July 9 (Tue)

15:00-17:00

Registration

17:20-17:30 Opening Remarks: Keiji TANAKA

Plenary Lecture

	Chair: Keiji TANAKA
	New insights into biochemical mechanisms of the 26S proteasome
17:30-18:30	Department of Cell Biology, Harvard Medical School, USA
	Alfred L. GOLDBERG

18:40-20:00

Welcome Reception

July 10 (Wed)

7:00-8:50

Breakfast

Session A: Basic Mechanisms of the Proteasome

Chairs: Wolfgang P. BAUMEISTER, Keiji TANAKA		
9:00-9:30	A-1	Recent advances in structural studies of the 26S proteasome
		Department of Structural Biology, Max-Planck-Institute of Biochemistry, Germany
		Wolfgang P. BAUMEISTER
	A-2	Regulation of proteasome activity
9:30-10:00		Department of Cell Biology, Harvard Medical School, USA
		Daniel FINLEY
10:00-10:30	A-3	BAG6 is essential for selective elimination of aggregation- prone defective proteins
		Department of Biological Sciences, Tokyo Metropolitan University, Japan
		Hiroyuki KAWAHARA

10:30-11:00

Coffee Break

11:00-11:15	A-4 (41)	Proteasomal degradation resolves competition between cell polarization and cellular wound healing Graduate School of Medical Science, Nagoya City University, Japan
		Keiko KONO
11:15-11:30	A-5 (35)	Structure and function of moyamoya disease-associated AAA+ ATPase/ubiquitin ligase mysterin Faculty of Life Sciences, Kyoto Sangyo University, Japan
		Daisuke MORITO
11.30-12.00	A-6	In-depth analysis of cellular dynamism of the proteasome Tokyo Metropolitan Institute of Medical Science, Japan
11.30-12.00		Keiji TANAKA
12:00-12:30	A-7	Regulation of eukaryotic proteasome assembly
		Department of Molecular Biophysics and Biochemistry, Yale University, USA
		Mark HOCHSTRASSER

Number in parenthesis: poster number

12:30-14:00

Lunch

Session B: Regulation of Ubiquitylation

Chairs: Raymond DESHAIES, Shigetsugu HATAKEYAMA			
		Regulation of culling-RING ubiquitin ligases	
14:00-14:30	B-1	California Institute of Technology, USA	
		Raymond DESHAIES	
14:30-15:00	B-2	Ubiquitin-dependent regulation of ASK1 stress signaling in cell death	
		Laboratory of Cell Signaling, Graduate School of Pharmaceutical Sciences, The University of Tokyo, Japan	
		Hidenori ICHIJO	
15:00-15:30	B-3	Functional analysis of F-box proteins in vivo	
		School of Medicine, Tohoku University, Japan	
		Keiko NAKAYAMA	

15:30-16:00

Coffee Break

16:00-16:15	B-4 (46)	Real-time, label-free monitoring of poly ubiquitin chain formation with bacterial ubiquitin ligase
		Division of Bacterial Infection Biology, Institute of Medical Science, The University of Tokyo, Japan
		Minsoo KIM
16:15-16:30	B-5 (19)	Recognition of K63-linked ubiquitin chains by the Ankrd13 family of UIM-bearing proteins regulates endocytosis of plasma membrane proteins Department of Biological Sciences, Tokyo Institute of Technology, Japan
		Masayuki KOMADA
16:30-17:00	B-6	Regulation of cellular functions by TRIM proteins Department of Biochemistry, Hokkaido University, Japan
		Shigetsugu HATAKEYAMA
17:00-17:30	B-7	The HRD ubiquitin ligase: managing protein quality control and ubiquitylation to maintain protein homeostasis in the secretory pathway Cancer Department, Max-Delbrück-Center for Molecular Medicine, Germany
		Thomas SOMMER

Number in parenthesis: poster number

17:30-19:00	Dinner	
19:00-21:00	Poster Session [I]	

July 11 (Thu)

7:00-8:50

Breakfast

Keynote Lecture		
	Chair: Kazuhiro IWAI	
9:00-9:40	The ubiquitin proteolytic system-from basic mechanisms thru human diseases and on to drug development	
	Faculty of Medicine, Technion-Israel Institute of Technology, State of Israel	
	Aaron J. CIECHANOVER	

Session C: Diverse Roles of Ubiquitylation		
Chairs: Kazuhiro IWAI, Stefan JENTSCH		
		Linear polyubiquitination: a new regulator of NF-κB activation
9:40-10:10	C-1	Department of Molecular and Cellular Physiology, Graduate School of Medicine, Kyoto University, Japan
		Kazuhiro IWAI
		Structural basis of ubiquitin chain recognition
10:10-10:40	C-2	Department of Structural Biology, Stanford School of Medicine, Stanford University, USA
		Soichi WAKATSUKI

10:40-11:10

Coffee Break

11:10-11:25	C-3 (20)	Stabilization of mouse CRY proteins by FBXL21 is critical for circadian oscillation of the biological clock. ~Molecular characterization of FBXL21 in circadian clockwork~ Department of Biophysics and Biochemistry, Graduate School of Science, The University of Tokyo, Japan
		Arisa HIRANO
11:25-11:40	C-4 (31)	Functional analysis of budding yeast SCFYlr224w E3 ligase Graduate School of Science, Nagoya University, Japan Takumi KAMURA
11:40-12:10	C-5	Ubiquitin-fold modifier 1 as a transcriptional regulator Department of Biological Sciences, Seoul National University, Korea Chin Ha CHUNG
12:10-12:40	C-6	Relevance of substrate-selective and protein group SUMOylation for nuclear activities Department of Molecular Cell Biology, Max-Planck-Institute of Biochemistry, Germany Stefan JENTSCH

Number in parenthesis: poster number

12:40-14:00

Lunch

Session D: Pathophysiological Roles of the Proteasome

		Chairs: Ron R. KOPITO, Shigeo MURATA
14:00-14:30	D-1	Presentation of unique peptide repertoire on MHC class I by the thymoproteasome
		Graduate School of Pharmaceutical Sciences, The University of Tokyo, Japan
		Shigeo MURATA
		Immunoproteasomes in the regulation of protein
14:30-15:00	D-2	homeostasis
		Institute of Biochemistry, Charité-Universitätsmedizin Berlin, Germany
		Peter-Michael KLOETZEL
15:00-15:30	D-3	Immunoproteasomes and human diseases
		Department of Immunology, The University of Tokushima, Japan
		Koji YASUTOMO

15:30-16:00

Coffee Break

16:00-16:15	D-4 (32)	A missense mutation in Psmb11 impairs thymoproteasome assembly and T cell development Department of Immunology and Pathology, National Center for Global Health and Medicine, Japan
		Takeshi NITTA
16:15-16:30	D-5 (43)	Loss of MHC class II ubiquitination negatively regulates dendritic cells Laboratory of integrative infection immunology, Showa Pharmaceutical University, Japan
		Satoshi ISHIDO
16:30-17:00	D-6	Regulation of leaf organ size and gene silencing by plant proteasome
		Research Faculty of Science, Hokkaido University, Japan
		Junji YAMAGUCHI
17:00-17:30	D-7	The role of protein aggregation and ubiquitylation in the pathogenesis of Huntingtons disease Department of Biology, Stanford University, USA
		Ron R. KOPITO

Number in parenthesis: poster number

17:30-19:00

Dinner

19:00-21:00

Poster Session [II]

July 12 (Fri)

7:00-8:50

Breakfast

Session E: Ubiquitin and Diseases

Chairs: Keiichi NAKAYAMA, Michele PAGANO		
9:00-9:30	E-1	Control of cell proliferation by SCF ubiquitin ligases and its relevance in human cancers
		Howard Hughes Medical Institute and Department of Pathology, NYU Cancer Institute, New York University School of Medicine, USA
		Michele PAGANO
9:30-10:00	E-2	Fbw7 is essential for maintenance of quiescence and function of cancer stem cells
		Department of Molecular and Cellular Biology, Medical Institute of Bioregulation, Kyushu University, Japan
		Keiichi NAKAYAMA
		Identification of a primary target of thalidomide's complex biological effects
10:00-10:30	E-3	Graduate School of Bioscience and Biotechnology, Tokyo Institute of Technology, Japan
		Hiroshi HANDA

10:30-11:00

Coffee Break

11:00-11:15	E-4 (48)	Coordinated functions of BRCA1, Claspin and HERC2 in DNA damage response and cell cycle Department of Translational Oncology, St. Marianna University Graduate School of Medicine, Japan Tomohiko OHTA
11:15-11:30	E-5 (7)	Identifying HECT E3 ubiquitin ligase substrate pairs by mechanism based approach Department of Chemistry, Northwestern University, USA Sungjin PARK

11:30-12:00	E-6	Regulation of cullin-RING ligases by viral deneddylases Department of Cell and Molecular Biology, Karolinska Institutet, Sweden
		Maria MASUCCI
		The Keap1-Nrf2 system for environmental response
12:00-12:30	E-7	Department of Medical Biochemistry, Tohoku University Graduate School of Medicine, Japan
		Masayuki YAMAMOTO

Number in parenthesis: poster number

12:30-12:45	Announcement
12:45-12:50	Closing Remarks: Kazuhiro IWAI
12:50-13:30	Lunch