



2024 JSPS-NIH Forum

March 1, 2024

Japan Society for the Promotion of Science Fogarty International Center, National Institutes of Health



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DATE 3:00 pm - 7:30 pm March 1, 2024 (EST) *[Japan Standard Time 日本時間] 5:00 am - 9:30 am March 2, 2024

VENUE Hybrid (The Lawton Chiles International House (Building 16)) In-person Registration (Invitation Only) Online Registration from here

*Time and Format are subject to change based on speakers and regulation on COVID-19

3:00pm - 3:25pm (EST)	Opening Remarks
5:00am - 5:25am (JST)	Dr. Junji URAKAWA, Director, JSPS Washington Office
	Mr. Koji ARIBAYASHI, Science Counselor, Embassy of Japan in the USA
	Dr. Yoh-suke MUKOYAMA, Chair of Review Panel / Senior Investigator, NHLBI,
	NIH
	Dr. Michael GOTTESMAN, Chief, Laboratory of Cell Biology, NCI/CCR, NIH

Lecture Session 1 (Speaker from Japan)	Theme: Neuroscience
3:25pm – 4:10pm (EST) 5:25am – 6:10am (JST)	Invited Speaker (1) 45 min. including Q & A Dr. Motoyuki ITOH , Professor, Graduate School of Pharmaceutical Sciences, Chiba University Lecture title: Investigating Age-Induced Alterations in Brain Function and Regeneration Using Zebrafish Model
4:10pm - 4:20pm (EST) 6:10am - 6:20am (JST)	Break
Lecture Session 2 (Speakers from NIH)	
4:20pm - 5:00pm (EST) 6:20am -7:00am (JST)	Invited Speaker (2) 20 min. including Q & A Dr. Yuko YASUI , (NIH/NIDA) [E] Lecture title: Sigma-1 receptor plays a critical role in cocaine-induced synaptic AMPA receptor plasticity in the VTA dopamine neurons Invited Speaker (3) 20 min. including Q & A Dr. Yuta KOUI , (NIH/NHLBI) [F] Lecture title: NGF-TrkA-PI3K signaling promotes sensory hypersensitivity in diabetes
5:00pm - 5:10pm (EST) 7:00am - 7:10am (JST)	Break
Flash Talk Session	

5:10pm - 5:35pm (EST) 7:10am - 7:35am (JST)	Presentations on Research Plan at NIH by JSPS-NIH Fellows (KAITOKU-NIH) Awarded in FY2023 (8 Fellows) 3 mins for each
5:35pm - 5:45pm (EST) 7:35am - 7:45am (JST)	Closing Remarks Dr. Peter Kilmarx, Acting Director, Fogarty International Center, NIH Dr. Nina F. SCHOR, Deputy Director for Intramural Research, NIH
5:45pm - 7:30pm (EST) 7:45am - 9:30am(JST)	Networking (only for in-person participants)

Investigating Age-Induced Alterations in Brain Function and Regeneration Using Zebrafish Model

Motoyuki ITOH

Professor Graduate School of Pharmaceutical Sciences Chiba University



Abstract

Zebrafish, with a lifespan of 3-5 years, share about 70% homology with the mammalian genome and retain over 80% of disease proteins. Our lab uses zebrafish for aging research, comparing their aging processes with mammals. We studied age-related changes in metabolism, learning ability, and the cerebrovascular system. Metabolic aging in zebrafish is evident at 10-14 months, and a decline in learning ability is observed at 15 months. A positive correlation was found between cerebral blood flow and learning ability in 14-month-old zebrafish, similar to aged humans. This suggests that aging symptoms appear in the brain and metabolic rate in zebrafish after 14 months. Unlike mammals, fish, especially zebrafish, have a higher regenerative capacity, including remarkable brain regeneration abilities, considered an anti-aging mechanism. While zebrafish show reduced learning ability with age, the relationship between their regenerative ability and age-related functional decline is unclear. Recent research is focused on elucidating this link, aiming to gain insights into overcoming age-related brain vulnerability in mammals.

Motoyuki ITOH

Biography

Education

1991 Osaka University, Pharmaceutical science, Japan Bachelor

1993 Osaka University, Pharmaceutical science, Japan Master

1998 Osaka University, Medical science, Japan Ph.D

Positions and Employment

1998-1999 Osaka University, Japan Postdoctoral fellow

1999-2003 LMG/NICHD, USA Postdoctoral fellow

2003-2012 Nagoya University, Japan Associate professor

2012- Chiba University, Japan Professor

Sigma-1 receptor plays a critical role in cocaine-induced synaptic AMPA receptor plasticity in the VTA dopamine neurons

Yuko YASUI Staff Scientist National institute on Drug Abuse NIH



Abstract

Sigma-1 receptor (Sig1R) is a ligand-operated chaperone that regulates a variety of cellular functions through interaction with proteins including receptors and ion channels. Sig1R has been implicated in cocaine addiction since Sig1R ligands attenuate cocaine-induced behaviors. Cocaine causes neuroadaptation in the mesocorticolimbic dopamine system which is essential for reward and motivational process. This system connects the ventral tegmental area (VTA) to the major projected areas, nucleus accumbens and prefrontal cortex, where alpha-amino-3-hydroxy-5methyl-4-isoxazolepropionic acid receptors (AMPARs) play an important role in cocaine-induced synaptic plasticity and long-lasting behavioral effects of cocaine. Single cocaine exposure in vivo induces long-term potentiation in the VTA dopamine neurons at least partly through insertion of GluR2-lacking AMPAR. However, the molecular mechanisms of this process are not fully understood. In this study, we found that a single cocaine injection increased synaptic GluR2-lacking AMPAR in the VTA dopamine neurons of wild-type mice as previously reported, but not in Sig1R knockout mice. Moreover, the expression of GluR2 and GluR3 was decreased on the surface and increased in the endosomes with cocaine treatment in wild-type but not in Sig1R knockout mice. Together, we demonstrate that Sig1R controls cocaine-induced formation of GluR2-lacking AMPAR by altering the intracellular distributions of GluR2 and GluR3. This study suggests Sig1R can be a therapeutic target for cocaine addiction.

Yuko YASUI

Biography

Gifu University, Gifu, Japan Ph.D., Pharmaceutical Sciences, March 2012

National Institutes of Health (NIH), National Institute on Drug Abuse (NIDA), Baltimore, MD Postdoctoral Research Fellow, October 2012-Septemebr 2018

National Institutes of Health (NIH), National Institute on Drug Abuse (NIDA), Baltimore, MD Research fellow, November 2018-August 2020

National Institutes of Health (NIH), National Institute on Drug Abuse (NIDA), Baltimore, MD Staff Scientist, August 2020-present

NGF-TrkA-PI3K signaling promotes sensory hypersensitivity in diabetes

Yuta KOUI

Postdoctoral Fellow Laboratory of Stem Cell and Neuro-Vascular Biology, Cell and Developmental Biology Center, National Heart, Lung, and Blood Institute National Institutes of Health



Abstract

Diabetic patients often develop small fiber neuropathy in the early stage of diabetes, including obesity and prediabetes. Patients with small fiber neuropathy exhibit dysfunction in skin sensory nerves and blood vessels, leading to severe burning or shooting pain. However, how symptoms are sequentially established and which signaling mechanisms are involved in sensory hypersensitivity remain unclear. Using a novel pain behavior assay and ex vivo live tissue calcium imaging, we demonstrated that diet-induced obesity (DIO) model mice develop painful symptoms and sensory hypersensitivity at 22 weeks-of-age, and exhibit reduced painful symptoms and hypersensitivity at 30 weeks-of-age. Our immunohistochemical analysis revealed sensory axon degeneration in the skin epidermis at 30 weeks-of-age but not at 22 weeks-of-age, indicating that hypersensitivity is established before the onset of axon degeneration. We also observed hyperpermeability in capillary vessels from 22 weeks-of-age onward, suggesting that vascular damage may play a role in the development of sensory dysfunction in the DIO skin. At the mechanistic level, enhanced NGF expression was observed in skin epidermal keratinocytes of the DIO mice at 22 weeks-of-age. Local treatment with anti-NGF neutralized antibody or a small compound inhibitor for PI3K, a downstream component of NGF signaling, suppresses hypersensitivity in the DIO skin. This suggests that keratinocyte-derived NGF induces hypersensitivity in the DIO skin. Collectively, NGF locally regulates sensory activities in diabetes and could be a potential therapeutic target for diabetic small fiber neuropathy.

Yuta KOUI

Biography

Education and position of employment

2014-2019: Master student, Ph.D. Candidate, and Postdoctoral Fellow supervised by Dr. Atsushi Miyajima, Laboratory of Cell Growth and Differentiation. Institute for Quantitative Biosciences, The University of Tokyo, Japan. 2019-present: Postdoctoral Fellow supervised by Dr. Yosuke Mukoyama, Laboratory of Stem Cell and Neuro-Vascular Biology. National Heart, Lung, and Blood Institute (NHLBI), National Institutes of Health (NIH), USA.

Fellowship and Award

2016-2019: Research fellow of Japanese Society of the Promotion of Science (JSPS)

2020-2022: Research Fellowship for Japanese Biomedical and Behavioral Researchers at NIH (JSPS)

2022-2023: Uehara Postdoctoral Fellowship Award (The Uehara Memorial Foundation)

2022-present: Lenfant Fellowship Award (National Heart, Lung, and Blood Institute)

Research Plan at NIH by JSPS-NIH Fellows Awarded in FY2023

	Name	page
1	TAMURA Atsumi	10
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6	WADA Yoichi	15
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Atsumi Tamura , M.D., Ph.D.



Effect of mutated Hras expression and deletion of Nkx2-1 on thyroid carcinogenesis in mice





Aim: To understand whether the loss of *Nkx2-1* promotes carcinogenesis expressing mutated *Hras* in the thyroid of model mouse

- 1) To determine the incidence of thyroid tumors between different genotype groups
- 2) To identify metabolites in serum, specific to thyroid tumors and genotypes in mice metabolomics analysis
- 3) To analyze the metabolites of human clinical serum samples

Chiori Tabe





2023 Ph.D. Morphological features of bronchiectasis in patients with non-tuberculous mycobacteriosis and interstitial pneumonia. BMC Res Notes. 2022 Jul 26;15(1):263.



2020.4-

1. The detection of drug-resistant mutation in anaplastic lymphoma kinase (ALK) positive lung cancer patients treated with ALK-TKIs.



2. The monitoring of the medication-induced changes in the intestinal microbiota in epidermal growth factor receptor (EGFR) positive lung cancer patients treated with EGFR-TKIs.



Identification and characterization of proteins associated with regions of extrachromosomal DNA regulating oncogene expression in small cell lung cancer cells



- Knockdown experiments using siRNA / shRNA
- Knockout experiments by genome editing
- Overexpression

Analysis of:

- Cell proliferation (in vitro / in vivo)
- Metastasis (in vitro / in vivo)

Biochem Biophys Res Commun 439(1):132-6,2013)

DNA analysis RNA analysis Protein analysis

Reverse crosslink

Keisuke Fukutomi

Mar 2011 Apr 2011-Mar 2017	M.D. , Osaka University Faculty of Medicine, Osaka, Japan Clinical training in Gastroenterology and Hepatology Osaka National Hospital and Osaka University Hospital, Osaka, Japan
Apr 2017-Nov 2021	Ph.D., Osaka University Graduate School of Medicine, Osaka, Japan
 Innate immune resp under treatment with B cell responses in Development of an host immune response 	ponse of HBV-infected hepatocytes h Capsid Assembly Modulator and peg-IFNα patients with chronic HBV infection HBsAg-transgenic mouse model to study nses Fukutomi K et al. <i>Hepatol Commun</i> 2022
Feb 2022- present	Postdoctoral Fellow, Immunology Section, Liver Diseases Branch, NIDDK, NIH Section Chief: Dr. Barbara Rehermann
Immunopathogene	sis of perinatal flares of hepatitis in pregnant women with chronic HBV infection

Future: Conduct translational research in immunology.

A clinical and immunological pilot study to restore HBV-specific immune responses in chronic hepatitis B virus infection

Background:

- HBV-specific immune responses are required to recover from acute and chronic HBV infection
- Long-term exposure to HBV antigens results in exhaustion of HBsAg-specific T cells

The problem:

- HBsAg levels exceed HBV virion levels by a factor of 100,000.
- · Conventional anti-HBV drugs block HBV replication but not HBsAg production from cccDNA & integrated DNA

Approach:

• Block HBs Ag production with siRNA treatment prior to antiviral treatment





Leo Yamada, M.D., Ph.D.





Lab in NIH: Laboratory of Human Carcinogenesis/NCI

Education & Training

- Surgeon (in Gastroenterology)
- 2012: M.D., Fukushima Medical University
- 2016-2020: Ph.D., Fukushima Medical University

Gastric Cancer (2021) 24	60.71		
https://doi.org/10.1007,	s10120-020-01094-0		
ORIGINAL ARTI	LE		

Selective sensitivity of EZH2 inhibitors based on synthetic lethality in ARID1A-deficient gastric cancer

Leo Yamada¹ - Motonobu Saito¹ ⁽¹⁾ - Aung Kyi Thar Min¹ - Katsuharu Saito¹ - Mai Ashizawa¹ - Koji Kase¹ -Shotaro Nakajima¹² - Ilisashi Onozawa¹ - Hirokazu Okayama¹ - Hisahito Endo¹ - Shotaro Fujita¹ -Wataru Sakawoto¹ - Zencihico Saze¹ - Tomoyuki Momma¹ - Kosaku Mimura¹³ - Shinji Oiku¹ - Koji Kono¹

Previous Affiliation in Japan:

- 1. Department of Gastrointestinal Tract Surgery, Fukushima Medical University
- 2. Laboratory of Molecular Carcinogenesis, National Cancer Center Research Institute

After JSPS-NIH

• A physician scientist for translational research.

Investigation of p53 Isoform-Based Therapeutic Strategy for Physiological and Pathological Aging.

Background

 Δ 133p53 α inhibits premature cellular senescence and extends cellular lifespan in progeria cells *in vitro*.

- Therapeutic potential in CD8⁺T cells and brain astrocytes as well.
- Non-oncogenic and non-mutagenic, encouraging clinical translation.

Hypothesis

 $\Delta133p53\alpha$ inhibits premature aging phenotypes and extends lifespan in a progeria mouse model in vivo.

Specific Aim

Investigate the effects of Δ 133p53 α on aging phenotypes and lifespan in the transgenic mice.





PI: Dr. Curtis C. Harris

Symptoms

Growth impairment Cardiovascular disease Skeletal dysplasia

Lipodystrophy Alopecia in and nail defects

Hutchinson-Gilford Progeria Syndrome

LmnaG609G

(Progeria mouse model)

Skin and nail defec Joint contractures

Reona Okada, M.D. Ph.D.

- Pediatric oncologist (2015-2022)
- Ph.D. work in the Seki Lab at Chiba University Graduate School of Medicine (2019-2021): Identification of oncogenes in various cancers using microRNA signatures. Dissertation: Regulation of oncogenic targets by miR-99a-3p (passenger strand of miR-99aduplex) in head and neck squamous cell carcinoma
- Postdoc in the Nguyen Lab in the Pediatric Oncology Branch, NCI (2022-present): Immunotherapy development for various pediatric cancers

GPC2-CAR T-cells engineered with NFAT-inducible membrane-tethered IL-15/IL-21 exhibit enhanced activity against neuroblastoma (under review)

• Post-fellowship goals: building a translational research program at a university in Japan

Orthotopic patient-derived xenografts (PDXs) of adrenocortical carcinoma (ACC) to accelerate drug discovery

ACC natient

1. ACC is a <u>rare</u> cancer of the adrenal gland without many therapy options for advanced disease



2. We established the largest collection of ACC PDXs (n=6)

Characterization of the PDXs

- Histology (immunohistochemistry)
- Flow cytometry
- RNA sequencing
- Whole exome sequencing
- Proteomics
- Surface proteomics
- Serum sample analysis
- 4. ACC PDXs are pan-sensitive to XPO1 inhibition



- Cytotoxicity of eltanexor in vivo (ongoing)
- Mechanism studies of eltanexor in ACC (pending proteomic and genetic study)
- Efficacy of combinatorial therapy B7H3-CAR + eltanexor (ongoing)

J Clin Endocrinol Metab (2014) Illustrations created with BioRender.com.

Yoichi Wada M.D., Ph.D.

Board Certified Pediatrician, Supervisor of Pediatrics (Japan Pediatric Society) Board Certified Medical Geneticist (Japan Society of Human Genetics) Specialty: Inborn Errors of Metabolism

- 2023 Present Metabolic Medicine Branch, Organic Acid Research Section, NHGRI Laboratory of Dr. Charles P. Venditti, Visiting Fellow
- 2019 2023 Tohoku University Hospital, Assistant Professor
- 2015 2019 Tohoku University Graduate School of Medicine
- 2010 2015 Kurashiki Central Hospital, Junior and Senior Resident

Previous studies

Galactosemia type IV (galactose mutarotase deficiency)

- Identification as a disease entity (MIM#618881) Wada et al, Genet Med 2018
- Prevalence estimation Iwasawa, Kikuchi, Wada, et al., Mol Genet Metab 2019
- Drug repurposing Wada et al, JIMD 2022
- Phenotypic and genetic clarification Mikami-Saito, Wada et al, under review
- Others also based on clinical questions as a pediatrician
- Establishment of mouse model of lanosterol synthase deficiency
- Rapid detection tests for blood Phenylalanine

Future direction after NIH-KAITOKU

- Gene therapy development for patients with Inborn Errors of Metabolism







Wada et al, PLOS Genet 2020 Wada et al, MGM rep 2023

Nuclease free genome editing to treat cobalamin B class methylmalonic acidemia

Cobalamin B class methylmalonic acidemia (cblB)

- · caused by biallellic mutations in MMAB gene
- characterized by metabolic instability, chronic renal disease, and neurological complications
- high mortality and morbidity
- there is no treatment to restore the metabolic condition

We are planning to insert normal *MMAB* gene into *Alb* locus without nuclease in model mouse of cblB





	2024 Q1	2024 Q2	2024 Q3	2024 Q4	2025 Q1	2025 Q2	2025 Q3	2025 Q4
Colony expansion	+	+						
Treatment: short term analysis		+	+					
Treatment: long term analysis			+	+	+	+		
Abstract							+	+
Manuscript								+



Project: Identification of Pathological Variants in Inflammatory Diseases of Unknown Cause



Yuta Myojin M.D., Ph.D.

-2021.6 Department of Gastroenterology, Osaka University

Research:

- 1. Cancer associated fibroblast (CAFs) and cancer cell interaction in hepatocellular carcinoma(HCC) -Autophagy of CAFs promotes tumor proliferation via several cytokine production.
- 2. Biomarker research in TKI treatment of HCC
- -The candidate marker discovered in mice screening was validated with clinical samples.

2021.7- NCI / Thoracic and GI Malignancies Branch / Dr. Greten's lab

Main Research : Correlative study of clinical trial of HCC patients treated with immunotherapy

My plan after leaving NIH

As a physician scientist, I'd like to conduct a clinical trials and translational research in Japan.

Correlative study of clinical trial of HCC patients treated with immunotherapy



Biopsy samples in pre-post treatment

=>Whole exon sequencing/ Bulk RNA sequencing/ 37-plex immunohistochemistry(CODEX)/single cell RNA sequencing Blood samples in pre-post treatment

=>PBMC : Spectro-cytometry(2 panel, 42 targets in total)/ Plasma : Multiplex cytokine panel

<Plan>

- 1. Collect samples of clinical trial and conduct the experiments with multiple modality
- 2. CODEX data => annotate cells and analyze the cellular neighborhood
- 3. scRNAseq data => validate the findings above by cell-cell interaction analysis + DEG analysis
- 4. BulkRNAseq data => validate the data in 1 and 2 by GSEA analysis and immune cell composition analysis
- 5. WES/Blood data => analyze the change pre and post/ response => combine with results from 1/2/3
- 6. Mice orthotopic liver tumor model => validate the hypothesis found in the above analysis