



Asteria Benchtop Kit for Single-cell RNA-seq & Cytonaut Cloud Software for Single-cell Data Analysis

Gain control of your experiments right now without the need for microfluidics equipment

Save the hassles and expense of a major equipment purchase and start your single-cell experiment today

Easily expand your skill set and lab's workflow with our simple kit and intuitive software

Allocate your time to experiments, not on purchasing microfluidics equipment or waiting for a slot to open in the Core Lab

sensitive hydrogel.

Gain confidence in your results with biological and technical replicates of up to 10,000 cells

Get your answers 24/7/365 using our cloudbased software and bypass the need for your own high performance computing infrastructure

Take control of single-cell data exploration with intuitive and powerful software tools

Iterate your experiments to find your answers and publish faster

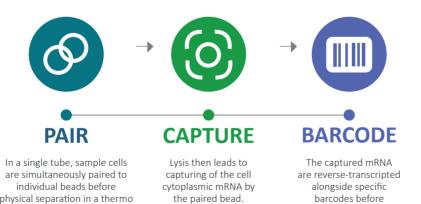
amplification.

Advanced Single-cell Analysis in a Straightforward Benchtop Kit

o Profile up to four samples of 10,000 cells with each kit

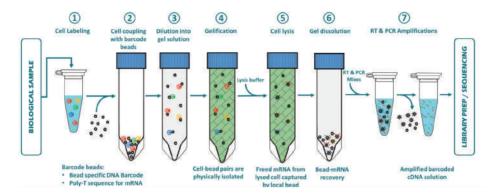
- o Use the simple, benchtop protocol for direct single-cell barcoding
- o Only basic laboratory skills and equipment are needed, no need for microfluidics instruments
- o As early as two hours, safely stop and store samples and then restart your experiment later
- o Experience high mRNA capture and gene sensitivity with low multiplet rates

Scipio's Novel Approach to Single-cell cDNA Preparation



At the completion of the benchtop protocol, the individually barcoded cDNA from up to 10,000 cells is ready for library preparation and next generation sequencing.

The Elegance of the Asteria Single-cell RNA-seq Workflow Take advantage of the simple benchtop protocol powered by RevGel-seq[™] Technology



The entire workflow can be done on the workbench with basic laboratory equipment

Single-cell Data Insights at your Fingertips with Cytonaut Cloud Software *Powerful for a bioinformation & intuitive for a biologist*

Rely on our intuitive software with the stepby-step guided experience and demo data designed for a fast learning curve for biologists new to single-cell data analysis

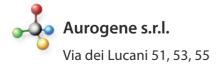
Assess the robustness of your data by checking dedicated quality indicators related to sequencing, sample preparation, and information retrieval

Run and analyze anytime with 24-hour online access to secure Cloud-based software to identify clusters and sub-clusters of cell subtypes and analyze differential gene expression Keep your data and results safe thanks to substantial computing infrastructure engineered on Trusted Cloud Principles

Dive deep into the rich data with our sophisticated software or use the interoperable data export feature for analysis with open source tools

Easily obtain publication-ready datasets and striking visuals thanks to a powerful interactive visualization functionality with high parametrization

Identify high-fidelity cell subtypes and perform accurate transcript quantification



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