# **Consumables conceived by biologists for biologists**

# **Flexomics Workflow:**



# 8 sites / 8-sample consumables

Consumable Dimensions: 25x75mm – compatible with standard microscope slide holder

Number of wells: per site / sample: 77,784

Well layout: 208 subsections of  $\sim$ 16x24 layouts mimicking 384 well-plate format with individual ID fiducial for easy site/well navigation Well geometry: circles and hexagons available – 30µm diameter and 35µm depth

Spacing between sites: 9mm - compatible with multi-channel pipettes and automated liquid handlers

Well mapping: well pre-loaded with optically encoded barcoded capture beads (~1 million barcodes available)



#### **Cellular assay**

Live cell enabled: gas permeable for easy incubation in standard CO<sub>2</sub> incubators Type of assays: immunofluorescence-based assays such as antibody staining, secretion assays, proliferation assays

Assay working volume: 100µl recommended (as low as 50µl), 500µl reagent reservoirs

## Imaging

Microscope compatibility: compatible with most inverted fluorescent microscopes

(microscopes tested include: Nikon Ti2-E, Evos FL Auto 2, Leica DMi8, Keyence BZ-X700, Zaber Nucleus) Magnification: 2x-40x compatible

**Brighfield / Fluorescence**: brightfield and standard fluorescent channel enabled (e.g. DAPI, CFP, FITC, YFP, RFP/Cy3, mCherry, Cy5, Cy7)

## Sequencing

Site/Sample "2"

**scRNA-seq**: 3'end RNA-seq – standard 2x100 short-read sequencing, hybrid-capture compatible for targeted

RNA-seq

**Isoform analysis**: compatible with ONT and PacBio sequencing

# Analysis

**Data processing**: Imaging and sequencing data processed at single-well resolution

**Visualization**: Browser-based FLXplorer analysis platform for visualization and integration of both imaging and sequencing data at single-cell level

(filtering, sequencing and/or fluorescence-based clustering, differential gene expression, per well sequencing and imaging mapping)





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