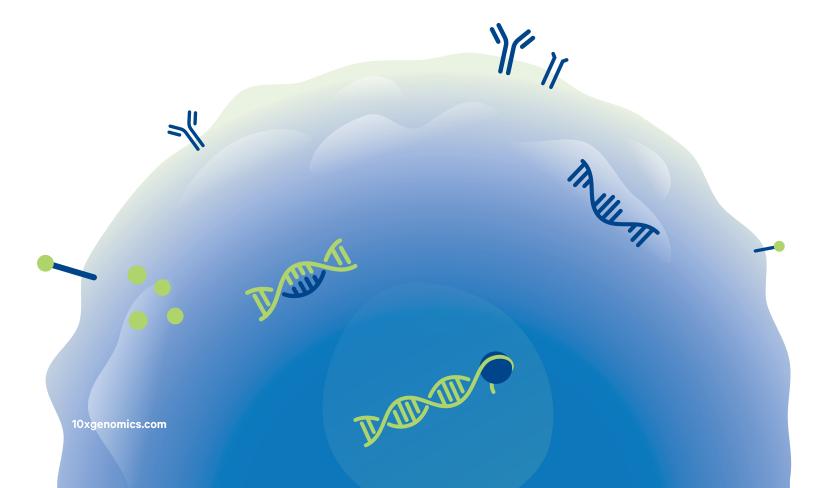


Single Cell Buyer's Guide

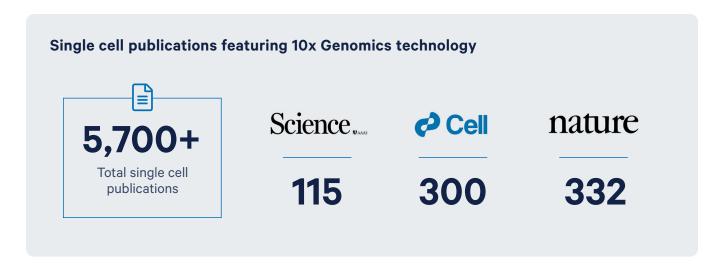
Gene Expression
Immune Profiling
Epigenomic Profiling
Protein Expression



Introduction

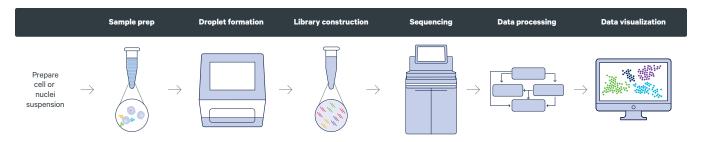
Why single cell sequencing?

Cell heterogeneity is a key contributor to biological complexity that is often masked by bulk techniques, like RNA sequencing (RNA-seq) or microarray analysis. Rather than providing an average snapshot of how cells work, single cell sequencing technology gives researchers the ability to more fully characterize tissue heterogeneity, identify rare cell types, and dissect molecular mechanisms cell by cell. This high level of resolution leads to groundbreaking insights, as evidenced by over 6,000 publications featuring 10x Genomics technology, the majority of which use single cell approaches.



How does it work?

Single cell sequencing starts with a suspension of single cells or nuclei. Droplets containing a single cell or nucleus are generated using advanced microfluidics with a Chromium instrument. Library construction follows a streamlined workflow. After sequencing, data can be processed and analyzed using our integrated data analysis pipelines and visualization software—intuitive enough for novice users yet powerful enough for the experts. Support resources provide guidance for the entire workflow, from sample preparation to data analysis.



Tackle your single cell experiments confidently with personalized support and detailed guidance across an end-to-end workflow.

Single cell product offerings

10x Genomics provides robust solutions that let you analyze gene and protein expression, perform comprehensive immune phenotyping, and map regions of open chromatin, all at single cell resolution.

Gene Expression ←

Perform molecular and cellular characterization of cells at scale with single cell RNA-sequencing for whole transcriptome gene expression.

Immune Profiling •-

Gain a comprehensive view of the immune system with paired, full-length receptor sequences from T and/or B cells, surface protein expression, antigen specificity, and gene expression, all from a single cell.

Epigenomic Profiling •

Uncover hidden insights with single cell epigenomic profiling using the assay for transposase-accessible chromatin, ATAC-seq, alone or in combination with single cell RNA-seq to simultaneously measure chromatin accessibility and gene expression.

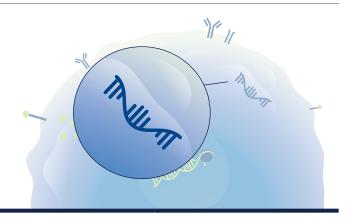
Protein Expression •

Reveal cell phenotypes and uncover functional information at single cell resolution with simultaneous measurement of hundreds of cell surface proteins and gene expression.



Single Cell Gene Expression

Characterize cellular heterogeneity, explore mechanisms driving development and disease, and trace cell lineages with 3'-, 5'-, or probe-based transcriptome sequencing at single cell resolution for hundreds to hundreds of thousands of cells.



At-a-glance comparison

	Single Cell Gene Expression Flex	Single Cell Gene Expression	Single Cell Immune Profiling
Description	A probe-based solution for whole transcriptome gene expression, starting from fresh or fixed samples, including FFPE	A versatile solution providing 3' gene expression with multiomic capabilities	Enhanced immune profiling combining 5' gene expression with multiomic profiling
Additional multiomic capabilities	Cell Surface Protein	Cell Surface Protein CRISPR Screening	Cell Surface Protein Antigen Specificity
		Citis it screening	BCR/TCR sequencing CRISPR Screening
Extensions	Built-in sample mulitplexing	Sample multiplexing Automated workflow	Automated workflow
Throughput	Singleplex: 10,000 cells per channel; up to 80,000 cells per chip	Standard: 10,000 cells per channel; up to 80,000 cells per chip; 140,000 cells per chip with 3' CellPlex High: 20,000 cells per channel; up to 320,000 cells per chip; up to 730,000 cells per chip with 3' Cellplex	Standard: 10,000 cells per channel; up to 80,000 cells per chip
	Multiplex: 128,000 cells per channel; up to 1,024,000 cells per chip		High: 20,000 cells per channel; up to 320,000 cells per chip
Sample compatibility	Cells or nuclei	Cells or nuclei	Cells
	Tested on diverse sample types, including cell lines, primary cells, and dissociated fresh or fixed tissue, including FFPE	Tested on diverse sample types, including cell lines, primary cells, and dissociated fresh tissue	Tested on diverse sample types, including cell lines, primary cells, and dissociated fresh tissue
Species compatibility	Human and mouse	Demonstrated on multiple species, including human, mouse, and rat	Demonstrated on multiple species, including human and mouse
			Amplification of human or mouse TCR/BCR sequences
Instrument compatibility	Chromium X/iX	Chromium X/iX Chromium Connect	Chromium X/iX Chromium Connect
Analysis software	Cell Ranger	Cell Ranger	Cell Ranger
	Cloud Analysis	Cloud Analysis	Cloud Analysis
Visualization software	Loupe Browser	Loupe Browser	Loupe Browser Loupe V(D)J Browser
Resources	Product sheet	Product sheet	Product sheet
	Kits and part numbers	Kits and part numbers	Kits and part numbers

Choosing the right gene expression assay for your experiment

When should I use the Single Cell Gene Expression Flex assay?

We recommend this for everyone running single cell experiments with human or mouse samples who is looking to measure gene expression alone or simultaneously with cell surface protein expression.

Single Cell Gene Expression Flex is our most sensitive, cost-effective, and flexible assay. Key benefits include:

- Streamlined processing provides compatibility for fresh or PFA-fixed tissues, cells, or nuclei, plus access to FFPE samples
- Built-in multiplexing capabilities provide the flexibility to run 1–128 samples or up to 1 million cells in a single run
- Highly sensitive probe-based chemistry combined with sample multiplexing provides our lowest cost per sample

When should I use other 10x gene expression assays?

Single Cell Gene Expression ←

- 3' gene expression in non-human or non-mouse samples
- Automated workflows on the Chromium Connect instrument

Single Cell Immune Profiling .___

- High-resolution immune cell characterization: 5' gene expression & full length VDJ sequences
- CRISPR screening: 5' gene expression & CRISPR perturbations
- 5' gene expression in non-human or non-mouse samples
- Automated workflows on the Chromium Connect instrument

Single Cell Multiome Gene Expression + ATAC ←

• Epigenomics applications: 3' gene expression and chromatin accessibility in the same single cell



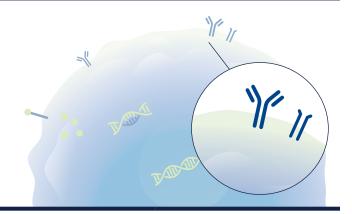






Single Cell Immune Profiling

Reveal the full diversity of the immune system to understand clonal expansion, accelerate immunotherapy, and investigate disease states.

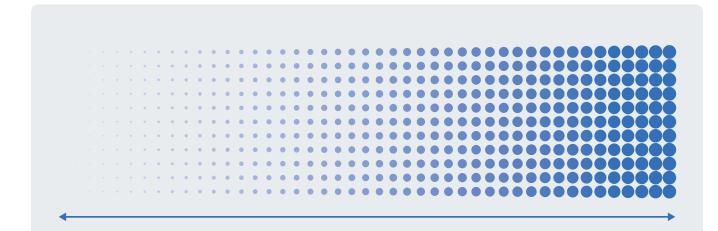


At-a-glance comparison

	Immune Receptor Profiling	Barcode Enabled Antigen Mapping (BEAM)	Single Cell 5' Gene Expression
Description	Perform BCR/TCR sequencing to obtain paired, full-length receptor sequences from T cells and/or B cells with complete isotype resolution	Perform multiplexed antigen screening to obtain comprehensive cellular profiles of antigen-specific B-cell (BEAM-Ab) and T-cell (BEAM-T) clonoypes	Study innate and adaptive immunity by combining single cell 5' gene expression with multiomic profiling
Additional multiomic	Antigen Specificity	BCR/TCR sequencing	Antigen Specificity
capabilities	Gene Expression	Gene Expression	BCR/TCR sequencing
	Cell Surface Protein	Cell Surface Protein	Cell Surface Protein
	CRISPR Screening		CRISPR Screening
Extensions	Automated workflow	-	Automated workflow
Throughput	Standard or high	Standard or high	Standard or high
Sample compatibility	Cells	Cells	Cells or nuclei
	Tested on diverse sample types, including cell lines, primary cells, and dissociated fresh tissue Amplification of human or mouse TCR/	Tested on diverse sample types, including PBMCs, splenocytes, lymph node aspirates, and enriched B or T cells	Tested on diverse sample types, including cell lines, primary cells, and dissociated fresh tissue Demonstrated on multiple species,
	BCR sequences	Amplification of human or mouse TCR/ BCR sequences	including human and mouse
	Demonstrated on multiple species, including human and mouse	Compatible with human and mouse	
Analysis software	Cell Ranger	Cell Ranger	Cell Ranger
	Cloud Analysis		Cloud Analysis
Visualization software	Loupe Browser	Loupe Browser	Loupe Browser
	Loupe V(D)J Browser	Loupe V(D)J Browser	Loupe V(D)J Browser
Resources	Product sheet	Product sheet	Product sheet
	Kits and part numbers	Kits and part numbers	Kits and part numbers

Adapt the scale you need for immune cell profiling

From clonotyping studies to large-scale antibody and TCR discovery projects, Single Cell Immune Profiling lets you scale up easily. Select the kit and instrument that enable your projects, from discovery to translational research.



Standard scale

- Process 500–10,000 cells per sample; up to 8 samples per chip
- Explore adaptive and innate immune cell diversity
- Identify and characterize rare cell types and biomarkers
- Analyze tissue microenvironments, disease progression, and drug immune response
- Characterize immune response to infection by measuring clonal expansion and immune cell phenotypes

Kit: Chromium Single Cell Immune Profiling

Instrument: Chromium X, Chromium iX, Chromium Connect

Large scale

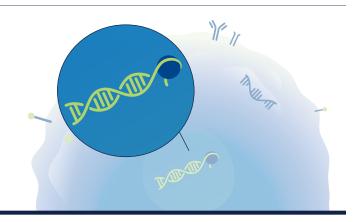
- Process 2,000–20,000 cells per sample; up to 16 samples per chip
- Perform large-scale antibody and TCR discovery for novel antigens
- Detect ultra-rare cell populations with increased throughput and multiomic readouts
- Perform large-scale time course experiments for clonal tracking
- Achieve economy of scale for immune repertoire profiling with simultaneous capture of gene expression and cell surface protein

Kit: Chromium Single Cell Immune Profiling HT

Instrument: Chromium X

Single Cell Epigenomic Profiling

Unmask epigenomic profiles, explore gene regulatory interactions, and resolve cell lineage relationships to transform your understanding of biology and uncover hidden insights.

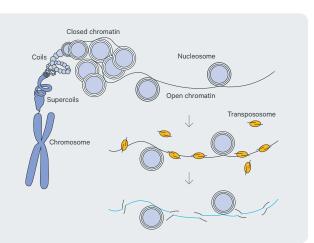


At-a-glance comparison

	Single Cell Multiome ATAC + Gene Expression	Single Cell ATAC
Description	Simultaneously profile single cell 3' gene expression and open chromatin from the same nucleus	Profile chromatin accessibility with single cell resolution
Chromatin accessibility profiling	Yes	Yes
Transcriptional profiling	Yes	No
Throughput	Standard	Standard
Sample & species compatibility	Nuclei	Nuclei
	Demonstrated with cell lines, primary cells, cryopreserved samples, and fresh- and	Demonstrated with cell lines, primary cells, cryopreserved samples, and fresh- and flash-frozen tissue
	flash-frozen tissue	Demonstrated on multiple species, including human and
	Demonstrated on multiple species, including human and mouse	mouse
Analysis software	Cell Ranger ARC	Cell Ranger ATAC
	Cloud Analysis	
Visualization software	Loupe Browser	Loupe Browser
Resources	Product sheet	Product sheet
	Kits and part numbers	Kits and part numbers

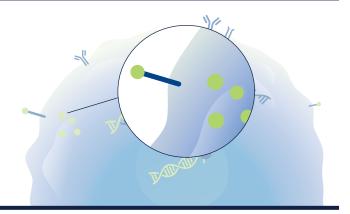
Why ATAC-seq?

The assay for transposase-accessible chromatin (ATAC) is a sequencing method that surveys the physical structure of the genome by identifying regions of open chromatin, where regulatory sequences are accessible to DNA-binding proteins. Single cell ATAC-seq can provide information about gene regulatory elements, including their cell-type specificity and binding site motifs.



Single Cell Protein Expression

Comprehensively resolve cell types and phenotypes, validate your single cell RNA-seq, and unravel cell signaling pathways to explore molecular mechanisms or transform drug discovery.



At-a-glance comparison

	Single Cell Gene Expression Flex	Single Cell Gene Expression	Single Cell Immune Profiling
Description	A probe-based solution for whole transcriptome gene expression and cell surface protein expression	A flexible multiomic solution providing single cell 3' gene expression and cell surface protein profiling	Enhanced immune profiling combining single cell 5' gene expression with cell surface protein profiling
Additional	_	_	Antigen Specificity
multiomic capabilities			BCR/TCR sequencing
Extensions	Built-in sample multiplexing	Sample multiplexing	-
Throughput	Singleplex: 10,000 cells per channel; up to 80,000 cells per chip	Standard: 10,000 cells per channel; up to 80,000 cells per chip; 140,000 cells	Standard: 10,000 cells per channel; up to 80,000 cells per chip
	Multiplex: 128,000 cells per channel;	per chip with 3' CellPlex	High: 20,000 cells per channel; up to
	up to 1,024,000 cells per chip	High: 20,000 cells per channel; up to 320,000 cells per chip; up to 730,000 cells per chip with 3' Cellplex	320,000 cells per chip
Sample compatibility	Cells	Cells	Cells
	Tested on diverse sample types, including cell lines, primary cells, and dissociated fresh or PFA-fixed tissue	Tested on diverse sample types, including cell lines, primary cells, and dissociated fresh tissue	Tested on diverse sample types, including cell lines, primary cells, and dissociated fresh tissue
Species compatibility	Human and mouse	Demonstrated on multiple species, including human and mouse	Demonstrated on multiple species, including human and mouse Amplification of human or mouse TCR/
			BCR sequences
Instrument compatibility	Chromium X/iX	Chromium X/iX	Chromium X/iX
		Chromium Connect	Chromium Connect
Analysis software	Cell Ranger	Cell Ranger	Cell Ranger
	Cloud Analysis	Cloud Analysis	Cloud Analysis
Visualization software	Loupe Browser	Loupe Browser	Loupe Browser
			Loupe V(D)J Browser
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Instrumentation that works for you

Key to our single cell sequencing technology is the ability to generate tens of thousands of single cell partitions, each containing an identifying barcode for downstream analysis. Each of the Chromium instruments uses advanced microfluidics to perform single cell partitioning and barcoding in a matter of minutes.

Next-generation technology for ultimate single cell flexibility

The Chromium X Series is designed with modern functionality plus enhanced hardware technology to offer the most optimized approach to single cell studies, with the option to match the scale of single cell research at any level and enable routine high-throughput experiments. All of our single cell assays are supported on Chromium X Series instruments.

Learn more about the Chromium X Series





Automate your single cell workflows

Chromium Connect provides automation of the full single cell gene expression and immune profiling workflows, including cell partitioning and library construction, letting you go from single cell suspensions to up to eight sequencing-ready libraries with walk-away convenience.

Learn more about Chromium Connect

Resources from 10x Genomics

We are dedicated to helping you get the most out of your 10x Genomics system by offering multiple helpful resources:

Technology brochure

Discover the power of single cell partitioning and learn how Next GEM technology enables integrated analysis of single cells at massive scale.

Learn more →

Support

Visit the support site for documentation, software, and datasets that will help you get the most out of your 10x Genomics products.

Learn more →

10x Genomics library

Learn more about what you can do with our single cell solutions from our collection of product literature, application notes, and scientific posters.

Learn more \longrightarrow

10x Genomics publications

Get inspired by more than 4,500 peer-reviewed publications and search papers by product, research area, or sample type.

Learn more \longrightarrow

10x Genomics compatible products

Access our list of key partner products that have been certified compatible to work with our various solutions.

Learn more \longrightarrow

10x Blog

Keep up to date with the 10x Genomics Blog, where you'll find everything from tips and tricks to the latest 10x news.

Learn more →

