent mutations. Overall, the comparison of the clinical profiles suggested a gradient of severity between the three major mutations with differences in intellectual ($p = 0.029$) and motor ($p = 0.039$) disabilities being statistically significant. For patients with epilepsy, age at onset of seizures was earlier for patients with either p.Glu815Lys or p.Gly947Arg mutation, compared to those with p.Asp801Asn mutation ($p < 0.001$). With regards to the five mutation clusters, some clusters appeared to correlate with certain clinical phenotypes. No statistically significant clinical correlations were found between patients with and without *ATPIA3* mutations.

CONCLUSIONS:

Our results, demonstrate a highly variable clinical phenotype in patients with AHC that correlates with certain mutations and possibly clusters within the *ATPIA3* gene. Our description of the clinical profile of patients with the most frequent mutations and the clinical picture of those with less common mutations confirms the results from previous studies, and further expands the spectrum of genotype-phenotype correlations.

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Academic Qualifications (most current date first)

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French inter-University Diploma of Epileptology. 2005 University Henri Poincaré, Nancy 1, France.

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From March 2003 to January 2004. Paediatric Hospital of Aghia Sophia, Athens, Greece.

From 2000 to 2002. Resident in paediatrics Paediatric hospital of Aghia Sophia, Athens, Greece.

From May 1996 to October 1998. Resident in paediatrics General Hospital of Kalamata, Greece.

From October 1993 to October 1995. General Practitioner. Following Greek legislation, all doctors are obliged, after their degree, to work for at least one year in a rural area. Public Medical Center of Areopolis, Greece.

Peer Reviewed Publications

- 1: Rheims S, Herbillon V, Villeneuve N, Auvin S, Napuri S, Cances C, Berquin P, Castelneau P, Nguyen The Tich S, Villega F, Isnard H, Nabbout R, Gaillard S, Mercier C, Kassai B, Arzimanoglou A; investigators of the Paediatric Epilepsy REsearch NETwork (PERENE). ADHD in childhood epilepsy: Clinical determinants of severity and of the response to methylphenidate. *Epilepsia*. 2016 Jul;57(7):1069-77.
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- 11: Bizec CL, Nicole S, **Panagiotakaki E**, Seta N, Vuillaumier-Barrot S. No Mutation in the SLC2A3 Gene in Cohorts of GLUT1 Deficiency Syndrome-Like Patients Negative for SLC2A1 and in Patients with AHC Negative for ATP1A3. *JIMD Rep*. 2014;12:115-20.
- 12: Heinzen EL, Swoboda KJ, Hitomi Y, Gurrieri F, Nicole S, de Vries B, Tiziano FD, Fontaine B, Walley NM, Heavin S, **Panagiotakaki E**; European Alternating Hemiplegia of Childhood (AHC) Genetics Consortium; Biobanca e Registro Clinico per l'Emiplegia Alternante (I.B.AHC) Consortium; European Network for Research on Alternating Hemiplegia (ENRAH) for Small and Medium-sized Enterprise (SMEs) Consortium, Fiori S, Abiusi E, Di Pietro L, Sweney MT, Newcomb TM, Viollet L, Huff C, Jorde LB, Reyna SP, Murphy KJ, Shianna KV, Gumbs CE, Little L, Silver K, Ptáček LJ, Haan J, Ferrari MD, Bye AM, Herkes GK, Whitelaw CM, Webb D, Lynch BJ, Uldall P, King MD, Scheffer IE, Neri G, Arzimanoglou A, van den Maagdenberg AM, Sisodiya SM, Mikati MA, Goldstein DB. De novo mutations in ATP1A3 cause alternating hemiplegia of childhood. *Nat Genet*. 2012 Sep;44(9):1030-4.

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- 19: Loudianos G, Lovicu M, Solinas P, Kanavakis E, Tzetis M, Manolaki N, **Panagiotakaki E**, Karpathios T, Cao A. Delineation of the spectrum of Wilson disease mutations in the Greek population and the identification of six novel mutations. *Genet Test*. 2000;4(4):399-402.

Session 2-2

Transcriptome analysis and clinical database/biorepository of Alternating Hemiplegia of Childhood patients with *ATPIA3* mutations for potential therapeutic discovery

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Mutations in the *ATPIA3* gene have been identified to be prevalent in patients with alternating hemiplegia of childhood (AHC). Over the past twenty years, we have made important contributions towards understanding AHC by establishing a clinical database and linked biorepository. This resource now contains data from approximately 300 affected probands. Our biorepository contains ~350 biological samples from probands including DNA, RNA, plasma, serum, whole blood, and cell lines, as well as CNS tissues from autopsy in 5 patients. Of 300 patients with an AHC phenotype, ~55% (164/300) have confirmed *ATPIA3* mutations with a total of 40 unique mutations. The majority of mutations are in exons 17, 18, and 21 with the most frequent mutations being: p.Asp801Asn (D801N) 20% (61/300), p.Glu815Lys (E815K) 14% (42/300), and p.Gly947Arg (G947R) 3.7% (11/300). Other *ATPIA3* mutations are present in 17% (51/300) and no *ATPIA3* mutations are present in ~12% (35/300) of affected individuals.

To date, no large scale transcriptomic data exist for human-derived AHC samples. The Connectivity Map (CMap) is a database of transcriptional signatures from cell lines treated with chemical and genetic reagents. When paired with the high-throughput L1000 by using a pattern-matching algorithm, the CMap links compounds with disease phenotypes. These CMAP/L1000 tools have helped identify several drug classes or compounds with potential activity for other treatment indications. Examples include the antihelminthic drug parabendazole as an inducer of osteoclast differentiation, the triterpene celastrol as a leptin sensitizer, compounds targeting COX2 and ADRA2A as potential treatments for diabetes, and small molecule therapeutics for skeletal muscular atrophy and spinal muscular atrophy. These tools have also helped to generate new therapeutic hypotheses for the treatment of inflammatory bowel disease and cancer. The present work leverages the rich tissue and phenotypic resources in our database/repository and expertise of the microarray analysis and deep sequencing resources at the Broad Institute/MIT. Using the CMap/L1000 tools, we performed a pilot experiment with 31 whole blood RNA samples from AHC patients and 89 healthy control samples. Our preliminary results revealed differences in the dynamic range of fold-expression achieved per gene in AHC samples when compared to controls. The gene ontology pathways most represented in the AHC gene expression profiles included genes involved in the circulatory system, vascular and embryonic development, as well as in cell proliferation pathways.

Following completion of testing and analyses of the current dataset, we hope to generate hypotheses and perform additional studies in patient derived samples including blood, cell lines, and brain. Identification of unique gene profiles in AHC could help identify novel pathways involved in disease pathogenesis and compounds that could ameliorate the disease phenotype.

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Education

| | | | |
|-------------|-----------------|-------------------|--|
| 07/82-06/86 | BA | Biology | Washington University, St. Louis, MO |
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| 07/86-06/90 | MD | Medicine | Feinberg School of Medicine, Northwestern University, Chicago, IL |
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Publications

1. Heinzen EL, **Swoboda KJ**, Hitomi Y, Gurrieri F, Nicole S, de Vries B, Tiziano FD, Fontaine B, Walley MN Heavin S, Panagiotakaki E, European Alternating Hemiplegia of Childhood, Genetics Consortium, Biobanca e Registro Clinico per l'Emiplegia Alternate (I.A.AHC) Consortium, European Network for

- Research on Alternating Hemiplegia (ENRAH) for small and medium size enterprises (SMEs) Consortium, Fiori S, Abiusi E, DiPietro L, Sweney MT, Newcomb TM, Viollet L, Huff C, Jorde LB, Reyna SP, Murphy KJ, Shianna KV, Gumbs CE, Little L, Silver K, Ptacek LJ, Haan J, Ferrari MD, Bye AM, Herkes GK, Whitelaw CM, Webb D, Lynch BJ, Uldall P, King MD, Scheffer IE, Neri G, Arzimanoglou A, Van den Maagdenberg AM, Sisodiya SM, Mikati MA, Goldstein DB. (07/29/2012). De Novo mutations in ATP1A3 cause alternating hemiplegia of Childhood. *Nat Genet*. 2012; 44(9), 1030-1034.
2. Flanigan KM, Ceco E, Lamar KM, Kaminoh Y, Dunn DM, Mendell JR, King WM, Pestronk A, Florence JM, Mathews KD, Finkel RS, **Swoboda KJ**, Gappmaier E, Howard MT, Day JW, McDonald C, McNally EM, Weiss RB; United Dystrophinopathy Project. LTBP4 genotype predicts age of ambulatory loss in Duchenne Muscular Dystrophy. *Ann Neurol*. 2012 Nov 26. doi: 10.1002/ana.23819. [Epub ahead of print] PMID: 23440719
 3. Rothwell E, Anderson RA, **Swoboda KJ**, Stark L, Botkin JR. Public attitudes regarding a pilot study of newborn screening for spinal muscular atrophy. *Am J Med Genet A*. 2013 Apr; 161(4): 679-86. doi: 10.1002/ajmg.a.35756. Epub 2013 Feb 26. PMID: 23443997
 4. Kobayashi DT, Shi J, Stephen L, Ballard KL, Dewey R, Mapes J, Chung B, McCarthy K, **Swoboda KJ**, Crawford TO, Li R, Plasterer T, Joyce C; Biomarkers for Spinal Muscular Atrophy Study Group, Chung WK, Kaufmann P, Darras BT, Finkel RS, Sproule DM, Martens WB, McDermott MP, De Vivo DC; Pediatric Neuromuscular Clinical Research Network, Walker MG, Chen KS. SMA-MAP: a plasma protein panel for spinal muscular atrophy. *PLoS One*. 2013;8(4): e60113. doi: 10.1371/journal.pone.0060113. Epub 2013 Apr 2. PMID: 23565191
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 6. Cano SJ, Mayhew A, Glanzman AM, Krosschell KJ, **Swoboda KJ**, Main M, Steffensen BF, Bérard C, Girardot F, Payan CA, Mercuri E, Mazzone E, Elsheikh B, Florence J, Hynan LS, Iannaccone ST, Nelson LL, Pandya S, Rose M, Scott C, Sadjadi R, Yore MA, Joyce C, Kissel JT; International Coordinating Committee for SMA Clinical Trials Rasch Task Force. Rasch Analysis of Clinical Outcome Measures in Spinal Muscular Atrophy. *Muscle Nerve*. 2013 Jul 8. PMID: 23836324
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Session 2-3

Cardiac phenotype in *ATP1A3* related-syndromes

S Balestrini, M Mikati, R Vavassori, M McLean, A Brashear, C Miller, R Samoes, J Novy, E de Grandis, E Veneselli, M Stagnaro, F Ragona, T Granata, N Nardocci, C Zucca, C Fons, J Campistol, I Scheffer, G Hollingsworth, E Pangiotakaki, A Arzimanoglou, I Carrilho, S Groppa, A Potic, QS Padiath, V Brankovic, R Pons, K Dzieżyc, M Mazurkiewicz-Beldzińska, J Pilch, K Vezyroglou, H Cross, JP Kaski, SM Sisodiya

Alternating hemiplegia of childhood (AHC) is a rare neurodevelopmental disorder with significant phenotypic diversity, caused in ~80% of cases by mutations in the *ATP1A3* gene. This gene encodes the catalytic alpha-3 subunit of the Na⁺/K⁺ ATPase exchange pump. Known outcomes range from life into adulthood, with comparatively little disability, to premature mortality from sudden death, including sudden unexpected death in epilepsy (SUDEP). We recently demonstrated that many patients with AHC have ECG changes, varying from repolarisation abnormalities, J-wave or J-point changes, to periods of asystole, with an increasing prevalence with age. The ECG abnormalities were dynamic, reflecting characteristics of inherited cardiac channelopathies, and suggesting, along with the paroxysmal neurological features, periodic systemic decompensation in *ATP1A3*-expressing organs. Given these findings, systematic cardiac investigation is warranted in this condition, as cardiac arrhythmic morbidity and mortality are potentially preventable (with implantation of a cardiac pacemaker or defibrillator). Also, certain drugs should be avoided in view of the increased risk of precipitating serious arrhythmias. We are now carrying out a second study enlarging the cohort (113 cases in total so far) and including cases with other *ATP1A3*-related phenotypes: cerebellar ataxia, areflexia, pes cavus, optic atrophy, and sensorineural hearing loss (CAPOS); and rapid-onset dystonia–parkinsonism (RDP). In this study we are investigating the cardiac phenotype further, collecting data also on prolonged ECG monitoring, echocardiography and other cardiological evaluation tests. This will validate our previous findings and give more insight on the phenotype-genotype correlation in AHC and other *ATP1A3*-associated syndromes.

Simona BALESTRINI

WORK EXPERIENCE

From 1.8.2014 ongoing. Clinical Research Associate at the Department of Clinical and Experimental Epilepsy, University College of London (UCL). Main topic of research: genetics of epilepsies, exome and genome analysis, genotype-phenotype association studies, transcranial magnetic stimulation.



From 23.2.2017 ongoing. Honorary Consultant Neurologist, at University College London Hospital (UCLH) NHS Foundation Trust.

From 1.7.2016 to 30.11. 2016. Locum Consultant Neurologist – Special Interest in Epilepsy, at University College London Hospital (UCLH) NHS Foundation Trust.

From 30.6.2009 to 4.7.2014. Residence in Neurology at Polytechnic University of Marche (Ancona, Italy), with focus on epilepsy, neuro-oncology, cerebrovascular disease, epidemiological research.

EDUCATION AND TRAINING

From 1.11.2013 to 1.3.2017. Phd student at Polytechnic University of Marche (Ancona, Italy). Main topic of research: genetics of drug-resistant epilepsy. Final thesis on ‘Biomarkers of Sudden Unexpected Death in Epilepsy (SUDEP)’.

On 4th Jul 2014. Completion of Medical Residency in Neurology at Polytechnic University of Marche (Ancona, Italy), with thesis on “Electrical Stimulations of Parietal Lobe: Stereo-EEG Study in Patients with Drug-Resistant Focal Epilepsy” (50/50 with honour).

From Jan to Dec 2013. Clinical Research Associate at the Department of clinical and experimental Epilepsy, University College of London (UCL) Main topics: clinical research projects about genetic basis of epilepsy, phenotype-genotype correlation, genetic mechanisms of SUDEP.

On 24th Jul 2008. Degree in Medicine and Surgery at Polytechnic University of Marche with thesis on "Progression of Carotid Atherosclerosis and Zinc Omeosthesis" (110/110 with honour).

PUBLICATIONS

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Session 3-1

New causative gene(s) in AHC?

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As not all patients with alternating hemiplegia of childhood (AHC) carry a mutation in the coding regions of the *ATPIA3* gene, a gene hunt was initiated aimed at identifying a second gene for this disabling disorder. Data of exome sequencing on genomic DNA of over 25 trio's (that is DNA of the patient and both parents) was performed to search primarily for *de novo* pathogenic mutations. Even with this large number of samples, the search for additional AHC genes is challenging, not in the least because of clinical heterogeneity among *ATPIA3*-negative cases, many being classified as atypical. The status of the gene hunt will be discussed on behalf of the Consortium. Approaches to screen for possible mutations in *ATPIA3* in presumed *ATPIA3*-negative cases will also be discussed. The gene hunt suggests that there may not be a second major gene after all. To unravel the (molecular) pathophysiology of AHC, it may be fruitful to learn from developments, with respect to disease pathways involved and functional assays to assess brain dysfunction, in disorders with resemblance to AHC.

Arn M.J.M. VAN DEN MAAGDENBERG



▪ Present position

Professor, Departments of Human Genetics and Neurology,
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▪ Education

1981 - 1988 Biology, Catholic University of Nijmegen, the Netherlands

1989 - 1993 Fellow in Human Genetics, Leiden University Medical
Centre, the Netherlands

1993 PhD, Leiden University Medical Centre, the Netherlands

▪ Academic appointments

1993 - 1998 Postdoctoral Fellow, Department of Cell Biology & Histology, University of
Nijmegen, The Netherlands

1998 - 2005 Postdoctoral Fellow, Department of Human Genetics, Leiden University Medical
Centre, The Netherlands

2005 - 2011 Associate Professor, Departments of Human Genetics & Neurology, Leiden
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2011 - Full professor, Departments of Human Genetics & Neurology (LUMC). Chair:
“Molecular and Functional Neurogenetics”

▪ Publications (relevant selection of in total 240 peer-reviewed publications)

1. Heinzen E, Swoboda K, Hitomi Y, **van den Maagdenberg A**, Sisodiya S, Mikati M, Goldstein D. De novo mutations in ATP1A3 cause alternating hemiplegia of childhood. *Nature Genetics* 2012;44:1030-1034.
2. Freilinger T, Anttila V, de Vries B, **van den Maagdenberg AM**; International Headache Genetics Consortium. Genome-wide association analysis identifies susceptibility loci for migraine without aura. *Nature Genetics* 2012;44:777-782.
3. Heinzen E, Arzimanoglou A, Brashear A, **van den Maagdenberg A**, Vilsen B; ATP1A3 Working Group. Distinct neurological disorders with ATP1A3 mutations. *Lancet Neurology* 2014;13:503-514.
4. Ferrari M*, Klever R*, Terwindt G, Ayata C, **van den Maagdenberg A**. Migraine pathophysiology: lessons from mouse models and human genetics. *Lancet Neurology* 2015;14:65-80.
5. Eising E, Shyti R, 't Hoen PAC, **van den Maagdenberg AM**. Cortical Spreading Depression Causes Unique Dysregulation of Inflammatory Pathways in a Transgenic Mouse Model of Migraine. *Molecular Neurobiology* 2017;54:2986-2996.

Session 3-2

Biochemical consequences of *ATP1A3* mutations

Jan B. Koenderink

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Alternating Hemiplegia of Childhood (AHC) is a neurodevelopmental disorder caused by de novo mutations in *ATP1A3*, the gene encoding the $\alpha 3$ subunit of Na,K-ATPase. Na,K-ATPase regulates electrogenic transport by exporting three Na⁺ ions in exchange for two K⁺ ions across the cellular membrane by hydrolysis of one adenosine triphosphate molecule. Maintaining these ionic gradients has been shown to be of great importance in a variety of cellular functions, including neuronal activity.

Depending on the location of the mutation in *ATP1A3* differences in AHC development have been observed. Moreover, some mutations in *ATP1A3* do not cause AHC, but rapid-onset dystonia-parkinsonism (RDP) and cerebellar ataxia, areflexia, pes cavus, optic atrophy, and sensorineural hearing loss (CAPOS).

The functional consequences of many AHC mutations have been characterized using over-expression in different mammalian cells, insect cells or *Xenopus laevis* oocytes. In addition, also several functional characteristics have been analyzed: Electrophysiological activity, ATPase activity, cell survival, ouabain binding, and phosphorylation capacity. A loss of function has been observed in most assays. A detailed analysis has been performed only for some of the many different mutations detected.

Many fundamental questions still remain, but what did we learn from these functional assays? Is it important to continue detailed functional assays for all mutations identified? Which knowledge gaps still exist and should be addressed?

Jan Koenderink

Jan is assistant professor in biochemical toxicology at the Department of Pharmacology and Toxicology, Radboud University Nijmegen Medical Centre, The Netherlands. His research interest is focussed on membrane transport proteins and their ligands (substrates and inhibitors). The interaction between membrane transporters (uptake and efflux) and novel and existing human drugs, as well as the multidrug resistance transport proteins (ABC transporters) of parasites (e.g. *Mycobacterium tuberculosis* and *Plasmodium falciparum*) are investigated. In addition, the molecular activity and interactions of the sodium pump (Na,K-ATPase) are studied. The studies include the expression of recombinant transport proteins and their molecular / biochemical and pharmacodynamic characterization.



Publications

Alternating Hemiplegia of Childhood mutations have a differential effect on Na(+),K(+)-ATPase activity and ouabain binding. Weigand KM, Messchaert M, Swarts HG, Russel FG, **Koenderink JB**. *Biochim Biophys Acta*. 2014 Jul;1842(7):1010-6.

Na(+),K(+)-ATPase isoform selectivity for digitalis-like compounds is determined by two amino acids in the first extracellular loop. Weigand KM, Laursen M, Swarts HG, Engwerda AH, Prüfert C, Sandrock J, Nissen P, Fedosova NU, Russel FG, **Koenderink JB**. *Chem Res Toxicol*. 2014 Dec 15;27(12):2082-92.

Vital and dispensable roles of Plasmodium multidrug resistance transporters during blood- and mosquito-stage development. Rijpma SR, van der Velden M, Annoura T, Matz JM, Kenthirapalan S, Kooij TW, Matuschewski K, van Gemert GJ, van de Vegte-Bolmer M, Siebelink-Stoter R, Graumans W, Ramesar J, Klop O, Russel FG, Sauerwein RW, Janse CJ, Franke-Fayard BM, **Koenderink JB**. *Mol Microbiol*. 2016 Jul;101(1):78-91.

Moxifloxacin Is a Potent In Vitro Inhibitor of OCT- and MATE-Mediated Transport of Metformin and Ethambutol. Te Brake LH, van den Heuvel JJ, Buaben AO, van Crevel R, Bilos A, Russel FG, Aarnoutse RE, **Koenderink JB**. *Antimicrob Agents Chemother*. 2016 Nov 21;60(12):7105-7114.

The role of efflux pumps in TB treatment and their promise as a target in drug development: unraveling the black box. Te Brake LH, de Knecht GJ, de Steenwinkel JE, van Dam TJ, Burger DM, Russel FG, van Crevel R, **Koenderink JB**, Aarnoutse RE. *Annual Review of Pharmacology and Toxicology*, 2017, *In press*.

Session 3-3

ATP1A3 neurological disease mutations affecting Na⁺ binding: Structural and functional perspectives and rescue of compromised function continued

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Several neurological phenotypes derive from ATP1A3 mutations. The effects of several of these mutations on Na⁺,K⁺-ATPase function have been studied *in vitro* [1]. The high-resolution crystal structures of the Na⁺,K⁺-ATPase provide a solid basis for understanding functional defects of the ATP1A3 disease mutants and of the corresponding ATP1A1 mutants. Na⁺ binds at three sites, I, II, and III, of which I and II overlap with the K⁺ binding sites, whereas site III is unique and Na⁺ specific [2]. In several ATP1A3 disease mutants the compromised function can be traced to disturbance of the Na⁺ specific binding site III, although some ATP1A3 disease mutants show disruption of Na⁺ binding at sites I and II. Furthermore a limited number of the ATP1A3 mutations occur in the cytoplasmic P domain near the catalytic site. We have recently characterized the functional disturbance caused by some of these mutations.

A secondary mutation was found to rescue the defective Na⁺ binding at site III caused by RDP/AHC mutation D923N [1, 3]. A perspective is that it may be feasible to develop an efficient pharmaceutical mimicking the rescuing effect, which optimally would rescue the compromised function of a variety of ATP1A3 mutants. In our continued work on the rescuing effect we have studied combinations of the rescuing mutation with mutations in sites I and II and additional ATP1A3 mutations.

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Present Position

Professor, Department of Biomedicine, Faculty of Health, Aarhus University, Aarhus, Denmark

Education

1984 Masters degree (cand.scient.) in Biology, Aarhus University
1995 D.M.Sc. (dr.med.), Aarhus University



Academic Appointments

1986-1988 Research Assistant, Department of Physiology, Aarhus University
1988-1989 Visiting Scientist at Professor D.H. MacLennan, University of Toronto, Canada
1988-1992 Assistant Professor, Department of Physiology, Aarhus University
1992-2006 Associate Professor, Department of Physiology, Aarhus University
2006-Present Full professor, Department of Biomedicine, Aarhus University

Publication (119 articles in international journals, H-index 41)

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Session 4-1

Development of new treatments in the *Myshkin* mouse model of AHC

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Alternating hemiplegia of childhood (AHC; OMIM: 614820) is a rare neurodevelopmental disorder that manifests as episodic hemiplegia starting in the first 18 months of life, with a spectrum of persistent motor, movement and cognitive deficits that become progressively more apparent with age. Flunarizine reduces the severity, duration, or frequency of hemiplegic attacks in some patients, while two AHC patients have shown marked improvements in symptoms when treated with a ketogenic diet. However, the complexity and severity of AHC make it imperative that new therapeutic options be explored.

Heterozygous missense mutations of the *ATP1A3* gene, encoding the neurone-specific Na⁺,K⁺-ATPase (NKA) α 3 subunit, have been identified as the primary cause of AHC as well as rapid-onset dystonia-parkinsonism (RDP) and CAPOS syndrome. Towards gaining a fuller understanding of the neurological effects of NKA α 3 dysfunction, we developed the *Myshkin* mouse model, which has an I810N mutation identical to that present in AHC.

Heterozygous *Myshkin* mice move with a paretic, tremulous gait that becomes transiently more severe after stress. Other phenotypic abnormalities include decreased body mass, neuronal hyperexcitability, increased susceptibility to epileptic seizures, motor dysfunction and cognitive impairment. Genetic (*Atp1a3* transgene), pharmacological (rostafuloxin), and dietary (medium-chain fatty acids) approaches to the treatment of AHC-related phenotypes in the *Myshkin* model will be presented.

Myshkin mice carrying a wildtype *Atp1a3* transgene that confers a 16% increase in brain-specific total NKA activity show significant phenotypic improvements compared with non-transgenic *Myshkin* mice. Chronic treatment with rostafuloxin (PST-2238), a compound which selectively displaces endogenous ouabain from the NKA, reduces the risk-taking behaviour of *Myshkin* mice. The potential therapeutic effects of a diet rich in medium-chain fatty acids will also be discussed.

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- 1999 PhD, Dept. of Genetics & Microbiology, University of Liverpool, UK
- 1998-2002 Postdoctoral Fellow, Dept. of Zoology, University of Oxford, UK
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Academic Appointments

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Relevant Publications

1. **CLAPCOTE, S.J.**, Duffy, S., Xie, G., Kirshenbaum, G., Bechard, A.R., Rodacker Schack, V., Petersen, J., Sinai, L., Saab, B.J., Lerch, J.P., Minassian, B.A., Ackerley, C.A., Sled, J.G., Cortez, M.G., Henderson, J.T., Vilsen, B. & Roder, J.C. (2009). Mutation I810N in the $\alpha 3$ isoform of Na⁺,K⁺-ATPase causes impairments in the sodium pump and hyperexcitability in the CNS. *Proceedings of the National Academy of Sciences of the USA* 106: 14085-14090.
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3. Kirshenbaum, G.S., **CLAPCOTE, S.J.**, Petersen, J., Vilsen, B., Ralph, M.R. & Roder, J.C.(2012). Genetic suppression of agrin reduces mania-like behavior in Na⁺,K⁺-ATPase $\alpha 3$ mutant mice. *Genes, Brain and Behavior* 11: 436-443.

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Session 4-2

Hypothermia-induced dystonia and abnormal cerebellar activity in a mouse model with a single disease-mutation in the sodium-potassium pump

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The neurological spectrum associated with mutations in the *ATP1A3* gene, encoding the α_3 isoform of the Na^+/K^+ -ATPase, is complex and still poorly understood. To elucidate the disease-specific pathophysiology, we examined a mouse model harboring the mutation D801Y, which was originally found in a patient with Rapid onset Dystonia Parkinsonism, but recently, also in a patient with Alternating Hemiplegia of Childhood. We found that this model exhibited motor deficits and developed dystonia when exposed to a drop in body temperature. Cerebellar *in vivo* recordings in awake mice revealed irregular firing of Purkinje cells and their synaptic targets, the deep cerebellar nuclei neurons, which was further exacerbated and evolved into abnormal high-frequency burst firing during dystonia. The development of specific neurological features within the *ATP1A3* mutation spectrum, such as dystonia, are thought to reflect the functional consequences of each mutation, thus to investigate the consequence of the D801Y mutations we characterized mutated D-to-Y Na^+/K^+ -ATPases expressed in *Xenopus* oocytes. These *in vitro* studies showed that the D-to-Y mutation abolishes pump-mediated Na^+/K^+ exchange, but still allows the pumps to bind Na^+ and become phosphorylated, trapping them in conformations that instead support proton influx.

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Publications (selected)

*Isaksen, Kros, Vedovato, Holm, Gadsby, Khodakhah and **Lykke-Hartmann**, *Hypothermia-induced dystonia in an RDP mouse model*.

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Session 4-3

Study of Knock-in model mouse; Development of new treatments

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Alternating hemiplegia of childhood is a serious disorder that together with its related disorders is becoming increasingly recognized. Recent advances in animal models of AHC including the knock in mouse models of the two most common mutations have improved the understanding of the underlying pathophysiology of AHC and are suggesting possible therapeutic interventions and potential translational research. For example studies in the D801Y model showed improvement in cognitive deficits using clonazepam, in the E815K model showed that Flunarizine acutely decreased the number of hemiplegia and dystonia spells, but caused after its discontinuation no chronic deterioration or protection suggesting that the detrimental effects observed in some patients after Flunarizine discontinuation are due to withdrawal of its beneficial effects of the drug. Finally the D801N model is being used for screening of medications as related to the underlying physiology. In that model we have demonstrated increased excitability, increased predisposition to spreading depression, and decreased GABAergic inhibition as demonstrated in our lab. The Deepening understanding of the physiology is bound to continue to help in choosing future candidate drugs.

Mohamad A. Mikati M.D., is the Wilburt C. Davison Professor of Pediatrics, Professor of Neurobiology, and Chief of the Division of Pediatric Neurology. Dr. Mikati's clinical research has centered on characterization and therapy of pediatric epilepsy and neurology syndromes, describing several new pediatric neurological entities with two carrying his name (POSSUM syndromes # 3708 and 4468), developing novel therapeutic strategies for epilepsy and related disorders particularly Alternating Hemiplegia of Childhood, and applying cutting edge genetic and Magnetic Resonance Imaging techniques to drug resistant pediatric epilepsy. In the laboratory he has elucidated mechanisms of seizure related neuronal injury, particularly those related to the ceramide pathway, and demonstrated neuroprotective effects of several agents including erythropoietin. Most recently he has concentrated his laboratory research on the pathophysiology of ATP1A3 dysfunction in the brain as model for epilepsy and of Alternating Hemiplegia of Childhood.



He has more than 210 peer reviewed publications, 300 abstracts 40 chapters one book and two booklets. He also has more than 6,214 citations in the literature with an h-index of 46 and an i-10index of 127. Dr. Mikati has written chapters on epilepsy and related disorders in the major textbooks of Pediatrics and Pediatric Neurology including Swaiman's Pediatric Neurology and Nelson's Pediatrics.

Before joining Duke in 2008 he had completed his M.D. and Pediatric training at the American University of Beirut, his Neurology at the Massachusetts General Hospital, his Neurophysiology at Boston Children's Hospital and had been on the Faculty at Harvard as Director of Research in the Epilepsy Program at Boston Children's Hospital and then as Professor and Chairman, Department of Pediatrics, Founder and Director of the Adult and Pediatric Epilepsy Program at the American University of Beirut.

Dr. Mikati has had several international leadership roles including being President of the Union of the Middle Eastern and Mediterranean Pediatric Societies, on the Standing Committee of the International Pediatric Association, Officer of the International Child Neurology Association, Member of the Pediatric Content and Scientific Program Committees, Child Neurology Society International Affairs Committee Consultant to UNICEF, WHO, and the American Board of Pediatrics, and being one of only two Pediatric Neurologists, initially chosen worldwide, on the WHO advisory committee for the International Classification of Disease being the head of the neuromuscular group in that committee. He is also a co-founder of the International Alternating Hemiplegia of Childhood Research Consortium and Deputy Scientific Coordinator of the Consortium. He is also a member of the Planning Committee for the International ATP1A3 Disease Group. He has received several national and international honors including, among others, Merritt Putnam American Epilepsy Society Fellowship Award, Harvard Community Health Plan Peer Recognition Award, Debs Research Award, Hamdan Award for contributions to Medicine, Hans Zellweger Award for contributions to Pediatric Neurology, Patient Choice Award, American University of Beirut Alumni Achievement and Service Award, and the Michael Frank Award for research and lifetime contributions to the field of Pediatric Neurology. His name consistently appears in America's Top Doctors Series.

Mikat MA, authored and co-authored Alternating Hemiplegia of Childhood Publications

1. Masoud M, Lyndsey Prange L, CPNP, Wuchich J, Hunanyan A, **Mikat MA**. Diagnosis and Treatment of Alternating Hemiplegia of Childhood. *Current Treatment Options in Neurology*, 2017 Feb;19(2):8
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 12. **Mikati MA**, Fischman A. PET Scan Findings in Alternating Hemiplegia of Childhood. In: *Alternating Hemiplegia of Childhood*. Eds: Aicardi J, Andermann F. New York: Raven. 1995:109-14.
 13. **Mikati MA**, Maguire H, Barlow CF, Ozelius L, Breakefield XO, Klauck SM, Korf B, O'Tuama SL, Dangond F. A syndrome of autosomal dominant alternating hemiplegia: clinical presentation mimicking intractable epilepsy; chromosomal studies; and physiologic investigations. *Neurology.* 1992 Dec;42(12):2251-7. PubMed PMID: 1361034.

Session 5-1

Treatment with adenosine- 5'- triphosphate for AHC

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The primary treatment for AHC has been anti-epileptic drugs (AEDs) and while none of these have been reported effective on non-epileptic paroxysmal attacks, some reports indicate their success as prophylactics. Flunarizine, a non-selective calcium ion channel, is commonly used in long term treatment. It has been reported to be effective in reducing the frequency and intensity of plegic attacks. There is still no gold standard treatment for patients with AHC,

The principal cause of AHC is an abnormality of ATPase pump which is caused by mutation in the *ATP1A3* gene. ATPase pump is the catalytic component of the active enzyme, which catalyzes the hydrolysis of Adenosine-5'-triphosphate (ATP) coupled with the exchange of sodium and potassium ions across the plasma membrane. This action establishes and maintains the electrochemical gradients of sodium and potassium ions. These gradients are essential for osmoregulation, the sodium-coupled transport of a variety of organic and inorganic molecules, and the electrical excitability of nerves and muscles. Some of AHC causing mutations has been reported to reduce ATP pump activity. Therefore, this function of ATPase pump supports the hypothesis that supplementation of ATP may improve the dysfunction of Na⁺/K⁺ ATPase to alleviate symptoms.

In an innovative therapy done with the approval of the IRB commission, a 7-year-old boy with a de novo splicing mutation (c. 2542+1G>A) in *ATP1A3* was treated for two years with oral ATP supplements. Therapy outcome was evaluated through the follow-up of improvement of hemiplegic episodes and psychomotor development. Side-effects were regularly monitored to insure patient safety. With the dosage of ATP administration increased, he showed significantly less frequency and shorter duration of hemiplegic episodes. Treatment with ATP was correlated with a marked amelioration of alternating hemiplegia of childhood episodes, and psychomotor development has improved. The maximum dose of oral administration of ATP reached 25 mg/kg per day. ATP therapy was well tolerated without complaint of discomfort and side effects. The 2-year follow-up outcome of ATP therapy for alternating hemiplegia of childhood was successful. This therapy is now considered one of the most useful treatments for AHC.

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Academic appointments

7/2001-3/2007 Undergraduate research in molecular pathology, Pediatrics, Fukuoka University
4/2007-3/2010 Graduate research in the Research Center for the Molecular Pathomechanisms of Epilepsy, Fukuoka University
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4/2013-8/2013 Research Fellow in Central Research Institute for the Molecular Pathomechanisms of Epilepsy, Fukuoka University
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Publications

1. **Ishii A.**, Watkins JC., Chen D., Hirose S., Hammer MF.. Clinical implications of SCN1A missense and truncation variants in a large Japanese cohort with Dravet syndrome. *Epilepsia*. 2016 Dec 1.
2. **Ishii A**, Kang JQ., Schornak CC., Hernandez CC., Shen W., Watkins JC., Macdonald RL., Hirose S.. A de novo missense mutation of GABRB2 causes early myoclonic encephalopathy. *Journal of Medical Genetics*. 2016 Oct 27:jmedgenet-2016.
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Session 5-2

Loss of function *ATP1A3* mutations differentiated by pre-steady-state analysis

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³ Equal contribution

Mutations in *ATP1A3* have been increasingly implicated in various neurological disorders. So far, loss of forward cycling function of the Na⁺/K⁺ ATPase is a common pathological feature of *ATP1A3* mutants. However, there are no clinically available compounds that enhance the function of the Na⁺/K⁺ ATPase. We aimed to elucidate effects of pathological *ATP1A3* mutations on the structure-function relationships to aid discovery of mechanism based therapeutics. The biophysical properties of two mutants Q140H and E818K were compared to the human wildtype (WT) $\alpha 3$ subunit by expression in *X.laevis* oocytes and recording using two-electrode voltage clamp. The voltage dependence of the forward cycling rate was significantly altered for the E818K mutant. Analysis of the pre-steady-state current relaxations, that reflect charge displacements associated with cation binding/debinding, revealed previously undocumented functional features. We observed a significant hyperpolarizing shift in the voltage dependence of displacement charge for E818K compared with WT. In contrast, whereas the voltage dependence of Q140H was similar to WT, the rate of current relaxation was significantly slower as compared to the WT or E818K. Pre-steady-state relaxation analysis can offer novel insights into how sodium ion translocation may be distinctly altered in mutants associated with different *ATP1A3* disorders. Furthermore, it provides a robust readout that complements the steady-state pump activity and may reveal novel targeted therapeutic mechanisms.

Professor Steven Petrou PhD FAHMS

Academic Qualifications

1984 B.Sc. Hons University of Melbourne

1992 Ph.D. University of Melbourne

Postdoctoral Training

1992-1994 Samuel A. Levine Research Fellow of the American Heart

Association. Department of Physiology, University of Massachusetts Medical School, USA

1995 Research Fellow. Worcester Foundation of Experimental Biology, Massachusetts, USA



Academic Positions

1998-1999 National Heart Research Fellow. Department of Physiology, University of Melbourne

2000-2001 Research Fellow. Department of Physiology, University of Melbourne

2001 Research Fellow. Max Planck Institute for Experimental Medicine and Research, Heidelberg, Germany

2001 Senior Research Fellow. Department of Physiology,

2008 Associate Professor, The University of Melbourne

2002-2009 Head of the Laboratory of Ion Channels and Human Disease, HFI

2009-2011 Associate Director, Florey Neuroscience Institute

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2013 Appointed to Department of Electrical Engineering, The University of Melbourne

2014 Professor, The University of Melbourne

2015 Professor of Translational Neuroscience (continuing), Department of Medicine, The University of Melbourne

Representative Publications

1. Oliver KL, Franceschetti S, Milligan CJ, Muona M, Mandelstam SA, Canafoglia L, Boguszevska-Chachulska AM, Korczyn AD, Bisulli F, Di Bonaventura C, Ragona F, Michelucci R, Ben-Zeev B, Straussberg R, Panzica F, Massano J, Friedman D, Crespel A, Engelsen BA, Andermann F, Andermann E, Spodar K, Lasek-Bal A, Riguzzi P, Pasini E, Tinuper P, Licchetta L, Gardella E, Lindenau M, Wulf A, Møller RS, Benninger F, Afawi Z, Rubboli G, Reid CA, Maljevic S, Lerche H, Lehesjoki AE, **Petrou S**, Berkovic SF. Myoclonus epilepsy and ataxia due to KCNC1 mutation: Analysis of 20 cases and K(+) channel properties. *Ann Neurol*. 2017 May;81(5):677-689.
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Session 5-3

Modeling Alternating Hemiplegia of Childhood using Patient Derived Stem Cells

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Alternating Hemiplegia of Childhood (AHC) is an episodic neurological disorder that was recently discovered to be caused to mutations in the *ATPIA3* gene. *ATPIA3* encodes the neuronal specific $\alpha 3$ subunit of the sodium-potassium adenosine triphosphatase (Na,K-ATPase) pump. Patients with AHC present in early childhood, often with triggered unilateral episodes of weakness, unilateral nystagmus, seizures, dystonia, developmental delay, and ultimately intellectual disability. Throughout the world, most patients with AHC have heterozygous *ATPIA3* mutations with D801N, E815K, or G947R missense mutations being the most common. While animal models of AHC have been developed, the complexity of human neural development make mechanistic analyses difficult and laborious using rodent models. This is due to the possible involvement of multiple neuronal lineages such as GABAergic, glutamatergic, and dopaminergic neurons as well as overlapping patterns of expression of $\alpha 1$, $\alpha 2$, and $\alpha 3$ subunits and potential compensation in cells with *ATPIA3* mutations.

In collaboration with the laboratories of Drs. Kathy Swoboda and Al George, we have made numerous lines of induced pluripotent stem cells (iPSCs) from patients with AHC including those with known D801N, E815K, or G947R mutations in the *ATPIA3* gene. These lines have been extensively characterized including verification of pluripotency status, karyotype, and confirmation of mutations in the *ATPIA3* gene. We have also used these iPSCs to optimize neuronal differentiation protocols to various neuronal lineages and to also assess the expression of $\alpha 1$, $\alpha 2$, and $\alpha 3$ subunits using immunoblotting. Ongoing experiments using fluorescent dyes will determine relative concentrations of sodium and calcium as well as ion flux in neuronal cultures differentiated from AHC patient iPSCs. Finally, we are actively making isogenic iPSCs using CRISPR/Cas9 technology to either correcting *ATPIA3* mutations in AHC patient-derived iPSC lines or introduce *ATPIA3* mutations to existing control lines. This approach will allow us to control for genetic variations that may introduce experimental variability. However, the same genetic variations may also underlie variability in disease severity seen in individual patients and lead to the identification of modifier genes that can be exploited in the future as potential druggable targets.

This research was generously supported by the Alternating Hemiplegia of Childhood Foundation as well as the Association Française de l'Hémiplégie Alternante.

Ess, Kevin C., M.D., Ph.D

Gerald M. Fenichel Chair in Neurology; Associate Professor



EDUCATION/TRAINING

| INSTITUTION AND LOCATION | DEGREE | Completion Date | FIELD OF STUDY |
|---|----------|-----------------|--------------------------|
| University of Cincinnati, Cincinnati, OH | B. M. | 06/1989 | Music Performance |
| Children's Hospital Research Foundation, University of Cincinnati, Cincinnati, OH | Ph.D. | 12/1996 | Developmental Biology |
| University of Cincinnati, Cincinnati, OH | M.D. | 06/1998 | Medicine |
| University of Colorado, Denver, CO | Intern | 06/1999 | Pediatrics |
| St. Louis Children's Hospital/Washington University, St Louis, MO | Resident | 06/2002 | Child Neurology |
| St. Louis Children's Hospital/Washington University | Fellow | 06/2004 | Clinical Neurophysiology |
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A. Personal Statement

I have dedicated my career to understanding the genetic control of brain development and how aberrations in developmental processes lead to epilepsy, and autism. For the past 11 years at Vanderbilt, my research has focused on tuberous sclerosis complex (TSC) as patients with this disease have prominent brain malformations, white matter disease and a very high prevalence of epilepsy and autism. Resulting from mutations in either the *TSC1* or *TSC2* genes, this disorder involves dysregulation of the mTOR kinase with striking signaling abnormalities of both the mTORC1 (increased) as well as the mTORC2 (decreased) signaling pathways. We have also been interested in a recently defined disorder, alternating hemiplegia of childhood (AHC), a devastating neurodevelopmental disorder due to mutations in the *ATPIA3* gene. To study abnormal developmental processes in TSC and AHC, we have utilized diverse model systems including transgenic mouse, zebrafish, and human induced pluripotent stem cells (iPSCs). Principally employing iPSCs, our basic and translational research approaches to TSC and AHC should culminate in advanced knowledge about pediatric neurological disorders and hopefully lead to the development of novel and more effective therapies.

B. Positions and Honors

Positions and Employment

| | |
|-------------|--|
| 2002 – 2006 | Founder and Director, Tuberous Sclerosis Clinic, St. Louis Children's Hospital |
| 2004 – 2006 | Instructor, Department of Neurology, Washington University, St. Louis, MO |
| 2006 – 2014 | Assistant Professor, Department of Neurology, Vanderbilt University, Nashville, TN |

| | |
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| 2006 – 2014 | Assistant Professor, Department of Pediatrics, Vanderbilt University, Nashville, TN |
| 2006 – present | Investigator, Vanderbilt Kennedy Center for Research on Human Development. |
| 2006 – present | Founder and Director, Tuberous Sclerosis Clinic, Vanderbilt Children’s Hospital |
| 2009 – 2014 | Assistant Professor, Department of Cell and Developmental Biology, Vanderbilt University |
| 2009 – present | Member, Vanderbilt Center for Stem Cell Biology, Nashville, TN |
| 2011 – 2014 | Assistant Professor, Department of Biological Sciences, Vanderbilt University |
| 2014– present | Associate Professor, Pediatrics, Neurology, Cell & Dev. Biology, Biological Sciences |
| 2002 – 2006 | Founder and Director, AHC Clinic, Vanderbilt Children’s Hospital |

C. Contribution to Science

1. Animal Models of Tuberous Sclerosis Complex

My career focus was initiated during a post-doctoral fellowship in the lab of Dr. David Gutmann at Washington University in St. Louis. In my own lab, it has since encompassed the generation of various mouse as well as zebrafish models of TSC. We have tested various hypotheses about the contribution of specific lineages of the brain, that when deficient in either the *Tsc1* or *Tsc2* genes, may underlie neurologic disease.

Relevant peer-reviewed publications:

1. Kim, S. H., Speirs, C. K., Solnica-Krezel, L. and **Ess, KC**. Zebrafish model of tuberous sclerosis complex reveals cell-autonomous and non-cell-autonomous functions of mutant tuberin. *Disease Models Mechanisms* 4: 255-67 (2011). PMID: 20959633 PMCID: PMC3046101
2. Fu C, Cawthon C, Clinkscales W, Bruce A, Winzenburger P and **Ess, KC**. GABAergic Interneuron Development and Function is Modulated by the *Tsc1* Gene. *Cerebral Cortex* 22: 2111-2119 (2011). PMID: 22021912 PMCID: PMC3412444
3. Carson R, Van Nielen D, Winzenburger P and **Ess, KC**. Neuronal and glia abnormalities in *Tsc1*-deficient forebrain and partial rescue by rapamycin. *Neurobiology of Disease* 45: 369-80 (2012). PMID: 21907282 PMCID: PMC3225598
4. Robert Carson, Nathaniel Kelm, Kathryn West, Mark Does, Mark, Cary Fu, Grace Weaver, Eleanor McBrier, Mark Grier, and **Ess KC**, Hypomyelination following deletion of *Tsc2* in oligodendrocyte precursors". *Annals Clinical and Translational Neurology*. 58(3):440-52 (2015). PMID:25818646

2. Models of human diseases using induced pluripotent stem cells (iPSCs)

Over the past several years, the laboratory efforts have included generation and characterization of human iPSCs. These are typically made from patients at Vanderbilt who have neurological disorders with a known genetic etiology such as TSC and other mTOR related disorders.

Relevant peer-reviewed publications:

1. Neely MD, Litt M, Tidball A, Li G, Aboud A, Hopkins C, Chamberlin R, Hong C, **Ess, KC**, and Bowman A. DMH1, a highly selective small molecule BMP inhibitor promotes neurogenesis of hiPCSs: Comparison of PAX6 and SOX1 Expression during Neural Induction. *ACS Chemical Neuroscience* 3:482-491 (2012) PMID: 22860217 PMCID: PMC3400384
2. Asad A Aboud; Andrew M Tidball; Kevin K Kumar; M D Neely; Bingying Han; **Ess, KC**; Charles C Hong; Keith M Erikson; Peter Hedera. PARK2 patient neuroprogenitors show increased mitochondrial sensitivity to copper. *Neurobiol Dis.* (2014) Oct 12;73C:204-212. doi: 10.1016/j.nbd.2014.10.002. PMID: 25315681
3. Thomas LR, Wang Q, Grieb BC, Phan J, Foshage AM, Sun Q, Olejniczak ET, Clark T, Dey S, Lorey S,

Alicie B, Howard GC, Cawthon B, **Ess KC**, Eischen CM, Zhao Z, Fesik SW, Tansey WP. Interaction with WDR5 Promotes Target Gene Recognition and Tumorigenesis by MYC. *Mol Cell*. (2015) Mar 25. pii: S1097-2765(15)00142-2. PMID:25818646

4. Tidball AM, Bryan MR, Uhouse MA, Kumar KK, Aboud AA, Feist JE, **Ess KC**, Neely MD, Aschner M, Bowman AB. A novel manganese-dependent ATM-p53 signaling pathway is selectively impaired in patient-based neuroprogenitor and murine striatal models of Huntington's disease. *Hum Mol Genet*. (2015) Apr 1;24(7):1929-44. PMID: 25489053

3. Non-TSC Models of human diseases

Our research focus on TSC has led to collaborations with other investigators as TSC/mTOR signaling pathway and the reagents we created (mouse, zebrafish, iPSCs) have broad connections to many fields of research and medicine.

Relevant peer-reviewed publications:

1. Aboud, A, Tidball, A, Kumar, K, Neely, D, **Ess KC**, Erikson, KM and Bowman, AB, “Genetic risk for Parkinson's disease correlates with alterations in neuronal manganese sensitivity between two human subjects” *Neurotoxicology* 33:1443-49 (2012). PMID: 23099318
2. Boglev, Y, Badrock, P, Trotter, A, Du, Q, Richardson, E, Parslow, A, Markmiller, S, Hall, N, de Jong-Curtain, T, Ng, A, Verkade, H, Ober, E, Field, H, Shin, D, Shin, C, Hannan, K, Hannan, R, Pearson, R, Kim, S-H, **Ess, KC**, Lieschke, G, Stainier, D and Joan K. Heath, “Autophagy Induction is a Tor and Tp53-independent Cell Survival Response in a Zebrafish Model of Disrupted Ribosome Biogenesis”. *PLoS Genetics*. doi: 10.1371/journal.pgen.1003279. (2013). PMID:23408911
3. Kim, SH, Scott, SA, Bennett, MJ, Carson, RP, Brown, HA, and **Ess, KC**. “Multi-organ Abnormalities and mTORC1 Activation in Zebrafish Model of Multiple Acyl-CoA Dehydrogenase Deficiency *PLoS Genetics* (2013 Jun;9(6) PMID: 23785301
4. Chun YW, Balikov DA, Feaster TK, Williams CH, Sheng CC, Lee JB, Boire TC, Neely MD, Bellan LM, **Ess KC**, Bowman AB, Sung HJ, Hong CC. Combinatorial polymer matrices enhance in vitro maturation of human induced pluripotent stem cell-derived cardiomyocytes. *Biomaterials*. 2015 Oct;67:52-64. PMID: 26204225.

Complete list of published work is in:

<http://www.ncbi.nlm.nih.gov/myncbi/browse/collection/41636033/?sort=date&direction=ascending>

Session 5-4

Modeling Alternating Hemiplegia of Childhood in Induced Pluripotent Stem Cells

Alfred L. George, Jr,
Northwestern University, USA

Alternating hemiplegia of childhood (AHC) is one notable phenotype among a heterogeneous spectrum of disorders associated with mutations in *ATP1A3*. We have investigated the electrophysiological properties of human neurons generated from AHC patient-specific induced pluripotent stem cells (iPSCs) to ascertain functional disturbances underlying this neurological disease. We will present data collected on neurons differentiated from two subjects with the common mutation G947R in which we succeeded in measuring ouabain-sensitive electrogenic outward current ('pump current'), resting membrane potential and evoked action potential firing behavior. We will also present data demonstrating the consequences of a splice site mutation (c.2542T>C) associated with AHC. Our findings are consistent with impaired ion transport function of neuronal Na⁺/K⁺-ATPase but highlight uncertainties regarding the importance of haploinsufficiency versus dominant-negative mechanisms. The iPSC-neuron model offers opportunities to investigate the pathogenesis of *ATP1A3*-associated disorders and serve as a cellular platform for testing therapeutic strategies.

Alfred L. GEORGE, Jr.



• Present Position

Magerstadt Professor and Chairman, Department of Pharmacology,
Director, Center for Pharmacogenomics, Northwestern University
Feinberg School of Medicine, Chicago IL USA

• Education:

1982 M.D., University of Rochester School of Medicine, Rochester NY USA
1982-86 Internship and Residency in Medicine, Vanderbilt University, Nashville TN USA
1986-87 Clinical Fellowship, Department of Medicine, University of Pennsylvania
1987-88 Visiting Postdoctoral Fellow, Institut Suisse de Recherches
Experimentales sur le Cancer, Lausanne, Switzerland
1988-92 Research Fellow, Department of Medicine, and Department of Biochemistry and Biophysics,
University of Pennsylvania, Philadelphia
1992-95 Assistant Professor, Department of Medicine and Pharmacology, Vanderbilt University
1995-98 Associate Professor of Medicine and Pharmacology, Vanderbilt University
1999-2014 Professor of Medicine and Pharmacology, Vanderbilt University
1999-2014 Founder and Chief, Division of Genetic Medicine, Vanderbilt University
2014-present Magerstadt Professor and Chairman, Department of Pharmacology,
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• Representative Publications

George, A.L., Jr., Crackower, M.A., Abdalla, J.A., Hudson, A.J., and Ebers, G.C. Molecular basis of Thomsen's disease (autosomal dominant myotonia congenita). *Nature Genetics* 1993;3:305-310.

Bennett, P.B., Jr., Yazawa, K., Makita, N., and **George, A.L., Jr.** Molecular mechanism of an inherited cardiac arrhythmia. *Nature*, 1995;367:683-685.

Lossin, C., Wang, D.W., Rhodes, T.H., Vanoye, C.G., and **George, A.L., Jr.** Molecular basis for an inherited epilepsy. *Neuron*, 2002;34:877-884.

Crotti, L., Johnson, C.N., Graf, E., De Ferrari, G.M., Cuneo, B.F., Ovadia, M., Papagiannis, J., Feldkamp, M.D., Rathi, S.G., Kunic, J.D., Pedrazzini, M., Wieland, T., Lichtner, P., Beckmann, B., Clark, T., Shaffer, C., Benson, D.W., Kääh, S., Meitinger, T., Strom, T.M., Chazin, W.J., Schwartz, P.J., and **George, A.L., Jr.** Calmodulin mutations associated with recurrent cardiac arrest in infants. *Circulation*, 2013;127:1009-17.

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Poster Presentations

| No. | 1st Author | Title |
|------|--------------|---|
| P- 1 | Y Nakamura | A de novo p.Arg756Cys mutation in <i>ATP1A3</i> causes a distinctive phenotype with prolonged paralysis and involuntary movements triggered by fever |
| P- 2 | L Kremer | <i>De novo</i> mutations in <i>ATP1A3</i> as differential diagnosis for suspected mitochondrial disease patients |
| P- 3 | M Kubota | A 20-year-old male with the right-dominant dystonia and bulbar palsy associated with <i>ATP1A3</i> gene mutation |
| P- 4 | M Soleimani | De novo missense variants in <i>ATP1A3</i> with Alternating Hemiplegia of Childhood associated with childhood-onset schizophrenia: Report of 2 cases |
| P- 5 | F Gurrieri | Unusual clinical presentation of familial hemiplegic attacks in two first cousins: possible oligogenic inheritance of AHC-related phenotypes? |
| P- 6 | F Nery | Pathologic Findings in Alternating Hemiplegia of Childhood (AHC) Patients |
| P- 7 | E De Grandis | Alternating Hemiplegia of Childhood: a magnetic resonance spectroscopy, tractography and functional 3T study exploring ictal and interictal disease mechanisms |
| P- 8 | A Vilamala | EEG Biofeedback for Relaxation using Deep Neural Networks |
| P- 9 | N Fedosova | Application of EPR for description of the ouabain-binding site of Na,K-ATPase |
| P-10 | R Ragno | Towards the construction of a Structure-based protocol to select potential <i>ATP1A3</i> ligands through virtual screening procedures |
| P-11 | K Ikeda | Perinatal seizure and defective respiratory rhythm generation in <i>Atp1a3</i> knockout homozygous knockout mice |
| P-12 | FD Tiziano | Characterization of a human neuroblastoma model of AHC: progress report |
| P-13 | C Simmons | <i>ATP1A3</i> Splice Site Mutation Produces Haploinsufficiency in Alternating Hemiplegia of Childhood |
| P-14 | J Snow | An iPSC-Derived Neuronal Model for Determining the Pathophysiology of <i>ATP1A3</i> Mutations in Alternating Hemiplegia of Childhood |
| P-15 | E Akkuratov | Mice model of rapid-onset dystonia-parkinsonism |

P-1

A de novo p.Arg756Cys mutation in *ATP1A3* causes a distinctive phenotype with prolonged paralysis and involuntary movements triggered by fever

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Recently, p.Arg756Cys mutation in *ATP1A3* has been proven to cause a unique form of *ATP1A3*-related disorders, characterized by prolonged paralysis and involuntary movements triggered by febrile illness. Dard R, et al. expressed this phenotype using a term relapsing encephalopathy with cerebellar ataxia (RECA). We herein report an additional patient with a p.Arg756Cys mutation as the fourth case.

The patient is a 4-year-old boy, who has shown from 9 months of age recurrent paroxysmal prolonged motor functional deterioration triggered by febrile illness, followed by gradual recovering. He showed developmental delay, being able to control his head at 5 months, and sat unassisted at 10 months. He began to show movement disorders and ataxia at 1 year 3 months. Whole-exome sequencing detected a de novo p.Arg756Cys mutation.

The current patient shared the common core symptoms with the previously reported patients of RECA carrying a p.Arg756Cys mutation, confirming the distinctive clinical entity that results from the specific missense mutation.

P-2

***De novo* mutations in *ATPIA3* as differential diagnosis for suspected mitochondrial disease patients**

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Mitochondrial disorders are the most common group of inborn errors of metabolism affecting 1 in 5,000 live births. They present with a large variety of symptoms and are generally suspected upon a severe disease course, often involving multiple tissues. For childhood-onset mitochondrial disease, most patients present with neurological symptoms. While metabolic markers as elevated lactate and biochemical findings like impairment of respiratory chain complex deficiencies can further strengthen the assumption of a mitochondrial disorder, these indicators are unspecific and differential diagnosis needs to be considered. WES on 540 suspected mitochondrial disease patients identified a *de novo* mutation in *ATPIA3* in two unrelated patients. Both patients presented with a strikingly similar phenotype including childhood-onset ataxia, epilepsy, dysarthria, hearing impairment and encephalopathic crisis upon febrile illness. While for childhood-onset mitochondrial disorders a recessive mode of inheritance is suspected, *de novo* mutations are often not investigated. The examples show that *ATPIA3*-related disease mimics mitochondrial disorders and are therefore likely to be missed in molecular diagnostics. It also shows that a screen for *de novo* mutations should always be considered in suspected mitochondrial disease patients.

A 20-year-old male with the right-dominant dystonia and bulbar palsy associated with *ATP1A3* gene mutation

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Rapid-onset dystonia-parkinsonism (RDP) is a rare autosomal dominant movement disorder with variable clinical course caused by a mutation of *ATP1A3* (Na⁺/K⁺ -ATPase α 3 subunit gene). We here report the first Japanese case with RDP.

This right-handed 20-year-old male had noticed writing difficulty and right-sided facial myoclonus at 13 years of age. Later, dystonic posture and twisting movement of right-sided extremities progressed gradually, resulted in gait disturbance. Because he received MR vaccination 1 month before the onset, he suspected of immune-mediated encephalitis, but IVIG therapy failed to recover. One and a half years later from onset, swallowing difficulty and dysarthria became apparent. Trihexyphenidyl, L-dopa, clonazepam, carbamazepine, and baclofen were not effective for his movement disorders. He showed rigidity in extremities, but no tremor. There was no spasticity or diurnal fluctuation of his dystonia. There was no abnormal findings in brain MRI and SPECT. There was no abnormality in CSF-HVA, 5-HIAA, bipterin and neopterin. After the introduction of botulinum toxin i.m. to right-sided extremities at 15 years of age, his symptoms were stabilized, although severe dystonic features remained. The next-generation sequencing revealed *de novo* *ATP1A3* mutation (c.2477C>T, p.Ala826Val).

Although RDP is characterized by sudden onset of asymmetrical dystonia and parkinsonism, our patient's symptoms progressed slowly and later stabilized. Currently he is engaged in light work with job support. As Brashear et al. (1996) already reported, the clinical course of RDP may be quite variable.

P-4

De novo missense variants in *ATP1A3* with Alternating Hemiplegia of Childhood associated with childhood-onset schizophrenia: Report of 2 cases

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Childhood-Onset Schizophrenia (COS) is a rare (prevalence 1/40 000) but major mental disorder characterized by a wide spectrum of symptoms, including delusions, hallucinations, disorganisation of speech, negative symptoms, and cognitive deficits. Several publications have shown that some polymorphisms are associated with COS. Identification of rare variants is just beginning, regarding CNV and de novo mutations.

Here we report on two COS cases associated with Alternating Hemiplegia of childhood (AHC), a rare disease characterised by repeated episodes of hemiplegia that alternately affects one side of the body. These comorbidities have guided the genetic exploration and two de novo deleterious missense variants (c.2438.T>C and c.2443G>A) in *ATP1A3* were identified.

Both patients exhibited early (<13 years old) and severe psychotic signs with high scores on the standard clinical scales. AHC diagnosis was made at an early age for both patients (< 20 months old). Treatment response to antipsychotics was poor as it is often the case for COS.

To our knowledge, only one case report in a young boy is available, these two cases perfectly underlines the need to seek for all possible genetic conditions in children and adolescents presenting atypical psychotic symptoms and in a clinical point of view, on the importance of an early identification of psychiatric symptoms and disorders in rare organic disease like AHC.

Keywords: psychosis, genetic, mutation, early-onset schizophrenia, hemiplegia

P-5

Unusual clinical presentation of familial hemiplegic attacks in two first cousins: possible oligogenic inheritance of AHC-related phenotypes?

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We report on two first cousins presenting with different clinical manifestations within the AHC phenotypic spectrum. The first patient is a 30 y/o female who shows typical features of classical AHC, including early onset hemiplegic attacks, intellectual disability, movement disorders, and seizures. *ATP1A3* molecular analysis tested negative, whereas she bears an apparently *de novo* variant in a novel gene which is a strong candidate for AHC. Her first cousin (daughter of a maternal uncle) is a female of similar age who was healthy until 25 years, when she experienced the onset of severe migraine, suddenly followed by a persisting hemiplegic episode, lasting a few months. Brain MRI was normal. After the recovery from the first hemiplegic attack, she remained healthy for some years. She has recently had a similar episode of migraine followed by hemiplegia. *ATP1A3* as well as the hemiplegic migraine genes (*ATP1A2*, *CACNA1A*, *SCN1A* and *SLC2A1*) tested negative.

To understand the unusual presentation of a similar phenotype in the two cousins, for which a potential shared pathogenic variant would segregate through at least three unaffected subjects, we have undertaken the study of their whole exome (currently under bioinformatic analysis). A simple autosomal dominant model seems unlikely, unless we take into account a high lack of penetrance: rather, we hypothesize that the two cousins share a single or a few variants that alone are not sufficient to induce the pathological phenotype. Additional inherited or *de novo* variants could explain the low penetrance and the marked phenotypic discrepancies between the two patients.

An update of the ongoing genetic studies will be presented.

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Pathologic Findings in Alternating Hemiplegia of Childhood (AHC) Patients

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Alternating hemiplegia of childhood (AHC) is a rare and severe neurodevelopmental syndrome characterized by recurrent hemiplegic episodes and distinct neurological manifestations. The majority of individuals with AHC have a de novo mutation in *ATPase alpha subunit 3 (ATPIA3)* gene. Controversy remains as to whether patients with AHC develop consistent structural neuropathological changes that may contribute to disease progression or the acquisition of fixed neurologic deficits over time.

In this present study, detailed autopsies were performed, including the brains, on three AHC affected subjects carrying mutations in the *ATPIA3* gene, including Exon 17 c.2401G>A; p. D801N, Exon 18 c.2423C>T; p. P808L, and Exon 18 c.2443G>A; p. E815K. Pulmonary edema or congestion was present in all three, with scattered foci of dystrophic calcification present in the lung and kidneys of the patient with E815K mutation. Significant cardiac abnormalities were present in 2 of the 3 patients including thickened ventricular walls and endomyocardium. Examination of brain tissue from these three subjects revealed consistent cerebellar atrophy with granular and Purkinje cell loss, gliosis, and patchy disorganization of the cortical layer. Some Purkinje cells were found in the deeper white matter in two subjects with the P808L and E815 mutations—suggesting possible abnormal neuronal migration and development. The subject with D801N mutation additionally had diffuse acute and chronic ischemic neuronal injury, with shrunken pyknotic neurons and significant cortical gliosis. The subject with the P808L mutation had mild vacuolization in the cerebral white matter, cortex and lateral thalamus, pigmented macrophages in perivascular spaces and rare binucleated neurons in the cortex and hippocampus suggesting subtle developmental abnormalities. The visual cortex had very large and disorganized neurons in the third layer. The subject with the E815K mutation had moderate gliosis of the dentate nucleus, hippocampal sclerosis and astrogliosis, and microglial nodules throughout the spinal cord.

In aggregate, our findings revealed severe cerebellar atrophy in AHC subjects, predominantly in the superior vermis; mild atrophy of hippocampus, predominantly in the dentate gyrus; and diffuse gliosis throughout the brain. Curiously, individuals affected with Rapid-onset Dystonia-Parkinsonism (RDP) who carry different *ATPIA3* gene mutations also have mild cerebellar and hippocampus atrophy. However, in contrast to RDP, we did not observe atrophy of the amygdala or substantia nigra. More studies are needed to confirm the unique neuropathologic findings related to brain maldevelopment in these well-defined AHC subjects by comparing them to age- and sex-matched controls. Such pathologic data are essential to underscore the clinical studies and further our understanding of the pathogenesis of AHC.

P-7

Alternating Hemiplegia of Childhood: a magnetic resonance spectroscopy, tractography and functional 3T study exploring ictal and interictal disease mechanisms

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Background: Alternating Hemiplegia of Childhood (AHC) is a very severe and intractable disorder, characterized by a combination of paroxysmal movement disorders and permanent neurological features. Despite the recent discovery of *ATPIA3* mutations in 75% of cases, there is little knowledge about the disease pathophysiology and no proven effective therapeutic options. Moreover, although the use of Magnetic Resonance (MR) techniques in neurological diseases has rapidly grown, long-term MR imaging studies and spectroscopy, tractography and functional studies are lacking in this group of patients.

Aim of the study: Aim of the project is to explore brain pathways and metabolic changes involved in AHC and to describe long-term brain morphological alterations in older patients using a 3T MR unit.

Methods: 8 genetically confirmed AHC patients (6 males, 2 females, age range 10 – 25 years) recruited through the Italian Biobank and Clinical Registry for AHC (www.ibahc.org) underwent MRI with spectroscopy, tractography and functional resting state study. Findings have been collected and reviewed by 2 expert neuroradiologists.

Results: Cerebellar atrophy has been shown in 5/8 patients. Spectroscopy showed slightly reduced cerebellar N-acetylaspartate, used as a potential indicator of neuronal injury/loss, in 6 out of 7 patients. Six patients completed the functional MRI resting state study. Comparison of functional MRI data with normal controls is in progress.

In AHC, multiple stress factors may lead to paroxysmal and permanent movement disorders. In particular, it could be hypothesized a relationship between the role of brainstem and cerebellum in sensory motor and cognitive processing and the site distribution of defective Na-K ATPase pump in AHC disease. Functional 3T MRI analysis will better define the involvement of cerebellum and other brain structures in disease pathophysiology and will give the rationale for the use of newer and more efficacious molecules.

EEG Biofeedback for Relaxation using Deep Neural Networks

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Background: Alternating Hemiplegia of Childhood (AHC) is a neurodevelopmental condition characterised for transient episodes of partial or total body paralysis. The attacks are usually relieved by relaxation or sleep. EEG biofeedback is a non-invasive technique that has been successfully used for driving subjects to relaxation states, by increasing their measured EEG alpha band power.

Methods: We are currently developing a data processing pipeline to analyse EEG spectrograms using successful artificial intelligence techniques for vision, known as deep neural networks. These networks can extract more informative representations of the subject's cognitive processes than simple alpha band power, providing better feedback to the subject. The system relies on commercial EEG caps connected to portable devices, where the system is deployed. This setting allows its use not only in medical environments, but also at patient's home, making it suitable for regular use.

Results: Preliminary results on sleep stage scoring demonstrate the suitability of the proposed method to analyse EEG spectrograms, obtaining better results as compared to those systems using discrete band power.

Conclusions: We think that this tool might help AHC patients to relieve their hemiplegic episodes by teaching them to bring their emotions to relaxation. Study on this subset of the population is the purpose of future research work.

Acknowledgements: This project has received funding from the European Union's Horizon 2020 research and innovation programme under the Marie Skłodowska-Curie grant agreement No 659860.

P-9

Application of EPR for description of the ouabain-binding site of Na,K-ATPase

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In the long run, the project aims at development of new derivatives of cardiotonic steroids (CTS) with increased selectivity towards individual isoforms of the Na,K-ATPase. The Na,K-ATPase is a target for a number of pharmacological agents from the CTS family, but, due to the enzyme's ubiquity, the inhibitors in use have many adverse effects. A diminished systemic toxicity of the compounds may be acquired by an increase in selectivity towards individual enzyme isoforms. Since the isoform-related differences are found within the cavity leading to the CTS-binding site, we shall supplement the existing description of crystallized enzyme-CTS complexes with information valid under physiological conditions. For this purpose, we shall collect Electron Paramagnetic Resonance (EPR) spectra of our newly synthesized spin-labelled CTS with varying spacer arms specifically bound to the enzyme and correlate the mobility of the label to the distance between the label and the steroid core of CTS. The degree of the motional restriction of the spin label reports on the radius of the cavity at the chosen distance from the steroid core. At present, we screen a number of CTS derivatives for their binding properties to Na,K-ATPase in order to find a core molecule with largest specific signal. We also have several reporting nitroxide groups with different sensitivities to conformational re-arrangements within or in the vicinity of the CTS binding site. Mapping of the CTS-binding cavity and marking out its boundaries under physiological conditions will define the spatial position of the sugar units (known to improve affinity and contribute to isoform selectivity). The preliminary results will be presented.

P-10

Towards the construction of a Structure-based protocol to select potential *ATPIA3* ligands through virtual screening procedures

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The maintenance of the proper Na⁺ gradient across the plasma membrane is essential for the function and survival of all eukaryotic cells. Ion gradients are maintained by the Na,K-ATPases of which alpha α subunit carries out the ion transport and ATP hydrolysis.

Mutations in ion-transporting proteins severely affect their ability to traffic ions properly, perturbing the delicate ion gradient balance.

ATPIA3 encodes for an alpha subunit that contributes to the maintenance of Na⁺ and K⁺ concentrations across neuron membranes. *ATPIA3* mutations have been indicated as responsible for the vast majority of alternating hemiplegia of childhood (AHC) cases and other related conditions, including DYT12, CAOS and CAPOS.

Due to the lack of any structural data, we have applied homology modeling techniques to build three-dimensional structures of wild type (wt) and mutated *ATPIA3* forms. Starting from the primary sequence of the wt *ATPIA3* protein, and those bearing D801Y and E815K variants, we generated several homology models that were structurally optimized through molecular mechanics.

Molecular dynamics (MD) techniques have been applied to inspect, at the atomic level, how each point mutation could affect the ion flow through *ATPIA3*. Details of the MD runs and structural insights for future drug design protocol will be reported.

This work was supported by AISEA.

Perinatal seizure and defective respiratory rhythm generation in *Atp1a3* knockout homozygous knockout mice

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A neuron-specific human $\alpha 3$ subunit isoform of the sodium pump, *ATP1A3*, plays an important role in neuronal excitability. Its point and deletion mutations have been recognized in diverse neurological disorders, such as alternating hemiplegia of childhood (AHC), apnea, and severe infantile epileptic encephalopathy often appear shortly after birth. To get insight into the pathophysiology of the disorders and to understand the functional roles of the sodium pump $\alpha 3$ subunit in the brain *in vivo* during this period of development, we examined the phenotype of *Atp1a3* knockout homozygous mouse fetuses (*Atp1a3*^{-/-}). Immediately after extraction from the uterus and release from the placenta, about half of *Atp1a3*^{-/-} showed small spontaneous movements such as mouth opening, responded to pinching with slight body movement, and made an effort to breath. Within several seconds to several minutes, they started to develop atypical seizure with extension of both upper limbs anteroinferiorly and with tongue protrusion. Triggers of seizure seemed to be the body movements such as response to pinching stimulation or the breathing effort they made themselves. The epileptic seizures usually continued for several minutes. During and after the seizure attack, the *Atp1a3*^{-/-} did not respond to pinching, show spontaneous body movements, or breathe on their own. The other *Atp1a3*^{-/-} fetuses showed complete absence of spontaneous body movements, no responses to pinching, and no breathing movements from the very beginning at birth. To investigate the locus of hyperexcitability associated with seizure, we examined c-Fos expression in the brains of *Atp1a3*^{-/-} and found a significantly increased number of c-Fos-expressing cells in various regions of the brains, with unique distribution in the cerebellum, when compared with wild-type littermates (*Atp1a3*^{+/+}). We also found various abnormal respiratory rhythms produced in the brainstem of *Atp1a3*^{-/-}. The data suggest that *Atp1a3* plays a critical role in neural function during development, survival at perinatal period, and its deficiency causes hyperexcitability in the fetal brain.

Reference Ikeda, K et al. *Brain Research*, in press, 2017

P-12

Characterization of a human neuroblastoma model of AHC: progress report

Tiziano FD¹, Piacentini R.², Novelli A.¹, Diano F.¹, Rinelli M.¹, Cocco S.², Di Pietro L.¹, Ripoli C.², Neri G.¹, Grassi C.², Ragno R.³, Gurrieri F.¹

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We are currently refining the characterization of a cellular model of AHC, based on SH-SY5Y neuroblastoma. SH-SY5Y have been stably transfected with constructs expressing the wild type form or four different *ATP1A3* variants: E815K, D801N, D801Y, G947R. The rationale behind the choice of this model is related to the fact that SH-SY5H cells display a neuronal-like phenotype and express the endogenous *ATP1A3* gene.

ATP1A3 cDNAs have been cloned into the pIRES-eGFP eukaryotic expression vector. After linearization, cell lines have been permanently transfected with the constructs. The electrophysiological characterization has shown in E815K cells an accumulation of Na⁺ and Ca²⁺, as well as the reduction of the membrane resting potential.

The differentiation protocol of naïve SH-SY5Y cells with retinoic acid (RA) followed by neurobasal medium /B27 supplement (NB/B27) led to the development of neuron-like cells showing Ca²⁺ transients, trains of action potentials, and the expression of MAP2 and NeuN neuronal markers. To characterize the neuronal lineage that we are obtaining upon our differentiation protocol we are currently performing the whole transcriptome analysis of undifferentiated and differentiated (with RA only or with RA-NB/B27) cell lines.

Upon differentiation protocol, cell lines bearing *ATP1A3* mutations started dying two days after the medium switch from RA to NB/B27. By immunocytofluorescence, we demonstrated hyperexpression of activated Caspase 2 in AHC-like cells, suggesting the activation of apoptosis. We are currently planning to perform the transcriptome analysis of mutated cells in order to identify deregulated genes due to the presence of mutated *ATP1A3*.

Further, we are starting a medium throughput screening of candidate compounds to the treatment of AHC, identified by virtual screening of molecules binding to the mutated *ATP1A3* alleles. Candidate compounds will be tested for their effect in: 1) promoting Na⁺ and Ca²⁺ scavenging and 2) revert the death phenotype.

This work was supported by AISEA.

***ATPIA3* Splice Site Mutation Produces Haploinsufficiency in Alternating Hemiplegia of Childhood**

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AHC is most often associated with heterozygous *ATPIA3* missense mutations. There are no reports of overt loss-of-function mutations such as truncating frameshifts or premature stop codons associated with AHC. However, there have been cases associated with intronic splice site variants.

We investigated the effects of a recurrent intronic splice site mutation (c.2542+2T>C), which disrupts the canonical GT dinucleotide at the start of intron 18, on *ATPIA3* mRNA splicing in human neurons generated from patient-specific induced pluripotent stem cells (iPSCs). We differentiated cortical excitatory neurons from iPSCs from a proband with a clinical history of seizures and alternating hemiplegia presenting in early childhood and a healthy sex-matched control subject. We performed reverse transcription-polymerase chain reaction (RT-PCR) on 25 day post differentiation neurons using primers embedded in exons 17 and 20. We observed a single band of expected size in RT-PCR reactions performed on control neurons but two distinct amplicons in reactions from RNA isolated from AHC iPSC-derived neurons. Sequencing demonstrated that the mutation causes mis-splicing of *ATPIA3* mRNA with an in-frame inclusion of 66 bp from the adjacent intron and a predicted 22 amino acid insertion in a transmembrane domain of the protein. This prediction was consistent with immunoblot experiments performed on HEK-293 cells overexpressing recombinant *ATPIA3* engineered with the 22 amino acid insertion, which demonstrated a higher molecular mass than WT protein. Structural modeling indicated that the insertion would affect the ion binding site as well as interactions with the lipid bilayer and beta-subunit thus destabilizing the protein. Immunoblotting of differentiated neurons showed lower amounts of the normal mass *ATPIA3* protein with no larger products suggesting that the mutant protein is likely degraded in neurons.

Our findings indicate that a splice site mutation promotes mis-splicing and *ATPIA3* haploinsufficiency in neurons. Electrophysiology studies are underway to assess the cellular consequences associated with this mutation.

An iPSC-Derived Neuronal Model for Determining the Pathophysiology of *ATPIA3* Mutations in Alternating Hemiplegia of Childhood

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Vanderbilt University Department of Pediatrics and Department of Cell and Developmental Biology

Alternating Hemiplegia of Childhood (AHC) is a rare genetic neurodevelopmental disease caused by heterozygous missense mutations in the *ATPIA3* gene. *ATPIA3* encodes the neuronal specific $\alpha 3$ subunit of the sodium-potassium adenosine triphosphatase (Na,K-ATPase) pump. This protein complex is crucial for many cellular functions including setting the resting membrane potential, regulating ion homeostasis, and generating action potentials. AHC patients display unique symptoms beginning in early childhood, including episodes of weakness or paralysis often triggered by stress, abnormal eye movements, seizures, painful dystonia, developmental delay, and intellectual disability. It is now known that the majority of these individuals possess one of three heterozygous missense mutations in the *ATPIA3* gene, resulting in D801N, E815K, or G947R mutants of the $\alpha 3$ Na,K-ATPase subunit. However, mechanisms underlying symptoms in patients remain poorly understood and there are no empirically proven treatments for AHC.

The intricate relationships of the *ATPIA3* genotype and AHC phenotype make induced pluripotent stem cell (iPSC) technologies ideal for the investigation of disease mechanisms via creation of patient-specific disease models.

For our studies, we have generated iPSCs from patients with the three most common mutations in AHC. These iPSCs are differentiated into cortical neurons to test our hypothesis that mutant $\alpha 3$ decreases Na,K-ATPase function in a dominant-negative manner. We further hypothesize that mutant $\alpha 3$ protein perturbs normal neurodevelopment and results in neuronal depolarization that is exacerbated during cellular stress. Initial studies have focused on characterizing Na,K-ATPase subunit expression in developing cortical neurons. Results indicate that neurons derived from all three major *ATPIA3* mutations have increased levels of $\alpha 3$ protein and decreased levels of the $\alpha 1$ subunit compared to iPSCs. Expression during neuronal differentiation has similar temporal dynamics to control cells. Ongoing studies are exploring the neurodevelopmental consequences of mutant $\alpha 3$ protein by analyzing synaptic formation, neuronal maturation, and lineage specification. Fluorescent indicators of sodium and calcium concentration in patient-specific iPSC-derived cortical neurons are being used to investigate ion homeostasis in the presence of *ATPIA3* mutations under both physiological and stressed conditions.

Future studies will involve single-cell and multi-electrode electrophysiological analyses, along with testing for lineage-specific consequences of mutant $\alpha 3$ expression in GABAergic iPSC-derived neurons.

Finally, we are increasing the potential impact of our findings by using CRISPR/Cas9 techniques to create isogenic controls by correcting *ATPIA3* mutations in AHC patient-derived iPSC lines.

This research was generously supported by the Alternating Hemiplegia of Childhood Foundation as well as the Association Française de l'Hémiplégie Alternante.

Mice model of rapid-onset dystonia-parkinsonism

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Rapid-onset dystonia-parkinsonism (RDP) is the first identified disease manifestation of ATP1A3 mutation. It is characterized by abrupt onset of several symptoms, including generalized dystonia, severe bradykinesia and gait instability. The most common mutation observed in RDP patients, T613M, is located close to the nucleotide-binding domain. We have recently created a knock-in mouse model of T613M ATP1A3 using CRISPR/Cas technology. Homozygous mice are lethal. Heterozygous mice are born normally, but weigh less and are more hyperactive than their wild type (WT) littermates. During elevated plus-maze test heterozygous mice spend twice as much time in open space as WT. Mice memory has been tested with passive avoidance test and yielded similar results in heterozygous and WT mice. We observed seizures during ketamine anesthesia in heterozygous but not in WT mice. Recent studies have indicated that the neuronal sodium pump plays an important role for control of motor activity and gait. We have identified two heterozygous mice, that had obvious problems moving their hind legs and unable to walk normally. Thus we have in these ongoing studies of heterozygous ATP1A3 mice found several of the phenotypic characteristics observed in patients with ATP1A3 mutations.

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An Introduction of the Illustrator



Ms. Miya Tamura. She was born in Japan in 1969. She suffered from repeated hemiplegic attacks since she was 8 months of age. Although she went to several hospitals, she had not been diagnosed for 14 years. Finally, she was diagnosed as having Alternating Hemiplegia of Childhood in 1984 at National Center of Neurology and Psychiatry (NCNP) by Dr. Norio Sakuragawa. Her *ATPIA3* mutation was confirmed as p.D801N through gene analysis in 2012. She has fun in drawing pictures of animals with crayons.





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