2020年度 国際共同研究 採択課題一覧

課題 番号	研 究 課	国際共同研究員氏名	国際共同研究員所属	所属機関 所在国	蛋白質研究所 担当研究室
1	Development of a new method to conjugate the defensin peptide to the carrier protein P64K using a MeOGly strategy		Center for Genetic Engineering and Biotechnology	Cuba	蛋白質有機化学研究室
2	Peptide Quantum Dot conjugate as new-age theranostics	SHARMA ROHIT KUMAR	Panjab University	India	蛋白質有機化学研究室
3	Computational modelling of EML4-ALK signaling pathway	g SAMPSON IOSIFINA	University of Leeds	United Kingdom	細胞システム研究室
4	Analysis of cell cycle dynamics by integrati of mathematical-experimental approach	on KHOLODENKO BORIS	Systems Biology Ireland, University College Dublin	Ireland	細胞システム研究室
5	Structure and Dynamics of Musashi 2 protein apo and RNA-bound form	in CHUGH JEETENDER	Indian Institute of Science Education & Research	India	機能構造計測学研究室
6	Structural insight into cold adaptation mechanism of FK506-binding protein from psychrophilic bacteria	BUDIMAN CAHYO	Universiti Malaysia Sabah	Malaysia	機能構造計測学研究室
7	Solid-state NMR Studies on bone and other nanomaterials	RAMAMOORTHY AYYALUSAMY	University of Michigan	USA	機能構造計測学研究室
8	Structural study of dUMP hydroxymethylases f bacteriophage	rom SONG HYUN KYU	Korea University	Korea	超分子構造解析学研究室
9	Structural and functional research on the survival-essential factors from bacterial pathogens for the development of novel antibiotics which induces suicide effect(Pha V)	LEE BONG-JIN	Seoul National University	Korea	超分子構造解析学研究室
10	Crystallographic fragment screening and structure determination for anticancer targe proteins	t KIM HYOUN SOOK	National Cancer Center	Korea	超分子構造解析学研究室
11	Structures of Hpt and the C-terminal received domain of P. aeruginosa	r CHEN CHUN-JUNG	National Synchorotron Radiation Research Center	Taiwan	超分子構造解析学研究室
12	Structural analysis of human pattern recogni receptors	tion HWANG KWANG YEON	Korea University	Korea	超分子構造解析学研究室
13	Drug screen strategy targeting RpoS against bacterial antibiotic resistance	SAQIB UZMA	Indian Institute of Technoligy Indore	India	超分子構造解析学研究室