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Table 1. Time course of cell viability after bleomycin 30 micro gram/ml treatment.

	24 hours after mean(95%CI)	48 hours after mean(95%CI)	72 hours after mean(95%CI)
Control	75 (43-106)	63 (47-78)	60 (48-72)
BLM (30 μ g/ml)	66 (34-97)	49 (35-64)	35 (24-46)

Initially, 6×10^5 of lymphocytes were seeded on 25 cm² flask.

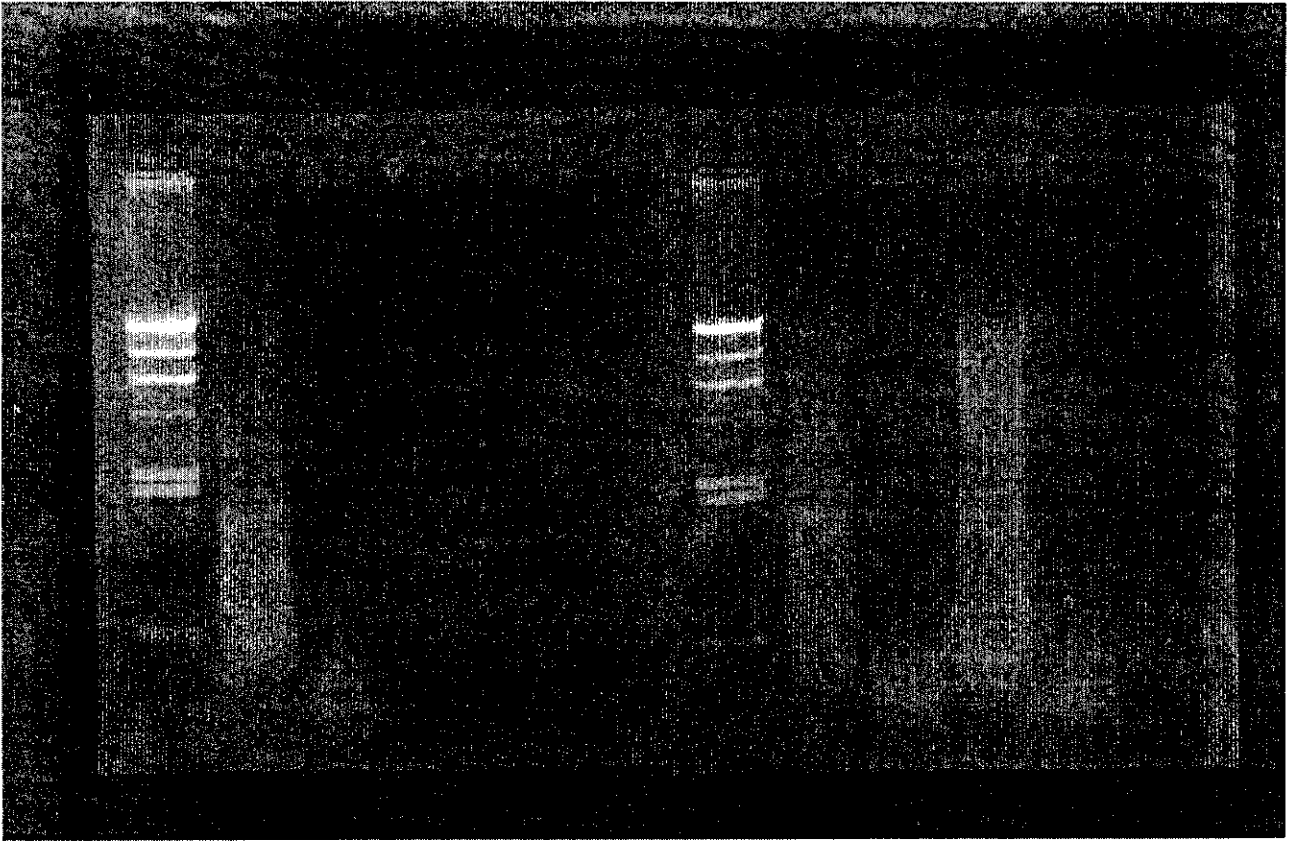


Figure 1. Gel Electrophoresis of cRNA.

(Left)

1. size marker λ / Hind III
2. cRNA of control cells
3. after fragmentation

(Right)

1. size marker λ / Hind III
2. cRNA of 7.5 micro g/ml bleomycin treatment cells
3. 2. after fragmentation
4. cRNA of 30 micro g/ml bleomycin treatment cells
5. 4. after fragmentation

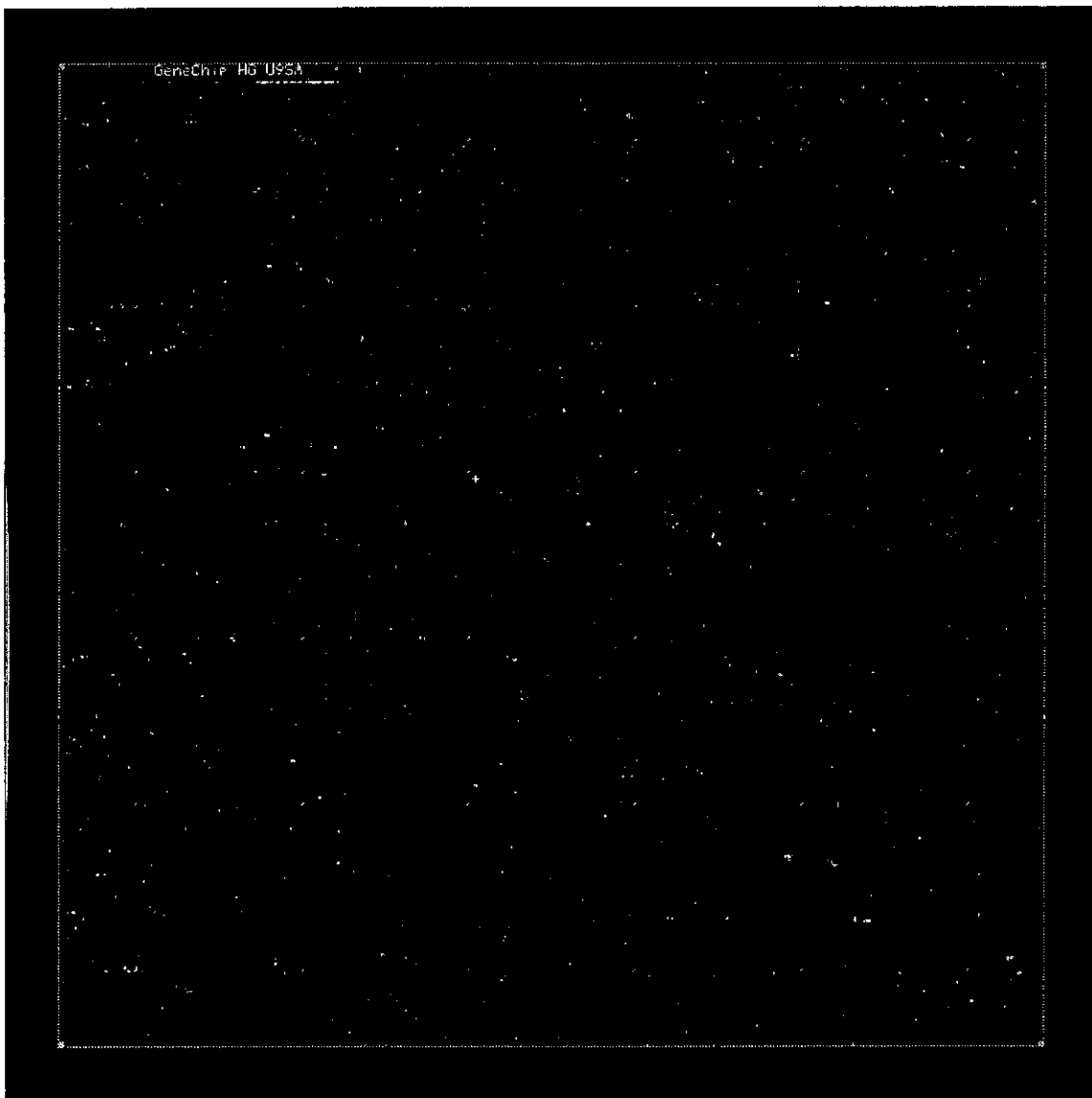


Figure 2.
Scanning Image of Control Cells.

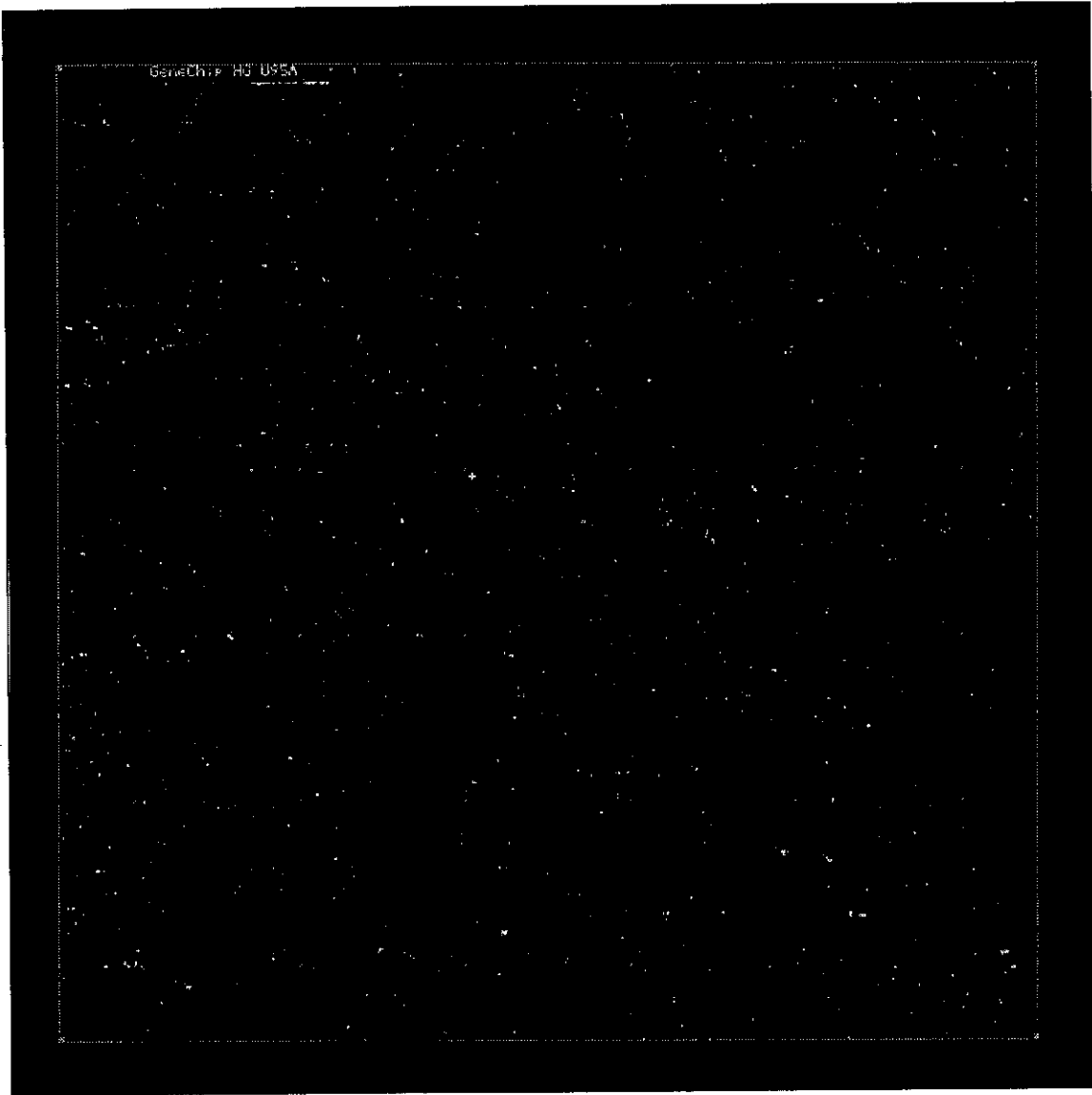


Figure 3.

Scanning Image of 7.5 micro gram/ml BLM Treated Cells.

Blm3.dat

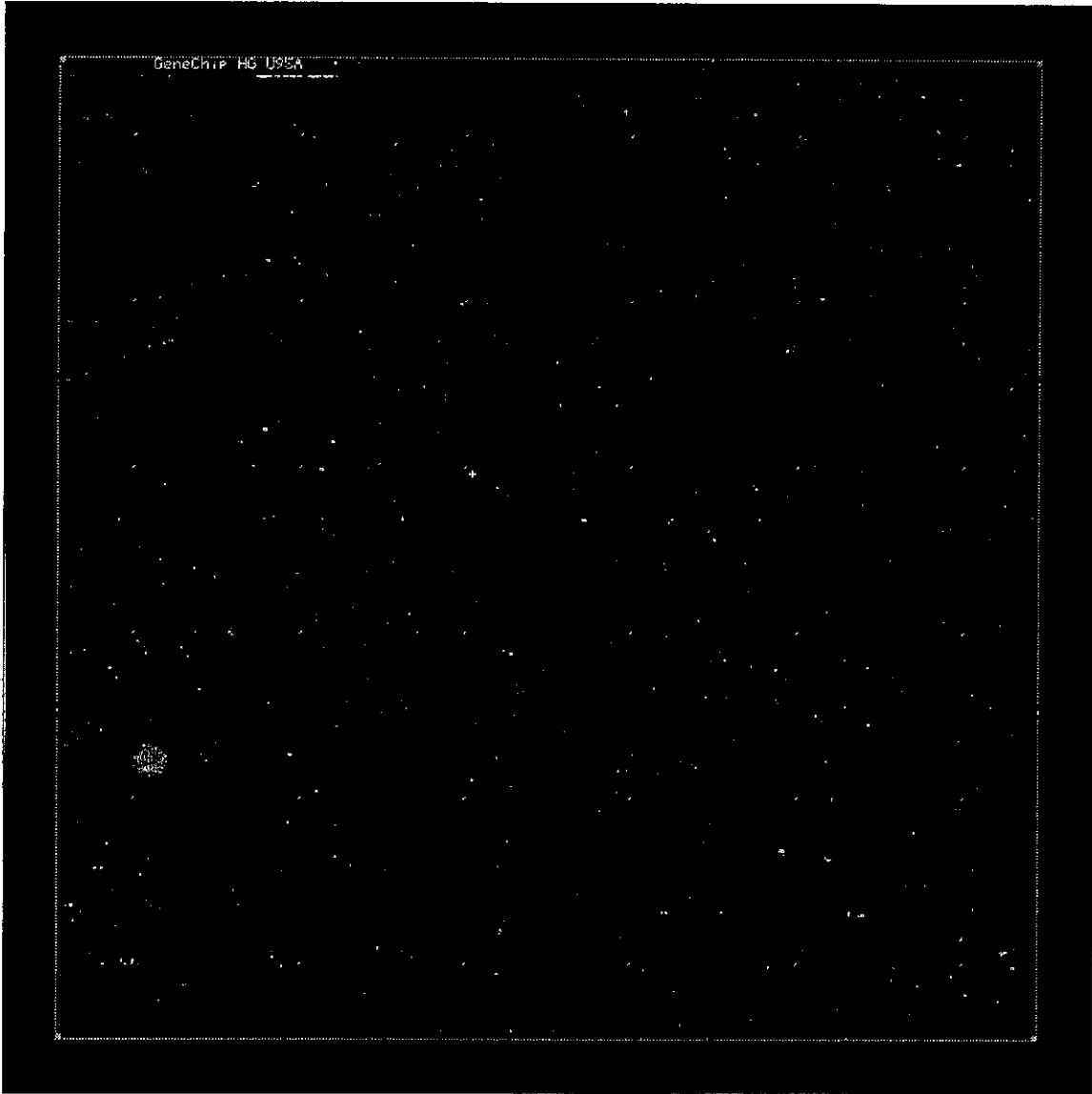


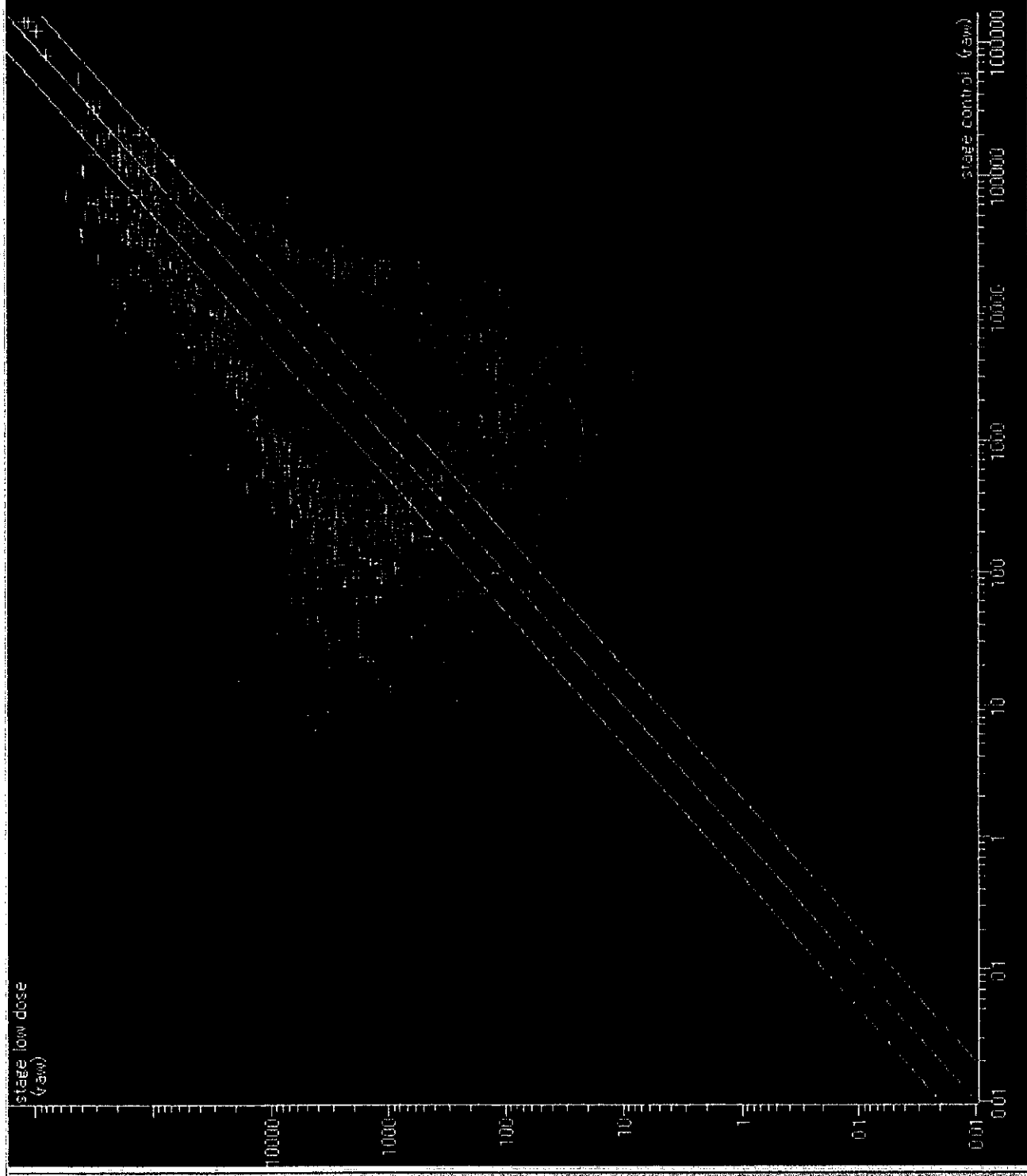
Figure 4.

Scanning Image of 30 micro gram/ml BLM Treated Cells.

- シーンリスト
- 実験
 - bleo-treat series 1
 - Default Interpretation
 - stage control
 - stage high dose
 - stage low dose
 - All experiments
 - bleo-treat series
 - test
- シーンツリー
- 実験ツリー
- クラスファイケーション
- Pathways
- 外部プログラム
- アニメーション
- Drawn Genes

Figure 5.

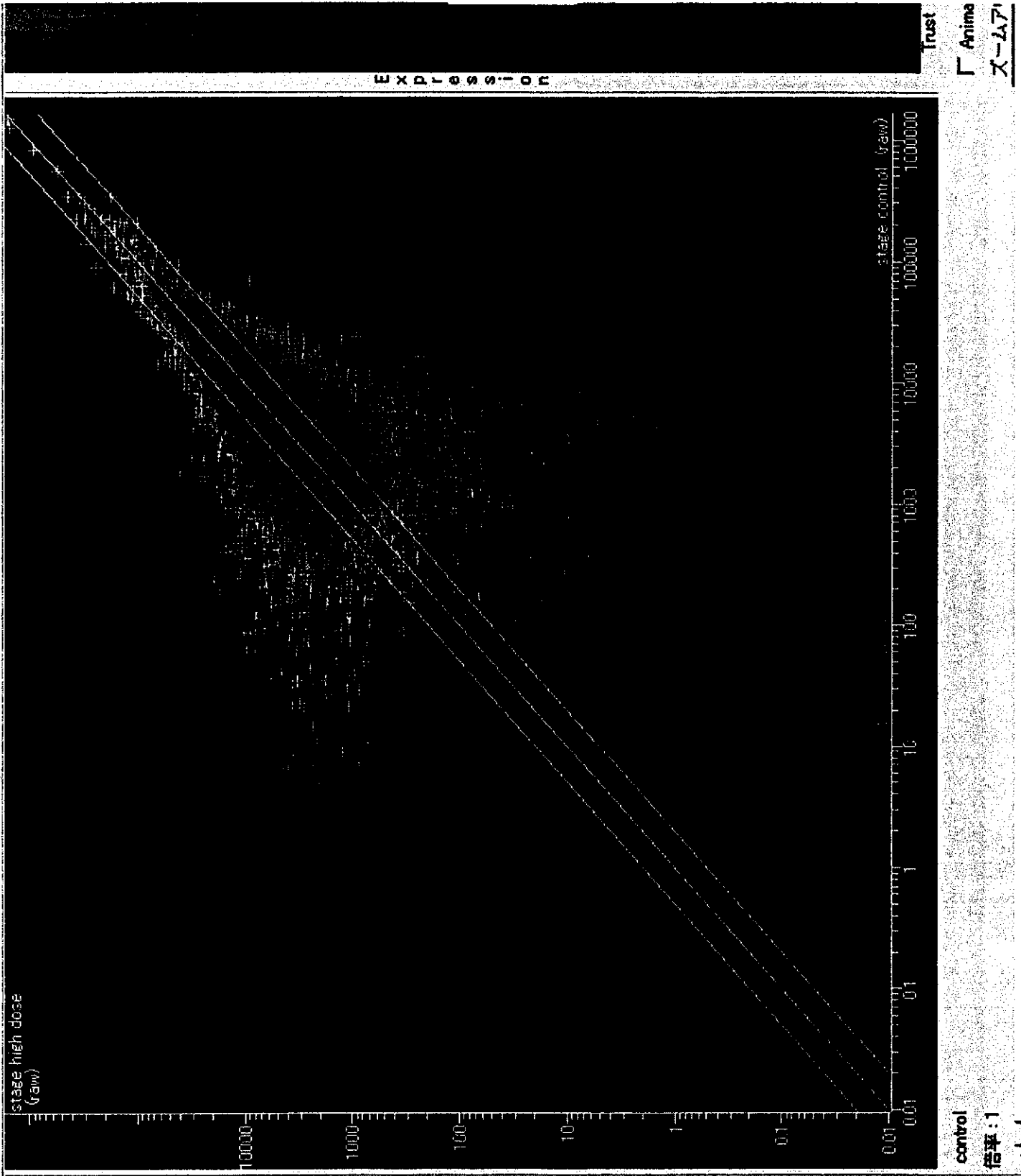
Scatter Plot. Control (x-axis) vs 7.5 micro gram/ml BLM (y-axis)



EXPRESSION

Trust
 Animat
 ZoomArc

control
 倍率: 1



- シーンリスト
- 実験
 - bleo-treat series 1
 - Default interpretation
 - stage control
 - stage high dose
 - stage low dose
 - All experiments
 - bleo-treat series
 - test
- シーンズリー
- 実験ツリー
- クラシフィケーション
- Pathways
- 外部プログラム
- アニメーション
- Drawn Genes

Figure 6.
Scatter Plot. Control (x-axis) vs 30 micro gram/ml BLM (y-axis)

Table 2-1.

Probe Set Name	Abs Call	Diff Call	Avg Diff Change	Fold Change
37463_r_at	A	NC	-293551.6	-35.8
34301_r_at	A	NC	-263839.1	-20.3
31947_r_at	A	NC	-35406.8	-11.4
40675_r_at	A	MD	-29240.9	-9.6
36440_at	A	NC	-28947.8	-9.5
36547_r_at	A	NC	-27796.3	-9.2
1152_i_at	P	NC	-27027.7	-8.9
35387_r_at	A	NC	-24857	-8.3
32468_f_at	A	NC	-23515.2	-7.9
39624_at	A	MD	-22979.6	-7.7
32484_at	A	MD	-22132.7	-7.5
40186_at	A	MD	-21811.1	-7.4
36848_r_at	A	NC	-21190.6	-7.2
34549_g_at	A	NC	-20575.7	-7
124_i_at	A	NC	-20259.2	-6.9
33568_at	A	NC	-19105.7	-6.6
1187_at	M	NC	-18417	-6.4
37785_at	A	NC	-17721.7	-6.2
36284_at	A	NC	-17082.2	-6
37877_at	A	NC	-17058.9	-6
31500_at	A	NC	-16264.9	-5.8
32031_at	A	NC	-16420.8	-5.8
40371_at	A	NC	-15933.5	-5.7
1704_at	A	NC	-17955.6	-5.6
35555_r_at	A	NC	-15576.9	-5.6
236_at	A	NC	-15449.7	-5.5
1493_r_at	A	NC	-14883.4	-5.4
32893_s_at	A	D	-14850.2	-5.4
34110_g_at	A	NC	-14950.8	-5.4
1111_at	A	NC	-14659.7	-5.3
32957_g_at	A	NC	-14545.7	-5.3
39560_at	A	MD	-14545.5	-5.3
35068_at	A	MD	-14236.3	-5.2
38569_at	A	NC	-14157.2	-5.2
32109_at	A	NC	-13847.7	-5.1
37455_at	A	NC	-13960.7	-5.1
38937_at	A	NC	-14001	-5.1
1663_at	A	NC	-13636.9	-5
31434_at	A	NC	-13628.8	-5
31705_at	A	NC	-13599.5	-5
31664_at	A	NC	-13288.9	-4.9
34094_i_at	A	NC	-13172	-4.9
35828_at	A	NC	-13365.9	-4.9
35917_at	A	NC	-13781.9	-4.9
37473_at	A	NC	-13337.8	-4.9
40944_at	A	NC	-13445.8	-4.9

Table 2-2.

Probe Set Name	Abs Call	Diff Call	Avg Diff Change	Fold Change
41566_at	A	MD	-13395.7	-4.9
110_at	A	NC	-12927.8	-4.8
31639_f_at	A	D	-12984.1	-4.8
34412_s_at	A	MD	-13085.2	-4.8
34949_at	A	NC	-12785.1	-4.8
36037_g_at	A	NC	-12811.6	-4.8
37993_at	A	NC	-12797.3	-4.8
38311_at	A	MD	-12855.9	-4.8
38535_at	A	NC	-12953.4	-4.8
40038_at	A	NC	-12927.9	-4.8
1944_f_at	A	NC	-12481.7	-4.7
33545_at	A	NC	-12699.4	-4.7
34365_at	A	MD	-12713.4	-4.7
38203_at	A	NC	-12580.3	-4.7
39668_at	A	NC	-12700.3	-4.7
754_s_at	A	NC	-12512.5	-4.7
2089_s_at	A	NC	-12371.7	-4.6
32108_at	A	NC	-12214.7	-4.6
34320_at	A	NC	-12241.7	-4.6
37369_s_at	A	NC	-12328.2	-4.6
38799_at	A	NC	-12367.5	-4.6
40963_at	A	NC	-12142.6	-4.6
1824_s_at	A	NC	-11926.1	-4.5
31663_at	A	MD	-11996.5	-4.5
36720_at	A	NC	-11794	-4.5
38787_at	A	NC	-12030.6	-4.5
39383_at	A	NC	-11972.5	-4.5
417_at	A	MD	-11906.9	-4.5
41748_at	A	D	-12001.2	-4.5
745_at	A	D	-11905.6	-4.5
32943_at	A	MD	-11421.5	-4.4
34126_at	A	D	-11534.7	-4.4
34555_at	A	NC	-11557.6	-4.4
35451_s_at	A	NC	-11553	-4.4
38963_i_at	A	NC	-19101.2	-4.4
39915_at	A	NC	-11728.7	-4.4
40753_at	A	NC	-11483.4	-4.4
41311_f_at	A	NC	-11637.8	-4.4
1561_at	A	NC	-11078.8	-4.3
31341_at	A	NC	-11242.5	-4.3
32614_at	A	D	-11120.4	-4.3
34494_at	A	NC	-11184	-4.3
39818_at	A	D	-11157.3	-4.3
41446_f_at	A	NC	-11404.2	-4.3
298_at	A	MD	-10848.7	-4.2
31340_at	A	D	-10945.9	-4.2

Table 2-3.

Probe Set Name	Abs Call	Diff Call	Avg Diff Change	Fold Change
31638_at	A	NC	-10843.5	-4.2
31825_at	A	NC	-11005.2	-4.2
32498_at	A	NC	-10813.3	-4.2
32894_at	A	NC	-10892.1	-4.2
33230_at	A	NC	-11015.3	-4.2
33627_at	A	NC	-10956.2	-4.2
34154_at	A	NC	-10739.4	-4.2
35381_at	A	NC	-10753.4	-4.2
36106_at	A	D	-10848.8	-4.2
36148_at	A	NC	-10770.2	-4.2
36787_at	A	NC	-10965.2	-4.2
40163_r_at	A	NC	-22527.8	-4.2
40703_at	A	NC	-10802.6	-4.2
40976_at	A	NC	-11015.8	-4.2
1607_at	A	MD	-10493.2	-4.1
2067_f_at	A	NC	-10532.1	-4.1
31484_at	A	NC	-10408.1	-4.1
31635_g_at	A	NC	-10482	-4.1
35191_at	A	NC	-10728.1	-4.1
36670_at	A	NC	-10663.2	-4.1
39458_s_at	A	D	-10494	-4.1
40227_at	A	NC	-10391	-4.1
40484_g_at	A	NC	-10501.1	-4.1
40794_at	A	NC	-10465.7	-4.1
41520_at	A	NC	-10601.4	-4.1
41775_at	A	NC	-10580.9	-4.1
991_g_at	A	NC	-10526.6	-4.1
31742_at	A	NC	-10165.8	-4
33698_at	A	NC	-10099.9	-4
33964_at	A	D	-10267.6	-4
34008_at	A	NC	-10277.3	-4
34429_at	A	NC	-10116	-4
34743_at	A	MD	-10334.2	-4
35691_r_at	A	NC	-10073.3	-4
35866_at	A	D	-10275.8	-4
36469_at	A	D	-10370.9	-4
37252_at	A	NC	-10081.3	-4
37529_at	A	D	-10195	-4
38885_at	A	D	-10324.5	-4
41497_at	A	NC	-10050.6	-4
1367_f_at	P	I	100943.8	4
31687_f_at	P	I	115625.5	4
329_s_at	P	I	13807.2	4
32969_r_at	A	NC	10134.5	4
36368_at	A	NC	10356.1	4
37077_at	A	NC	10207.8	4

Table 2-4.

Probe Set Name	Abs Call	Diff Call	Avg Diff Change	Fold Change
38504_at	A	NC	10223.1	4
39110_at	P	I	11622	4
39933_at	A	NC	10250.3	4
41203_at	A	MI	10209.4	4
41206_r_at	P	I	21080.1	4
41382_at	A	NC	10288.7	4
31481_s_at	P	I	10528.4	4.1
31932_f_at	P	I	10447.9	4.1
33908_at	A	NC	10599.4	4.1
34032_at	A	MI	10484.6	4.1
34707_at	M	NC	10498.9	4.1
35016_at	P	I	33282	4.1
36473_at	P	NC	10433.7	4.1
36938_at	P	I	10454.4	4.1
38081_at	P	I	10495.9	4.1
38789_at	A	I	10439.4	4.1
39947_at	A	NC	10453.7	4.1
895_at	P	I	10668.8	4.1
1518_at	A	NC	10842.9	4.2
256_s_at	P	I	121364.9	4.2
31492_at	A	I	10923.7	4.2
31815_r_at	A	NC	17940.8	4.2
34160_at	P	I	44874.2	4.2
36587_at	P	I	51069.6	4.2
36652_at	A	NC	10807	4.2
38121_at	P	I	10997.5	4.2
38733_at	P	MI	11013.7	4.2
38997_at	P	I	10976.1	4.2
1184_at	P	I	11363.6	4.3
1817_at	P	I	11239	4.3
31330_at	P	I	161863.5	4.3
31419_r_at	A	NC	38467.6	4.3
31719_at	P	I	11228.9	4.3
31955_at	P	I	43342.3	4.3
33178_at	A	NC	11130.9	4.3
33676_at	P	I	110220.5	4.3
33967_at	P	NC	11119.1	4.3
34351_at	A	NC	11164.2	4.3
41138_at	P	MI	12991.2	4.3
41485_at	P	I	11239.4	4.3
925_at	P	I	60679	4.3
1083_s_at	A	NC	11584.3	4.4
35175_f_at	P	I	14130.9	4.4
38589_i_at	A	MI	11499.3	4.4
41221_at	P	NC	11660.6	4.4
146_at	A	NC	11842	4.5

Table 2-5.

Probe Set Name	Abs Call	Diff Call	Avg Diff Change	Fold Change
1825_at	P	I	12031.7	4.5
33117_r_at	P	I	52774.1	4.5
33390_at	A	NC	12000.9	4.5
33987_at	M	NC	11971	4.5
32878_f_at	A	NC	12100.8	4.6
33143_s_at	A	I	12232.6	4.6
37649_at	A	NC	12113.1	4.6
39054_at	A	NC	12235.5	4.6
100_g_at	A	MI	12464.4	4.7
2090_i_at	A	NC	12678.8	4.7
31951_s_at	P	I	68234.5	4.7
33727_r_at	A	NC	16195.6	4.7
36209_at	P	I	12682.4	4.7
37307_at	P	NC	23687.7	4.7
38404_at	P	I	12541.5	4.7
41189_at	P	I	12444.1	4.7
41400_at	A	NC	12515.1	4.7
151_s_at	A	I	12856	4.8
32070_at	A	NC	12837.1	4.8
38088_r_at	A	NC	13037.3	4.8
505_at	A	NC	12830.5	4.8
868_at	A	I	13048.7	4.8
1404_r_at	P	I	13120.4	4.9
33425_at	A	NC	13218.4	4.9
1984_s_at	P	I	13552.9	5
32900_at	A	NC	13629.4	5
35083_at	P	I	250325.6	5
35823_at	A	I	13736	5
36330_at	A	NC	13649.5	5
36493_at	P	I	15585.2	5
37603_at	P	I	23925	5
38970_s_at	A	I	13694.6	5
40159_r_at	A	NC	13575.1	5
917_g_at	A	NC	13489.7	5
32316_s_at	P	I	14003.7	5.1
347_s_at	P	I	132536.4	5.1
34841_at	A	MI	14109.4	5.1
40480_s_at	P	I	13844.9	5.1
956_at	A	NC	13896	5.1
35117_at	A	MI	14322.2	5.2
36675_r_at	P	I	43562.8	5.2
37940_f_at	A	NC	14220.9	5.2
33485_at	P	I	46973.8	5.3
33500_i_at	P	I	26486.6	5.3
34826_at	P	I	14592.2	5.3
35298_at	P	I	14749.7	5.3

Table 2-6

Probe Set Name	Abs Call	Diff Call	Avg Diff Change	Fold Change
38061_at	P	I	71146.5	5.3
41164_at	P	I	24897.2	5.3
1346_at	A	MI	14872.1	5.4
31525_s_at	P	I	218352.5	5.4
32334_f_at	P	I	108562.5	5.4
34345_at	A	NC	15082.3	5.4
1161_at	P	MI	15463.8	5.5
34608_at	P	I	91001.3	5.5
33201_at	A	NC	-18970.1	5.6
39237_at	A	NC	-15712.5	5.6
32437_at	P	I	30183.7	5.7
40520_g_at	P	I	16008.9	5.7
1662_r_at	A	NC	16446	5.8
38661_at	A	MI	16385.6	5.8
39856_at	P	I	16202	5.8
38759_at	P	I	16863.2	5.9
39728_at	P	I	81729.1	5.9
39704_s_at	A	MI	16867.2	6
40155_at	A	I	17021.7	6
181_g_at	A	NC	17238.6	6.1
34645_at	P	I	81954.8	6.1
37145_at	P	I	17871.5	6.2
38745_at	P	I	17722.7	6.2
31546_at	P	I	41000	6.3
33984_at	A	I	18005.3	6.3
40253_at	A	NC	17978	6.3
935_at	P	I	18614.9	6.5
691_g_at	M	NC	18934.2	6.6
810_at	P	I	19007.5	6.6
32543_at	A	I	19286.9	6.7
38138_at	P	MI	19389.1	6.7
40874_at	P	I	19306.1	6.7
1506_at	P	I	19655.3	6.8
32166_at	P	NC	28628.1	6.8
32378_at	A	NC	19862.2	6.8
34495_r_at	A	NC	20349.7	7
40164_at	A	I	20394.1	7
31957_r_at	P	I	322667.3	7.1
33659_at	A	I	80715.1	7.1
34210_at	P	I	20732	7.1
37383_f_at	P	I	428958.5	7.1
34041_at	P	I	21554.3	7.3
33674_at	P	I	75933.7	7.4
1707_g_at	P	NC	22957.8	7.7
37121_at	P	I	23105.9	7.8
31907_at	P	I	46016.3	8.1

Table 2-7.

Probe Set Name	Abs Call	Diff Call	Avg Diff Change	Fold Change
34085_at	P	I	130166.3	8.2
33943_at	P	I	38389	8.3
40435_at	P	I	40538	8.4
37449_i_at	P	I	40364.3	8.5
38976_at	A	I	25887.5	8.6
37450_r_at	P	I	51843.4	8.7
33273_f_at	P	I	27244.1	9
37448_s_at	P	I	27436.6	9.1
40835_at	A	NC	27728	9.1
41165_g_at	P	I	42736.5	9.3
40370_f_at	P	I	29897.7	9.8
39119_s_at	P	I	30244.9	9.9
39166_s_at	A	NC	30694.4	10
40042_r_at	A	NC	31929.9	10.4
36786_at	P	I	33850.9	10.9
1288_s_at	P	I	345959.1	11.1
40887_g_at	P	I	255876	11.5
428_s_at	P	I	180066.3	12.2
32749_s_at	P	I	45487.7	12.3
201_s_at	P	I	191230.6	12.6
37759_at	P	I	56219.3	13.8
32321_at	P	I	45287	14.3
31870_at	P	I	46231.5	14.6
32318_s_at	P	I	153259.3	16.2
32487_s_at	P	I	168200.4	16.2
2016_s_at	P	I	160508.8	18.9
37421_f_at	P	I	62387.4	19.3
32794_g_at	P	I	75616.4	23.2
31385_at	P	I	185236.2	23.9
38524_at	A	NC	211658.7	63.1
1894_f_at	P	NC	240916.2	71.7
41624_r_at	A	NC	279838.7	83.1
39473_r_at	A	NC	304816.3	90.5
36546_r_at	A	NC	305109.5	90.6
32466_at	P	NC	533600.1	157.6
1008_f_at	P	I	934734	275.4

Table 3-1.

Probe Set Name	Diff Call	Fold Change	GenBank No.	Accession	Description
39560_at	MD	-5.3	H10776		unknown
38311_at	MD	-4.8	AF055012		unknown
34743_at	MD	-4	D63481		KIAA0147
41206_r_at	I	4	U05255		glycophorin HeP2
39110_at	I	4	AF070626		unknown
31687_f_at	I	4	M25079		sickle cell beta-globin
329_s_at	I	4			Nuclear Mitotic Apparatus Protein 1, Alt. Splice Form 2
1367_f_at	I	4	M26880		ubiquitin
31932_f_at	I	4.1	M90357		basic transcription factor 3a (BTF3a)
36938_at	I	4.1	U70063		acid ceramidase
38081_at	I	4.1	J03459		leukotriene A-4 hydrolase
31481_s_at	I	4.1	M92383		thymosin beta-10
895_at	I	4.1	L19686		macrophage migration inhibitory factor (MIF)
35016_at	I	4.1	M13560		Ia-associated invariant gamma-chain
256_s_at	I	4.2	M14199		laminin receptor (2H5 epitope)
36587_at	I	4.2	AI741756		cDNA clone similar to vascular proton pump subunit SFD beta isoform
38997_at	I	4.2	X96924		mitochondrial citrate transport protein
38121_at	I	4.2	X59892		IFN-inducible gamma2
34160_at	I	4.2	AB019529		OGG1 protein type 2c, partial cds
38733_at	MI	4.2	M30938		Ku (p70/p80) subunit
925_at	I	4.3	J03909		gamma-interferon-inducible protein (IP-30)
31330_at	I	4.3	M81757		S19 ribosomal protein
33676_at	I	4.3	X15940		ribosomal protein L31
31719_at	I	4.3	X02761		fibronectin (FN precursor)
1817_at	I	4.3	D89667		c-myc binding protein
41485_at	I	4.3	X02152		lactate dehydrogenase-A (LDH-A, EC 1.1.1.27)
31955_at	I	4.3	X65923		fau
41138_at	MI	4.3	M16279		MIC2
1184_at	I	4.3	D45248		proteasome activator hPA28 subunit beta

Table 3-2.

Probe Set Name	Diff Call	Fold Change	GenBank No.	Accession	Description
35175_f_at	I	4.4	X70940		elongation factor 1 alpha-2
1825_at	I	4.5	L33075		ras GTPase-activating-like protein (IQGAP1)
33117_r_at	I	4.5	AA977163		unknown
41189_at	I	4.7	Y09392		WSL-1R, WSL-S1 and WSL-S2 proteins
38404_at	I	4.7	M55153		transglutaminase (TGase)
31951_s_at	I	4.7	Z48501		polyadenylate binding protein II
36209_at	I	4.7	S78771		NAT=CpG island-associated gene
1404_r_at	I	4.9	M21121		T cell-specific protein (RANTES)
35083_at	I	5	AL031670		ferritin, light polypeptide-like 1
37603_at	I	5	X52015		interleukin-1 receptor antagonist
1984_s_at	I	5	X69549		rho GDP-dissociation Inhibitor 2
347_s_at	I	5.1	D14530		homolog of yeast ribosomal protein S28
40480_s_at	I	5.1	M14333		c-syn protooncogene
32316_s_at	I	5.1	X15183		90-kDa heat-shock protein
36675_r_at	I	5.2	J03191		profilin
38061_at	I	5.3	AI541256		unknown
33485_at	I	5.3	D23660		ribosomal protein
34826_at	I	5.3	L21936		succinate dehydrogenase flavoprotein subunit (SDH)
41164_at	I	5.3	X67301		IgM heavy chain constant region (Ab63)
35298_at	I	5.3	U54558		translation initiation factor eIF3 p66 subunit
33500_i_at	I	5.3	S71043		Ig alpha 2=immunoglobulin A heavy chain allotype 2
31525_s_at	I	5.4	J00153		alpha globin gene cluster on chromosome 16- zeta gene
32334_f_at	I	5.4	AB009010		polyubiquitin UbC
34608_at	I	5.5	X86174		SSX1
1161_at	MI	5.5	J04988		90 kD heat shock protein
36493_at	I	5.6	M33552		lymphocyte-specific protein 1 (LSP1)
40520_g_at	I	5.7	Y00638		leukocyte common antigen (T200)
32437_at	I	5.7	U14970		ribosomal protein S5
39856_at	I	5.8	AI708983		cDNA clone Ig kappa chain precursor V-III region

Table 3-3.

Probe Set Name	Diff Call	Fold Change	GenBank No.	Accession	Description
39728_at	I	5.9	J03909		gamma-interferon-inducible protein (IP-30)
38759_at	I	5.9	U97502		butyrophilin (BT3.3)
34645_at	I	6.1	X55715		40S ribosomal protein s3
38745_at	I	6.2	X76488		lysosomal acid lipase
37145_at	I	6.2	M85276		NKG5
31546_at	I	6.3	L11566		ribosomal protein L18 (RPL18)
935_at	I	6.5	L12168		adenylyl cyclase-associated protein (CAP)
810_at	I	6.6	U64105		guanine nucleotide exchange factor p115-RhoGEF
40874_at	I	6.7	AJ005259		EDF-1
38138_at	MI	6.7	D38583		calgizzarin
1506_at	I	6.8	D11086		interleukin 2 receptor gamma chain
34210_at	I	7.1	N90866		cDNA clone similar to CAMPATH-lymphocyte surface antigen
31957_r_at	I	7.1	M17886		acidic ribosomal phosphoprotein P1
37383_f_at	I	7.1	X58536		HLA class I locus C heavy chain
34041_at	I	7.3	U83171		macrophage-derived chemokine precursor (MDC)
33674_at	I	7.4	Z49148		ribosomal protein L29
37121_at	I	7.8	S69115		granulocyte colony-stimulating factor induced gene
31907_at	I	8.1	D87735		ribosomal protein L14
34085_at	I	8.2	Z26876		ribosomal protein L38
33943_at	I	8.3	L20941		ferritin heavy chain
40435_at	I	8.4	J03592		ADP/ATP translocase
37449_i_at	I	8.5	X04409		coupling protein G(s) alpha-subunit (alpha-S1)
37450_r_at	I	8.7	X04409		coupling protein G(s) alpha-subunit (alpha-S1)
33273_f_at	I	9	X57809		rearranged immunoglobulin lambda light chain
37448_s_at	I	9.1	X56009		GSA mRNA for alpha subunit of GsGTP binding protein
41165_g_at	I	9.3	X67301		IgM heavy chain constant region (Ab63)
40370_f_at	I	9.8	M90683		lymphocyte antigen (HLA-G1)
39119_s_at	I	9.9	AA631972		cDNA clone similar to NK4
36786_at	I	10.9	AL022721		60S Ribosomal Protein RPL10A

Table 3-4.

Probe Set Name	Diff Call	Fold Change	GenBank No.	Accession	Description
1288_s_at	I	11.1	J04617		elongation factor EF-1-alpha
40887_g_at	I	11.5	L41498		elongation factor 1-alpha 1 (PTI-1)
428_s_at	I	12.2	V00567		beta-2 microglobulin
32749_s_at	I	12.3	AL050396		unknown
201_s_at	I	12.6	S82297		beta 2-microglobulin {11bp deleted}
37759_at	I	13.8	U51240		lysosomal-associated multitransmembrane protein (LAPtm5)
32321_at	I	14.3	X56841		HLA-E
31870_at	I	14.6	X14046		leukocyte antigen CD37
32487_s_at	I	16.2	AI093511		cDNA clone similar to tau protein kinase CDK5/P20
32318_s_at	I	16.2	X63432		ACTB mRNA for mutant beta-actin (beta-actin)
2016_s_at	I	18.9	M64241		Wilms tumor-related protein (QM)
37421_f_at	I	19.3	AL022723		major histocompatibility complex, class I, F (CDA12)
32794_g_at	I	23.2	X00437		T-cell specific protein
31385_at	I	23.9	U14969		ribosomal protein L28
1008_f_at	I	274.5	U50648		interferon-inducible RNA-dependent protein kinase (Pkr)

Table 4.

Probe Set Name	Abs Call	Diff Call	Fold Change
124_i_at	A (コントロールでも A)	MD	-13.4
38203_at	A (コントロールでも A)	MD	-5.9
38356_at	A (コントロールでも A)	D	-5.8
373_at	A (コントロールでも A)	D	-5.4
35886_at	A (コントロールでも A)	MD	-5.2
34094_i_at	A (コントロールでも A)	MD	-5.1
33552_at	A (コントロールでも A)	D	-4.9
32990_at	A (コントロールでも A)	D	-4.9
36431_at	A (コントロールでも A)	D	-4.9
35318_at	A (コントロールでも A)	D	-4.8
32405_at	A	D	-4.7
32980_f_at	A (コントロールでも A)	D	-4.6
34716_at	A	D	-4.5
34209_at	A (コントロールでも A)	MD	-4.4
294_s_at	A (コントロールでも A)	D	-4.4
38236_at	A	D	-4.3
39615_at	A	D	-4.3
41282_s_at	A (コントロールでも A)	D	-4.3
38057_at	A (コントロールでも A)	MD	-4.2
33172_at	A (コントロールでも A)	D	-4.2
36035_at	A	D	-4.2
37844_at	A	D	-4.2
40209_at	A (コントロールでも A)	MD	-4.1
1207_at	A	D	-4.1
36535_at	A (コントロールでも A)	D	-4.1
33151_s_at	A	D	-4
37370_i_at	A (コントロールでも A)	MD	-4
31736_at	A (コントロールでも A)	D	-4
34630_s_at	A	D	-4
34412_s_at	A (コントロールでも A)	D	-4
1117_at	A (コントロールでも A)	D	-4
35139_at	A (コントロールでも A)	D	-4
38390_at	A (コントロールでも A)	D	-4
1161_at	P	I	4
40408_at	P	I	4.2
41189_at	P	I	4.3
36748_at	P	I	4.3
37344_at	P	I	4.8
34892_at	P	I	5
1008_f_at	P	MI	315.7