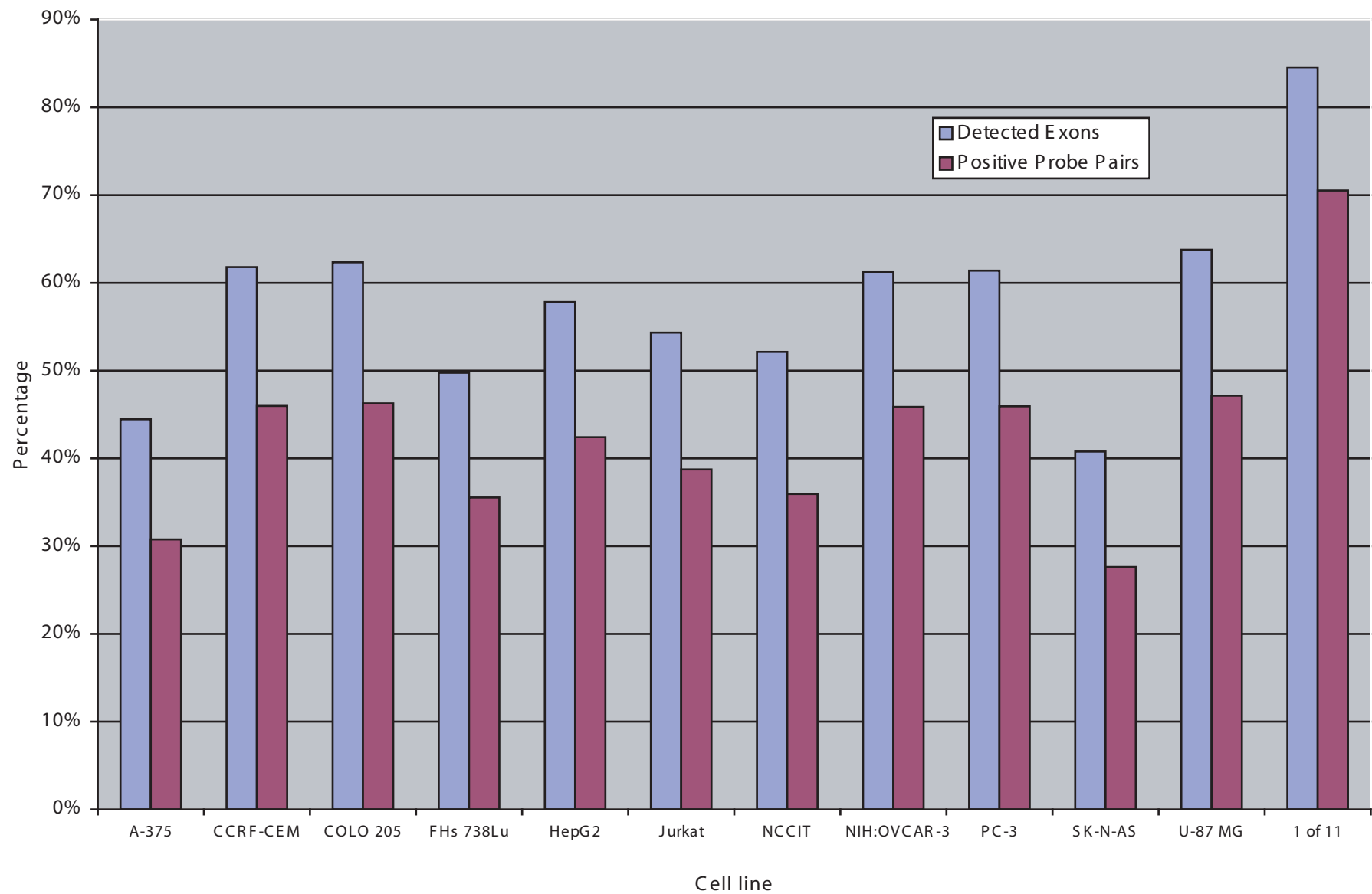
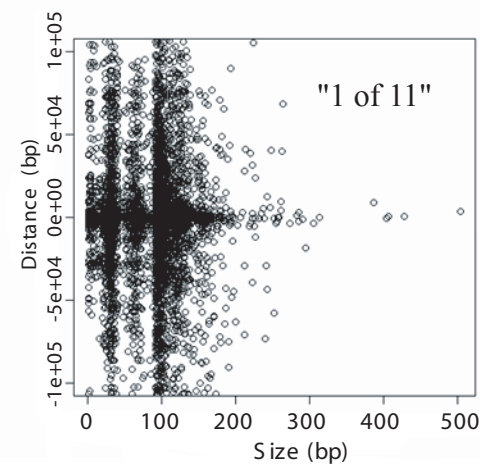
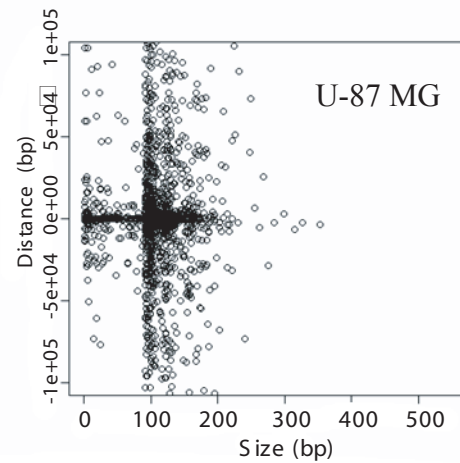
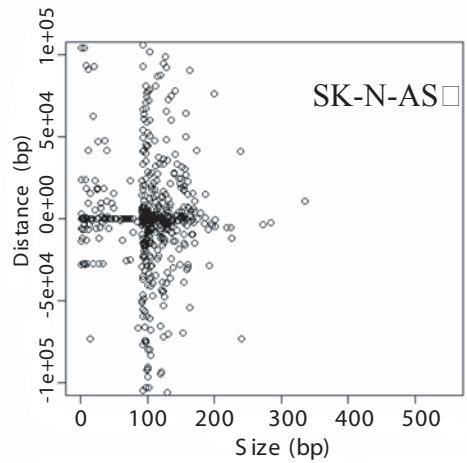
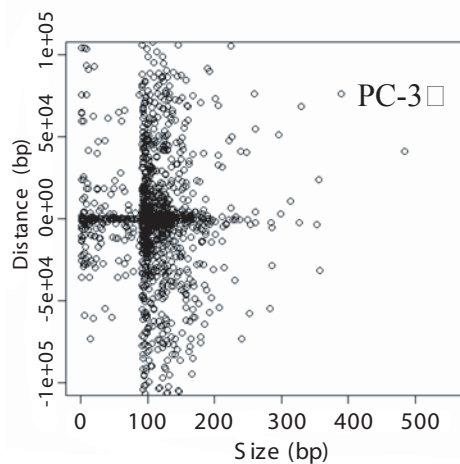
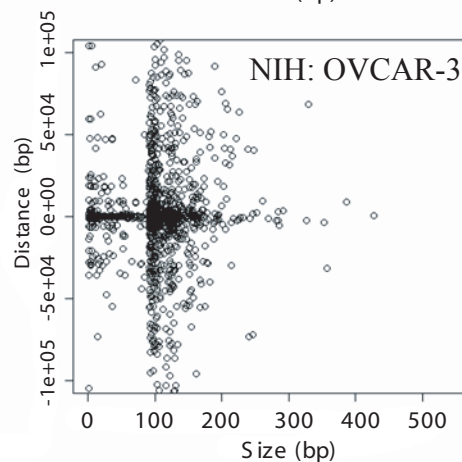
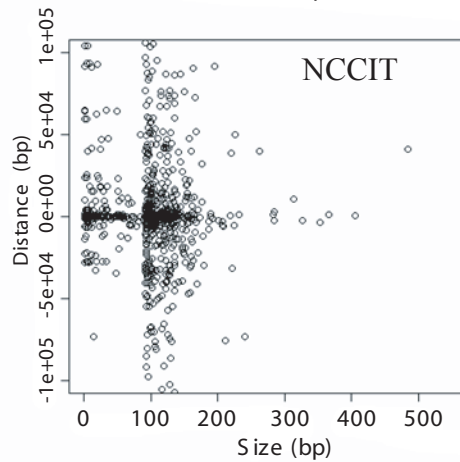
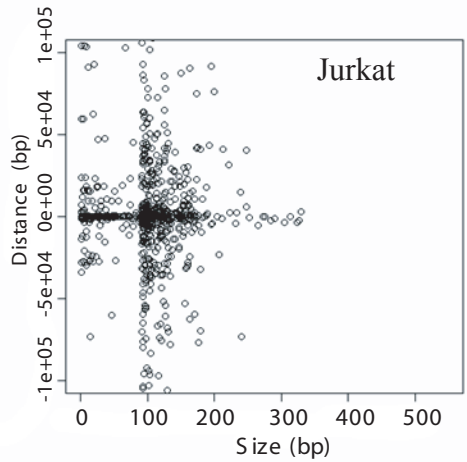
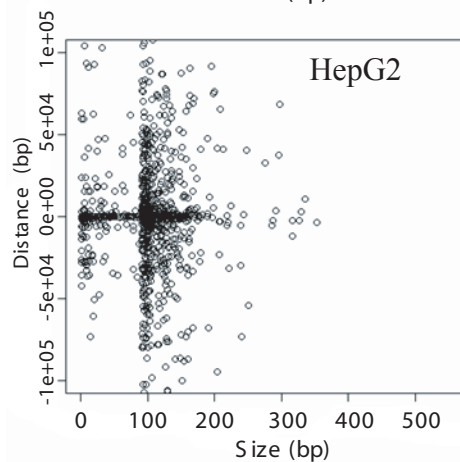
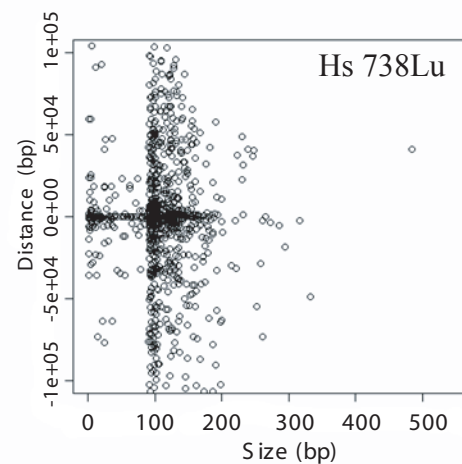
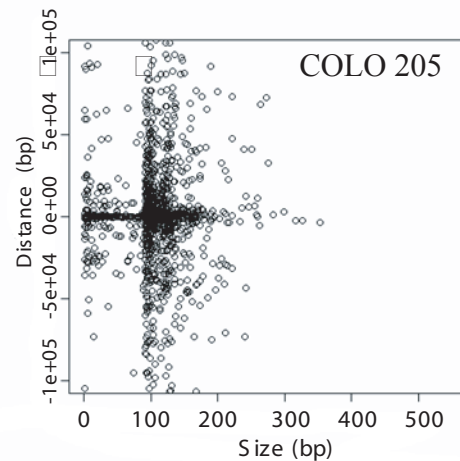
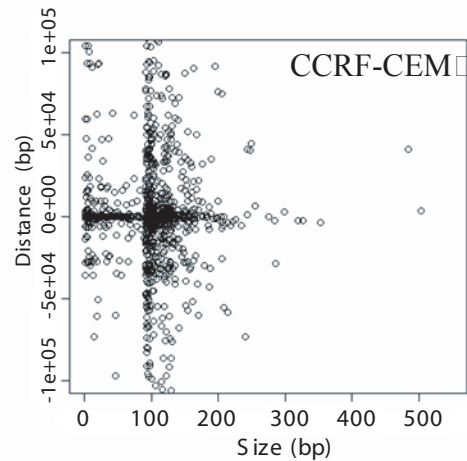
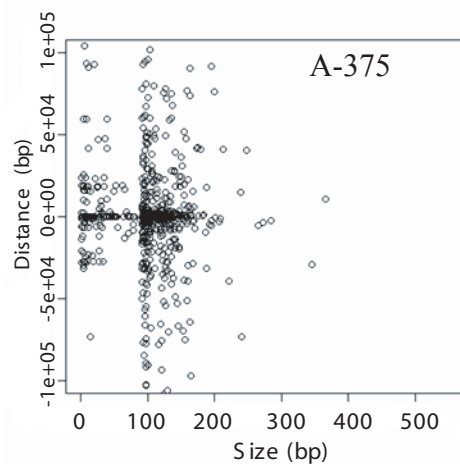


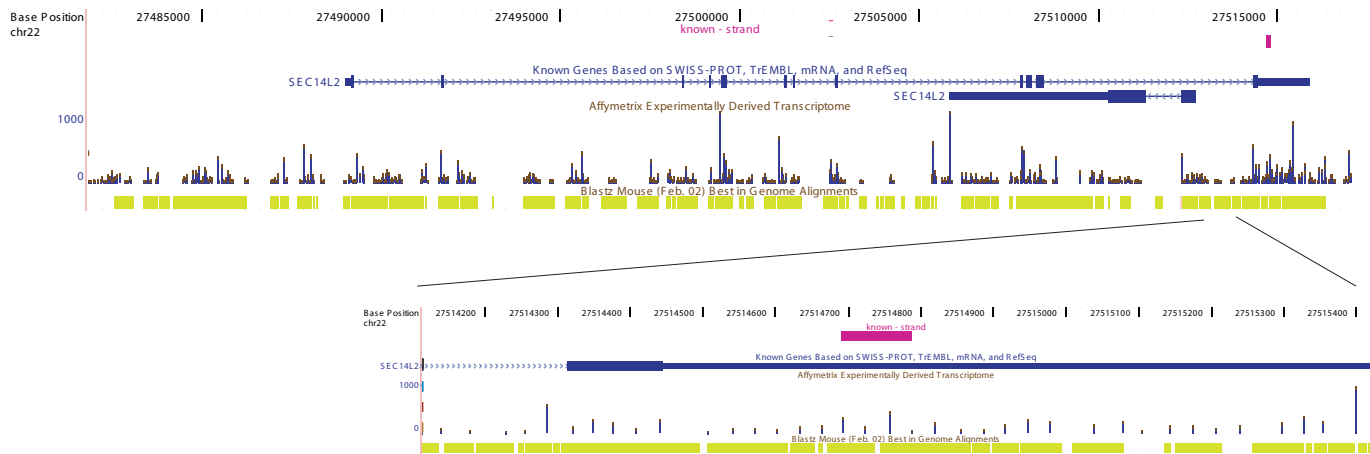


	A-375	CCRF-CEM	COLO 205	FHs 738Lu	HepG2	Jurkat	NCCIT	NIH:OVCA-3	PC-3	SK-N-AS	U-87 MG	1 of 11
Total # exons	5995	5995	5995	5995	5995	5995	5995	5995	5995	5995	5995	5995
Number of detected exons	2665	3705	3738	2983	3466	3257	3126	3670	3681	2445	3823	5068
Percent of exons detected	44.5%	61.8%	62.4%	49.8%	57.8%	54.3%	52.1%	61.2%	61.4%	40.8%	63.8%	84.5%

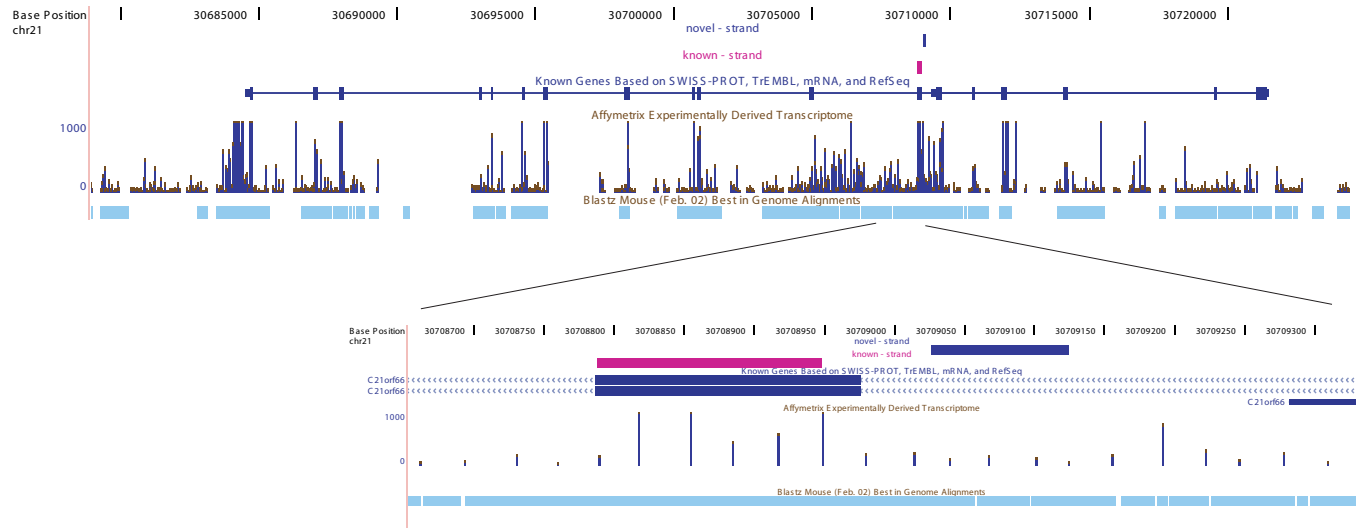
	A-375	CCRF-CEM	COLO 205	FHs 738Lu	HepG2	Jurkat	NCCIT	NIH:OVCA-3	PC-3	SK-N-AS	U-87 MG	1 of 11
Interrogation probe pairs	38407	38407	38407	38407	38407	38407	38407	38407	38407	38407	38407	38407
Positive probe pairs	11815	17661	17770	13653	16292	14878	13808	17617	17644	10610	18107	27088
Percent positive	30.8%	46.0%	46.3%	35.5%	42.4%	38.7%	36.0%	45.9%	45.9%	27.6%	47.1%	70.5%



A.



B.



C.

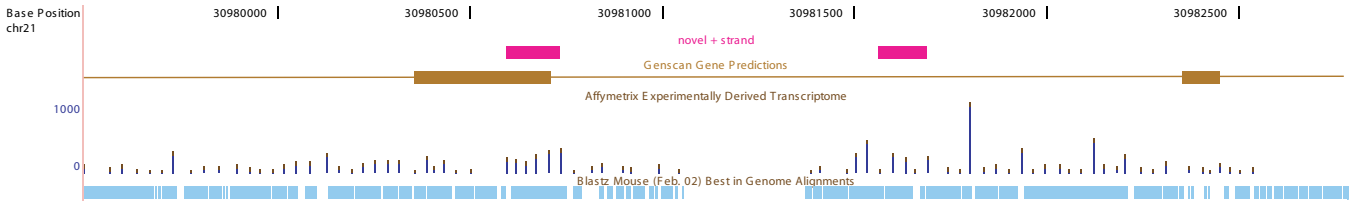


Table S1. Positive probe analysis summary by cell line.

Cell Line	PosProbes ^{1.}	MapProbes ^{2.}	% PosProbes ^{3.}
A-375	81685	1020643	8.00%
CCRF-CEM	110036	1020643	10.78%
COLO 205	121087	1020643	11.86%
FHs 738Lu	95159	1020643	9.32%
HepG2	108331	1020643	10.61%
Jurkat	92558	1020643	9.07%
NCCIT	89901	1020643	8.81%
NIH:OVCAR-3	117159	1020643	11.48%
PC-3	127531	1020643	12.50%
SK-N-AS	77021	1020643	7.55%
U-87 MG	134262	1020643	13.15%
Averages	104975	1020643	10.29%
"1 of 11"	263184	1020643	25.79%

1. PosProbes = positive probes (PM-MM) after applying wilcoxon test and the fixed threshold of 150
2. MapProbes = all interrogated probes on the array, includes probes that map to more than 1 location
The genomic coordinates associated with probes are transformed or mapped to those of newer versions of the genome based on exact alignment of probe sequence to genomic sequence. The analysis is then performed using these mapped probe coordinates.
3. %PosProbes = percent positive probes (PosProbes/MapProbes)

Table S2. False positive rates and sensitivities by chip type and cell line.

Cell Line	Chip Type ^{1.}	FPR ^{2.}	Bac. Sn. Lys ^{3.}	Bac. Sn. Phe ^{3.}	Bac. Sn. Dap ^{3.}	Bac. Sn. Thr ^{3.}	Bac. Sn. Trp ^{3.}
A-375	A	0.98	0.73	0.23	0.97	0.97	1.00
A-375	B	0.00	0.80	0.54	0.94	0.97	1.00
A-375	C	0.00	1.00	0.46	0.97	0.97	1.00
CCRF-CEM	A	0.98	1.00	0.77	1.00	0.97	1.00
CCRF-CEM	B	1.96	1.00	0.77	1.00	1.00	1.00
CCRF-CEM	C	0.98	1.00	0.69	1.00	0.97	1.00
COLO 205	A	2.94	0.93	0.46	1.00	0.97	1.00
COLO 205	B	2.94	0.93	0.46	0.97	0.97	1.00
COLO 205	C	1.96	1.00	0.46	0.97	0.97	1.00
FHs 738Lu	A	13.73	0.40	0.23	1.00	0.94	1.00
FHs 738Lu	B	0.98	0.47	0.38	0.97	0.91	1.00
FHs 738Lu	C	2.94	0.60	0.38	0.97	0.94	1.00
HepG2	A	3.92	1.00	0.77	1.00	1.00	1.00
HepG2	B	4.90	1.00	0.54	1.00	1.00	1.00
HepG2	C	1.96	1.00	0.62	0.97	1.00	1.00
Jurkat	A	1.96	0.67	0.46	0.97	0.97	1.00
Jurkat	B	0.00	0.80	0.62	0.97	0.94	1.00
Jurkat	C	0.00	0.93	0.38	0.97	0.97	1.00
NCCIT	A	2.94	0.60	0.38	1.00	0.91	1.00
NCCIT	B	2.94	0.73	0.54	1.00	0.97	1.00
NCCIT	C	3.92	0.73	0.46	1.00	0.91	1.00
NIH: OVCAR-3	A	3.92	0.80	0.54	1.00	0.97	1.00
NIH: OVCAR-3	B	1.96	0.93	0.46	1.00	0.97	1.00
NIH: OVCAR-3	C	0.00	0.87	0.38	1.00	0.91	1.00
PC-3	A	6.86	0.80	0.54	1.00	1.00	1.00
PC-3	B	3.92	0.93	0.46	0.97	0.97	1.00
PC-3	C	2.94	1.00	0.46	1.00	0.97	1.00
SK-N-AS	A	0.00	0.67	0.38	0.91	0.91	0.97
SK-N-AS	B	3.92	0.93	0.31	0.94	0.88	0.97

SK-N-AS	C	1.96	0.53	0.31	0.94	0.94	0.97
U-87 MG	A	3.92	0.87	0.46	1.00	1.00	1.00
U-87 MG	B	7.84	0.87	0.46	0.97	0.97	1.00
U-87 MG	C	4.90	0.93	0.46	1.00	0.97	1.00
MEDIAN		2.94	0.87	0.46	1.00	0.97	1.00

1. Chromosome 21 and 22 arrays contain uniformly spaced oligonucleotide probes (25-nucleotide oligomers) that interrogate on average every 35 base pairs comprised on 3 chips (A, B, or C). Genomic coordinates of chips are (NCBI version 30):
A – chr21: 11,000,000 – 31,412,500; B – chr21: 31,375,000 - 44,625,000; B – chr22: 12,500,000 – 22,750,000;
C – chr22: 22,660,000 – 47,748,250
2. Corresponding false positive rate at a fixed threshold of 150 determined from positive probes before generation of transfrags.
3. Probes were picked from the following *Bacillus subtilis* genes using identical probe selection rules: LYS 328-1344; PHE 2016-3331; THR 247-2231; DAP 1357-3196; TRP 1-2517. A polyadenylated transcript corresponding to a smaller portion of each five loci was generated to evaluate the sensitivity of the assay, while the bacterial region outside of the spiked regions was employed in determination of false positive rates. The regions of each gene that corresponds to the spiked transcripts are: LYS 817-1344; PHE 2852-3331; THR 1221-2231; DAP 1357-2493; TRP 1-1261. The control bacterial transcripts were spiked into human polyA+ RNA preparations before cDNA synthesis at the following concentrations (copies/cell): LYS and PHE - 3; DAP and THR - 10; and TRP - 30, assuming 300,000 different mRNA species in a human cell and the average size of a transcript to be 1300 nt.

Table S3. Transfrag analysis summary by cell line.

Cell Line	No. transfrags ^{1.}	Average length (bp)	bp coverage
A-375	1840	153	281171
CCRF-CEM	3372	160	540056
COLO 205	3616	154	557089
FHs 738Lu	2603	148	385295
HepG2	3028	155	468751
Jurkat	2552	159	406063
NCCIT	2392	153	366381
NIH:OVCAR-3	3578	154	552025
PC-3	3905	153	595700
SK-N-AS	1623	149	242609
U-87 MG	4110	148	608433
Averages	2965	153	454870
"1 of 11"	9001	154	1384484

1. No. transfrags = number of transfrags identified in indicated cell line after applying the fixed threshold (150), minimum length (90 bp), and maximum gap (40 bp) parameters.

Table S4. Gene profiles by cell line

1. Exon Profile = For each exon represented on the array, a "on" or "off" call was made based the percentage of positive probes. A "0" represents < 30% of the interrogating probes were called positive. A "1" represents > 30% of the interrogating probes were positive.

GenBank Acc. No.	Cell line	Exon Profile ¹ .
AB000099	A375	0
AB000099	CEM	0
AB000099	COLO	0
AB000099	FHS	0
AB000099	Hep	0
AB000099	Jurkat	0
AB000099	Nccit	0
AB000099	OVCAR	0
AB000099	PC3	0
AB000099	Sk	0
AB000099	U87	0
AB001535	A375	0001100001000010010001001010
AB001535	CEM	0110111001010011010011001010
AB001535	COLO	0100100001010010010001101011
AB001535	FHS	0000100001010000011001000000
AB001535	Hep	0100100000000001010001001010
AB001535	Jurkat	0100000000000010010001001010
AB001535	Nccit	0000100001010010011001001010
AB001535	OVCAR	0100100101010010011001101011
AB001535	PC3	0000100001000010010101101011
AB001535	Sk	0001110000000010000001001000
AB001535	U87	0000101001010011011001001010
AB002058	A375	010001101000
AB002058	CEM	110001100010
AB002058	COLO	110011100010
AB002058	FHS	010011100000

AB002058	Hep	010011100010	
AB002058	Jurkat	010010000010	
AB002058	Nccit	110011000000	
AB002058	OVCAR	110011100010	
AB002058	PC3	110011000010	
AB002058	Sk	110010101010	
AB002058	U87	111011111011	
AB002369	A375	0000000001010010000	
AB002369	CEM	1100110111110111111	
AB002369	COLO	0101110111110111111	
AB002369	FHS	0100010111011011011	
AB002369	Hep	0100110011010011111	
AB002369	Jurkat	0110010001010011010	
AB002369	Nccit	0000000111010011010	
AB002369	OVCAR	0111111111111111111	
AB002369	PC3	0100110111010111111	
AB002369	Sk	0100000001010010000	
AB002369	U87	0110110111011111011	
AB002374	A375	1111000100110	
AB002374	CEM	1111110111111	
AB002374	COLO	1111100111111	
AB002374	FHS	1111100111111	
AB002374	Hep	1111100111111	
AB002374	Jurkat	1111100101111	
AB002374	Nccit	1111100111111	
AB002374	OVCAR	1111110111111	
AB002374	PC3	1111100101110	
AB002374	Sk	1111000101110	
AB002374	U87	1111110111111	
AB002803	A375	11111	
AB002803	CEM	11111	
AB002803	COLO	11111	
AB002803	FHS	11111	

AB002803	Hep	11111	
AB002803	Jurkat	11111	
AB002803	Nccit	11111	
AB002803	OVCAR	11111	
AB002803	PC3	11111	
AB002803	Sk	11111	
AB002803	U87	11111	
AB006536	A375	000000001	
AB006536	CEM	000000001	
AB006536	COLO	000001101	
AB006536	FHS	000000000	
AB006536	Hep	000000001	
AB006536	Jurkat	000000001	
AB006536	Nccit	000000001	
AB006536	OVCAR	010000001	
AB006536	PC3	000000001	
AB006536	Sk	000000001	
AB006536	U87	000000001	
AB006630	A375	11011	
AB006630	CEM	11111	
AB006630	COLO	10111	
AB006630	FHS	10010	
AB006630	Hep	11111	
AB006630	Jurkat	11111	
AB006630	Nccit	10111	
AB006630	OVCAR	10111	
AB006630	PC3	10111	
AB006630	Sk	11001	
AB006630	U87	10111	
AB011111	A375	1110010010101011110111110010001111	
AB011111	CEM	11110111110011111111101110010110011	
AB011111	COLO	1111111111000011111111111010111011	
AB011111	FHS	00110101111001111111111110001010001	

AB011111	Hep	111111111101111110101011011100011
AB011111	Jurkat	11111011111111111011111011110111
AB011111	Nccit	11110111111011111111111011111111
AB011111	OVCAR	11111111110111111111111111111111
AB011111	PC3	111111111111111111111111011111111
AB011111	Sk	1111010110001011110101111011000111
AB011111	U87	1111111111100111111111111010110011
AB011114	A375	0000100000000000000010111110000
AB011114	CEM	01101110111111010001111011111100
AB011114	COLO	01001000000001000001110011111000
AB011114	FHS	000010000000000000001110011111000
AB011114	Hep	000010100000000000001110111111000
AB011114	Jurkat	01001110111000000001000111111000
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AB011114	OVCAR	00001000100001000011110011111000
AB011114	PC3	000010000000000000101110011111000
AB011114	Sk	000010000000000000000000111110000
AB011114	U87	00001000000001000101110011111000
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AB013885	FHS	0000001000
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AB014537	FHS	1

AB014537	Hep	1
AB014537	Jurkat	1
AB014537	Nccit	1
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AB014537	PC3	1
AB014537	Sk	0
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AB014545	CEM	01000100100111110010110001001110001011110
AB014545	COLO	00000100000101010010110001100111001011011
AB014545	FHS	00000100000010000000110001100010001010000
AB014545	Hep	00000000000100010010110001000110000010000
AB014545	Jurkat	01000000100101010010110101001110001010010
AB014545	Nccit	00000000000010010010110001000111001010000
AB014545	OVCAR	00010010000111110010110001100011001010010
AB014545	PC3	000000000000000010010110000001010001011010
AB014545	Sk	00000001000000000000110100000000000010010
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AB014553	CEM	001100
AB014553	COLO	000000
AB014553	FHS	000000
AB014553	Hep	001100
AB014553	Jurkat	000000
AB014553	Nccit	011110
AB014553	OVCAR	011110
AB014553	PC3	011110
AB014553	Sk	000000
AB014553	U87	011110
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AB014568	CEM	111110111111111
AB014568	COLO	111110110111111
AB014568	FHS	111110111111111

AB014568	Hep	111110111111111
AB014568	Jurkat	111010110110011
AB014568	Nccit	111010110110001
AB014568	OVCAR	111010101110111
AB014568	PC3	111010110111111
AB014568	Sk	110000110010011
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AB014585	A375	11011111011101
AB014585	CEM	11111111011111
AB014585	COLO	11111111011111
AB014585	FHS	11111111011111
AB014585	Hep	111111111111111
AB014585	Jurkat	111111111111111
AB014585	Nccit	111011110111110
AB014585	OVCAR	111111110111111
AB014585	PC3	111111110111111
AB014585	Sk	01001010010101
AB014585	U87	111111110111111
AB015282	A375	1001
AB015282	CEM	1111
AB015282	COLO	1111
AB015282	FHS	1101
AB015282	Hep	1111
AB015282	Jurkat	1111
AB015282	Nccit	1111
AB015282	OVCAR	1111
AB015282	PC3	1111
AB015282	Sk	0111
AB015282	U87	1111
AB015283	A375	01
AB015283	CEM	11
AB015283	COLO	11
AB015283	FHS	01

AB015283	Hep	11	
AB015283	Jurkat	11	
AB015283	Nccit	01	
AB015283	OVCAR	11	
AB015283	PC3	11	
AB015283	Sk	11	
AB015283	U87	01	
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AB017549	CEM	1101010	
AB017549	COLO	0111011	
AB017549	FHS	0100000	
AB017549	Hep	0101010	
AB017549	Jurkat	0101010	
AB017549	Nccit	0101010	
AB017549	OVCAR	0111011	
AB017549	PC3	0111011	
AB017549	Sk	0101000	
AB017549	U87	0101010	
AB018257	A375	111111101101001010100	
AB018257	CEM	111111101101101111101	
AB018257	COLO	11111111111111111101	
AB018257	FHS	111101101111111111101	
AB018257	Hep	111111101111101111101	
AB018257	Jurkat	111110101111101111100	
AB018257	Nccit	111111111110110111101	
AB018257	OVCAR	111111111111111111101	
AB018257	PC3	111111111111111111101	
AB018257	Sk	111111101101101011100	
AB018257	U87	111111101111111111101	
AB018310	A375	1111010011111011	
AB018310	CEM	11111111111111111	
AB018310	COLO	1111010001101011	
AB018310	FHS	1111010011111000	

AB018310	Hep	1111010011111001
AB018310	Jurkat	1111011111111111
AB018310	Nccit	1111010001111010
AB018310	OVCAR	1111010001101011
AB018310	PC3	1111010011101001
AB018310	Sk	0010000001111010
AB018310	U87	1111010011111010
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AB020337	CEM	00
AB020337	COLO	11
AB020337	FHS	00
AB020337	Hep	00
AB020337	Jurkat	00
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AB020626	CEM	100100
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AB020626	FHS	100000
AB020626	Hep	100000
AB020626	Jurkat	100000
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AB020626	PC3	100011
AB020626	Sk	101000
AB020626	U87	101011
AB020659	A375	00011101100000010101110000
AB020659	CEM	11011111111110010111110000
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AB020659	FHS	10010011101100010101110000

AB020659	Hep	111111111111011111111000
AB020659	Jurkat	11011101110000010111110000
AB020659	Nccit	1111111101110110101110000
AB020659	OVCAR	1111111111110110111110000
AB020659	PC3	11010111111010010111110000
AB020659	Sk	00011101100000010110110000
AB020659	U87	0101111111111010111110000
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AB020717	CEM	111001110101011010110000010110
AB020717	COLO	111001110101001000100001010110
AB020717	FHS	011001110101011000000000010010
AB020717	Hep	011001110101001000000000000010
AB020717	Jurkat	111001110101111010110000111110
AB020717	Nccit	011000110101001000100000000100
AB020717	OVCAR	111000110101011000000000010011
AB020717	PC3	111001110101111110100010111110
AB020717	Sk	101001110001001000100000000000
AB020717	U87	111001110101011000100000010010
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AB023061	CEM	1000000000
AB023061	COLO	1000000100
AB023061	FHS	1001001100
AB023061	Hep	0000000000
AB023061	Jurkat	1000000000
AB023061	Nccit	1001001100
AB023061	OVCAR	1001001101
AB023061	PC3	1000001100
AB023061	Sk	1000000000
AB023061	U87	1001001101
AB023144	A375	100011010010010
AB023144	CEM	100011010000010
AB023144	COLO	100001010000010
AB023144	FHS	100011000000000

AB023144	Hep	100011010000010
AB023144	Jurkat	100010000010010
AB023144	Nccit	100001000000000
AB023144	OVCAR	100001010000000
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AB023144	Sk	100000000010010
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AF007118	A375	100001000000010000000000
AF007118	CEM	000000000000001110000000
AF007118	COLO	001000100010000000000000
AF007118	FHS	000000100010100000000000
AF007118	Hep	000000100000000000000000
AF007118	Jurkat	0100000000000001000000000
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AF008655	COLO	0111
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AF008655	Hep	0111
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AF009039	FHS	011000111010101100000000001001
AF009039	Hep	011000111010100100000000000001
AF009039	Jurkat	111000111010111101011000011111
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AF012131	COLO	00001010
AF012131	FHS	00000110
AF012131	Hep	00101010
AF012131	Jurkat	00000110
AF012131	Nccit	00101110
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AF012872	COLO	11111111011111110111111111110111111111111111111101101
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AF012872	Hep	1111111111110011011111111110101111101101011111101101
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AF018081	Hep	111010110110111111111111111
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AF029689	Hep	000000001001
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AF035304	A375	0
AF035304	CEM	1
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AF035304	Nccit	1
AF035304	OVCAR	1
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AF035304	U87	1
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AF038175	CEM	00000
AF038175	COLO	01001
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AF038175	Hep	01000	
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AF040251	FHS	1	
AF040251	Hep	1	
AF040251	Jurkat	0	
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AF060862	U87	0000
AF060924	A375	0110
AF060924	CEM	1111
AF060924	COLO	1111
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AF060924	Hep	0110
AF060924	Jurkat	0101
AF060924	Nccit	1010
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AF064238	A375	00111011111100110111
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AF064238	Hep	01111011101110110111
AF064238	Jurkat	00110001100110000110
AF064238	Nccit	10100001000010000000
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AF067223	COLO	0100000001001000001
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AF067223	Hep	0110010001111001101	
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AF075458	Nccit	010000101000	
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AF075589	PC3	11	
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AF081535	Hep	1100110011111010100	
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AF142567	Hep	00001000000000000000101001100000011011
AF142567	Jurkat	00001000000000000000101000110000011000
AF142567	Nccit	0000100000010001000101101100000011001
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AF142567	PC3	0010100000110000000101101110000011011
AF142567	Sk	0010100000000000000101000000000011000
AF142567	U87	0000100000110011000101101110000011011
AF146568	A375	1011
AF146568	CEM	1111
AF146568	COLO	1111
AF146568	FHS	1011
AF146568	Hep	1011
AF146568	Jurkat	1011
AF146568	Nccit	1011
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AF146568	PC3	1111
AF146568	Sk	1011
AF146568	U87	1111
AF149785	A375	1111100
AF149785	CEM	1111110
AF149785	COLO	1111111
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AF149785	Hep	1111111
AF149785	Jurkat	1111100
AF149785	Nccit	1111111
AF149785	OVCAR	1111111
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AF151854	A375	001011110000011110111
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AF151854	COLO	011111110000010111111
AF151854	FHS	101111110001010111110

AF151854	Hep	001011110000010110111	
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AF156774	COLO	011101011	
AF156774	FHS	011110011	
AF156774	Hep	011111011	
AF156774	Jurkat	000000011	
AF156774	Nccit	000000010	
AF156774	OVCAR	011111011	
AF156774	PC3	011111111	
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AF156774	U87	011111011	
AF161350	A375	101110110	
AF161350	CEM	001100111	
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AF161350	Nccit	001100111	
AF161350	OVCAR	101110111	
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AF161350	U87	101110111	
AF165520	A375	1111	
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AF165520	Hep	1111	
AF165520	Jurkat	1111	
AF165520	Nccit	0100	
AF165520	OVCAR	0100	
AF165520	PC3	1111	
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AF165527	COLO	1	
AF165527	FHS	1	
AF165527	Hep	1	
AF165527	Jurkat	1	
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AF170562	Hep	0110011001001111111110110	
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AF173935	A375	0100100000	
AF173935	CEM	0101100001	
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AF173935	Hep	0000100000	
AF173935	Jurkat	01111111000	
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AF173935	OVCAR	0111100001	
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AF178532	Jurkat	100001111	
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AF178532	OVCAR	100001101	
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AF183411	Hep	011	
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AF183411	PC3	011	
AF183411	Sk	011	
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AF188276	Jurkat	10000111
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AF188276	OVCAR	10000101
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AF188277	Nccit	10000100
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AF190862	Hep	0101111111111111	
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AF197129	A375	1100011	
AF197129	CEM	1100111	
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AF201385	Hep	010001110111001	
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AF203032	Hep	10001	
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AF203032	OVCAR	10010	
AF203032	PC3	10010	
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AF207664	A375	001011111	
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AF211189	A375	00101000011000000010100010100001100	
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AF211189	Hep	000010000000000000010100110000001101
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AF212252	COLO	111111111
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AF212252	Jurkat	100001111
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AF217525	A375	001000001000100000000000000000011
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AF217525	Jurkat	000000001000100000000000000000011
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AF218085	A375	000000000	
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AF218085	Hep	000000000	
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AF218085	OVCAR	000000000	
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AF218085	Sk	101111100	
AF218085	U87	000000000	
AF220490	A375	0001100	
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AF220490	FHS	0101100	
AF220490	Hep	0001100	
AF220490	Jurkat	0000100	
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AF220490	OVCAR	0100100	
AF220490	PC3	0001100	
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AF221520	A375	01	
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AF228707	A375	11110	
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AF229058	A375	100110010110	
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AF231024	Hep	01000100000100000000100000000010
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AF231922	A375	0
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AF231922	Nccit	0	
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AF233436	A375	0000000101001000	
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AF237769	A375	111111000111	
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AF237769	Hep	100110000000	
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AF237772	Jurkat	011110111111	
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AF238328	CEM	0000110	
AF238328	COLO	0001110	
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AF238328	Hep	0001100	
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AF238328	Nccit	0000100	
AF238328	OVCAR	0101110	
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AF240775	A375	000000100011001001	
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AF257472	FHS	000100	
AF257472	Hep	000100	
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AF267865	Nccit	1	
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AF267865	PC3	1	
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AK056681	Nccit	011111111111
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AK056681	PC3	111111111111
AK056681	Sk	011111111110
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AK056738	A375	100000110000110001000
AK056738	CEM	110111111110111101111
AK056738	COLO	110100110100011001001
AK056738	FHS	110000110110000000001
AK056738	Hep	100100110100110001111
AK056738	Jurkat	110101111100111001111
AK056738	Nccit	110000110110011001001
AK056738	OVCAR	110100110110011001101
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AK056738	Sk	100000110100110001001
AK056738	U87	110100110110011001101
AK056895	A375	0
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AK056895	COLO	0
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AK056895	Jurkat	0
AK056895	Nccit	0
AK056895	OVCAR	0
AK056895	PC3	0
AK056895	Sk	0
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AK057029	A375	10001100110
AK057029	CEM	10011111011
AK057029	COLO	11001111110
AK057029	FHS	10001101010

AK057029	Hep	10001100010
AK057029	Jurkat	10001111110
AK057029	Nccit	11001101010
AK057029	OVCAR	11011101110
AK057029	PC3	10001101110
AK057029	Sk	10001111010
AK057029	U87	11001110010
AK057137	A375	0
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AK057137	COLO	0
AK057137	FHS	0
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AK057137	OVCAR	0
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AK057318	COLO	0
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AK057318	Jurkat	0
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AK057318	OVCAR	0
AK057318	PC3	0
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AK057349	A375	0010001
AK057349	CEM	0011000
AK057349	COLO	0011010
AK057349	FHS	0000010

AK057349	Hep	0010000	
AK057349	Jurkat	0010000	
AK057349	Nccit	0010000	
AK057349	OVCAR	0000010	
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AK057397	FHS	000000	
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AK057461	COLO	10110101	
AK057461	FHS	00100001	
AK057461	Hep	10100101	
AK057461	Jurkat	10000100	
AK057461	Nccit	00000000	
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AK058150	Nccit	00	
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AK074042	Hep	0	
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AK074042	Nccit	0	
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AK074091	FHS	010000110100011	
AK074091	Hep	110001100100011	
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AK074091	Nccit	110001110100111	
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AK074094	A375	00010	
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AK074094	Jurkat	00000
AK074094	Nccit	00101
AK074094	OVCAR	00110
AK074094	PC3	00010
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AK074162	COLO	11
AK074162	FHS	01
AK074162	Hep	11
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AK074162	Nccit	11
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AK074300	COLO	1
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AK074300	Nccit	0
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AK074445	Nccit	0	
AK074445	OVCAR	0	
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AL009266	COLO	11100100110	
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AL009266	Hep	10100010110	
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AL009266	OVCAR	10110110110	
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AL021683	Nccit	10	
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AL035364	Jurkat	000	
AL035364	Nccit	000	
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AL050075	Nccit	010	
AL050075	OVCAR	110	
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AL050173	COLO	00	
AL050173	FHS	10	
AL050173	Hep	11	
AL050173	Jurkat	01	
AL050173	Nccit	01	
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AL050173	Sk	00	
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AL050256	PC3	000	
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AL050257	Hep	0000
AL050257	Jurkat	0000
AL050257	Nccit	0100
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AL050258	Hep	11101010111100
AL050258	Jurkat	11101010111101
AL050258	Nccit	11101111101101
AL050258	OVCAR	11111111111101
AL050258	PC3	11111111111001
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AL050258	U87	11111011111100
AL050282	A375	111011111111
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AL050282	COLO	111111111111
AL050282	FHS	111111111111
AL050282	Hep	111111111111
AL050282	Jurkat	111111111111
AL050282	Nccit	111111111111
AL050282	OVCAR	111111111111
AL050282	PC3	111111111111
AL050282	Sk	111011111111
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AL050346	A375	0001100
AL050346	CEM	0001100
AL050346	COLO	0011100
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AL050346	Nccit	0001100	
AL050346	OVCAR	1011100	
AL050346	PC3	1001100	
AL050346	Sk	0001000	
AL050346	U87	1001100	
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AL079310	Jurkat	011111101111	
AL079310	Nccit	000101111111	
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AL080149	Hep	1001111110	
AL080149	Jurkat	1101111110	
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AL080187	FHS	0100011001100000	
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AL080197	Nccit	0	
AL080197	OVCAR	0	
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AL096746	Hep	0	
AL096746	Jurkat	0	
AL096746	Nccit	0	
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AL096780	Hep	01111111111	
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AL096780	PC3	01111111011	
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AL096880	Nccit	1111	
AL096880	OVCAR	1111	
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AL096881	Hep	000001001011	
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AL117466	Hep	011111111111011	
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AL117578	FHS	0	
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AL117578	Jurkat	0	
AL117578	Nccit	0	
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AL136871	A375	010
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AL137288	COLO	0	
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AL137288	Hep	0	
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AL137288	Nccit	0	
AL137288	OVCAR	0	
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AL137314	OVCAR	111111111111	
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AL137448	FHS	0	
AL137448	Hep	1	
AL137448	Jurkat	0	
AL137448	Nccit	0	
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AL137448	Sk	0	
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AL137499	A375	1001	
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AL137499	COLO	1100	
AL137499	FHS	0101	
AL137499	Hep	1000	
AL137499	Jurkat	1001	
AL137499	Nccit	1000	
AL137499	OVCAR	0000	
AL137499	PC3	0000	
AL137499	Sk	1000	
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AL137651	A375	0010	
AL137651	CEM	0000	
AL137651	COLO	1100	
AL137651	FHS	0100	

AL137651	Hep	1110
AL137651	Jurkat	0010
AL137651	Nccit	0000
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AL137672	COLO	00
AL137672	FHS	00
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AL138578	CEM	10000100
AL138578	COLO	11101111
AL138578	FHS	10100110
AL138578	Hep	11101111
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AL138578	Nccit	11100111
AL138578	OVCAR	10001010
AL138578	PC3	10101110
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AL157441	Hep	111111110000
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AL157441	Nccit	111111110000
AL157441	OVCAR	111111110100
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AL157480	Jurkat	111101
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AL157851	Hep	111011
AL157851	Jurkat	111111
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AL160111	A375	111111111
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AL161993	Jurkat	100
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AL355841	A375	1000000
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AL355841	COLO	0000110
AL355841	FHS	0000010

AL355841	Hep	1000000	
AL355841	Jurkat	1000000	
AL355841	Nccit	0000000	
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AL359401	A375	0010	
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AL359401	FHS	0010	
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AL359403	Jurkat	111	
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AL365511	Nccit	000	
AL365511	OVCAR	000	
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AL365513	Hep	0000100011	
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AL389951	FHS	11	
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AL389951	Jurkat	11	
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AL442096	Jurkat	0	
AL442096	Nccit	0	
AL442096	OVCAR	0	
AL442096	PC3	0	
AL442096	Sk	0	
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AL450314	Hep	0000100001	
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AL512711	Sk	01	
AL512711	U87	11	
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AL589867	CEM	01110001	
AL589867	COLO	01110101	
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AL589867	Hep	00010000
AL589867	Jurkat	01111100
AL589867	Nccit	01010000
AL589867	OVCAR	01110001
AL589867	PC3	01010001
AL589867	Sk	00010000
AL589867	U87	01110101
AL590888	A375	01110
AL590888	CEM	01110
AL590888	COLO	11111
AL590888	FHS	00100
AL590888	Hep	01110
AL590888	Jurkat	01110
AL590888	Nccit	01110
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AL590888	PC3	01110
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AL713640	A375	000000
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AL713640	COLO	110101
AL713640	FHS	100001
AL713640	Hep	100100
AL713640	Jurkat	000100
AL713640	Nccit	000000
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AL713698	A375	11111111
AL713698	CEM	11111111
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AL713698	Nccit	11111111
AL713698	OVCAR	10111111
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AL713701	Nccit	11000
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AL713710	Jurkat	1111111
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AY007132	Nccit	1	
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AY014878	Jurkat	00010	
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AY014879	Nccit	000
AY014879	OVCAR	000
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AY014906	Nccit	0000
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AY014914	CEM	000100
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AY016009	FHS	10111111

AY016009	Hep	10001100
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AY028896	COLO	1010001111111111111
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AY028896	Hep	1011111111111011111
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AY029208	A375	11101101001010100111
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AY035381	COLO	0	
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AY035383	Hep	0000
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AY035383	Nccit	0000
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AY039243	A375	0000010
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AY062123	CEM	011111111111
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AY062123	FHS	01000001101
AY062123	Hep	11011011111
AY062123	Jurkat	11011011111
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AY062123	OVCAR	11001011101
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AY077698	Hep	10001100
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AY077698	Nccit	10001000
AY077698	OVCAR	10000100
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AY077700	Hep	110000000010001001100100000010010000100000000
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AY077700	Nccit	1100000000000000000000100000000010000100000000
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D45906	Nccit	0111111101111010
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L21950	FHS	011

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L21950	Nccit	010	
L21950	OVCAR	010	
L21950	PC3	111	
L21950	Sk	011	
L21950	U87	111	
L25781	A375	1	
L25781	CEM	1	
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L25781	FHS	0	
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L25781	Jurkat	1	
L25781	Nccit	1	
L25781	OVCAR	0	
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L38562	CEM	1	
L38562	COLO	0	
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L38562	Hep	0	
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L38562	Nccit	0	
L38562	OVCAR	0	
L38562	PC3	0	
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L43092	COLO	0	
L43092	FHS	0	
L43092	Hep	0	
L43092	Jurkat	0	
L43092	Nccit	0	
L43092	OVCAR	0	
L43092	PC3	0	
L43092	Sk	0	
L43092	U87	0	
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L75823	CEM	111111111	
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L75823	Hep	111111111	
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L75823	Nccit	111111011	
L75823	OVCAR	111111111	
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L78790	CEM	1111111111111	
L78790	COLO	0100000001000	
L78790	FHS	0100000000100	
L78790	Hep	1111110010100	
L78790	Jurkat	1111111111110	
L78790	Nccit	0100000000100	
L78790	OVCAR	0110000000100	
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M12783	COLO	1001011	
M12783	FHS	1000011	
M12783	Hep	1100010	
M12783	Jurkat	0000000	
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M12849	Nccit	0000	
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M15395	Hep	11111100101100	
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M15533	COLO	11	
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M15533	Sk	11	
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M19730	COLO	1	
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M19730	Nccit	1	
M19730	OVCAR	1	
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M21535	Nccit	000000000	
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M26062	CEM	0011000100	
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M26062	Hep	0011000010
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M26062	Nccit	0011000110
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M27749	CEM	111
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M30474	CEM	11111
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M30474	Hep	11011	
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NM_012131	FHS	1	
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[illegible]

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U20980	FHS	0010000010110	
U20980	Hep	0110010011110	
U20980	Jurkat	0111010111111	
U20980	Nccit	1111010111111	
U20980	OVCAR	1111010111111	
U20980	PC3	0111010111111	
U20980	Sk	0110010011110	
U20980	U87	1111010111111	
U22526	A375	000010110101001011	
U22526	CEM	111111111111101111	
U22526	COLO	111111111111001111	
U22526	FHS	111111111111001111	

U22526	Hep	111111111111101111
U22526	Jurkat	111111110111101111
U22526	Nccit	111111111111001111
U22526	OVCAR	111111111111001111
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U22526	Sk	111011110111001101
U22526	U87	111111111111001111
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U35340	CEM	000000
U35340	COLO	000000
U35340	FHS	000000
U35340	Hep	000000
U35340	Jurkat	000000
U35340	Nccit	111000
U35340	OVCAR	000000
U35340	PC3	000001
U35340	Sk	000000
U35340	U87	100000
U44059	A375	1001
U44059	CEM	1001
U44059	COLO	1111
U44059	FHS	1011
U44059	Hep	1011
U44059	Jurkat	1001
U44059	Nccit	1010
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U44059	PC3	1011
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U44059	U87	1111
U45975	A375	0000000
U45975	CEM	0010000
U45975	COLO	1000000
U45975	FHS	0010000

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U45975	Jurkat	0000000	
U45975	Nccit	1010000	
U45975	OVCAR	1010100	
U45975	PC3	0010000	
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U49928	FHS	11000111111	
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U49928	Jurkat	11000101101	
U49928	Nccit	11000111101	
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U51269	A375	00000001000111000	
U51269	CEM	00001001000101001	
U51269	COLO	00011001010111000	
U51269	FHS	00011000010010000	
U51269	Hep	10011111000111001	
U51269	Jurkat	00000001000111001	
U51269	Nccit	00001111000111000	
U51269	OVCAR	10011111010011001	
U51269	PC3	00001001000011001	
U51269	Sk	00000001000111000	
U51269	U87	10001000010001001	
U52962	A375	001011000101011101001000000001100100100011010010	
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U52962	COLO	111111110100011101011001110001111101111111111111	
U52962	FHS	101011100100001101001000000001101101111101000101	

U52962	Hep	000011010100010101001100010001111101100111100110
U52962	Jurkat	001111110111111111111111010110111111111111111111
U52962	Nccit	00111111011111111111111101111111111111111111011111
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U53442	CEM	1111111111
U53442	COLO	0010010001
U53442	FHS	0111010001
U53442	Hep	0011010001
U53442	Jurkat	0011010101
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U53442	OVCAR	0110010001
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U59057	COLO	00110
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U59057	Hep	01110
U59057	Jurkat	01010
U59057	Nccit	00110
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U59057	PC3	00110
U59057	Sk	10010
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U59632	A375	10011100110
U59632	CEM	10010100111
U59632	COLO	00000100101
U59632	FHS	00110100110

U59632	Hep	00000100111
U59632	Jurkat	01011100110
U59632	Nccit	00101100111
U59632	OVCAR	00111110110
U59632	PC3	10111111111
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U59632	U87	00110110110
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U61083	CEM	11001
U61083	COLO	11001
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U61083	Hep	10010
U61083	Jurkat	10000
U61083	Nccit	00000
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U61084	A375	11111010
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U61084	Hep	11110010
U61084	Jurkat	11110000
U61084	Nccit	00000000
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U61500	A375	0111100010111100111101
U61500	CEM	1111101111111111111111
U61500	COLO	1111101111111111111111
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U61500	Hep	0111101010111111111111
U61500	Jurkat	0111101111111111111111
U61500	Nccit	0111101111111111111111
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U61500	Sk	0011000011111100111111
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U64444	CEM	1111111111
U64444	COLO	1011111111
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U64444	Hep	1011111101
U64444	Jurkat	1111111101
U64444	Nccit	1011111111
U64444	OVCAR	1011111111
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U65011	A375	11110
U65011	CEM	01000
U65011	COLO	00000
U65011	FHS	00000
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U65011	Jurkat	11110
U65011	Nccit	00000
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U65011	Sk	11110
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U66828	CEM	11111111111111111111
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U66828	Hep	000010110101110010
U66828	Jurkat	010011110111111111
U66828	Nccit	011010110001110110
U66828	OVCAR	001000000100110010
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U66828	Sk	010000000000111101
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U68727	A375	010010111110
U68727	CEM	11111101111
U68727	COLO	11001111111
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U68727	Hep	11001111111
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U68727	U87	11111101111
U71216	A375	0000
U71216	CEM	0001
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U71216	FHS	0100
U71216	Hep	0100
U71216	Jurkat	0000
U71216	Nccit	0101
U71216	OVCAR	0100
U71216	PC3	0100
U71216	Sk	0000
U71216	U87	0100
U74628	A375	1001110011
U74628	CEM	1001010011
U74628	COLO	0000010011
U74628	FHS	0011010011

U74628	Hep	0000010011	
U74628	Jurkat	0101110011	
U74628	Nccit	0010110011	
U74628	OVCAR	0011111011	
U74628	PC3	1011111111	
U74628	Sk	1011110011	
U74628	U87	0011011011	
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U75329	COLO	1111110111111	
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U75329	Hep	0101010100010	
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U75329	Nccit	0001010000010	
U75329	OVCAR	0100100010001	
U75329	PC3	1101110111111	
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U75330	A375	0100011010000100	
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U75330	Jurkat	0000010000000100	
U75330	Nccit	0100110000000100	
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U75330	PC3	0101010000000110	
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U78310	CEM	011111111111110	
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U78310	Hep	01111111111110
U78310	Jurkat	01111111111110
U78310	Nccit	01111111111110
U78310	OVCAR	11111111111110
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U78310	U87	11111111111110
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U79252	CEM	1
U79252	COLO	1
U79252	FHS	0
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U79252	Jurkat	1
U79252	Nccit	0
U79252	OVCAR	0
U79252	PC3	0
U79252	Sk	0
U79252	U87	0
U79280	A375	1
U79280	CEM	0
U79280	COLO	0
U79280	FHS	1
U79280	Hep	0
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U79280	PC3	0
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U79286	A375	0111111
U79286	CEM	1111110
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U79286	Nccit	1111110
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U79286	PC3	1111110
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U79754	CEM	001011000000111
U79754	COLO	111011011010110
U79754	FHS	101001000010010
U79754	Hep	000011000000110
U79754	Jurkat	000001010000100
U79754	Nccit	111111111110110
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U79754	PC3	000010010000010
U79754	Sk	000001000000000
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U80456	A375	00010000000
U80456	CEM	01010000000
U80456	COLO	11010110111
U80456	FHS	11010110110
U80456	Hep	01010000010
U80456	Jurkat	01010000000
U80456	Nccit	01010000000
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U80456	PC3	11111111111
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U82381	A375	000011000000
U82381	CEM	001011000001
U82381	COLO	111011011011
U82381	FHS	101001000010

U82381	Hep	000011000001	
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U82381	PC3	000010010000	
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U85431	A375	10011	
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U85431	Jurkat	01011	
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U93181	Hep	111011111111011011101101111101111
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U94352	Hep	00110011
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U94352	PC3	01010011
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U96394	A375	0
U96394	CEM	1
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X00474	COLO	110
X00474	FHS	010

X00474	Hep	100	
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X00474	PC3	111	
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X01147	FHS	00	
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X13967	A375	111	
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X14675	PC3	011
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X14676	A375	01
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X14789	Jurkat	0
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X15977	OVCAR	1
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X59656	CEM	111
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X68486	A375	1000	
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X69117	Hep	000101010111011001000
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X82209	A375	00
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X82209	Nccit	00	
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X82260	Hep	11111011111011	
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X85237	A375	111011111111110	
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X89887	A375	000100010110101011001000
X89887	CEM	111110111111111011111110
X89887	COLO	101110111111111111111111
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X89887	Hep	100110111111111011111110
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X89986	A375	0101
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X89986	COLO	1111
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X89986	Hep	1101	
X89986	Jurkat	1111	
X89986	Nccit	0101	
X89986	OVCAR	1001	
X89986	PC3	1111	
X89986	Sk	0100	
X89986	U87	1101	
X90980	A375	00	
X90980	CEM	10	
X90980	COLO	00	
X90980	FHS	00	
X90980	Hep	00	
X90980	Jurkat	10	
X90980	Nccit	00	
X90980	OVCAR	00	
X90980	PC3	00	
X90980	Sk	00	
X90980	U87	00	
X92715	A375	0111	
X92715	CEM	1111	
X92715	COLO	1111	
X92715	FHS	0111	
X92715	Hep	0101	
X92715	Jurkat	0111	
X92715	Nccit	0111	
X92715	OVCAR	1111	
X92715	PC3	1111	
X92715	Sk	0111	
X92715	U87	0111	
X95263	A375	010011110101101001110	
X95263	CEM	010011111111111011111	
X95263	COLO	000011101101101001110	
X95263	FHS	000011000101100001000	

[illegible]

X99584	Hep	111
X99584	Jurkat	111
X99584	Nccit	111
X99584	OVCAR	111
X99584	PC3	111
X99584	Sk	111
X99584	U87	111
Y00264	A375	1111111111111
Y00264	CEM	0000000010100
Y00264	COLO	1111111111111
Y00264	FHS	1111111111111
Y00264	Hep	1111111111111
Y00264	Jurkat	0000000010000
Y00264	Nccit	1111111111111
Y00264	OVCAR	1111111111111
Y00264	PC3	1111111111111
Y00264	Sk	1111111111111
Y00264	U87	1111111111111
Y00661	A375	0110010010000100111001
Y00661	CEM	1111111111111100111111
Y00661	COLO	1111111111111100111111
Y00661	FHS	1111111110100000111011
Y00661	Hep	1111111111111100111011
Y00661	Jurkat	111111111111110111011
Y00661	Nccit	1111111110111100111111
Y00661	OVCAR	1111111110111100111111
Y00661	PC3	1111111111111100111111
Y00661	Sk	1110110010000100111001
Y00661	U87	1111111110111100111111
Y00970	A375	00010
Y00970	CEM	00000
Y00970	COLO	00010
Y00970	FHS	00000

Y00970	Hep	00000
Y00970	Jurkat	00010
Y00970	Nccit	00010
Y00970	OVCAR	00010
Y00970	PC3	00010
Y00970	Sk	00010
Y00970	U87	00010
Y07593	A375	1000000
Y07593	CEM	0000000
Y07593	COLO	1000000
Y07593	FHS	0000000
Y07593	Hep	1111111
Y07593	Jurkat	0101010
Y07593	Nccit	1111111
Y07593	OVCAR	1111111
Y07593	PC3	1101000
Y07593	Sk	0101110
Y07593	U87	1000000
Y07846	A375	11011
Y07846	CEM	11111
Y07846	COLO	11011
Y07846	FHS	11011
Y07846	Hep	11010
Y07846	Jurkat	11111
Y07846	Nccit	11010
Y07846	OVCAR	11111
Y07846	PC3	11111
Y07846	Sk	11010
Y07846	U87	11111
Y07847	A375	000
Y07847	CEM	001
Y07847	COLO	001
Y07847	FHS	000

Y07847	Hep	000
Y07847	Jurkat	000
Y07847	Nccit	000
Y07847	OVCAR	001
Y07847	PC3	001
Y07847	Sk	001
Y07847	U87	001
Y10745	A375	10
Y10745	CEM	00
Y10745	COLO	00
Y10745	FHS	00
Y10745	Hep	10
Y10745	Jurkat	10
Y10745	Nccit	00
Y10745	OVCAR	00
Y10745	PC3	00
Y10745	Sk	10
Y10745	U87	00
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Y10776	CEM	000001
Y10776	COLO	000001
Y10776	FHS	000000
Y10776	Hep	000001
Y10776	Jurkat	000001
Y10776	Nccit	000000
Y10776	OVCAR	000000
Y10776	PC3	000001
Y10776	Sk	000001
Y10776	U87	000011
Y11392	A375	000100
Y11392	CEM	100110
Y11392	COLO	100101
Y11392	FHS	000101

Y11392	Hep	100100
Y11392	Jurkat	000100
Y11392	Nccit	100101
Y11392	OVCAR	100101
Y11392	PC3	100111
Y11392	Sk	000000
Y11392	U87	110101
Y12478	A375	11011
Y12478	CEM	11111
Y12478	COLO	11111
Y12478	FHS	01111
Y12478	Hep	11111
Y12478	Jurkat	11111
Y12478	Nccit	11111
Y12478	OVCAR	11111
Y12478	PC3	11111
Y12478	Sk	11011
Y12478	U87	11111
Y13492	A375	0011101111110011011
Y13492	CEM	0010000101001010010
Y13492	COLO	11111011111001010010
Y13492	FHS	1111111111101011111
Y13492	Hep	0111101110111011011
Y13492	Jurkat	0011000110011000010
Y13492	Nccit	1010000100001000000
Y13492	OVCAR	11111111111001011111
Y13492	PC3	1111111111101011111
Y13492	Sk	0001000100001000000
Y13492	U87	1111111111111011111
Y16186	A375	00
Y16186	CEM	01
Y16186	COLO	01
Y16186	FHS	10

Y16186	Hep	01	
Y16186	Jurkat	01	
Y16186	Nccit	00	
Y16186	OVCAR	00	
Y16186	PC3	01	
Y16186	Sk	00	
Y16186	U87	01	
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Z11695	Hep	1111111111	
Z11695	Jurkat	1111111110	
Z11695	Nccit	1111111110	
Z11695	OVCAR	1111111111	
Z11695	PC3	1111111111	
Z11695	Sk	1111111110	
Z11695	U87	1111111111	
Z17227	A375	1111011	
Z17227	CEM	1111111	
Z17227	COLO	1111111	
Z17227	FHS	1111111	
Z17227	Hep	1101111	
Z17227	Jurkat	1111111	
Z17227	Nccit	1000010	
Z17227	OVCAR	1111111	
Z17227	PC3	1111111	
Z17227	Sk	1101011	
Z17227	U87	1111111	
Z25423	A375	00	
Z25423	CEM	11	
Z25423	COLO	11	
Z25423	FHS	11	

Z25423	Hep	11
Z25423	Jurkat	11
Z25423	Nccit	11
Z25423	OVCAR	11
Z25423	PC3	11
Z25423	Sk	11
Z25423	U87	11
Z49989	A375	11110011011
Z49989	CEM	11001010010
Z49989	COLO	11001010010
Z49989	FHS	11101011111
Z49989	Hep	10111011011
Z49989	Jurkat	10011000010
Z49989	Nccit	10001000000
Z49989	OVCAR	01001011111
Z49989	PC3	11101011111
Z49989	Sk	10001000000
Z49989	U87	11111011111
Z97990	A375	0000101011000
Z97990	CEM	0000011001000
Z97990	COLO	0000001001001
Z97990	FHS	0000011001000
Z97990	Hep	0000101001000
Z97990	Jurkat	0000110001000
Z97990	Nccit	0000011001000
Z97990	OVCAR	0000011001001
Z97990	PC3	0000011001000
Z97990	Sk	0000101001000
Z97990	U87	0000011001000

Table S5. Experimentally validated novel regions.

Name	Start	End	Length	Longest ORF ¹	Acc. No.	dbEST_Id
Chr21-32	13532249	13532717	468	224	CF798425	20136229
Chr21-33	13573145	13573389	244	124	CF798426	20136230
Chr21-34	13112117	13112586	469	205	CF798427	20136231
Chr21-35	14000589	14000894	305	136	CF798428	20136232
Chr21-36	20756699	20757116	417	216	CF798429	20136233
Chr21-37	21035695	21036469	774	211	CF798430	20136234
Chr21-38	21130234	21130384	150	151	CF798431	20136235
Chr21-39	37595837	37597007	1170	469	CF798432	20136236
Chr21-40	37703737	37704257	520	480	CF798433	20136237
Chr21-41	37762535	37763019	484	214	CF798434	20136238
Chr21-42	37786389	37786628	239	212	CF798435	20136239
Chr21-43	37793687	37794002	315	275	CF798436	20136240
Chr21-44	37834234	37834524	290	259	CF798437	20136241
Chr21-45	38009278	38009633	355	128	CF798438	20136242
Chr21-46	38078621	38078999	378	259	CF798439	20136243
Chr21-47	38122379	38122534	155	105	CF798440	20136244
Chr21-48	38159580	38159800	220	148	CF798441	20136245
Chr21-49	38214744	38215386	642	235	CF798442	20136246
Chr21-50	42981570	42981897	327	328	CF798443	20136247
Chr21-51	42994508	42994855	347	217	CF798444	20136248
Chr21-52	43509473	43509974	501	344	CF798445	20136249
Chr21-53	43558305	43558600	295	259	CF798446	20136250
Chr21-54	43626860	43627238	378	379	CF798447	20136251
Chr21-55	43747072	43747514	442	433	CF798448	20136252
Chr22-14	21064484	21064911	428	307	CF798449	20136253
Chr22-15	22501144	22501474	331	295	CF798450	20136254
Chr22-16	23918058	23918261	204	165	CF798451	20136255
Chr22-17	34209281	34209713	433	355	CF798452	20136256
Chr22-18	15419554	15420051	498	360	CF798453	20136257
Chr22-19	35344202	35344808	607	319	CF798454	20136258
Chr22-20	35345199	35345532	334	334	CF798455	20136259
Chr22-21	37662387	37662780	394	386	CF798456	20136260
Chr22-22	39284594	39284803	210	159	CF798457	20136261
Chr22-23	43555031	43555394	363	260	CF798458	20136262
Chr22-24	43556735	43557113	378	379	CF798459	20136263
Chr22-25	46177453	46177838	385	375	CF798460	20136264
Chr22-26	46376450	46376631	181	182	CF798461	20136265
Chr22-27	46484726	46485203	478	259	CF798462	20136266
Chr22-28	46710711	46711159	449	274	CF798463	20136267
Chr22-29	17126473	17126742	270	202	CF798464	20136268
Chr22-30	18421277	18421618	342	342	CF798465	20136269

Chr22-31	16525822	16526489	667	610	CF798466	20136270
Chr22-32	16746580	16747170	590	511	CF798467	20136271
Chr22-33	16700390	16701169	779	445	CF798468	20136272
Chr22-34	16663572	16663979	407	408	CF798469	20136273
Chr22-35	16569847	16570242	395	396	CF798470	20136274
Chr22-36	16572090	16572591	501	259	CF798471	20136275
Chr22-37	44401517	44401846	329	166	CF798472	20136276
Chr22-38	44780193	44780473	280	194	CF798473	20136277
Chr22-39	45172041	45172339	298	241	CF798474	20136278
Chr22-40	45277896	45278535	639	349	CF798475	20136279
Chr22-41	45320687	45321613	926	535	CF798476	20136280
Chr22-42	32609718	32609944	226	157	CF798477	20136281
Chr22-43	30171596	30172195	599	283	CF798478	20136282
Chr22-44	34414001	34414604	603	346	CF798479	20136283
Chr22-45	24870996	24871394	398	390	CF798480	20136284
Chr22-46	41621808	41622224	416	386	CF798481	20136285
Chr22-47	43085598	43086106	508	400	CF798482	20136286
Chr22-48	36199847	36200359	512	247	CF798483	20136287
Chr22-49	36319938	36320429	491	433	CF798484	20136288
Chr22-50	36336780	36337066	286	287	CF798485	20136289
Chr22-51	36710439	36710919	480	481	CF798486	20136290
Chr22-52	36738851	36739509	658	442	CF798487	20136291
Chr22-53	36909766	36909951	185	147	CF798488	20136292
Chr22-54	36930827	36931125	298	296	CF798489	20136293
Chr22-55	37402848	37403150	302	157	CF798490	20136294
Chr22-56	37478099	37478560	461	181	CF798491	20136295
Chr22-57	37728943	37729266	323	160	CF798492	20136296
Chr22-58	38049485	38049857	372	196	CF798493	20136297
Chr22-59	38301254	38301590	336	254	CF798494	20136298
Chr22-60	38565658	38566195	537	352	CF798495	20136299
Chr22-61	38981341	38981624	283	284	CF798496	20136300
Chr22-62	27849091	27849289	198	199	CF798497	20136301
Chr22-63	27892375	27892715	340	286	CF798498	20136302
Chr22-64	28234525	28234719	194	195	CF798499	20136303
Chr22-65	28628149	28628410	261	192	CF798500	20136304
Chr22-66	43554887	43555434	547	271	CF798501	20136305
Chr22-67	14335764	14335909	145	146	CF798502	20136306
Chr22-68	18225097	18225803	706	274	CF798503	20136307
Chr22-69	20518824	20519137	313	215	CF798504	20136308
Chr22-70	20159838	20160306	468	379	CF798505	20136309
Chr22-71	20161136	20161594	458	223	CF798506	20136310

1. Longest ORF defined as a stretch of the corresponding genomic DNA without a stop codon. All coordinates were obtained from the NCBI version 30.