



2022 Accelerator Grant Awardees

	Project Title	Investigator(s) ¹	Institution(s) ²
01	Identifying rare variants in patients with epilepsy and ataxia using whole genome sequencing	Danielle Andrade (PI) Ryan Yuen Renato Munhoz Helio Teive	University Health Network The Hospital for Sick Children Univ Federal do Paraná, Brazil
02	Mapping genetic networks underlying autism spectrum disorder	Brenda Andrews (PI) Charles Boone Jason Moffat	University of Toronto
03	Developing a practical guide for conducting equitable genomics research with underrepresented and racialized populations	Yvonne Bombard (PI) Aisha Lofters Nav Persaud Melyssa Aronson Aaron Pollett Harriet Feilotter Jordan Lerner-Ellis	St. Michael's Hospital Women's College Hospital Mount Sinai Hospital Queen's University
04	Development of a novel approach for noninvasive prenatal CNV detection by establishing the human placental methylome in early pregnancy	Karen Chong (PI) Yiming Wang Rosanna Weksberg David Chitayat Elena Greenfeld Abdul Noor Trevor Pugh Hansen He Jack Brzezinski Sanaa Choufani Maian Roifman	Mount Sinai Hospital The Hospital for Sick Children University Health Network
05	Building capacity for genome sequencing in under- resourced settings: a global pilot project	Gregory Costain (PI) Ramses Badilla Nadirah Damseh Hanna Faghfoury Robin Hayeems Eriskay Liston Christian Marshall Roberto Mendoza-Londono	The Hospital for Sick Children Nat Children Hosp, Costa Rica Al-Makassed Hosp, Jerusalem University Health Network
06	Uncovering epigenetic regulators of genetic risk loci in osteoarthritis through deep phenotyping	Osvaldo Espin-Garcia (PI) Shabana Amanda Ali Andy Wong	University of Toronto Wayne State University University Health Network
	Personalized Medicine for Stroke Prevention: Implementing and Evaluating a Home-Delivery Pharmacogenetic Testing Service in Canada to Guide Antiplatelet Medication Selection for Patients with Ischemic Stroke and TIA	David Gladstone (PI) Mark Boulos Phavalan Rajendram Moni Lustig Aaron Goldman Andrew Demchuk Andrew Micieli	Sunnybrook Research Institute University of Toronto University of Calgary DNALabs Canada
	A chemical genomic map of therapeutic treatment of multi-drug resistant Staphylococcus aureus Whole genome-wide DNA methylation analyses of	Justin Nodwell (PI) Roman Melnyk Ivan Radovanovic (PI)	University of Toronto The Hospital for Sick Children University Health Network
09	arteriovenous malformations (AVMs): understanding the methylation profile of AVMs and its relation to both disease and behaviour		

		Whole genome sequencing of granulomatosis with	Katherine Siminovitch (PI)	Mount Sinai Hospital
1	10	polyangiitis (GPA) families to identify common and rare risk variants and provide more effective diagnostic and	Christian Pagnoux	
	10	risk variants and provide more effective diagnostic and		
		therapeutic strategies		
1		The utilization of cell-free DNA methylation signatures	Gelareh Zadeh (PI)	University Health Network
	11	in cerebrospinal fluid for diagnosis and early detection		
		of progression in glioblastomas and brain metastases		
1		Using Long-read Whole-genome Sequencing to Understand Treatment Resistance in Psychiatric Patients	Clement Zai (PI)	Centre for Addiction and Mental
			Arun Tiwari	Health
			Gwyneth Zai	University of California, San
			Joyce So	Francisco
			Jerry Warsh	
			James Kennedy	

¹ Bold indicates Lead or Co-Lead Principal Investigator (PI)
² Bold indicates Lead Institution