The 17th Asian Bioinformatics Consortium (ABC) Symposium

August 22nd to 23rd, 2019

Gui' an, Guizhou Province, China.

Program Version3.0 (August 14^h, 2019) The Asian Bioinformatics Consortium (ABC) Symposium (formerly known as the Japan-China-Korea Bioinformatics Training Course or Symposium) started in 2002 which aims at promoting bioinformatics research activity in the three Asian countries through scientific exchange, training future talents and exploring new translational biomedical strategies to improve human health.

History



Time

Registration	August 21st, 2019	
Symposium August 22nd to 23rd, 2019		
Departure August 24th, 2019		

Venue

Gui'an New Area, Guiyang, Guizhou Province, China.



Local Organizers and Sponsors

- · Shanghai Institute of Nutrition and Health (SINH), CAS
- Bio-Med Big Data Center, SINH, CAS
- Guizhou Science and Technology Information Center
- Guizhou Normal University
- Shanghai Center for Bioinformation Technology
- Shanghai Society for Bioinformatics
- Gui An Bio-Med Big Data Center, SIBS, CAS

Attendants

- One keynote speaker and nine speakers from each country, 30 in total.
- A keynote speaker from each country will give 35-min presentation.
- Remaining nine speakers from each country will give 20-min presentation.

Meeting Place

Conference Center, Guizhou Normal University (Hua Xi Campus) (贵州师范大学,花溪大学城,会议中心)

Accommodation



Hotel: Wan Yi Panorama Hotel (万宜丽景酒店:贵阳市花溪区清溪路 111 号) Address: No.111, Qing Xi Road, Hua Xi District, Guiyang, Guizhou Province. Tel: 86-0851-83962888

Cost and Facility

The attendance fee is free. Speakers should cover their own travel expenses.

• Three days' lodging will be provided by Bio-Med Big Data Center of Shanghai Institute of Nutrition and Health of CAS including August 21st, 22nd, 23rd.

Invited Speakers

📕 China		
Chen Runsheng	Professor, Institute of Biophysics, CAS	Topic: Noncoding sequences of genome and non coding RNA.
Zhao Guoping	Professor, Institute of Plant Physiology and Ecology, CAS.	Topic: Strive for international collaboration on the challenge of Bio-medical data — — From Genomics and through Biomes
Li Yixue	Professor, CAS-MPG Partner Institute for Computational Biology, Shanghai Institute of Nutrition and Health, CAS.	1
Xie Lu	Professor, Shanghai Center for Bioinformation Technology.	Topic:GeneexpressionproteomelandscapeofChineseHCCrevealsprognosticdeterminantsand therapeutic targets
Wang Zefeng	Professor, CAS-MPG Partner Institute for Computational Biology, Shanghai Institute of Nutrition and Health, CAS.	Topic: Increasing the human genome complexity at the RNA levels
Shen Tie	Associate Professor, Guizhou Normal University	Topic: Fluxomics: Big data for synthetic biology
Lv Hui	Professor, Shanghai Jiao Tong University	Topic: Joint analysis framework for multiomics and multidisease data
Wei Chaochun	Professor, Shanghai Jiao Tong University	Topic: HUPAN: a human pan-genome analysis pipeline
Shi Tieliu	Professor, East China Normal University	Topic:DeepLearning-BasedMulti-OmicsDataIntegrationRevealsTwoPrognosticSubtypesin

		High-Risk Neuroblastoma
Wang Sijia	Professor, CAS-MPG Partner Institute for Computational Biology, Shanghai Institute of Nutrition and Health, CAS.	Topic: Human physical traits and the phenome research
Liu Qi	Professor, Tongji University	Topic: Initi-IO: An Integrative and Intelligentized Immuno-Oncology Study Platform
💌 Japan		
Gojobori Takashi	Professor, KAUST, Saudi Arabia & NIG	Topic: Digital Health, AI, and Biomedical Genomics
Saitou Naruya	Professor, National Institute of Genetics	Topic:Introduction of YaponesianGenomeProjectwithspecial reference to modernhuman genome analyses
Yoko Satta	Professor, SOKENDAI Hayama	Topic: Detecting signals of positive selection by two dimensional site frequency spectrum (2DSFS): a case study of schizophrenia related SNPs
Nakamura Yasukazu	Professor, National Institute of Genetics	Topic: Chromosome-scale assemblies are coming back: DB updates and future plan
lturo Inoue	Professor, National Institute of Genetics	Topic: Viruses in the planet
Tadashi Imanishi	Professor, Okai University Medical School	Topic: Bioinformatics for the diagnosis of infectious diseases
Kosuke Hanada	Associate Professor, Kyushu Technology University	Topic: The origin and evolution of Japanese cultivated radishes
Masatoshi Matsunami	Assistant Professor,	Торіс:

	University of the Ryukyus Medical School	A comprehensive reference transcriptome resource for the Iberian ribbed newt Pleurodeles waltl, an emerging model for developmental and regeneration biology
Hideaki Kanzawa	Staff Scientist, National Museum of Science and Nature	Topic: History of Yaponesians inferred from ancient Yaponesian genome sequences
Kae Koganebuchi	Specially appointed assistant professor, University of the Ryukyus Medical School	Topic: Demographic history of the Ryukyu Islanders inferred from whole-genome sequencing
Korea		
Kim Sanguk	Professor, Pohang University of Science and Technology	Topic: Evolution of functional modules and regulatory networks: Understanding genotype-phenotype relationship
Park Ji Hwan	Senior Researcher, Korean Bioinformation Center	Topic:TranscriptomicandproteomicapproachesidentifyregulatoryNACnetwork ofArabidopsisleafsenescence
Choi Sun Shim	Professor, Kangwon National University	Topic:Functional relevance of firstintronrevealedbyenrichmentofepigeneticmodification signals
Lee Younghee	Assistant Professor, University of Utah, Asia Campus	Topic: Alternative splicing and human diseases
Rhee Je-Keun	Assistant Professor, Pusan National University	Topic: Genomic and epigenomic data analysis using probabilistic evolutionary learning

Park Kiejung	Specially Employed Professor, Sangmyung University	Topic: Development of a prediction system for positively selected genes against massive orthologous groups
Choe Yong-Kyung	Professor, Institut Pasteur Korea	Topic: Institut Pasteur Korea: role and works
Choi Inhee	Senior Researcher, Institut Pasteur Korea	Topic: Identification of TB Drug Targets from NGS
Jeong Choongwon	Assistant Professor, Seoul National University	Topic: The genetic history of admixture across inner Eurasia
Hwang Seungwoo	Principal Investigator, Korean Bioinformation Center	(Poster presentation only) Genome-wide DNA methylation profile of Korean CLL patients

Program

Talk time for keynote speakers is 35 min. Talk time for other speakers is 20min

Registration: Wednesday, August 21st , 2019 Wan Yi Panorama Hotel.	
13:00-18:00	Registration at Wan Yi Panorama Hotel.
18:30-20:00	Welcome Reception at Wan Yi Panorama Hotel.

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Day 1: Thursday, August 22 nd , 2019 Conference Center, Guizhou Normal University (Hua Xi Campus).				
08:50-09:00	:00 Introduction Li Yixue & Xie Lu			
	Session 1: Chaired by Wang Ze	feng		
09:00-09:35	Keynote Talk 1: Noncoding sequences of genome and non coding RNA.	Chen Runsheng Institute of Biophysics, CAS		
09:35-10:10	Keynote Talk 2: Strive for international collaboration on the challenge of Bio-medical data—— From Genomics and through Biomes	Zhao Guoping Institute of Plant Physiology and Ecology, CAS		
10:10-10:30	Talk 1: Detecting signals of positive selection by two dimensional site frequency spectrum (2DSFS): a case study of schizophrenia related SNPs	Yoko Satta SOKENDAI Hayama		
10:30-10:50	Talk 2: Joint analysis framework for multi-omics and multi-disease data	Lv Hui Shanghai Jiao Tong University		
10:50-11:10	Talk 3: Institut Pasteur Korea: role and works	Choe Yong-Kyung Institut Pasteur Korea		
11:10-11:30	Talk 4: Deep Learning-Based Multi-Omics Data Integration Reveals Two Prognostic Subtypes in High-Risk Neuroblastoma	Shi Tieliu East China Normal University		
11:50-13:00	1:50-13:00 Lunch at Howard Johnson Qunsheng Plaza Guizhou			
13:10-13:30	Taking Group Picture & Posters Preparation			
	Session 2: Chaired by Yixue	Li		
13:35-14:10	Keynote Talk 3:	Gojobori Takashi		

	Digital Health, AI, and Biomedical Genomics	KAUST, Saudi Arabia & NIG	
14:10-14:30	Talk 5: Transcriptomic and proteomic approaches to identify regulatory NAC network of Arabidopsis leaf senescenceImage: SenescencePark Ji Hwan Korean Bioinformation 		
14:30-14:50	Talk 6: Increasing the human genome complexity at the RNA levels	Wang Zefeng Shanghai Institute of Nutrition and Health, CAS.	
14:50-15:10	Talk 7: Alternative splicing and human diseases	Lee Younghee University of Utah, Asia Campus	
15:10-15:30	Talk 8: A comprehensive reference transcriptome resource for the Iberian ribbed newt Pleurodeles waltl, an emerging model for developmental and regeneration biology	Masatoshi Matsunami University of the Ryukyus Medical School	
15:30-16:00	Tea Break		
	Session 3: Chaired by Lv Hu	i	
16:00-16:20	Talk 9: HUPAN: a human pan-genome analysis pipeline	Wei Chaochun Shanghai Jiao Tong University	
16:20-16:40	Talk 10: The genetic history of admixture across inner Eurasia	Jeong Choongwon Seoul National University	
16:40-17:00	Talk 11: History of Yaponesians inferred from ancient Yaponesian genome sequences	• Hideaki Kanzawa National Museum of Science and Nature	
17:00-17:20	Talk 12: Development of a prediction system for positively selected genes against massive orthologous groups	Sangmyung University	
17:20-17:40	Talk 13: Demographic history of the Ryukyu Islanders inferred from whole-genome sequencing	Kae Koganebuchi University of the Ryukyus Medical School	
18:40-20:10	Dinner at Wan Yi Panorama Hotel, 6 th Floor		
20:30-21:30	1:30 Discussion Meeting about next year's Bioinformatics Symposium (Invited lecturers), Azalea Hall (杜鹃厅)8 th Floor at Wan Yi Panorama Hotel		

-	y, August 23 rd , 2019 Center, Guizhou Normal University (Hua Xi C	Campus).
	Session 4: Chaired by Xie Lu	
08:55-09:30	Keynote Talk 4: Evolution of functional modules and regulatory networks: Understanding genotype-phenotype relationship	Kim Sanguk Pohang University of Science and Technology
09:30-09:50	Talk 14: Human physical traits and the phenome research	Wang Sijia Shanghai Institute of Nutrition and Health, CAS.
09:50-10:10	Talk 15: Functional relevance of first intron revealed by enrichment of epigenetic modification signals	Choi Sun Shim Kangwon National University
10:10-10:30	Talk 16: Genomic and epigenomic data analysis using probabilistic evolutionary learning	Rhee Je-Keun Pusan National University
10:30-10:50	Talk 17: Viruses in the planet	• Ituro Inoue National Institute of Genetics
10:50-11:10	Talk 18: The origin and evolution of Japanese cultivated radishes	Kosuke Hanada Kyushu Technology University
11:30-12:50	Lunch at Howard Johnson Qunsheng Plaza Gu	izhou
13:10-13:40	Poster Presentation 2	
	Session 4: Chaired by Wang Sij	ia
13:40-14:00	Talk 19: Introduction of Yaponesian Genome Project with special reference to modern human genome analyses	Saitou Naruya National Institute of Genetics
14:00-14:20	Talk 20: Initi-IO: An Integrative and Intelligentized Immuno-Oncology Study Platform	Liu Qi Tongji University
14:20-14:40	Talk 21: Identification of TB Drug Targets from NGS	Choi Inhee Institut Pasteur Korea
14:40-15:00	Talk 22: Bioinformatics for the diagnosis of infectious diseases	Tadashi Imanishi Okai University Medical School

15:00-15:20	Tea Break	
15:20-15:40	Talk 23: Fluxomics: Big data for synthetic biologyShen Tie Guizhou Normal Univer	
15:40-16:00	Talk 24: Chromosome-scale assemblies are coming back: DB updates and future plan	• Nakamura Yasukazu National Institute of Genetics
16:00-16:20	Talk 25: Gene expression and proteome landscape of Chinese HCC reveals prognostic determinants and therapeutic targets	Xie Lu Shanghai Center for Bioinformation Technology
16:20-16:30	Ending Remark	
18:00-19:40	Dinner at Wan Yi Panorama Hotel, 6 th Floor. (If the participants have their own arrangement and do not need to take dinner in the hotel, please inform us in advance)	

	day, August 24 th , 2019 Check-out orama Hotel.
07:00-12:00	Check-out at Wan Yi Panorama Hotel

Travelling Guidance



We will arrange the pick-up at the airport for participants.

If someone need to go to the hotel by himself or herself, travelling by taxi is highly recommended than any other ways.

A single trip from Guiyang Longdongbao International Airport to Wan Yi Panorama Hotel takes 50 minutes and costs around 75 RMB.

Please contact Ms. HE Qinwen if you need any help during the travelling.



Wan Yi Panorama Hotel is about 10 kilometers from the Conference Center in Guizhou Normal University (Hua Xi Campus).

A shuttle bus will take all the participants to the conference center on 22nd and 23rd morning and take everyone back to the hotel after the symposium.



This is a guidance map of Guizhou Normal University (Hua Xi Campus). The conference center is located in the blue circle.

Shuttle Bus		
Date	Departure Time	Start & Terminal
August 22nd	08:00	Wan Yi Panorama Hotel——Conference Center in
August 22nd	08.00	Guizhou Normal University (Hua Xi Campus)
August 22nd	st 22nd 18:00	Conference Center in Guizhou Normal University (Hua
August 22110		Xi Campus) ——Wan Yi Panorama Hotel
Assessed OD rol	08:10	Wan Yi Panorama Hotel——Conference Center in
August 23rd		Guizhou Normal University (Hua Xi Campus)
August 22rd	August 23rd 16:45	Conference Center in Guizhou Normal University (Hua
August 2310		Xi Campus) ——Wan Yi Panorama Hotel
August 22rd	ust 23rd 20:00	Wan Yi Panorama Hotel——Guiyang Longdongbao
August 23rd		International Airport

Warm Reminder

1. The hotel check-in time is 14:00 and check-out time is no later than 12:00.

2. We'll pay for room costs (breakfasts included) . Other expenses including mini bar, laundry, and call, etc. would be kindly paid by yourself.

3. The internet access is free in the hotel room. Each room has its own wifi which

username will the room number.

4. We recommend that all participants should take out the travel insurance to cover themselves against accident or illness.

Contact

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