



McGill Applied Genomics Innovation Core (MAGIC)

Partners for Advancing Genomics Research



Cores and Supported Applications

CORES	ACTIVITY	TECHNOLOGY / ACTIVITY
Production Genomics	Whole Genome Sequencing Exome / Targeted Capture Sequencing Whole Genome Bisulfite Sequencing (WGBS) RNA-Seq / miRNA-Seq ChIP-Seq, ChIPmentation, ATAC-Seq Expression Arrays / Genotyping	Illumina NovaSeq Illumina NovaSeq Illumina NovaSeq Illumina NovaSeq Illumina NovaSeq Illumina iScan / Affymetrix GeneTitan
Advanced Genomics Technologies	10X Genomics Single-cell applications Long-reads sequencing SmartSeq2 Single-cell RNAseq Protein expression Hi-C	10X Genomics / Illumina NovaSeq / Complete Genomics Oxford Nanopore / PacBio Illumina NovaSeq 10X Genomics Visium Illumina NovaSeq
Biomarker	Spatial genomics Immunoassay-based detection of biomarkers High-throughput aptamer-based proteomics LC/MS (Liquid Chromatograph-Mass Spectrometry) GC/MS (Gas Chromatograph-Mass Spectrometry)	Nanostring GeoMx Olink Proteomics SomaLogic Shimadzu Shimadzu
Microbial Genomics	Amplicon sequencing and metagenomics	Illumina MiSeq
Canadian Centre for Computational Genomics	Bioinformatics support	Technology Development and Analysis

Contact us: pm.genome@mcgill.ca

