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## Massively parallel digital transcriptional profiling of single cells

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2250	Perturb-Seq: Dissecting Molecular Circuits with Scalable Single-Cell RNA Profiling of Pooled Genetic Screens. <b>2016</b> , 167, 1853-1866.e17		675
2249	Pooled CRISPR screening with single-cell transcriptome readout. <b>2017</b> , 14, 297-301		435
2248	Single cell sequencing: a distinct new field. <b>2017</b> , 6, 10		42
2247	From single-molecule detection to next-generation sequencing: microfluidic droplets for high-throughput nucleic acid analysis. <b>2017</b> , 21, 58		34
2246	Visualization and analysis of single-cell RNA-seq data by kernel-based similarity learning. <b>2017</b> , 14, 414-416		355
2245	Power analysis of single-cell RNA-sequencing experiments. <b>2017</b> , 14, 381-387		357
2244	Taking inventory of metastasis effectors. <b>2017</b> , 23, 275-276		
2243	Single-cell RNA-seq reveals new types of human blood dendritic cells, monocytes, and progenitors. <b>2017</b> , 356,		1176
2242	High-throughput Characterization of HIV-1 Reservoir Reactivation Using a Single-Cell-in-Droplet PCR Assay. <b>2017</b> , 20, 217-229		39
2241	Non-equivalence of Wnt and R-spondin ligands during Lgr5 intestinal stem-cell self-renewal. <b>2017</b> , 545, 238-242		209
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2239	Microfluidics as a Strategic Player to Decipher Single-Cell Omics?. <b>2017</b> , 35, 713-727		19
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