## Supplementary Figures



Supplementary Figure 1. Analysis of cell-identifying barcodes. a) Expression profiles for all detected cell-identifying barcodes were downsampled to between 20 and 500 unique molecules in regular increments ( y -axis of heatmap) and the distribution of number of molecules per read (x-axis) is
histogrammed across the heatmap. The series of histograms reveals two clear populations of barcodes. The barcodes in the population on the left shift continuously to the left and are associated with relatively few molecules per read compared to the population on the far right which have closer to one molecule per read. We take this small subpopulation of high-coverage barcodes to indicate the actual single cell RNA samples captured in our device. b) A single histogram from the heatmap in a) where the profiles have been downsampled to 100 reads per barcode. The high-coverage subpopulation is highlighted in red. c) Comparison between the number of unique cell-identifying barcodes predicted from the high-coverage subpopulation in a) and b) and the number of unique cell-identifying barcodes expected from imaging bead-cell pairs in the microwells in our device by fluorescence microscopy (Fig. 4c). Note that the number of cell-identifying barcodes in each lane was determined based on a single threshold for the whole data set with barcodes from all lanes pooled together (we didn't need to choose a different threshold for each lane to get these results).


Supplementary Figure 2. Heatmap display of downsampling analysis in Supplementary Fig. 1a on a lane-by-lane and barcode-by-barcode basis where the downsampled distributions are shown for all 960 cell-identifying barcodes in each lane after ordering the barcodes by total coverage. Green tic-marks indicate the predicted number of cell-identifying barcodes based on imaging cell-bead pairs in the device.


Supplementary Figure 3. Fluorescence image of a glass coverslip after single cell RNA capture, reverse transcription, and Sytox Orange staining using the scheme in Fig. 1a with an improperly or partially sealed microwell array demonstrating the rapid escape and uniform distribution of RNA by diffusion from individual cells into every printing site.


Supplementary Figure 4. Cell type separation from single cell RNA-Seq Experiment 2. a) t-SNE clustering of 247 single cell profiles based on differentially expressed genes color-coated by the lane-oforigin of each profile. Two clear spatial clusters form and each is exclusively associated with a specific cell type-exclusive lane. b) The same t-SNE clustering shown in a) but color-coated with a score indicating expression of the U87-specific genes vs. the WI-38-specific genes. The score is based on the relative rank-ordering of WI-38 and U87-specific genes in each cell (see Methods).

## Supplementary Tables

Table 1. List of key oligonucleotides used for barcoding and library preparation (not including the long list of cell-identifying barcodes that appear in the subsequent tables) for Experiment 1.

| Oligonucleotide Name | Oligonucleotide Sequence |
| :--- | :--- |
| Bead Capture Oligo (5'-dual biotinylated) | AGGTAAGGTAATACGACTCACTATAGGGGTTCAGAGT <br> TCTACAGTCCGACGATC |
| RT1 (Reverse Transcription Primer for Lane 1) | GCCTTGGCACCCGAGAATTCCANNNNNNNNCGTGATN <br> NNNNN |
| RT2 (Reverse Transcription Primer for Lane 2) | GCCTTGGCACCCGAGAATTCCANNNNNNNNACATCGN <br> NNNNN |
| RT3 (Reverse Transcription Primer for Lane 3) | GCCTTGGCACCCGAGAATTCCANNNNNNNNGCCTAAN <br> NNNNN |
| RT4 (Reverse Transcription Primer for Lane 4) | GCCTTGGCACCCGAGAATTCCANNNNNNNNTGGTCAN <br> NNNNN |
| RT5 (Reverse Transcription Primer for Lane 5) | GCCTTGGCACCCGAGAATTCCANNNNNNNNCACTGTN <br> NNNNN |
| RP1 (PCR Primer 1) | AATGATACGGCGACCACCGAGATCTACACGTTCAGAG <br> TTCTACAGTCCGA |
| RPI1 (PCR Primer 2) | CAAGCAGAAGACGGCATACGAGATCGTGATGTGACTG <br> GAGTTCCTTGGCACCCGAGAATTCCA |

Table 2. Oligonucleotide sequences used to generate the first set of barcoded beads (FBC) for combinatorial synthesis in Experiment 1.

| Oligonucleotide Name | Oligonucleotide Sequence |
| :--- | :--- |
| FBC_Oligo1 | CAGGTCAACCAGAGAGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo2 | CAGGTCAAAGTACGCGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo3 | CAGGTCGTTTGGCATGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo4 | CAGGTCAAGTGAGGTGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo5 | CAGGTCACGTTAGCTGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo6 | CAGGTCGTGCTAGAGGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo7 | CAGGTCGTCCTGTGTGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo8 | CAGGTCTCTACGGCAGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo9 | CAGGTCACAGGGCTTGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo10 | CAGGTCGTGCGTTATGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo11 | CAGGTCGGGTAAGTAGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo12 | CAGGTCTCCCTTAGGGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo13 | CAGGTCTTCTCACTCGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo14 | CAGGTCTCCCACTCTGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo15 | CAGGTCCGGTATACCGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo16 | CAGGTCAGGCATGTGGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo17 | CAGGTCCCCAGATTGGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo18 | CAGGTCTTCCCTTGAGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo19 | CAGGTCGTTGTACGAGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo20 | CAGGTCTGCTTGCAGGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo21 | CAGGTCGGCCTCATTGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo22 | CAGGTCAACAGCCTAGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo23 |  |


| FBC_Oligo24 | CAGGTCGATGCAATGGATCGTCGGACTGTAGAACTCTGAAC |
| :---: | :---: |
| FBC_Oligo25 | CAGGTCGAAGGAACGGATCGTCGGACTGTAGAACTCTGAAC |
| FBC Oligo26 | CAGGTCCAGCCACTTGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo27 | CAGGTCCTCTGCTTCGATCGTCGGACTGTAGAACTCTGAAC |
| FBC Oligo28 | CAGGTCGGCTTATGAGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo29 | CAGGTCCTAGTCCTCGATCGTCGGACTGTAGAACTCTGAAC |
| FBC Oligo30 | CAGGTCCTAGAGGAGGATCGTCGGACTGTAGAACTCTGAAC |
| FBC Oligo31 | CAGGTCAGCTTTACCGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo32 | CAGGTCGTCCATGAAGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo33 | CAGGTCCTCGAACCTGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo34 | CAGGTCCATTGTACGGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo35 | CAGGTCTTGAACGCTGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo36 | CAGGTCTACGTCATGGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo37 | CAGGTCAAGCCGTTAGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo38 | CAGGTCCGGACGTATGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo39 | CAGGTCTCGTTACCGGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo40 | CAGGTCATCCCCCATGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo41 | CAGGTCCAGACGATTGATCGTCGGACTGTAGAACTCTGAAC |
| FBC Oligo42 | CAGGTCATCGATCCCGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo43 | CAGGTCCCTGAGGATGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo44 | CAGGTCAGCTCTTTGGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo45 | CAGGTCGGAATACGGGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo46 | CAGGTCCTATCCTGGGATCGTCGGACTGTAGAACTCTGAAC |
| FBC Oligo47 | CAGGTCGGTTGTAGTGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo48 | CAGGTCGAACGTAGCGATCGTCGGACTGTAGAACTCTGAAC |
| FBC Oligo49 | CAGGTCGTCTATCGGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo50 | CAGGTCTACGAGTGGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo51 | CAGGTCTCATGTCGGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo52 | CAGGTCAAACACCCGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo53 | CAGGTCACTAGTCCGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo54 | CAGGTCCGAGGAATGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo55 | CAGGTCACAATGGCGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo56 | CAGGTCTAGGTCTCGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo57 | CAGGTCTCTGTGAGGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo58 | CAGGTCGGGATTGAGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo59 | CAGGTCAACTCTGGGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo60 | CAGGTCAAACGCGTGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo61 | CAGGTCTCCTACGAGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo62 | CAGGTCTAGCAGGTGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo63 | CAGGTCCCTGCATTGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo64 | CAGGTCGTGATGCAGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo65 | CAGGTCCGATTCAGGATCGTCGGACTGTAGAACTCTGAAC |
| FBC Oligo66 | CAGGTCAGGATGACGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo67 | CAGGTCAGGCCATAGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo68 | CAGGTCGCTTGCTTGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo69 | CAGGTCTCCCAAGTGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo70 | CAGGTCTCAAGGCAGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo71 | CAGGTCACGAGGTAGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo72 | CAGGTCGGAACGAAGATCGTCGGACTGTAGAACTCTGAAC |


| FBC_Oligo73 | CAGGTCAATCCCAGGATCGTCGGACTGTAGAACTCTGAAC |
| :--- | :--- |
| FBC_Oligo74 | CAGGTCCGATAAGGGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo75 | CAGGTCTATCGCGAGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo76 | CAGGTCCGCATAACGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo77 | CAGGTCGTGCAGTTGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo78 | CAGGTCAGAACGCTGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo79 | CAGGTCTAGAGGTCGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo80 | CAGGTCCTGTGATGGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo81 | CAGGTCTAGAGCCAGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo82 | CAGGTCCTTGATGCGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo83 | CAGGTCTTCGTGTCGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo84 | CAGGTCTATCTGCGGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo85 | CAGGTCTGGTAGGAGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo86 | CAGGTCCCTAGACAGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo87 | CAGGTCAGTCAACGGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo88 | CAGGTCAAGGGTGAGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo89 | CAGGTCCTTCACACGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo90 | CAGGTCAGGTTGCTGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo91 | CAGGTCACCCGAAAGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo92 | CAGGTCGAAAAGGGGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo93 | CAGGTCACTTCCCAGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo94 | CAGGTCTGCTGCATGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo95 | CAGGTCATTCCTGGGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo96 | CAGGTCCAGAACTCGATCGTCGGACTGTAGAACTCTGAAC |

Table 3. Oligonucleotide sequences used to generate the second set of barcoded beads (SBC) for combinatorial synthesis in Experiment 1.

| Oligonucleotide Name | Oligonucleotide Sequence |
| :--- | :--- |
| SBC_Oligo1 | AAAAAAAAAAAAAAAAAAAAAAAAAGGTGATACAGGTCAAAAAAAAAGATCG <br> TCGGACTGTAGAACTC |
| SBC_Oligo2 | AAAAAAAAAAAAAAAAAAAAAAAAATGAATGCCAGGTCAAAAAAAAAGATCG <br> TCGGACTGTAGAACTC |
| SBC_Oligo3 | AAAAAAAAAAAAAAAAAAAAAAAAATGCCAAACAGGTCAAAAAAAAAGATCG <br> TCGGACTGTAGAACTC |
| SBC_Oligo4 | AAAAAAAAAAAAAAAAAAAAAAAAAACAGAAGCAGGTCAAAAAAAAAGATCG <br> TCGGACTGTAGAACTC |
| SBC_Oligo5 | AAAAAAAAAAAAAAAAAAAAAAAAACACTGGACAGGTCAAAAAAAAAGATCG <br> TCGGACTGTAGAACTC |
| SBC_Oligo6 | AAAAAAAAAAAAAAAAAAAAAAAAACGATGATCAGGTCAAAAAAAAAGATCG <br> TCGGACTGTAGAACTC |
| SBC_Oligo7 | AAAAAAAAAAAAAAAAAAAAAAAAAGTGTCCACAGGTCAAAAAAAAAGATCG <br> TCGGACTGTAGAACTC |
| SBC_Oligo8 | AAAAAAAAAAAAAAAAAAAAAAAAATCCTCTTCAGGTCAAAAAAAAAGATCG <br> TCGGACTGTAGAACTC |
| SBC_Oligo9 | AAAAAAAAAAAAAAAAAAAAAAAAAGTGCAGTCAGGTCAAAAAAAAAAGATCG <br> TCGGACTGTAGAACTC |
| SBC_Oligo10 | AAAAAAAAAAAAAAAAAAAAAAAAAAAGGTAGACAGGTCAAAAAAAAAAGATCG <br> TCGGACTGTAGAACTC |

Table 4. List of key oligonucleotides used for barcoding and library preparation (not including the long list of cell-identifying barcodes that appear in the subsequent tables) for Experiment 2.

| Oligonucleotide Name | Oligonucleotide Sequence |
| :--- | :--- |
| Bead Capture Oligo (5'-dual biotinylated) | AGGTAAGGTAATACGACTCACTATAGGGGTTCAGAGT <br> TCTACAGTCCGACGATC |
| RT1 (Reverse Transcription Primer for Lane 1) | GCCTTGGCACCCGAGAATTCCANNNNNNNNCGTCATN <br> NNNNN |
| RT2 (Reverse Transcription Primer for Lane 2) | GCCTTGGCACCCGAGAATTCCANNNNNNNNTACCCAN <br> NNNNN |
| RT3 (Reverse Transcription Primer for Lane 3) | GCCTTGGCACCCGAGAATTCCANNNNNNNNGCCATTN <br> NNNNN |
| RT4 (Reverse Transcription Primer for Lane 4) | GCCTTGGCACCCGAGAATTCCANNNNNNNNGAGTACN <br> NNNNN |
| RT5 (Reverse Transcription Primer for Lane 5) | GCCTTGGCACCCGAGAATTCCANNNNNNNNAGAGTCN <br> NNNNN |
| RP1 (PCR Primer 1) | AATGATACGGCGACCACCGAGATCTACACGTTCAGAG <br> TTCTACAGTCCGA |
| RPI2 (PCR Primer 2) | CAAGCAGAAGACGGCATACGAGATACATCGGTGACTG <br> GAGTTCCTTGGCACCCGAGAATTCCA |

Table 5. Oligonucleotide sequences used to generate the first set of barcoded beads (FBC) for combinatorial synthesis in Experiment 2.

| Oligonucleotide Name | Oligonucleotide Sequence |
| :--- | :--- |
| FBC_Oligo1 | CAGGTCCTGATCGATGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo2 | CAGGTCGTGTAGACAGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo3 | CAGGTCCATTGTTCCGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo4 | CAGGTCCTTGACTACGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo5 | CAGGTCACCGTTTCGGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo6 | CAGGTCAAGGACCGTGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo7 | CAGGTCTCACTATGCGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo8 | CAGGTCCTGCAATGGGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo9 | CAGGTCTGAGTCGTCGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo10 | CAGGTCCTCACACTAGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo11 | CAGGTCTTACCCCCTGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo12 | CAGGTCCCAAGTAGAGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo13 | CAGGTCATAGCGCACGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo14 | CAGGTCTGACGTACGGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo15 | CAGGTCGTAGAGTTGGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo16 | CAGGTCTTTCTGGCGGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo17 | CAGGTCGGAATGTGTGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo18 | CAGGTCCTATGGAAGGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo19 | CAGGTCAAGTCCATGGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo20 | CAGGTCAGTACTTGGGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo21 | CAGGTCACAGGACTAGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo22 | CAGGTCACCAGGTAAGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo23 | CAGGTCGCATGAACCGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo24 | CAGGTCGTTGGTGTTGATCGTCGGACTGTAGAACTCTGAAC |


| FBC_Oligo25 | CAGGTCCCTTCAGACGATCGTCGGACTGTAGAACTCTGAAC |
| :---: | :---: |
| FBC_Oligo26 | CAGGTCCCTCTTGGTGATCGTCGGACTGTAGAACTCTGAAC |
| FBC Oligo27 | CAGGTCGGGAAAGTTGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo28 | CAGGTCAGCCAGAGTGATCGTCGGACTGTAGAACTCTGAAC |
| FBC Oligo29 | CAGGTCTCGCATCTGGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo30 | CAGGTCGATACGGCAGATCGTCGGACTGTAGAACTCTGAAC |
| FBC Oligo31 | CAGGTCTCGGCCAAAGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo32 | CAGGTCAGATTTCGCGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo33 | CAGGTCGACCCTCAAGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo34 | CAGGTCAGTCCACTCGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo35 | CAGGTCCAAACGATCGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo36 | CAGGTCGCCTAATAGGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo37 | CAGGTCGGCTACATCGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo38 | CAGGTCTATGAGCAGGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo39 | CAGGTCGGTAGTAACGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo40 | CAGGTCCGCGTATATGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo41 | CAGGTCTACTGGAGCGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo42 | CAGGTCAGGGAATCAGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo43 | CAGGTCATCCGAGATGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo44 | CAGGTCTCCCAAGCAGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo45 | CAGGTCGAGCCGTTTGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo46 | CAGGTCTGCTCTTACGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo47 | CAGGTCACGACTACCGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo48 | CAGGTCCAAGCAGCTGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo49 | CAGGTCGTATTCGCGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo50 | CAGGTCGCTCTGAAGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo51 | CAGGTCACGTAGTGGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo52 | CAGGTCATTGGGTCGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo53 | CAGGTCAACAGCACGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo54 | CAGGTCTCAGAGACGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo55 | CAGGTCGTGTGCTAGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo56 | CAGGTCGCAGTTGAGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo57 | CAGGTCTTAACGGGGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo58 | CAGGTCGCTCGATTGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo59 | CAGGTCACACCTGTGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo60 | CAGGTCAGACGGTTGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo61 | CAGGTCGCAAACCAGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo62 | CAGGTCGAGTATGGGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo63 | CAGGTCGGTCTTTCGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo64 | CAGGTCCATCTGCTGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo65 | CAGGTCTTCGCAAGGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo66 | CAGGTCTTGTGACGGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo67 | CAGGTCTGCATGACGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo68 | CAGGTCCAACGTGAGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo69 | CAGGTCTAGGCTTCGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo70 | CAGGTCTGGTAGGAGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo71 | CAGGTCTGCAGCTTGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo72 | CAGGTCCTGTACCTGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo73 | CAGGTCCGCAATGAGATCGTCGGACTGTAGAACTCTGAAC |


| FBC_Oligo74 | CAGGTCGATCCAAGGATCGTCGGACTGTAGAACTCTGAAC |
| :--- | :--- |
| FBC_Oligo75 | CAGGTCCACTTACGGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo76 | CAGGTCAACTAGGCGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo77 | CAGGTCACTAGCGTGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo78 | CAGGTCCGTTCGTTGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo79 | CAGGTCAGTCACGAGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo80 | CAGGTCCCTGTAACGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo81 | CAGGTCGTCCTCTTGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo82 | CAGGTCCAGCGAATGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo83 | CAGGTCATGGTTGGGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo84 | CAGGTCGAGGTTCTGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo85 | CAGGTCTACCTCGAGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo86 | CAGGTCTTCTGTGCGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo87 | CAGGTCGACAACTGGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo88 | CAGGTCCGACAACAGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo89 | CAGGTCTCGATACCGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo90 | CAGGTCCCATACTCGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo91 | CAGGTCATTCGCAGGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo92 | CAGGTCACCATAGGGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo93 | CAGGTCCGATCAAGGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo94 | CAGGTCACCTTGCTGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo95 | CAGGTCGACTCAGTGATCGTCGGACTGTAGAACTCTGAAC |
| FBC_Oligo96 | CAGGTCGTCAATCCGATCGTCGGACTGTAGAACTCTGAAC |

Table 6. Oligonucleotide sequences used to generate the second set of barcoded beads (SBC) for combinatorial synthesis in Experiment 2.

| Oligonucleotide Name | Oligonucleotide Sequence |
| :--- | :--- |
| SBC_Oligo1 | AAAAAAAAAAAAAAAAAAAAAAAAAATATGCGCAGGTCAAAAAAAAAGATCG <br> TCGGACTGTAGAACTC |
| SBC_Oligo2 | AAAAAAAAAAAAAAAAAAAAAAAAAAGGACATCAGGTCAAAAAAAAAGATCG <br> TCGGACTGTAGAACTC |
| SBC_Oligo3 | AAAAAAAAAAAAAAAAAAAAAAAAAGACTACGCAGGTCAAAAAAAAAGATCG <br> TCGGACTGTAGAACTC |
| SBC_Oligo4 | AAAAAAAAAAAAAAAAAAAAAAAAACTGAAACCAGGTCAAAAAAAAAGATCG <br> TCGGACTGTAGAACTC |
| SBC_Oligo5 | AAAAAAAAAAAAAAAAAAAAAAAAATAGGACCCAGGTCAAAAAAAAAGATCG <br> TCGGACTGTAGAACTC |
| SBC_Oligo6 | AAAAAAAAAAAAAAAAAAAAAAAAATAACGCACAGGTCAAAAAAAAAGATCG <br> TCGGACTGTAGAACTC |
| SBC_Oligo7 | AAAAAAAAAAAAAAAAAAAAAAAAACCCAACACAGGTCAAAAAAAAAGATCG <br> TCGGACTGTAGAACTC |
| SBC_Oligo8 | AAAAAAAAAAAAAAAAAAAAAAAAACGCATTTCAGGTCAAAAAAAAAGATCG <br> TCGGACTGTAGAACTC |
| SBC_Oligo9 | AAAAAAAAAAAAAAAAAAAAAAAAACATCTACCAGGTCAAAAAAAAAGATCG <br> TCGGACTGTAGAACTC |
| SBC_Oligo10 | AAAAAAAAAAAAAAAAAAAAAAAAATACGGATCAGGTCAAAAAAAAAGATCG <br> TCGGACTGTAGAACTC |

Table 7. Table of reagents and cost estimates per run. Costs associate with combinatorial bead synthesis are highlighted in gray.

| Reagent | Volume | Stock Volume | Price of Stock | Price per Run |
| :---: | :---: | :---: | :---: | :---: |
| SUPERaseIN (Ambion) | 19 uL | 500 uL | \$350.40 | \$13.32 |
| dNTPs (NEB) | 10 uL | 800 uL | \$44.80 | \$0.56 |
| HiScribe <br> (NEB) IVT Kit | 1 uL | 50 uL | \$169.60 | \$3.39 |
| MessageAamp II Kit (Ambion)* | 3 uL | 740 uL | \$3,668.00 | \$14.87 |
| PrimeScript (Clontech) $\quad$ RT | 5 uL | 200 uL | \$501.63 | \$12.54 |
| Phusion polymerase | 0.5 uL | 250 uL | \$336.00 | \$0.67 |
| Lane <br> primers (IDT) Barcode RT | 15 uL | 3155 uL | \$542.25 | \$0.86 |
| NHS beads (GE) | 3.25 uL | 25000 uL | \$155.80 | \$0.02 |
| Streptavidin (NEB) | 5 uL | 1000 uL | \$188.80 | \$0.94 |
| Dual-biotin anchor oligo (IDT) | 0.64 uL | 700 uL | \$225.75 | \$0.21 |
| $\begin{aligned} & \text { Klenow fragment } \\ & \text { exo- (NEB) } \end{aligned}$ | 1.5 uL | 200 uL | \$188.80 | \$1.42 |
| dNTPs (NEB) | 3.5 uL | 800 uL | \$44.80 | \$0.20 |
| FBC primers (IDT) | 0.96 uL (all 96) | 240,000 uL (all 96) | \$484.56 total | \$0.002 |
| SBC primers (IDT) | 0.66 uL (all 10) | 4000 uL (all 10) | \$340.00 total | \$0.67 |
| Experiment Costs |  |  |  | \$46.21 |
| Bead Costs |  |  |  | \$3.46 |
| Total Cost per Run |  |  |  | \$49.67 |
| Cost per Cell | 250-500 cells |  |  | \$0.10-\$0.20 |

* Only the second-strand synthesis reagents are used here.

