## Preliminary Programme\*\* (as at 14 August 2019 - \*\*may be subject to change)

Date	Session	Location
Sunday 1 December		St David Lecture Theatre Complex (unless otherwise stated)
1.00 pm	Registration Opens	
2.00 pm – 5.00 pm	Street Art Tour (optional)	
5.00 pm – 5.30 pm	Official Welcome /Mihi whakatau/ Kapa Haka	
5.30 pm – 6.15 pm	Keynote: David Levens	
	Center for Cancer Research, National Cancer Research	
	Institute, Maryland, USA	
	MYC and the principles and consequences of global	
	transcription amplification	
6.15 pm – 7.15 pm	Welcome Function: Drinks and Canapes	Trade Exhibition Area
Monday 2 Docombor		
Monday 2 December 8.00 am – 8.30 am	Registration	
8.30 am – 8.45 am	Welcome/Housekeeping	
8.45 am – 9.30 am	Keynote: Susan Clark	
6.45 dili = 9.50 dili	Garvan Institute of Medical Research, Sydney, Australia	
	Reprogramming of the cancer genome and epigenome in 3D	
	DNA methylation	
9.35 am – 9.50 am	Aniruddha Chaterjee, University of Otago, Dunedin, NZ TBA	
9.50 am – 10.05 am	Amy Osborne, University of Canterbury, NZ	
3.00 0.11 20.00 0.11	DNA methylation differences in environmental exposures	
10.05 am – 10.30 am	Presentations selected from submitted abstracts	
10.30 am – 11.00 am	Morning Tea	Trade Exhibition Area
	Chromatin control of transcription I	
11.00 am – 11.15 am	Ping Gao, University of Science & Technology of China TBA	
11.15 am – 11.30 am	Tracy Hale, Massey University, NZ	
11.15 am – 11.30 am	Heterochromatin Protein 1a interacts with parallel RNA and	
	DNA G-quadruplexes	
11.30 am – 11.45 am	JHA Sudhakar, Cancer Science Institute of Singapore	
	TIP60 keeps the endogenous oncogenic demons repressed	
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Date	Session	Location
Monday 2 December (continued)		St David Lecture Theatre Complex (unless otherwise stated)
11.45 am – 12.00 pm	Helen Fitzsimons, Massey University, NZ Investigating nuclear and cytoplasmic roles of HDAC4 in the brain	
12.00 pm – 12.15 pm	Chen Davidovich, Monash University, Melbourne, Australia Context-dependent master-regulator: molecular basis for the regulation of the histone methyltransferase PRC2	
12.15 pm – 1.00 pm	Presentations selected from submitted abstracts	
1.00 pm – 2.00 pm	Lunch	Trade Exhibition Area
	RNA and transcription	
2.00 pm – 2.15 pm	Paul Gardner, University of Otago, NZ  Design principles for mRNA-based fine-tuning of protein expression	
2.15 pm – 2.30 pm	Sarah Diermeier, University of Otago, NZ Long non-coding RNAs as regulators of gene expression	
2.30 pm – 2.45 pm	Xiaoyuan Song, University of Science & Technology of China The function of IncRNAs in brain aging	
2.45 pm – 3.00 pm	Robert Weatheritt, Garvan Institute of Medical Research, Sydney, Australia TBA	
3.00 pm – 3.15 pm	Xiangting Wang, Institute Transcriptional Control of Carbonic Anhydrase9 by A Pair of Long Noncoding RNAs	
3.15 pm – 3.30 pm	Presentations selected from submitted abstracts	
3.30pm – 4.00 pm	Afternoon Tea	
	Genome organisation and chromosome structure	
4.00 pm – 4.15 pm	Inkyung Jung, Korea Advanced Institute of Science and Technology  3D genome organization in colon cancer	
4.15 pm – 4.30 pm	Marnie Blewitt, Walter and Eliza Hall Institute of Medical Research, Melbourne, Australia Unexpected contributors to the establishment of X Inactivation	
4.30 pm – 5.00 pm	Presentations selected from submitted abstracts	
5.00 pm – 7.00 pm	Poster Session: Drinks and Nibbles	Trade Exhibition/Poster Area

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Date	Session	Location
Tuesday 3 December		St David Lecture Theatre Complex (unless otherwise stated)
	Transcription and cell growth/proliferation and energy	
8.30 am – 9.15 am	Keynote: Leonie Quinn The John Curtin School of Medical Research, The Australian National University, Canberra To bind RNA or DNA? The multifaceted functions of DNA/RNA binding proteins in stem cell fate, development and disease	
9.15 am – 9.30 am	Kate Hannan, The Australian National University, TBA	
9.30 am – 9.45 am	Kate Quinlan, University of New South Wales, Sydney, Australia Transcriptional regulation of adipose tissue energy expenditure	
9:45 am -10:00 am	Jerry Workman, Stowers Institute for Research, Missouri, USA Nutritional stress signaling through chromatin	
10.00 am – 10.30 am	Morning Tea	Trade Exhibition Area
	Student/Early Career Awards Session	
10.30 am – 12.15 pm	Presentations selected from submitted abstracts	
12.15 pm – 12.30 pm	Judging and Awards	
12.30 pm – 1.30 pm	Lunch	Trade Exhibition Area
	Features of prokaryotic transcription	
1.30 pm – 1.45 pm	Ranjan Sen, Centre for DNA Fingerprinting and Diagnostics, India Design of bacterial transcription terminator-inhibitory peptides by directed evolution	
1.45 pm – 2.00 pm	Mayuree Fuangthong, Chulabhorn Research Institute, Bangkok, Thailand tRNA modification and oxidative stress response in Pseudomonas aeruginosa	
2.00 pm – 2.15 pm	Kalai Mathee, International Forensic Research Institute, Florida, USA TBA	
2.15 pm – 2.30 pm	Changwon Kang, Korea Advanced Institute of Science and Technology  RNA polymerase retention on DNA for transcription reinitiation after intrinsic but not rho-dependent termination	
2.30 pm – 2.45 pm	Jaehoon Kim, Korea Advanced Institute of Science and Technology Regulation of Heat Shock-Responsive Gene Transcription by RNF20/40-Mediated eEF1BdL Monoubiquitylation	

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Date	Session	Location
Tuesday 3 December (continued)		St David Lecture Theatre Complex (unless otherwise stated)
2.45 pm – 3.00 pm	Gang Wang, Shanghai Institutes for Biological Sciences, Chinese Academy of Sciences Transcriptional Mediator subunit MED23 regulates inflammatory responses and liver fibrosis	
3.00 pm – 3.30 pm	Afternoon Tea	Trade Exhibition Area
3.30 pm – 3.45 pm	Vincent Harley, Hudson Institute of Medical Research, Melbourne Australia TBA	
3.45 pm – 4.00 pm	Nisanart Charoenlap, Chulabhorn Research Institute, Bangkok, Thailand Modulation of antibiotic resistances by oxidative stress responsive transcriptional factor inStenotrophomonas maltophilia	
4.00 pm – 5.00 pm	Presentations selected from submitted abstracts	
5.00 pm – 6.00 pm	SOC/IOC Meeting: 2021 Conference	
7.00 pm - late	Conference Dinner	Toitu Early Settlers Museum
Wednesday 4 December		
Wednesday 4 December	Chromatin control of transcription II	
9.00 am – 10.15 am	<b>Keynote:</b> Ed Seto, The George Washington Cancer Center, Washington, USA  Mechanisms and functions of HATs and HDACs: transcription and beyond	
10.15 am – 10.45 am	Morning Tea	
10.45 am – 11.00 am	Huafeng Zhang, University of Science & Technology of China TBA	
11.00 am – 11.15 am	Valakunja Nagaraja, Indian Institute of Science TBA	
11.15 am – 12.30 pm	Presentations selected from submitted abstracts	
12.30 pm – 1.00 pm	Farewell	
1.00 pm – 2.00 pm	Lunch	Trade Exhibition Area
2.00 pm onwards	Optional Tours: Monarch Peninsula Tour Taieri Gorge Railway Olveston Anatomy Museum	

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