

# Special lectures on biomedicine

Time : Wed. 15:30 – 17:00

Classroom : Meeting room No.3 of Auditorium and Activity Center

No.	Date	Speaker	Topic	Organizer
1	3/2	Dr. Chuan-Chin Chiao (National Museum of Natural Science)	Cephalopods as model animals for neuroscience research: historical perspectives and modern approaches	Institute of Neuroscience
2	3/9	Dr. Kai-Chien Yang (Department and Graduate Institute of Pharmacology, National Taiwan University)	Targeting ER Protein TXNDC5 as a Novel Therapeutic Approach against Cardiac and Organ Fibrosis	Department of Life Sciences and Institute of Genome Sciences
3	3/16	Dr. Charles Pin-Kuang Lai (Institute of Atomic and Molecular Sciences, Academia Sinica)		Institute of Biopharmaceutical Science
4	3/23	Dr. Eminy H.Y. Lee (Institute of Biomedical Sciences, Academia Sinica)		College of Life Sciences
5	3/30	Dr. Pei-Lin Cheng (Institute of Molecular Biology, Academia Sinica)		Institute of Biochemistry and Molecular Biology
6	4/13	Dr. Yan-Shen Shan (Institute of Clinical Medicine, National Cheng Kung University)	The impact of tumor microenvironment on pancreatic tumorigenesis: from bedside to bench	Institute of Microbiology & Immunology
7	4/20	Dr. Kuei-Sen Hsu (Department of Pharmacology, National Cheng Kung University / Vice Dean of School of Medicine)		College of Life Sciences / College of Medicine / Institute of Neuroscience
8	4/27	Dr. Hsing-Pang Hsieh (Biomedical Translation Research Center, Academia Sinica)	From Bench to Clinic: Translational Research to Startups	Department of Life Sciences and Institute of Genome Sciences
9	5/4	Dr. Steve R. Roffler (Institute of Biomedical Sciences, Academia Sinica /		Institute of Biopharmaceutical Science

		Ph.D. University of California, Berkeley)		
10	5/18	Dr. Li-Jung Juan (Genomics Research Center, Academia Sinica)		Institute of Biochemistry and Molecular Biology
11	5/25	Dr. Jhih-Wei Chu (Department of Biological Science & Technology, National Yang Ming Chiao Tung University )	On the Molecular Origin of Functional Properties in Gene Expression and Protein Allostery: Mechanism Uncovered by Molecular Simulation, Machine Learning, and Graph Theory	Institute of Microbiology & Immunology