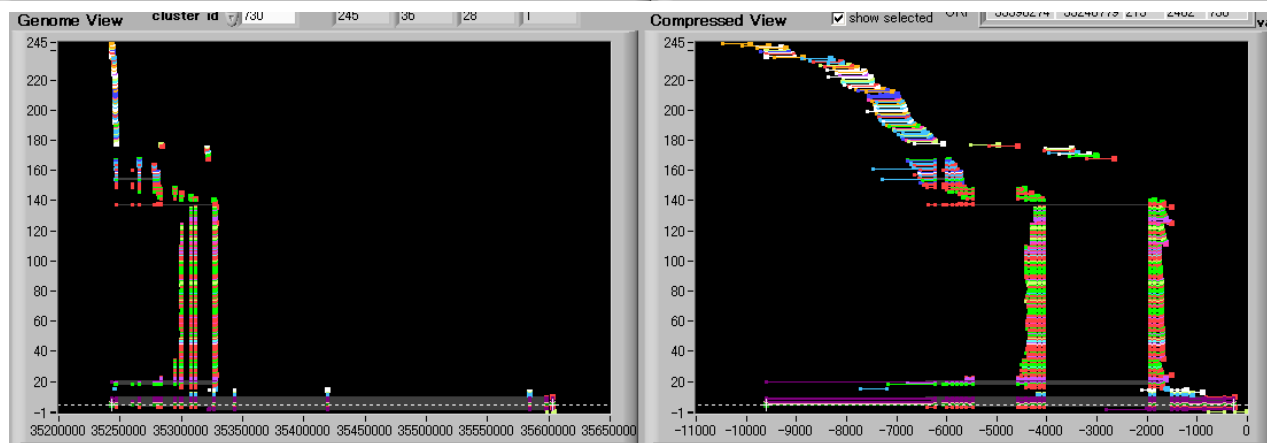
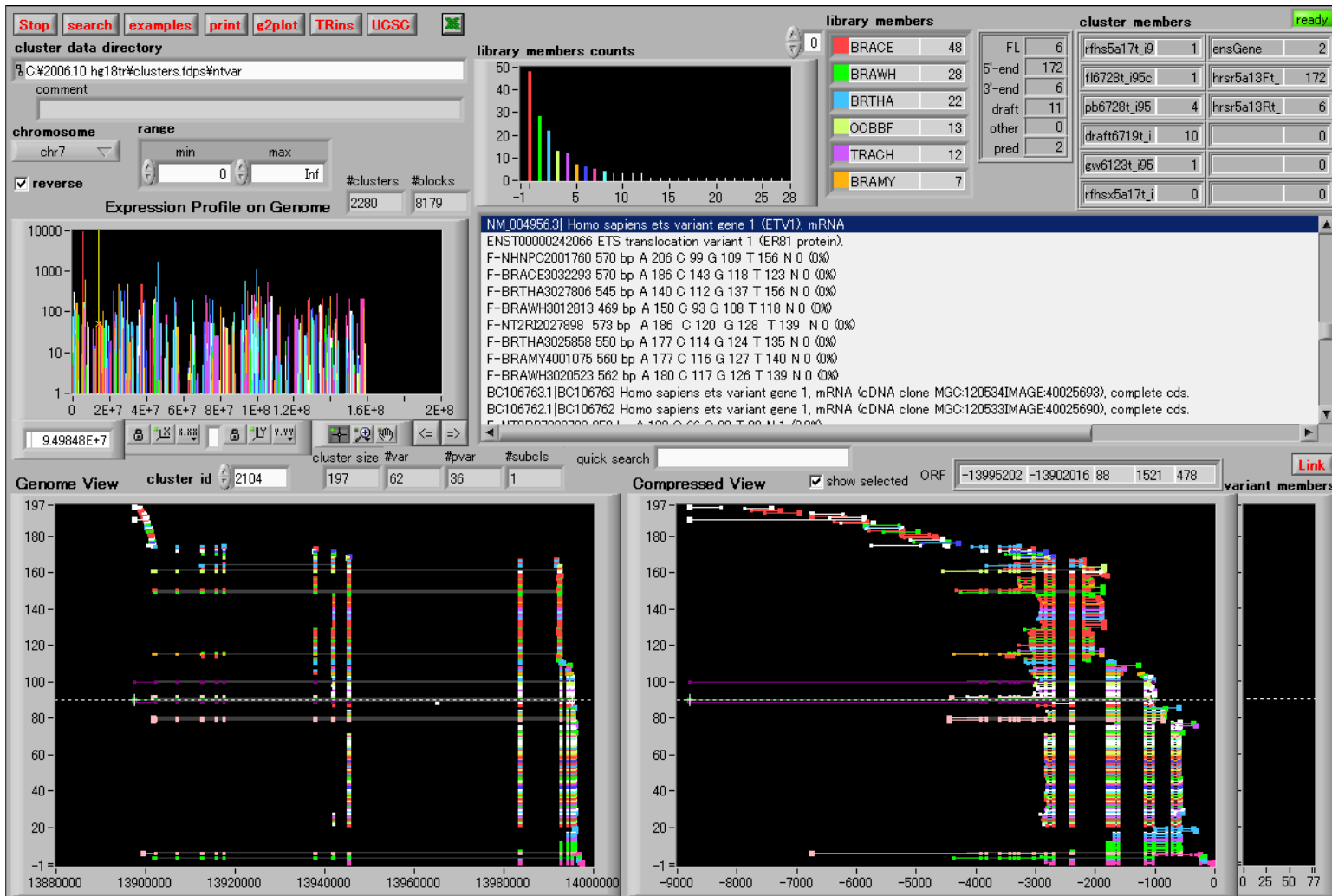


FLJ50539

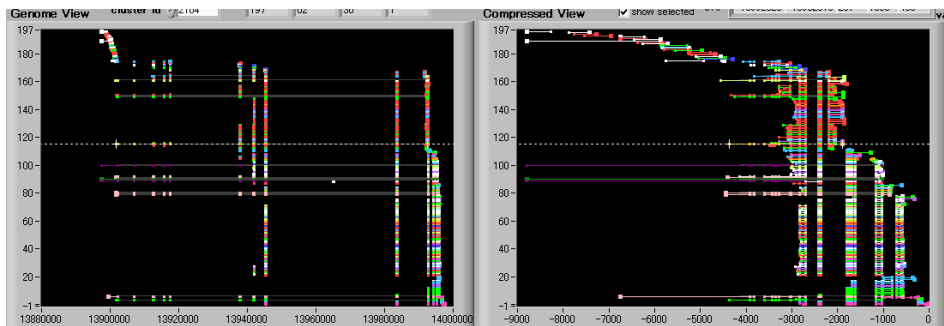


NM_004734.2

DCLK1_FLJ50539,
NM_004734.2

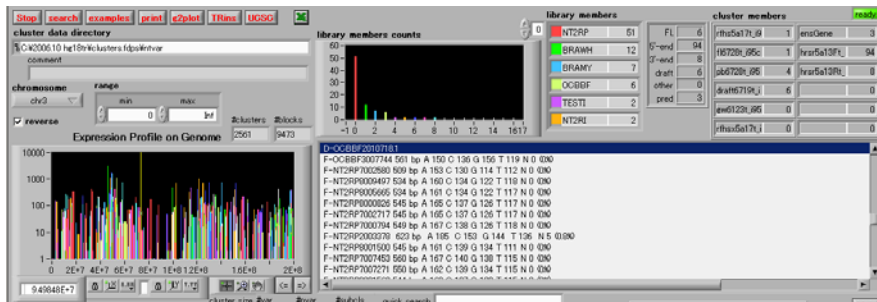


NM_004956.3



FLJ50494

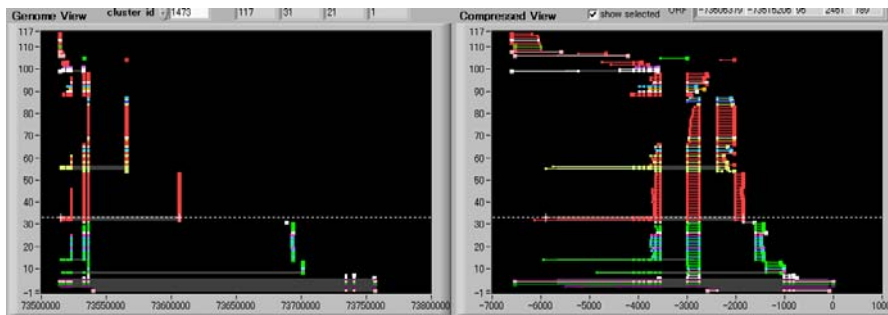
ETV1_NM_004956.3,
 FLJ50494



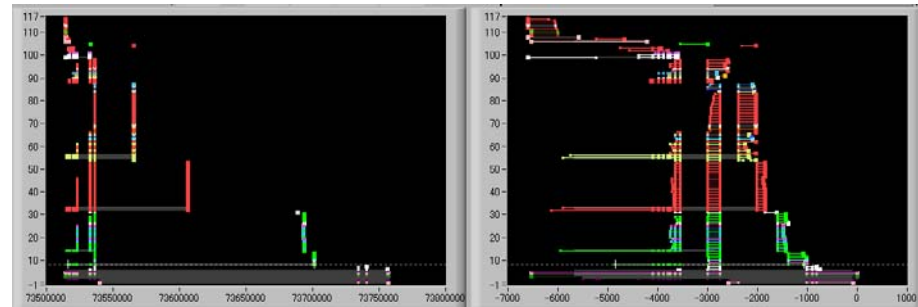
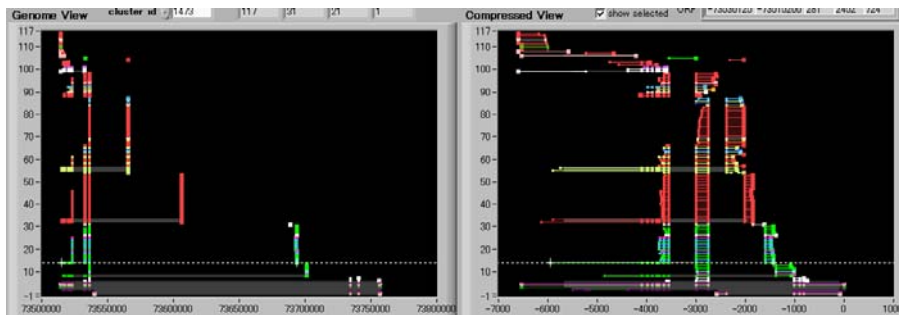
FLJ55043

FLJ97766

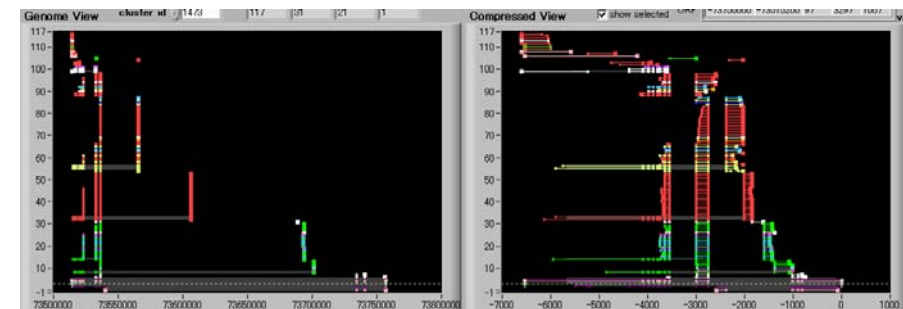
FLJ54746



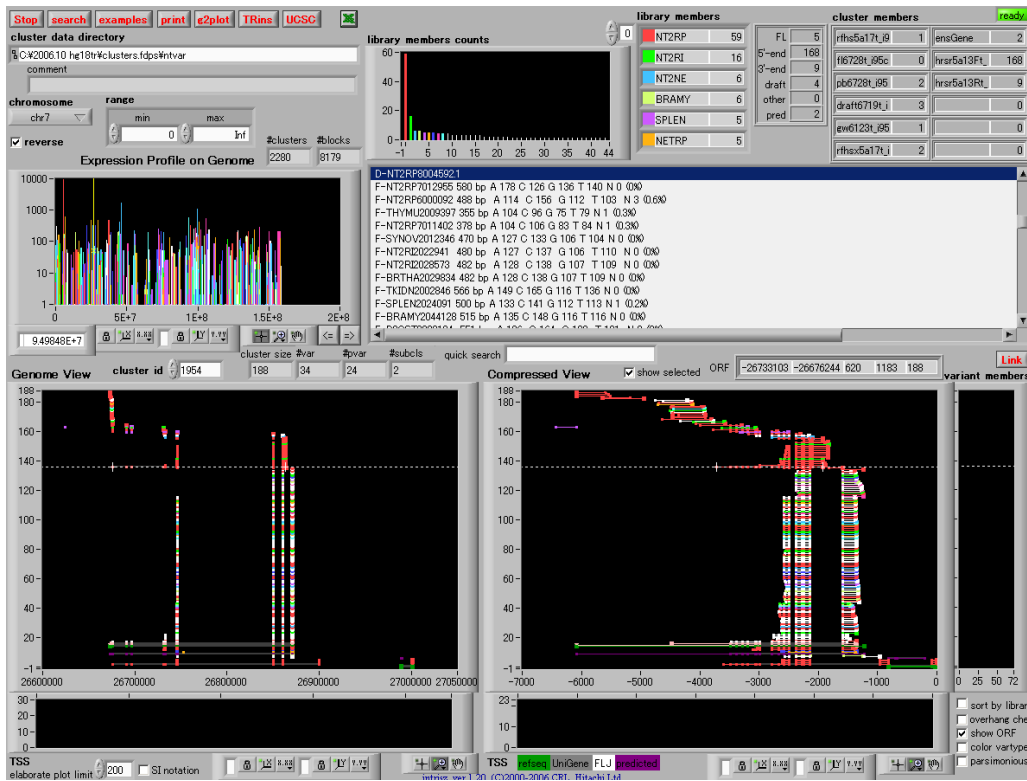
FLJ54738



NM_015009.1



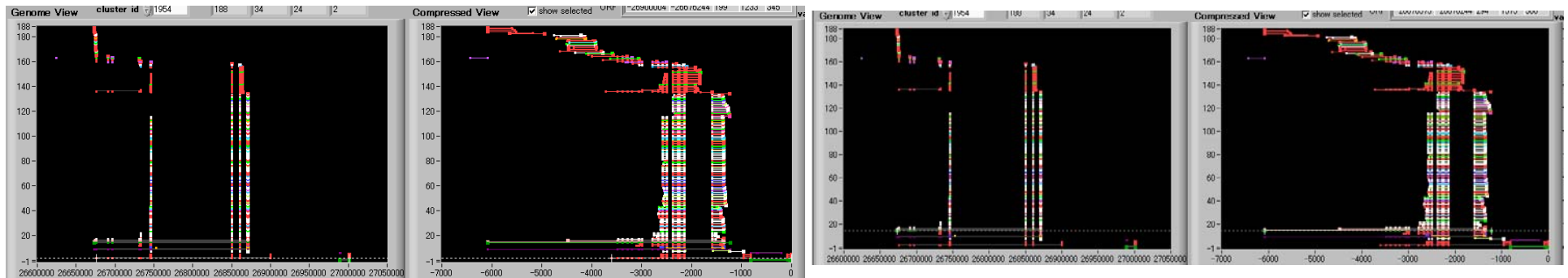
PDZRN3_FLJ55043,
FLJ54746, FLJ54738,
FLJ97766,
NM_015009.1



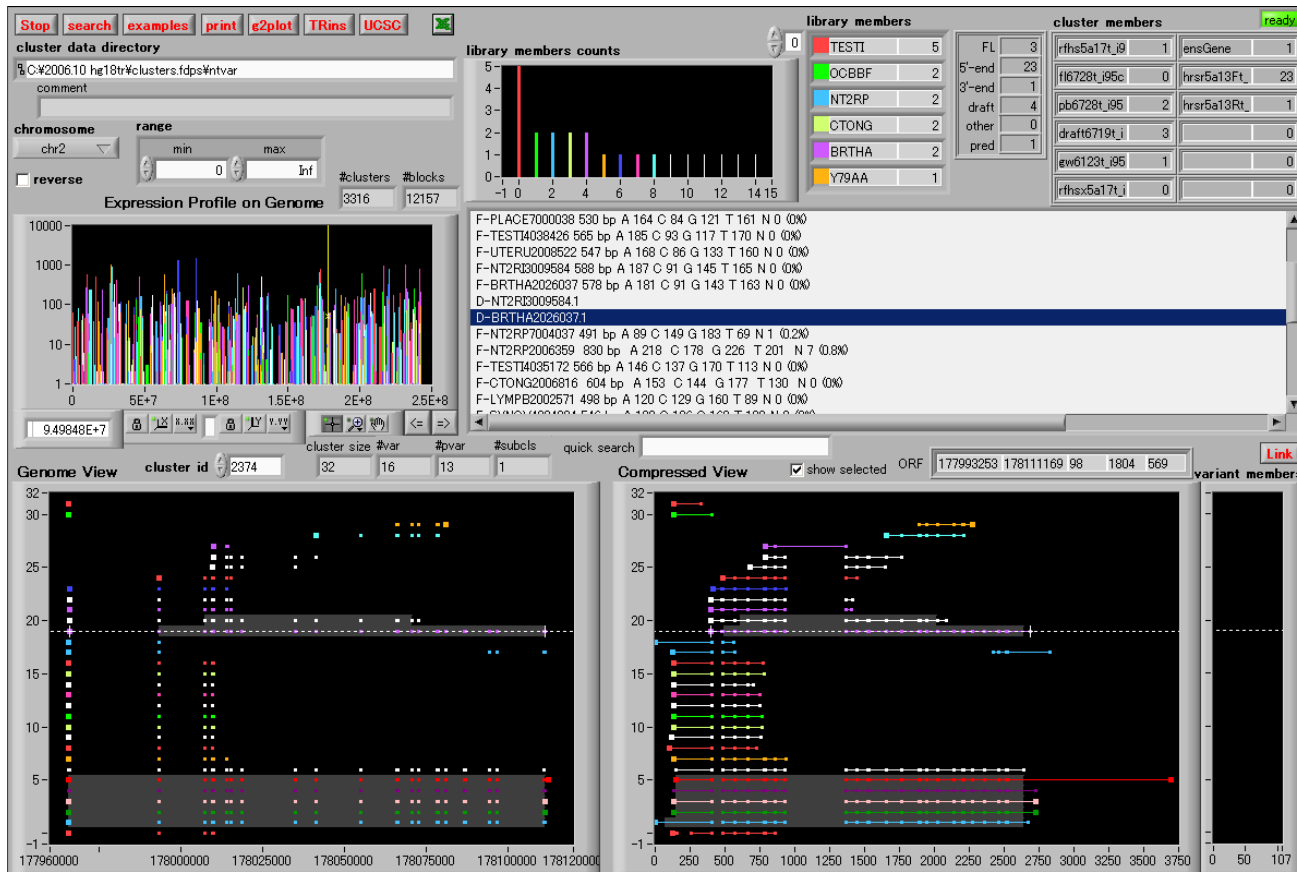
FLJ50630

FLJ51950

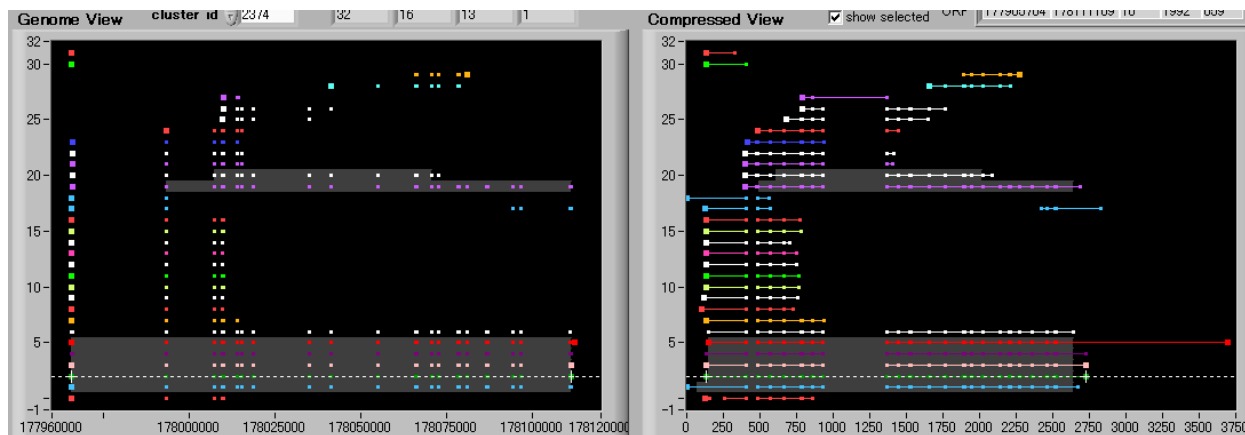
NM_003930.3



SKAP2_FLJ50630,
FLJ51950,
NM_003930.3

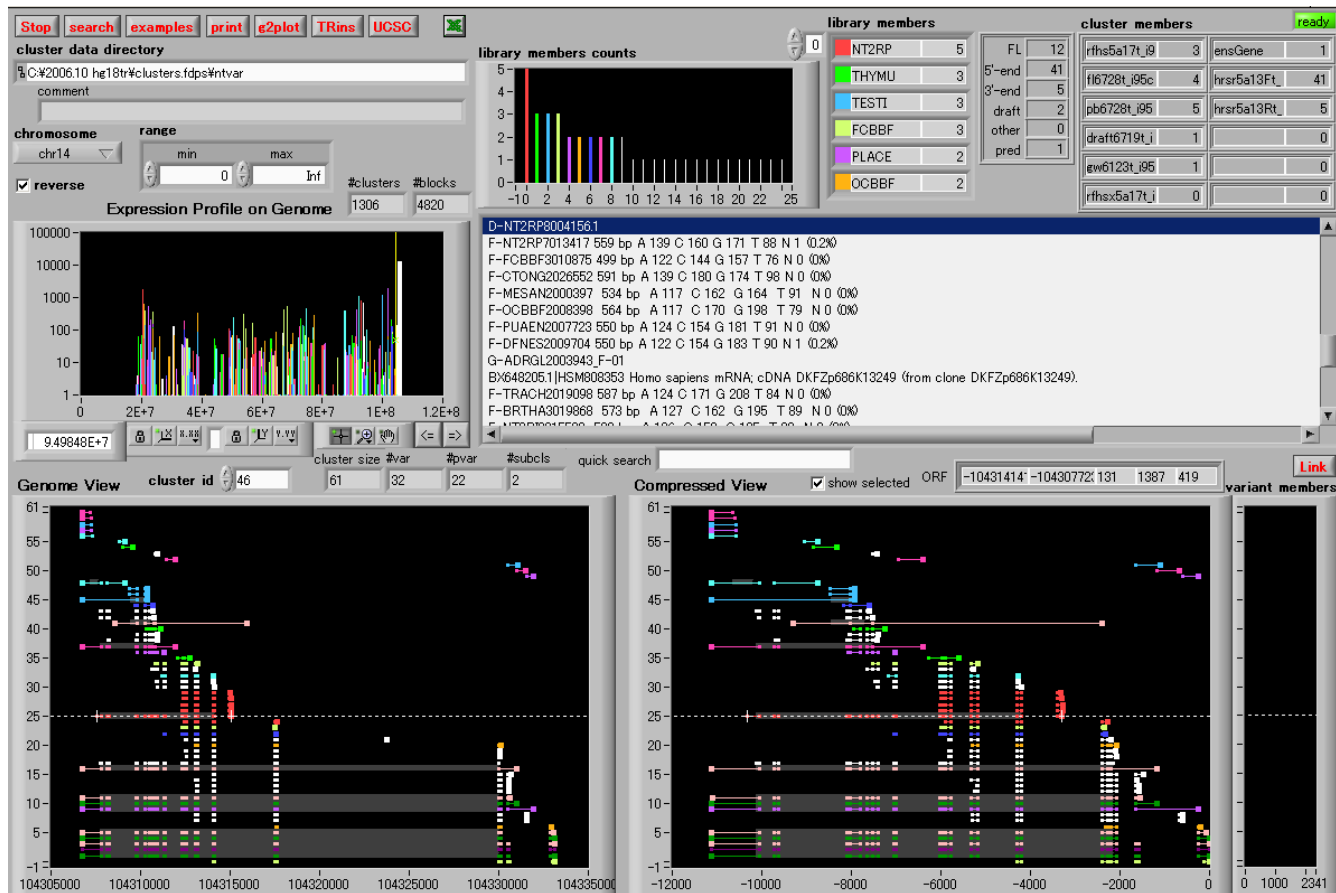


FLJ51873

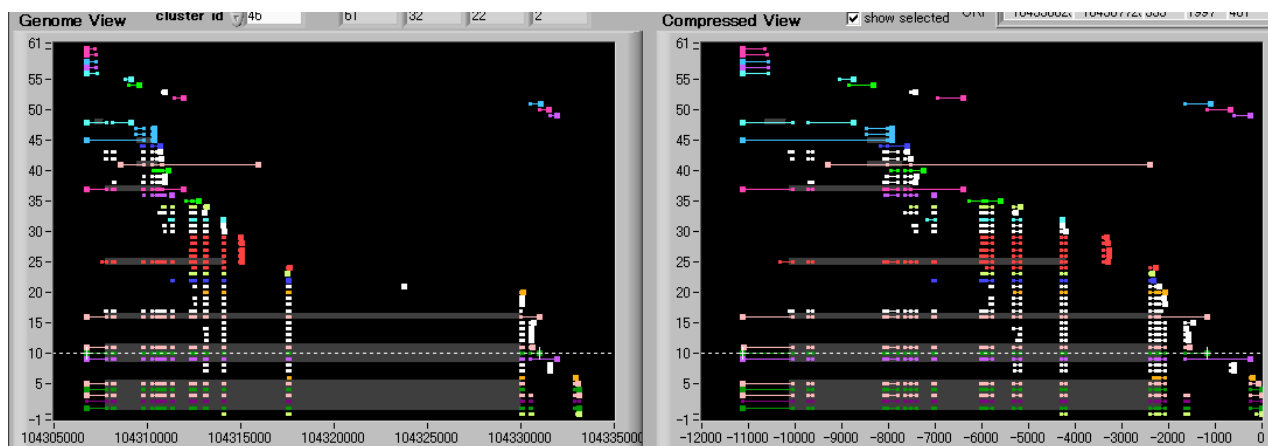


NM_003659.1

AGPS_FLJ51873,
NM_003659.1

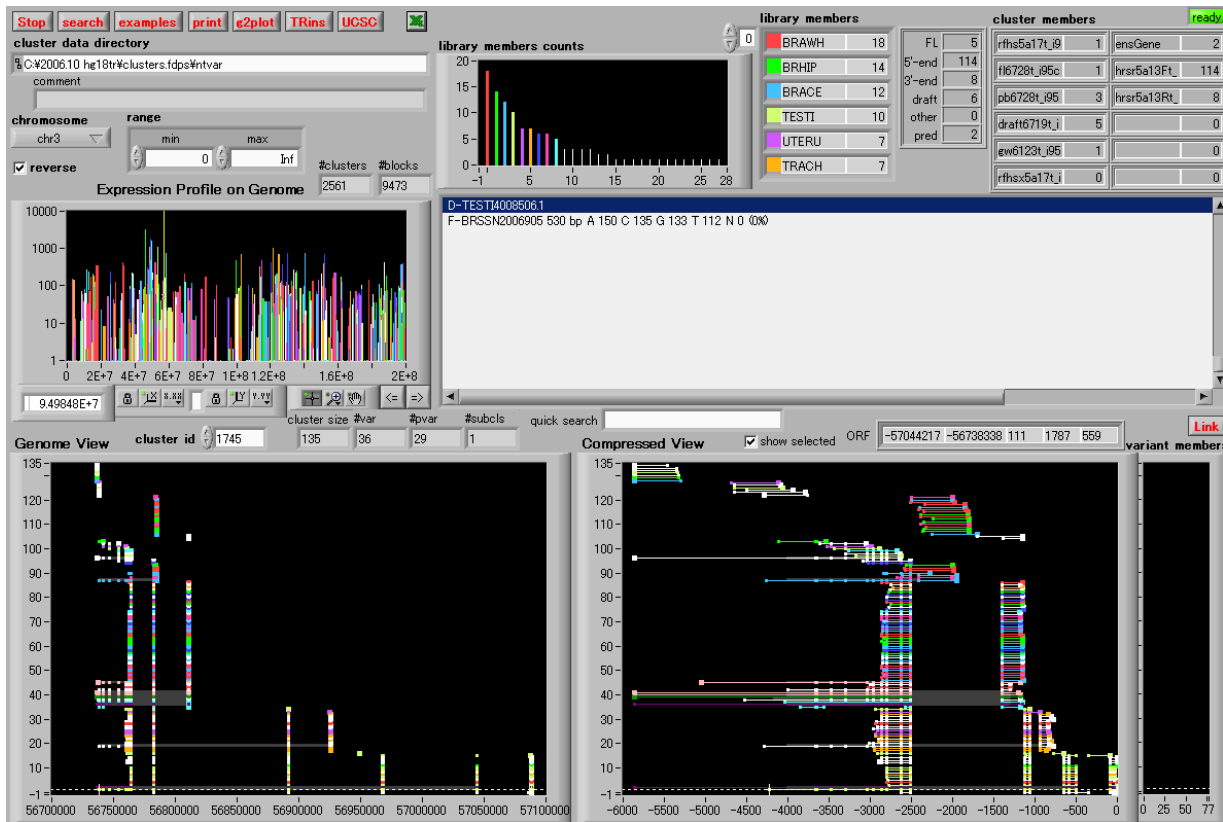


FLJ53606



NM_005163.2

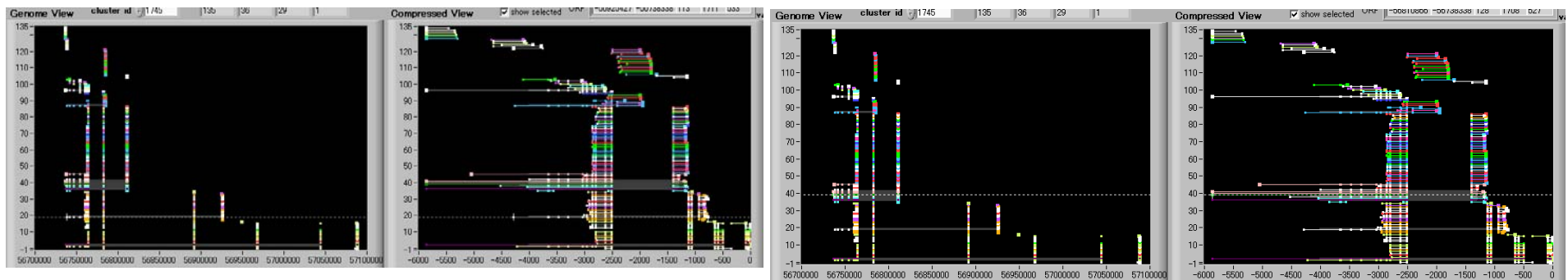
AKT1_FLJ53606,
 NM_005163.2



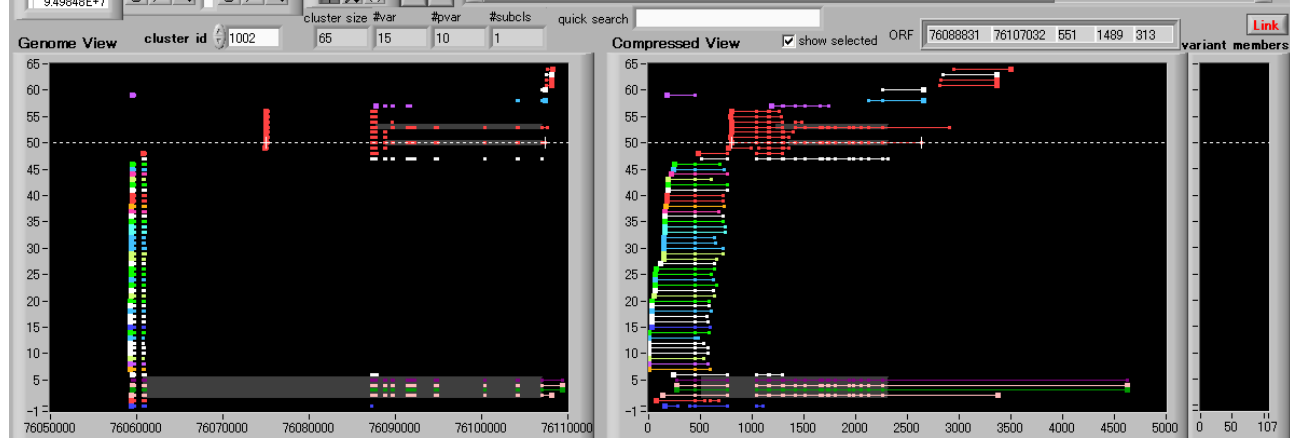
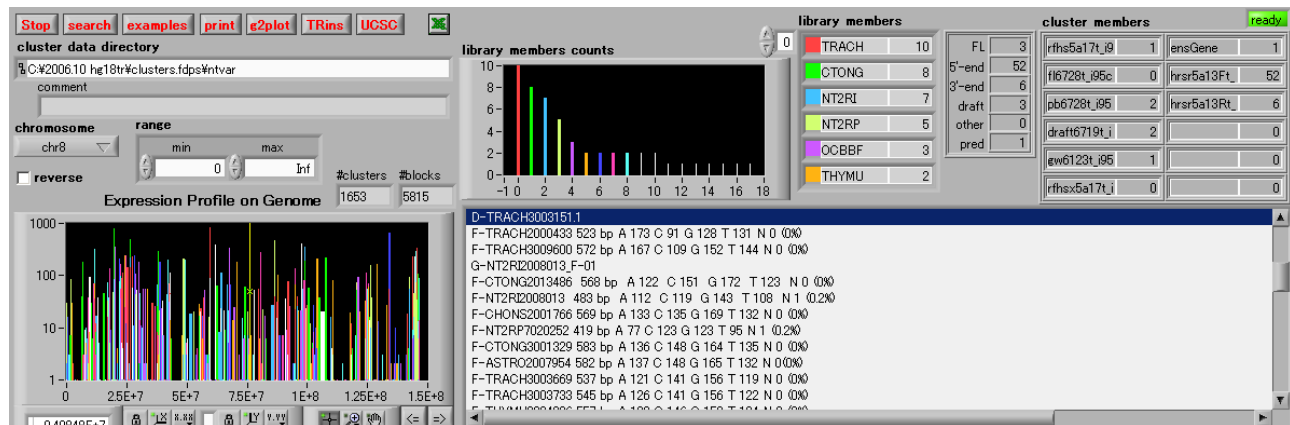
FLJ55856

NM_019555.1

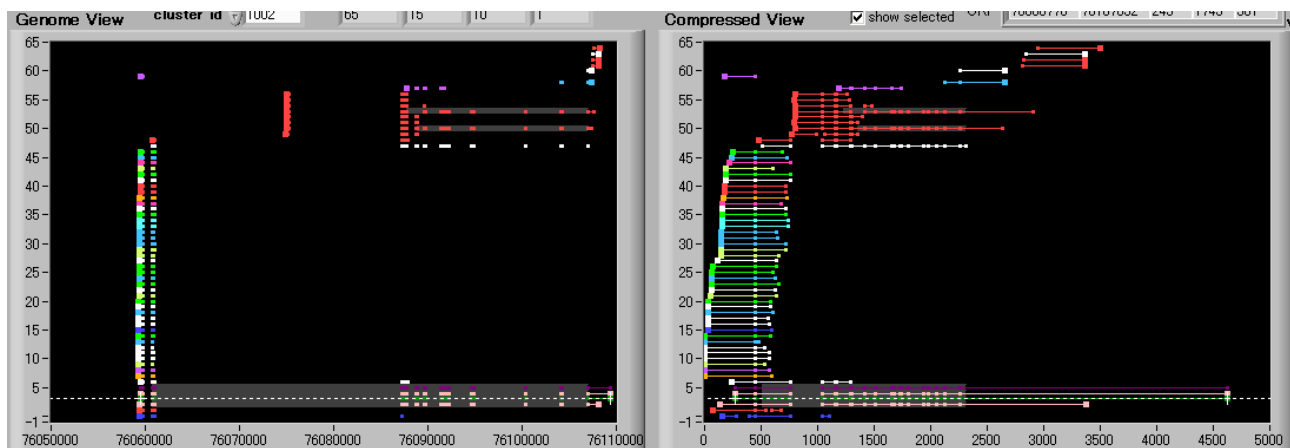
FLJ55591



ARHGEF3_FLJ55856,
FLJ55591,
NM_019555.1



FLJ57290



NM_031461.3

CRISPLD1_FLJ57290,
NM_031461.3

Stop search examples print e2plot TRins UCSC

cluster data directory
%C:\2006.10 hg18tr\clusters.fdp5#ntvar

chromosome chr2 range min max
reverse #clusters #blocks 3316 12157

Expression Profile on Genome

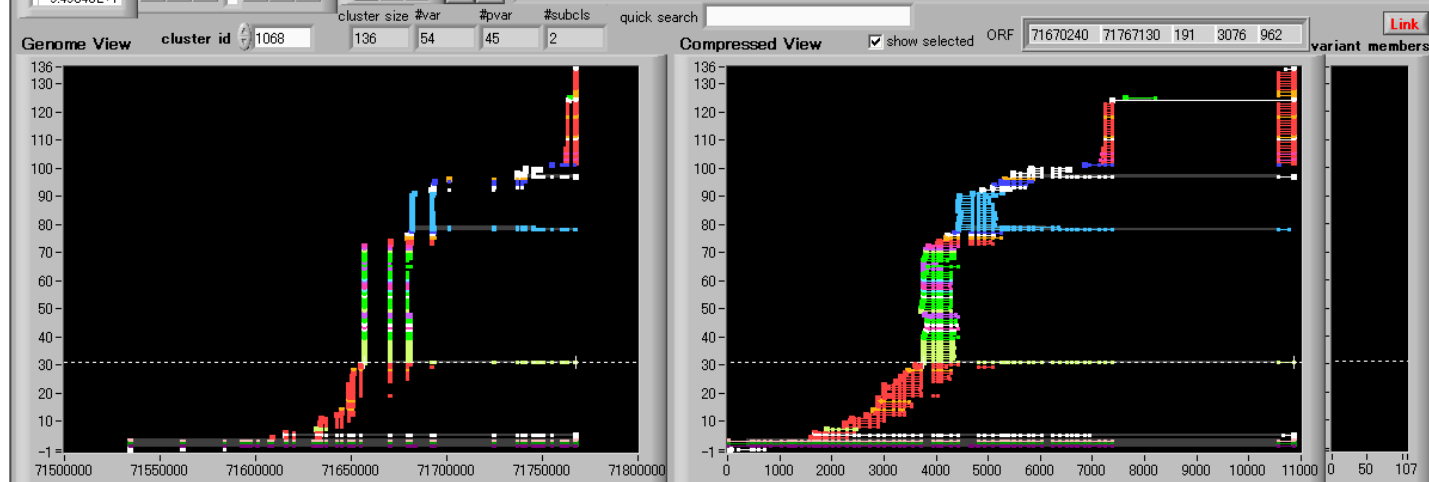
library members counts

library members	count
VESEN	24
BRTHA	18
THYMU	12
BRAWH	10
BRAMY	6
PLACE	5

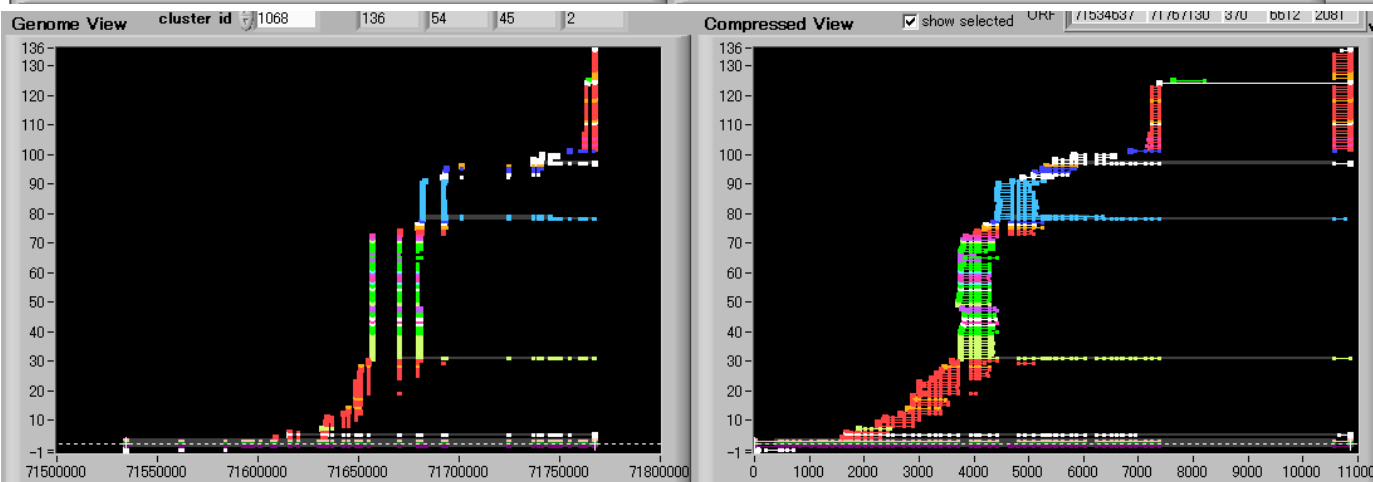
cluster members	count
rths5a17t_i9	1
fl6728t_i95c	2
pb6728t_i95	2
draft6719t_i	3
gw6123t_i95	0
rthsx5a17t_i	0

D-BRAWH3031931.1

Gene	Length (bp)	A	C	G	T	N	%GC
F-VESEN1000582	725	149	210	209	154	3	0.4%
F-VESEN1000394	837	180	254	227	169	7	0.8%
F-PLACE2000227	833	179	232	246	172	4	0.4%
F-VESEN1000058	789	166	226	232	161	4	0.5%
F-VESEN1000475	783	174	216	238	152	3	0.3%
F-VESEN2000459	778	171	220	236	148	3	0.3%
F-VESEN2000521	813	183	227	249	149	5	0.6%
F-VESEN2000285	791	176	210	259	143	3	0.3%
F-VESEN1000283	710	155	187	233	131	4	0.5%
F-VESEN2000493	760	170	203	249	135	3	0.3%
F-VESEN1000214	789	174	211	250	150	4	0.5%

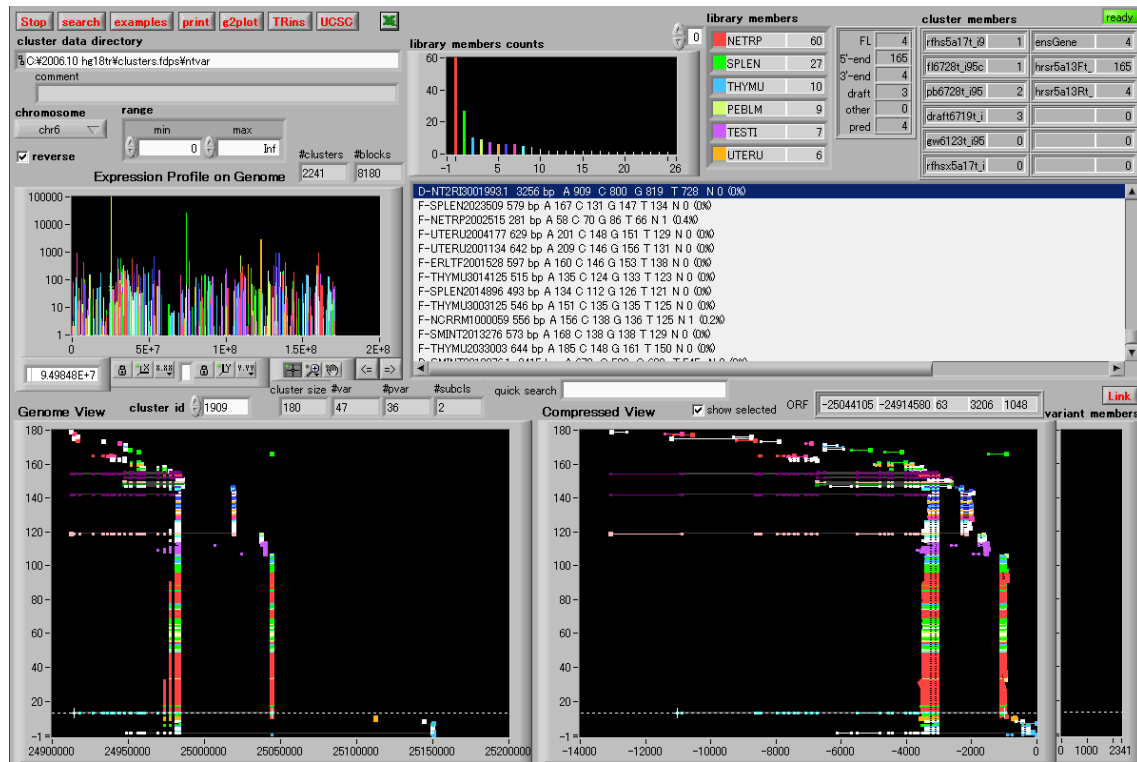


FLJ55344



NM_003494.2

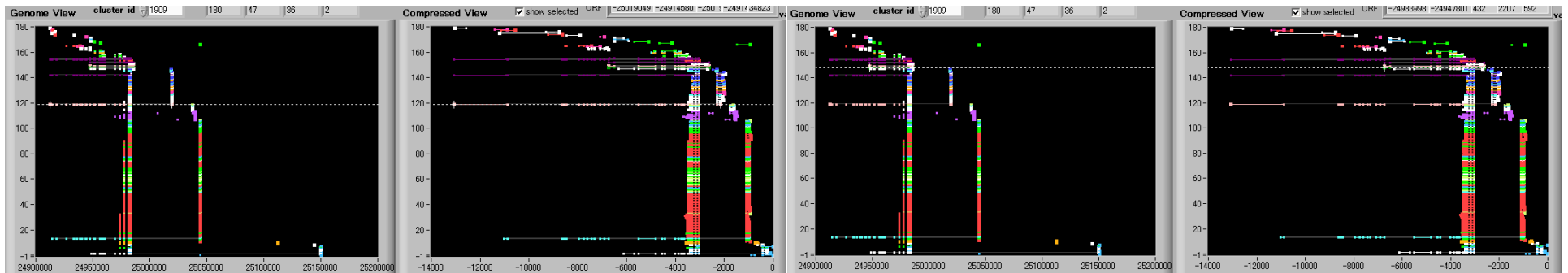
DYSF_FLJ55344,
NM_003494.2



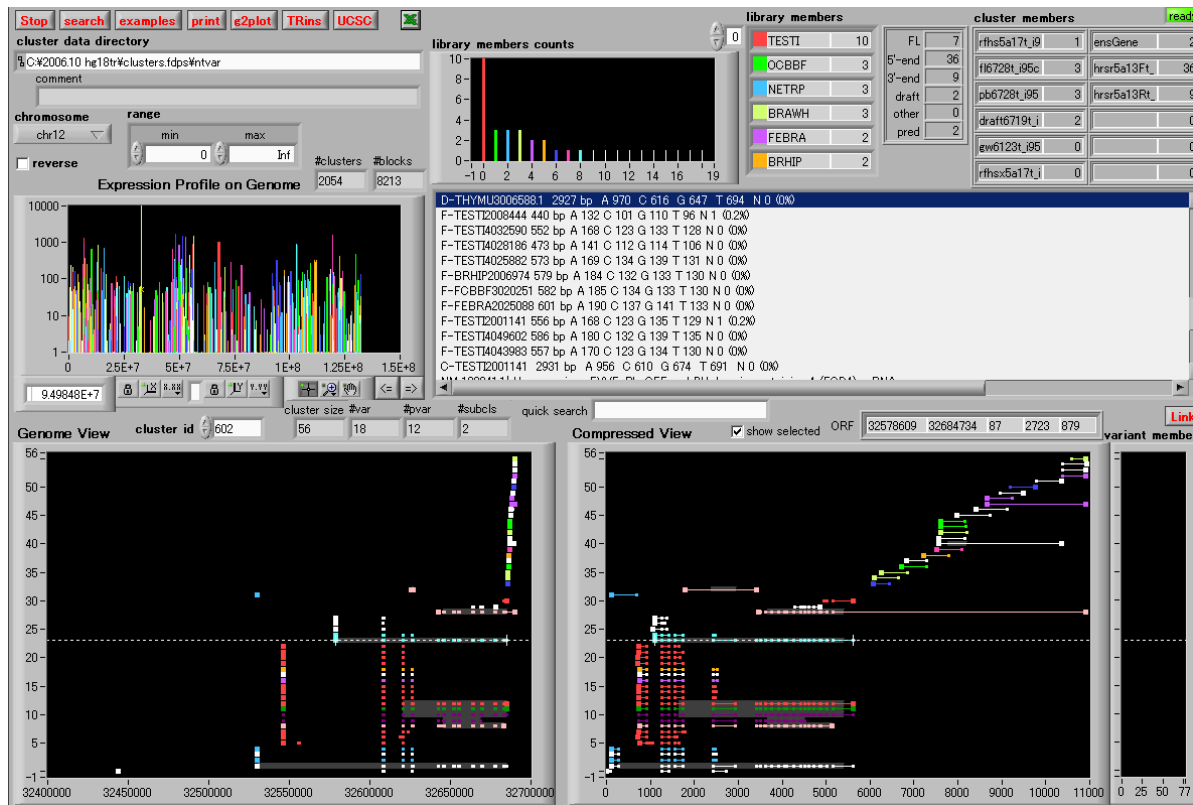
FLJ56137

AB002384.1

NM_015864.2



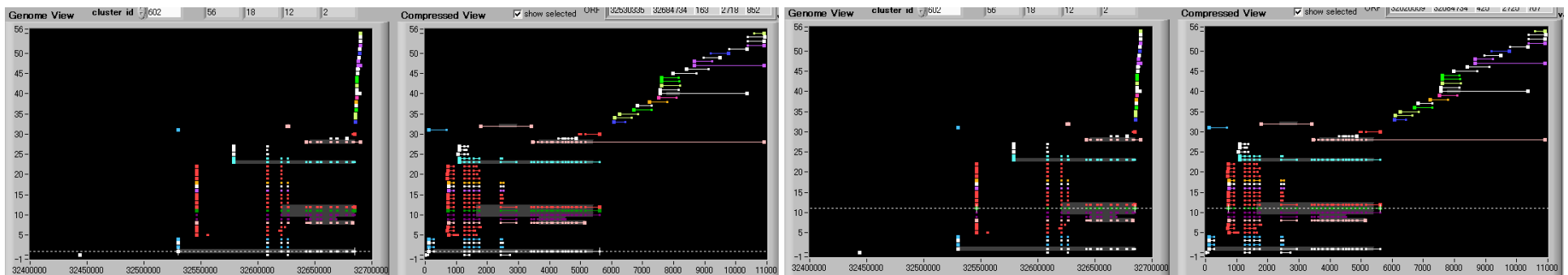
FAM65B_FLJ56137,
AB002384.1,
NM_015864.2



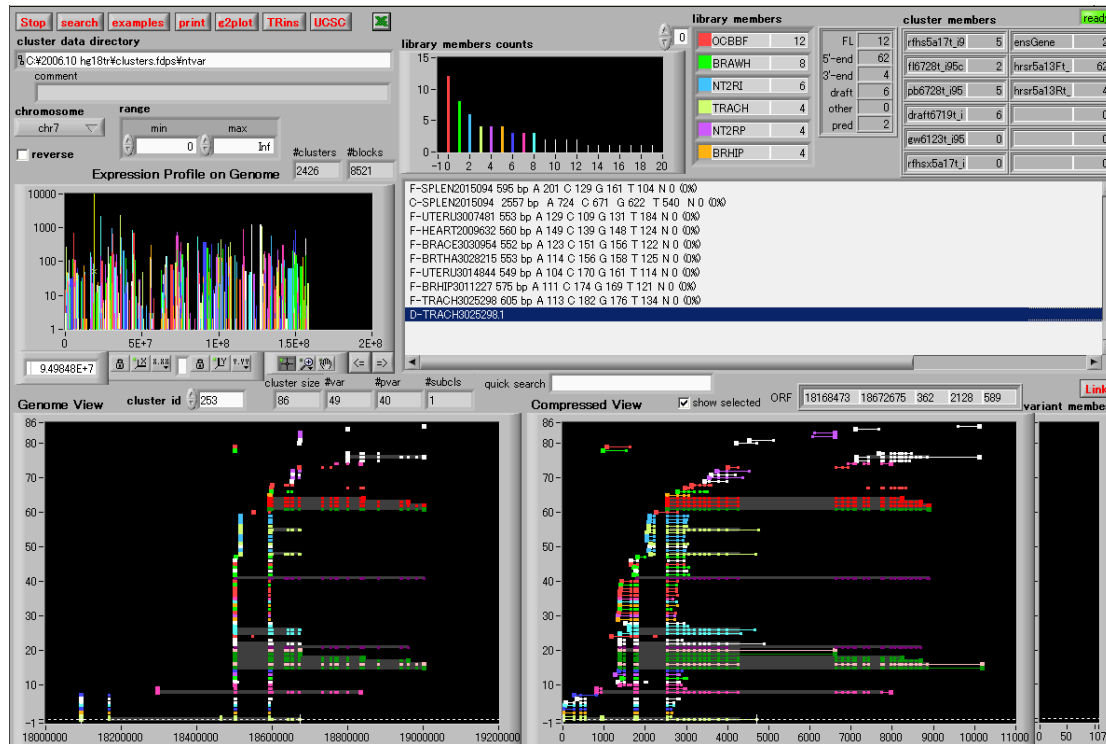
FLJ56188

FLJ55905

NM_139241.1



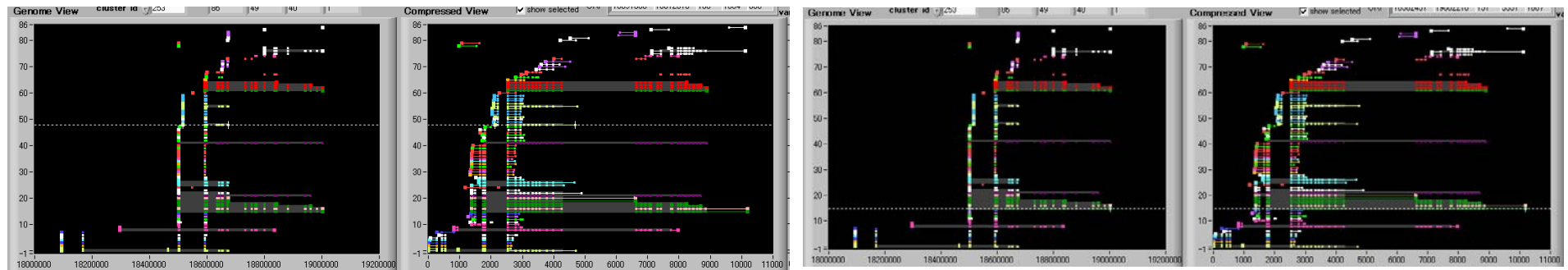
FGD4_FLJ56188,
 FLJ55905,
 NM_139241.1



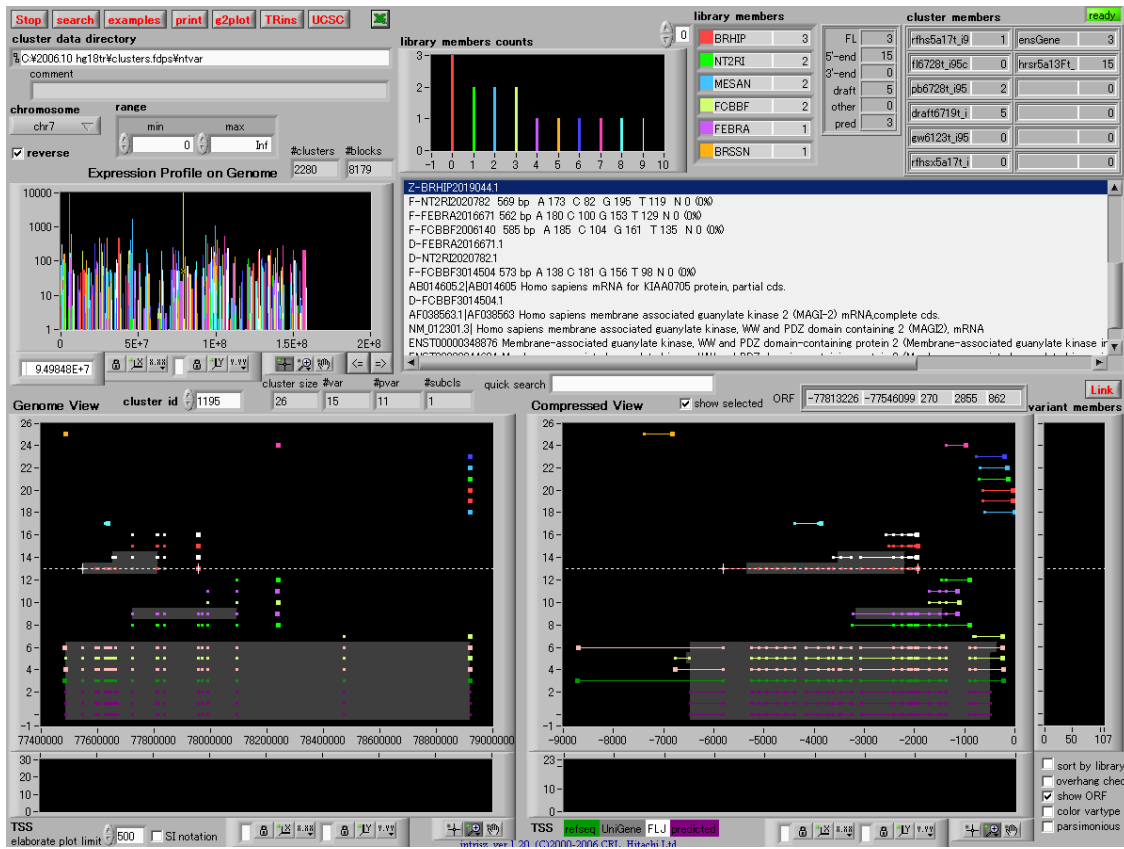
FLJ55607

FLJ54577

NM_178423.1



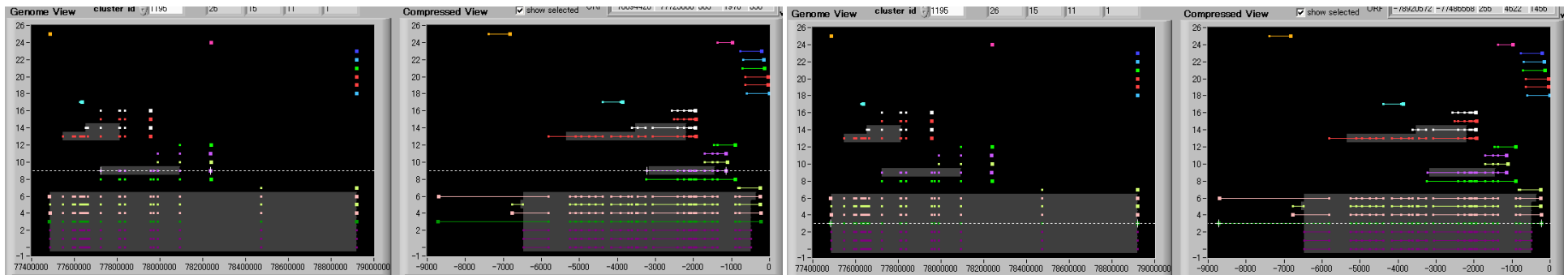
HDAC9_FLJ55607,
FLJ54577,
NM_178423.1



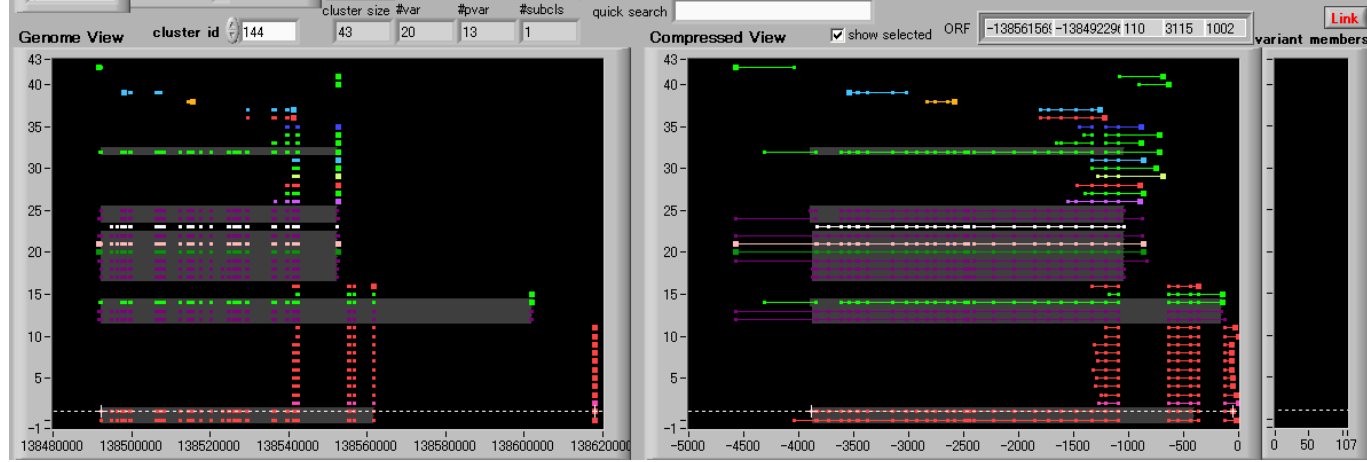
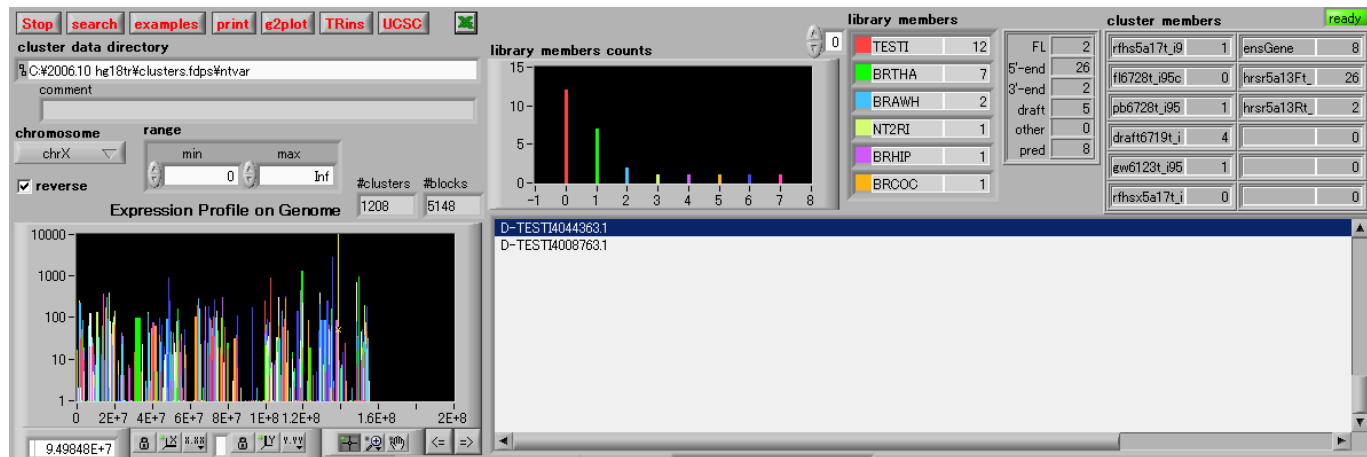
FLJ99053

FLJ50810

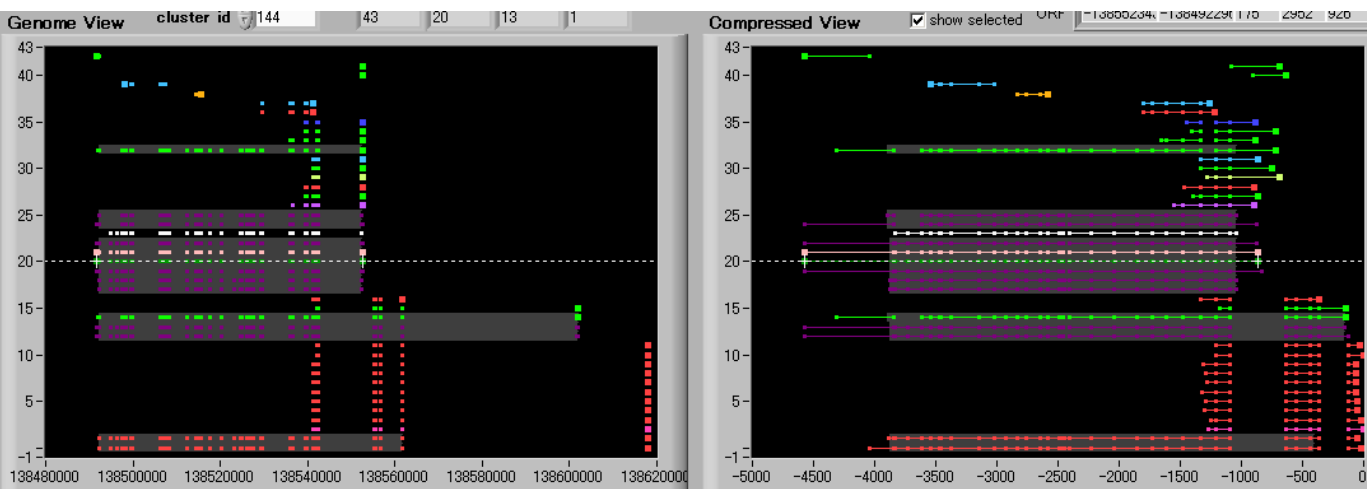
NM_012301.3



MAGI2_FLJ99053,
FLJ50810,
NM_012301.3

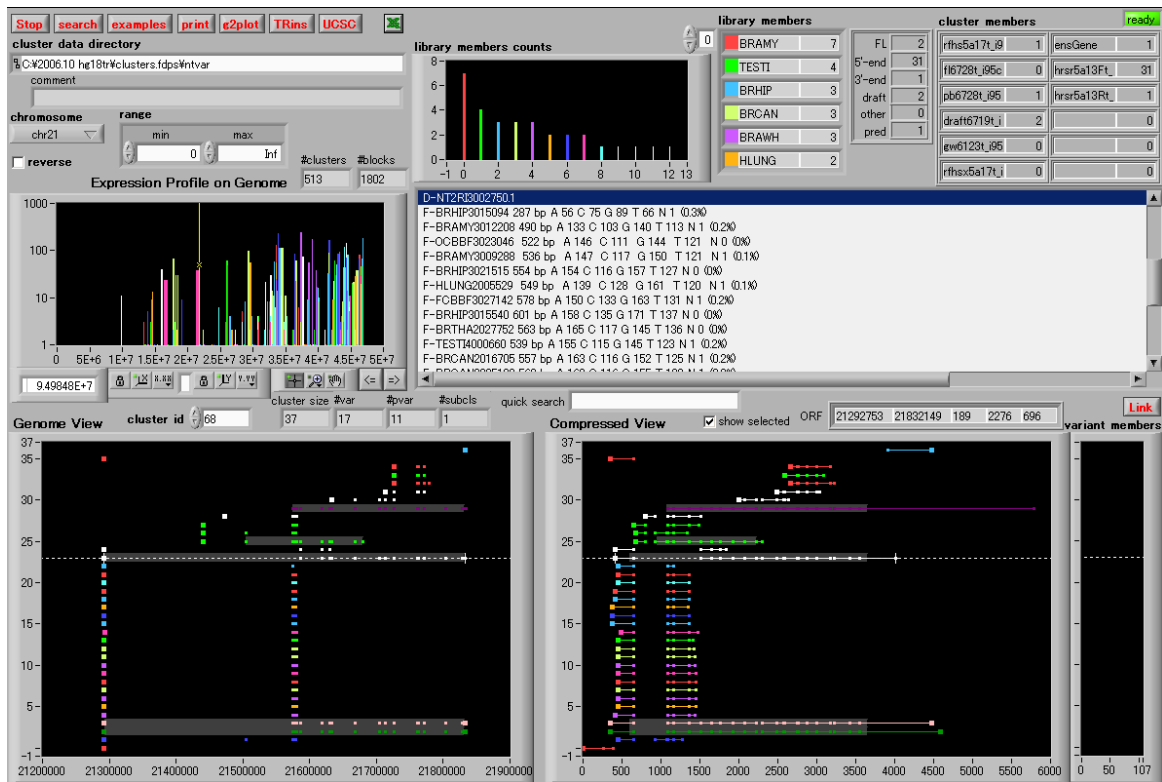


FLJ51685



NM_005369.2

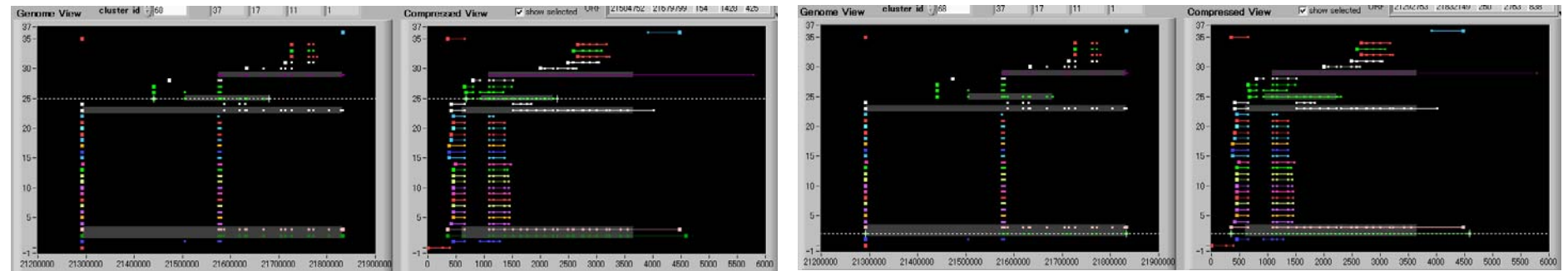
MCF2_FLJ51685,
NM_005369.2



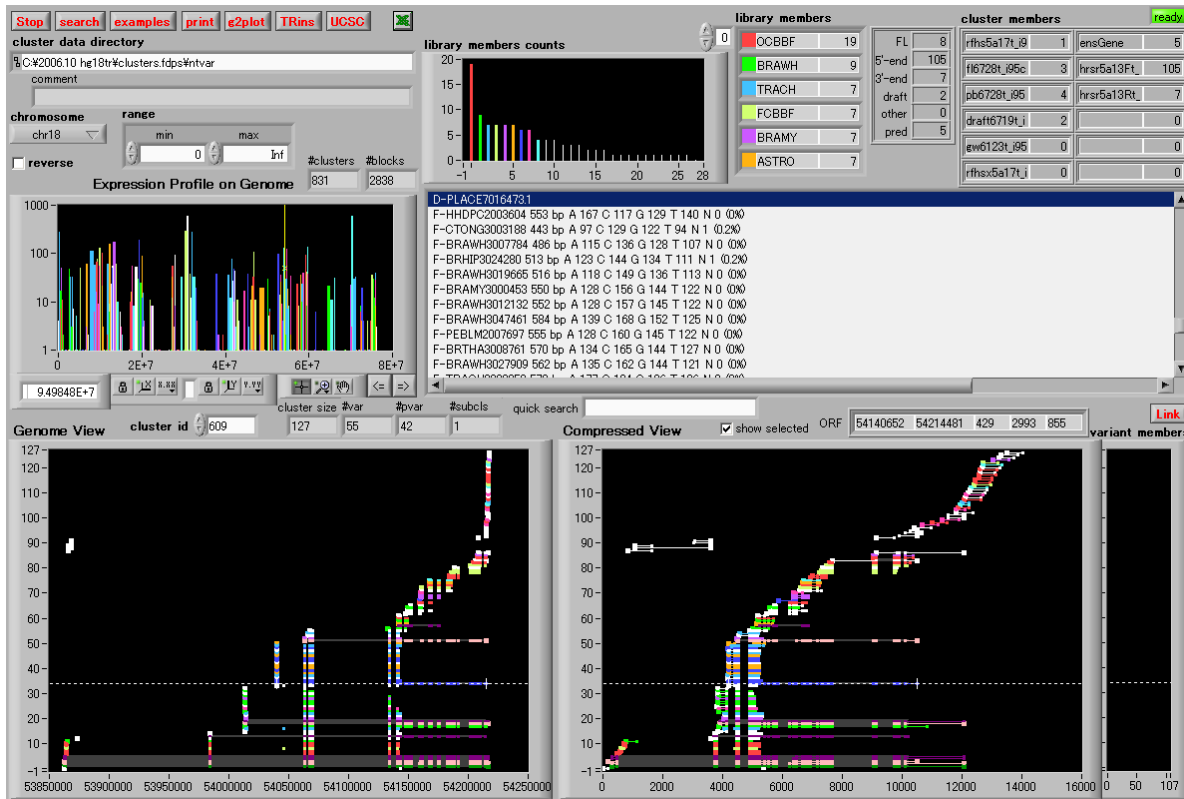
FLJ54289

FLJ53114

NM_004540.2



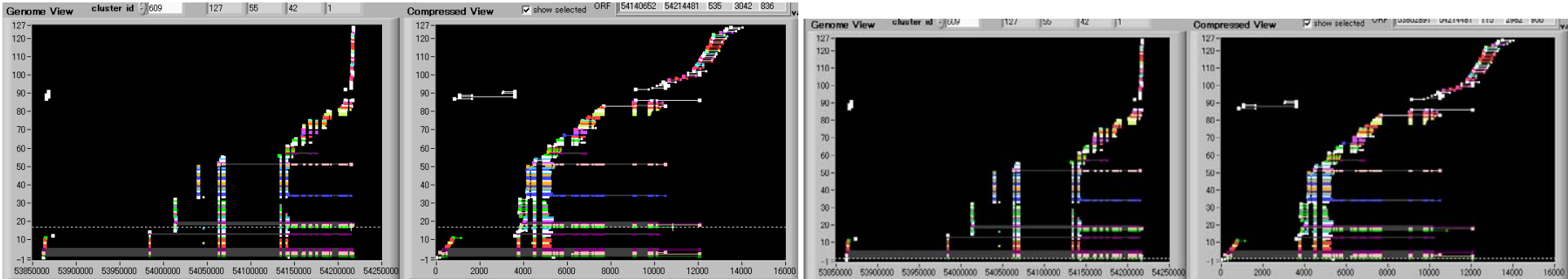
NCAM2_FLJ54289,
FLJ53114,
NM_004540.2



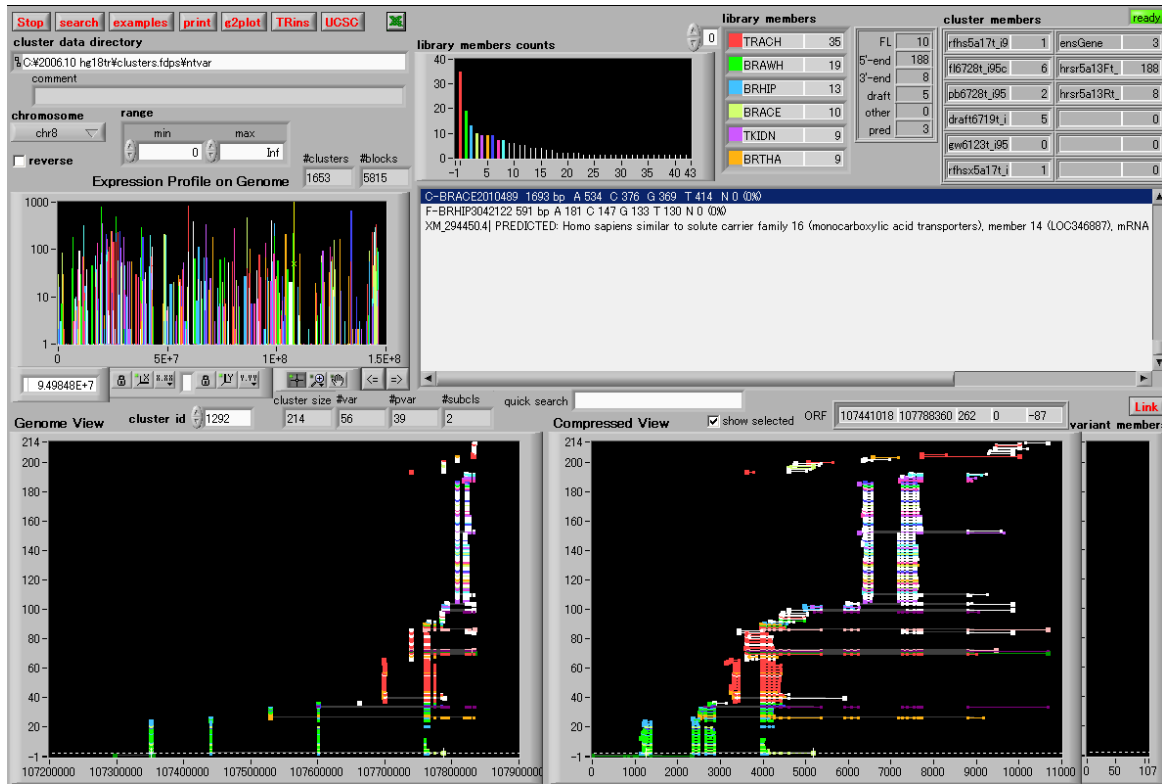
FLJ53199

FLJ61249

NM_015277.2



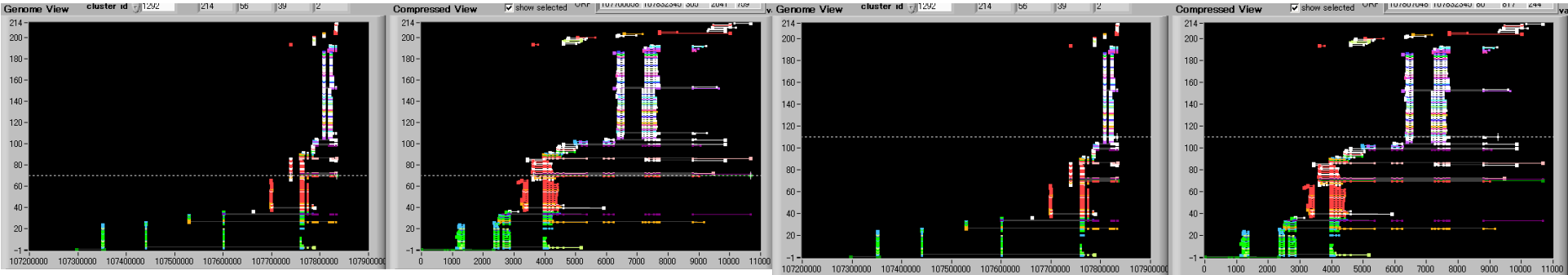
NEDD4L_FLJ53199,
FLJ61249,
NM_015277.2



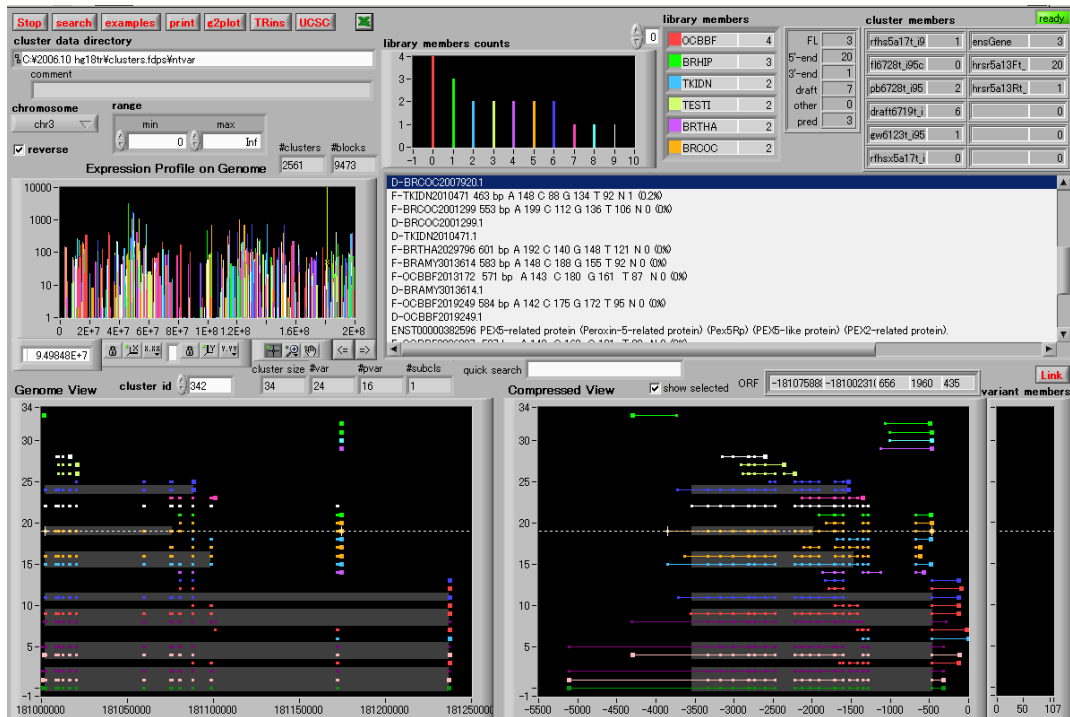
FLJ42450

NM_181354.3

FLJ55036



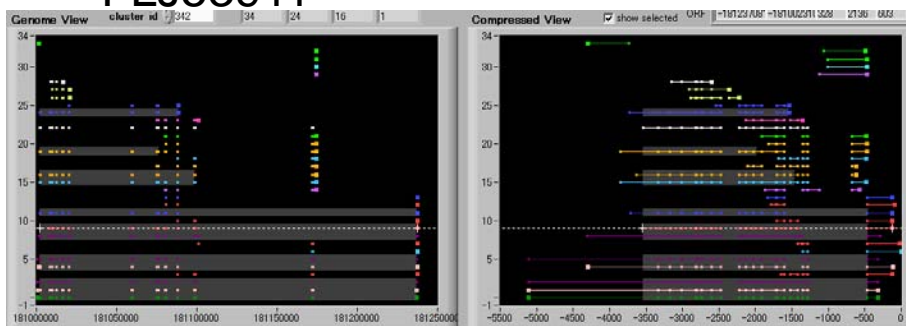
OXR1_FLJ42450,
NM_181354.3,
FLJ55036



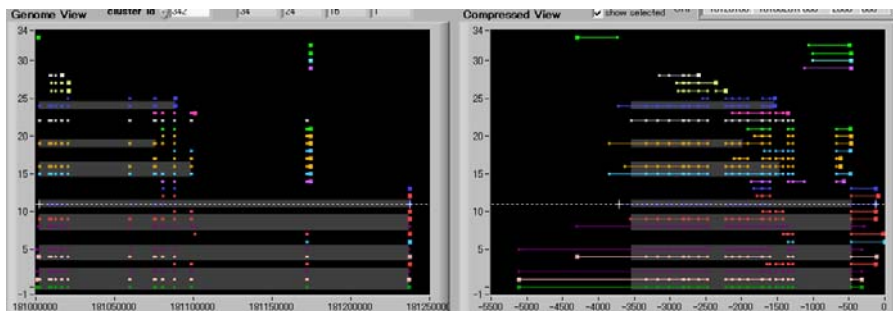
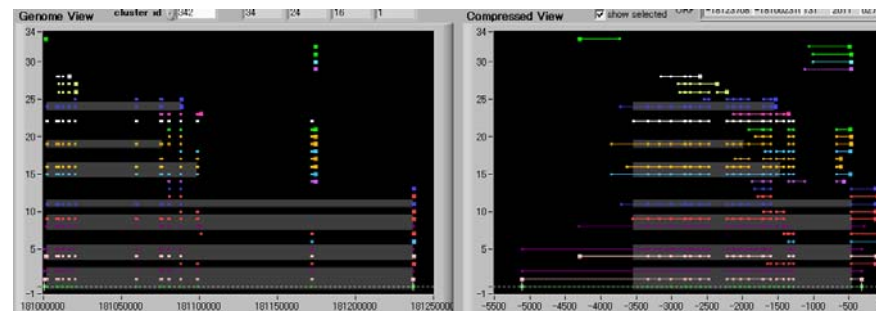
FLJ53911

FLJ50526

NM_016559.1



FLJ50489



PEX5L_FLJ50526,
FLJ53911, FLJ50489,
NM_016559.1

[Stop](#) [search](#) [examples](#) [print](#) [e2plot](#) [TRins](#) [UCSC](#)

cluster data directory
 C:\200610 hg18tr\clusters.fdp\ntvar
 comment

chromosome chr4 **range** min max
 reverse #clusters: 1901 #blocks: 6371

Expression Profile on Genome

library members counts

BRAWH	9
TESTI	6
BRTHA	6
BRAMY	3
NT2RP	2
BRHIP	2

library members

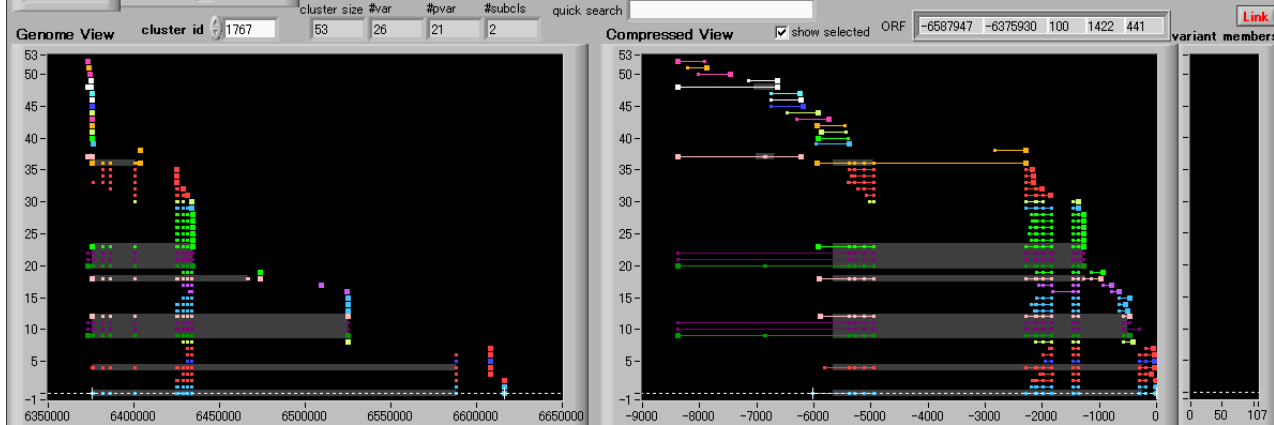
FL	8
5'-end	35
3'-end	4
draft	2
other	0
pred	4

cluster members

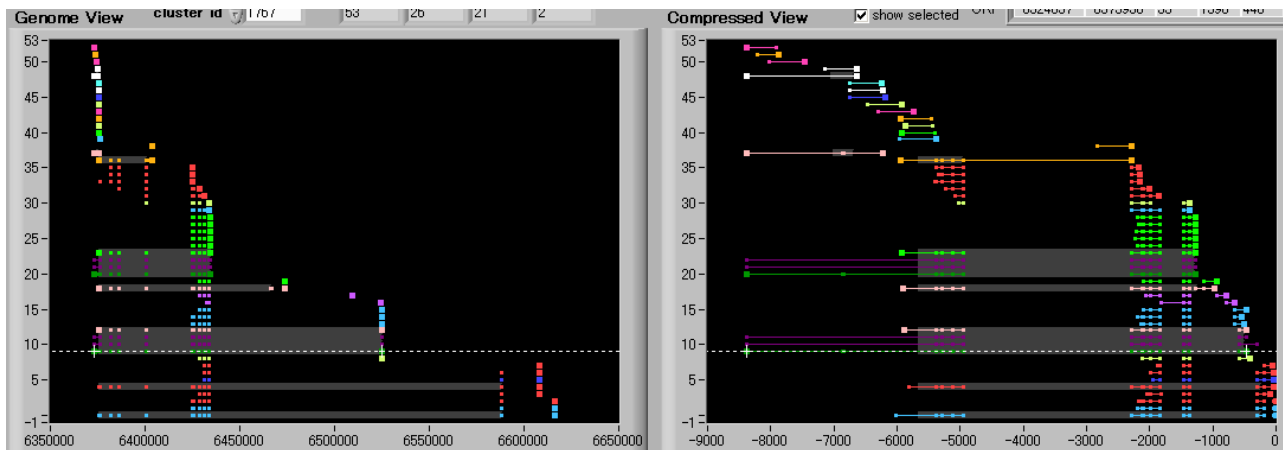
rffsx5a17t_i	2	ensGene	4
fl6728t_95c	3	hrr5a13Ft	35
pb6728t_95	3	hrr5a13Rt	4
draft6719t_i	2		0
gw6123t_95	0		0
rffsx5a17t_i	0		0

ENST00000264959 Serine/threonine-protein phosphatase 2A 55 kDa regulatory subunit B gamma isoform (PP2A, subunit B, B-gamma isoform) (A)
 ENST00000382599 Serine/threonine-protein phosphatase 2A 55 kDa regulatory subunit B gamma isoform (PP2A, subunit B, B-gamma isoform) (C)
 NM_020416.2 Homo sapiens protein phosphatase 2 (formerly 2A), regulatory subunit B (PR 52), gamma isoform (PPP2R2C), transcript variant 1, (M)

F-BRAMY3010518 548 bp A 147 C 153 G 164 T 84 N 0 (0%)
 F-BRAWH3028123 289 bp A 69 C 89 G 87 T 43 N 1 (0.3%)
 F-BRAWH3013275 517 bp A 141 C 144 G 143 T 89 N 0 (0%)
 F-BRCAN2017766 480 bp A 127 C 130 G 137 T 86 N 0 (0%)
 D-BRAWH3013275.1
 F-BRAWH3007753 542 bp A 155 C 151 G 144 T 92 N 0 (0%)
 F-BRAWH3048704 589 bp A 167 C 151 G 163 T 108 N 0 (0%)
 F-BRTHA3017732 610 bp A 174 C 160 G 164 T 112 N 0 (0%)
 D-BRTHA3017732.1

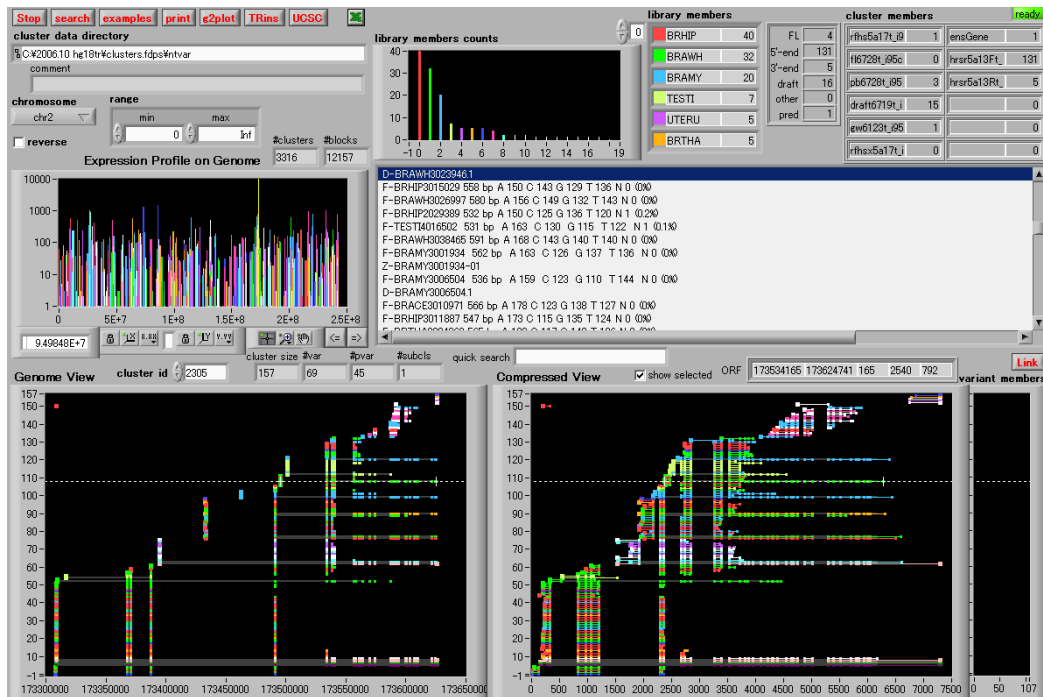


FLJ58008



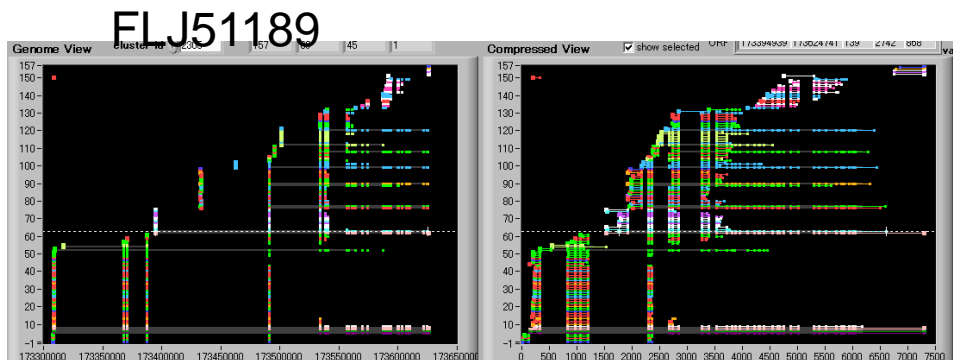
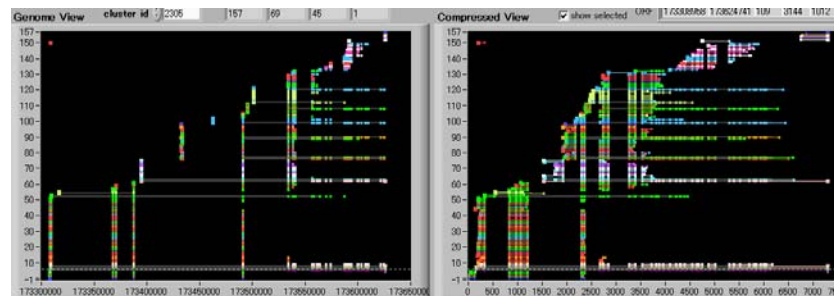
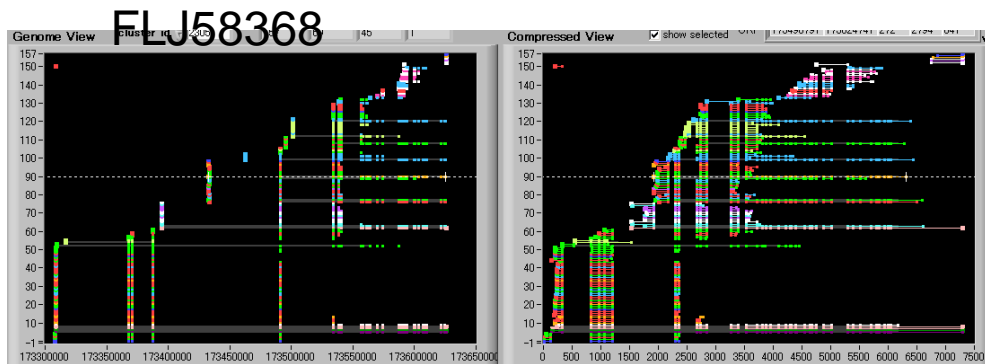
NM_020416.2

PPP2R2C_FLJ58008,
NM_020416.2

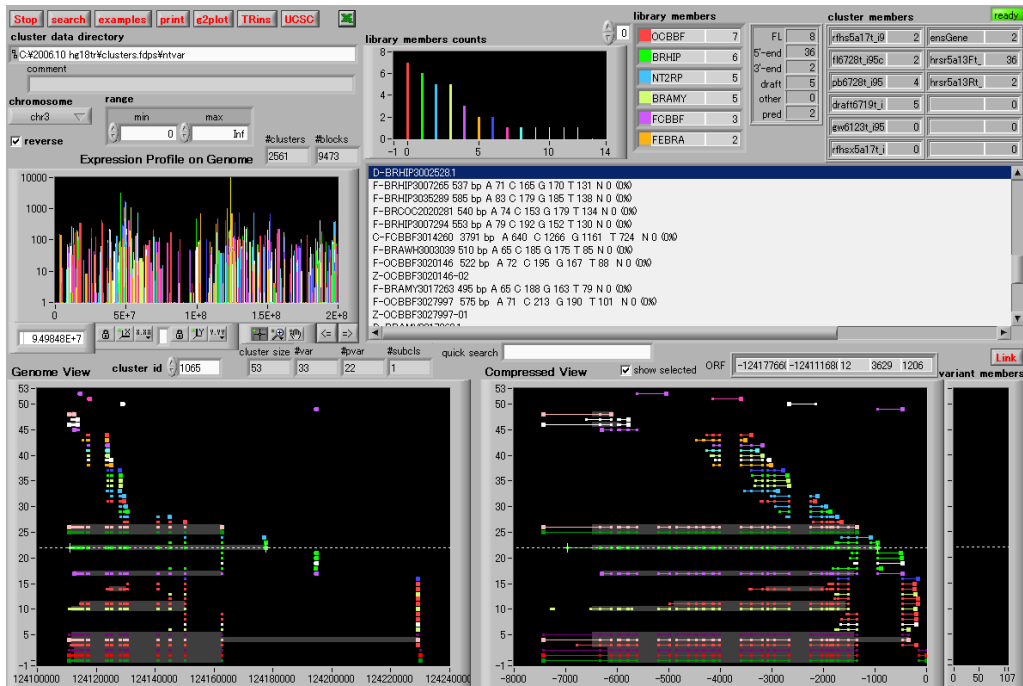


FLJ50956

NM_007023.1



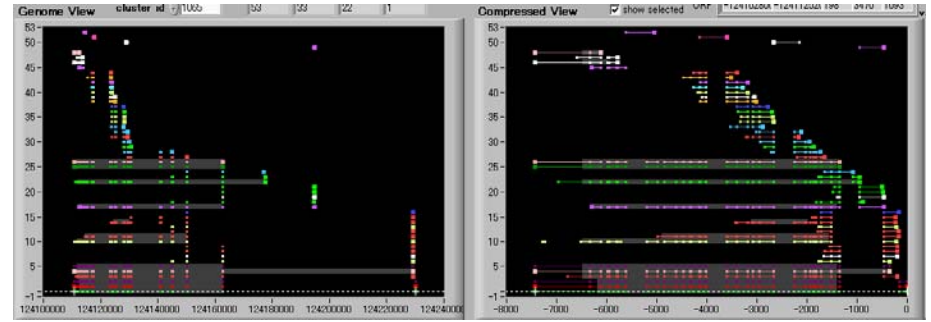
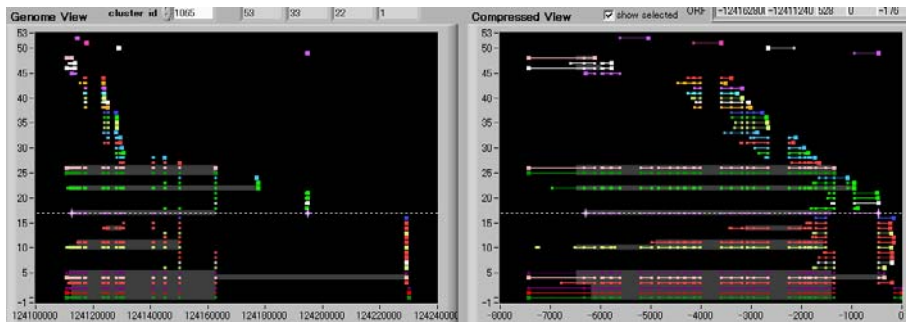
RAPGEF4_FLJ50956,
FLJ58368, FLJ51189,
NM_007023.1



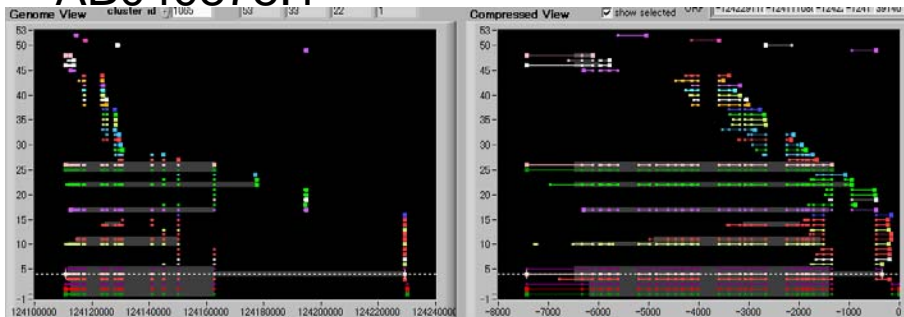
FLJ55460

NM_018987.1

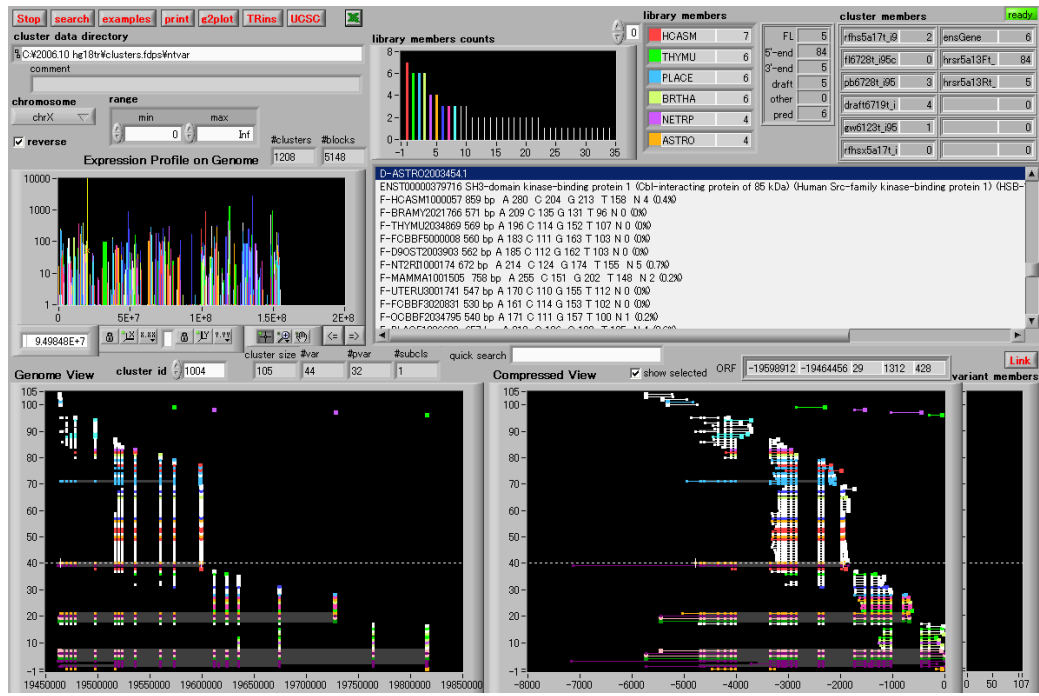
FLJ34162



AB040878.1



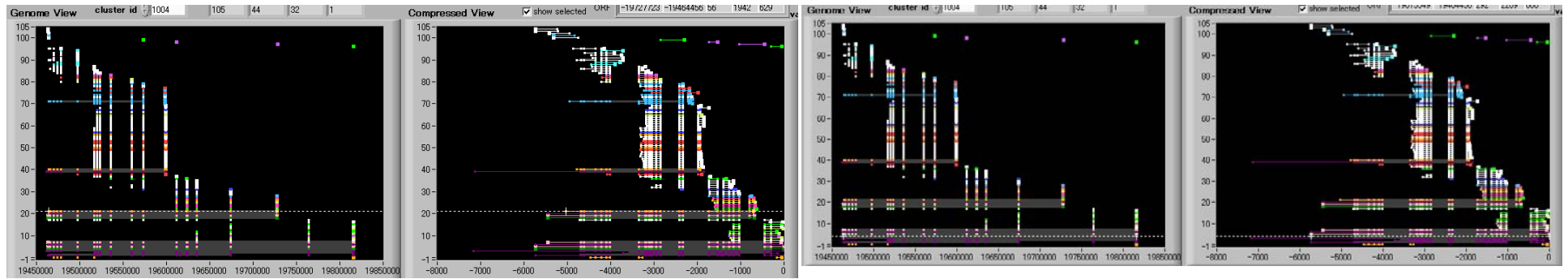
SEMA5B_FLJ55460,
FLJ34162,
AB040878.1,
NM_018987.1



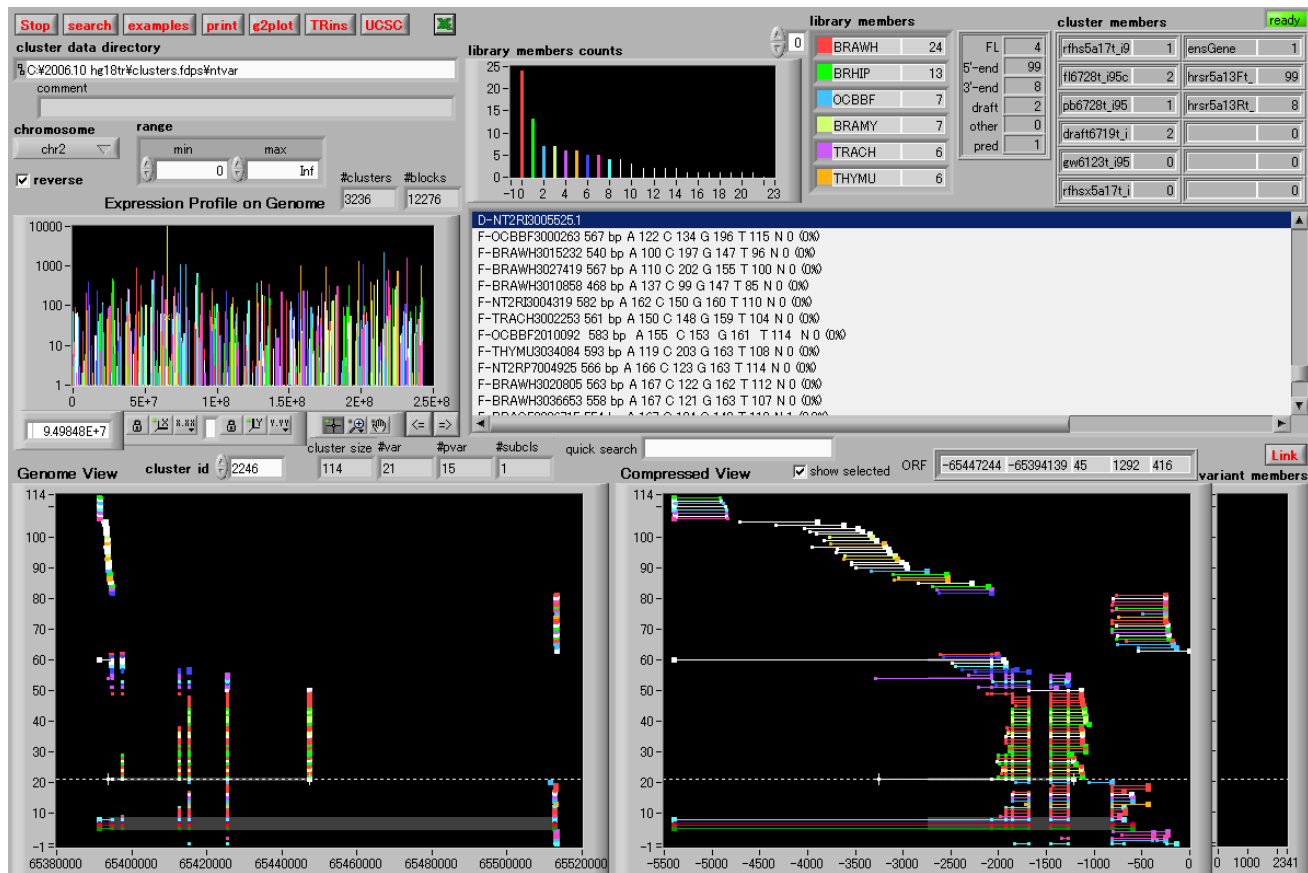
FLJ54612

FLJ54623

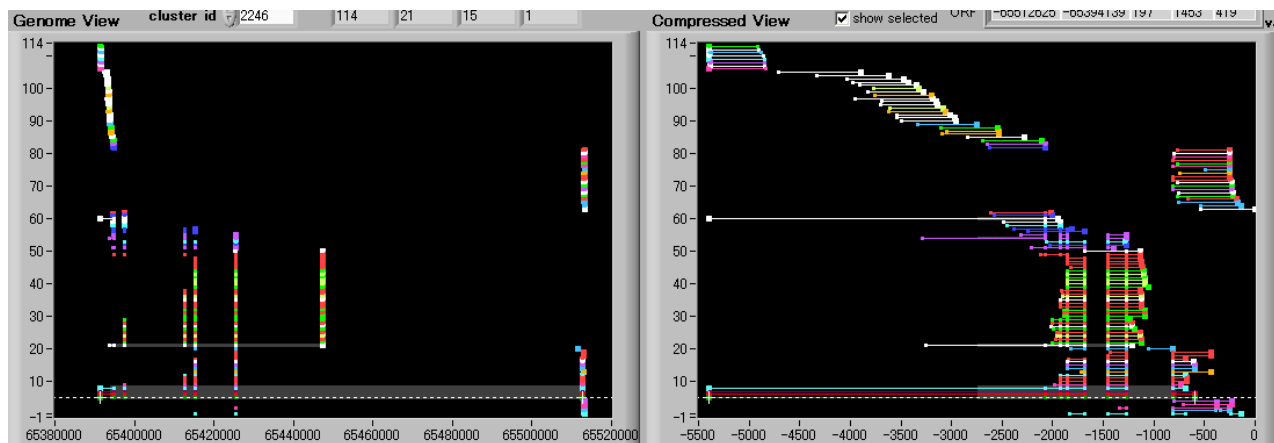
NM_031892.1



SH3KBP1_FLJ54612,
 FLJ54623, NM_031892.1

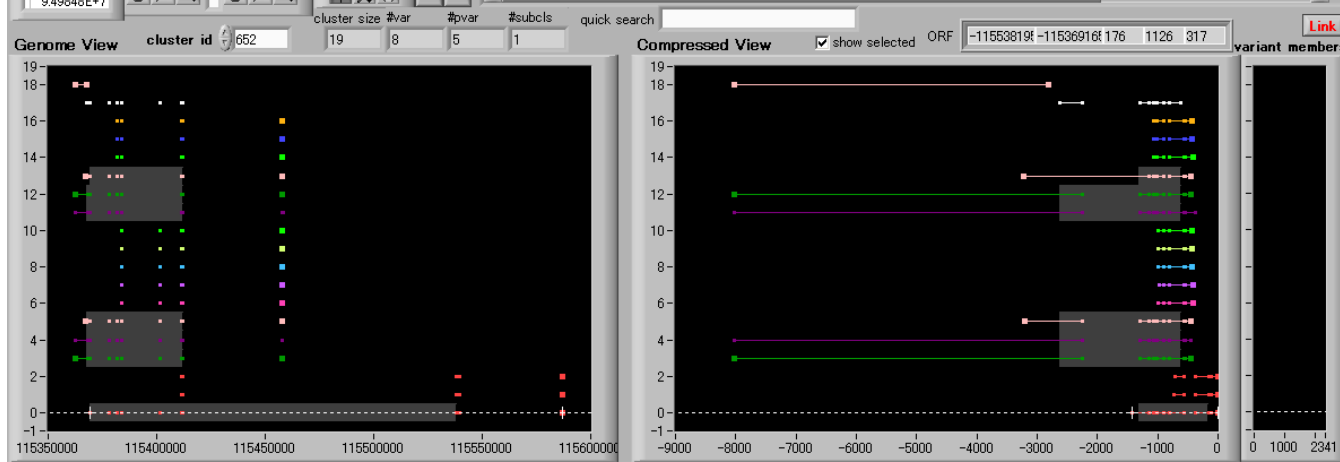
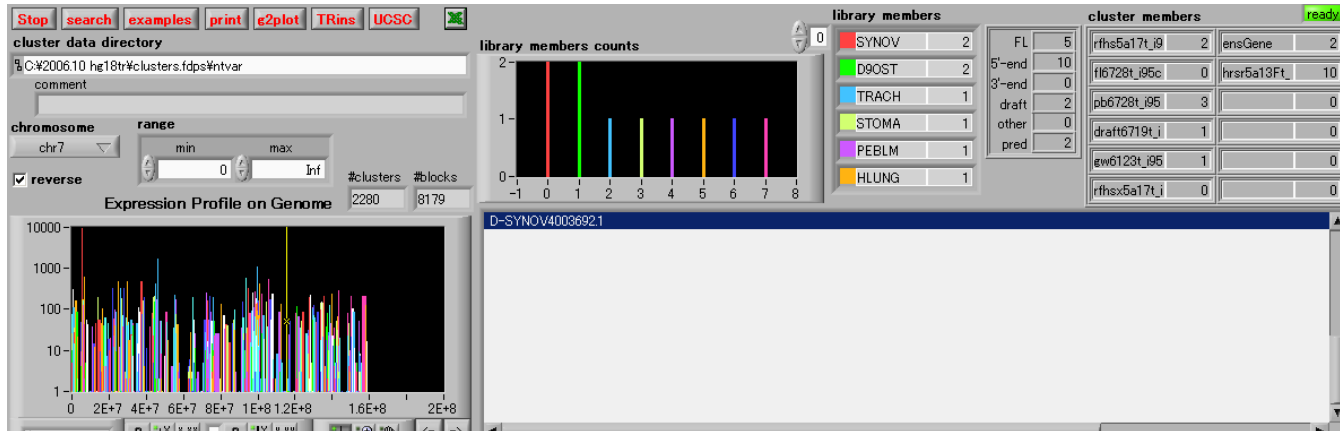


FLJ52731

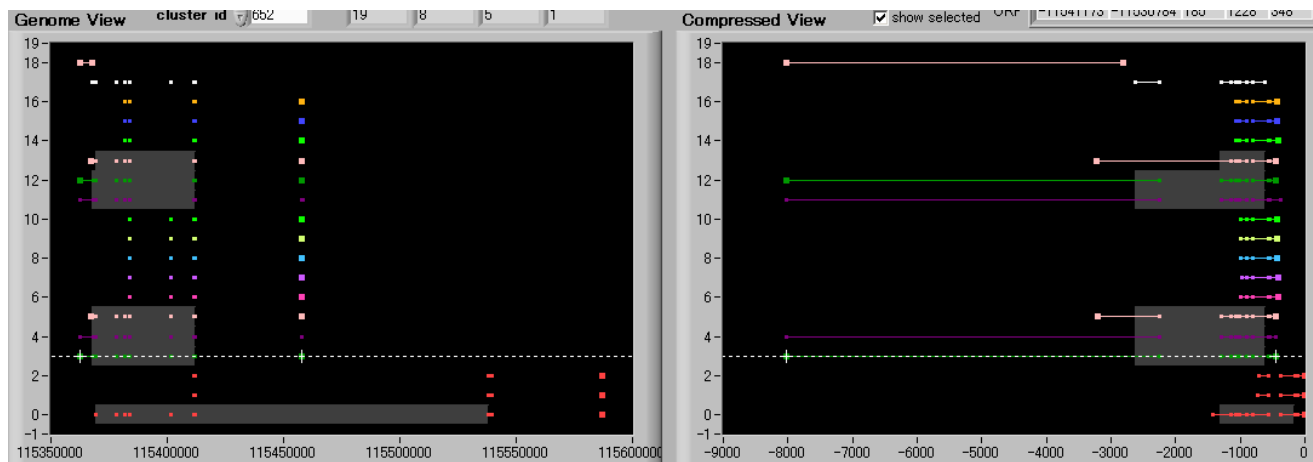


NM_181784.1

SPRED2_FLJ52731,
 NM_181784.1

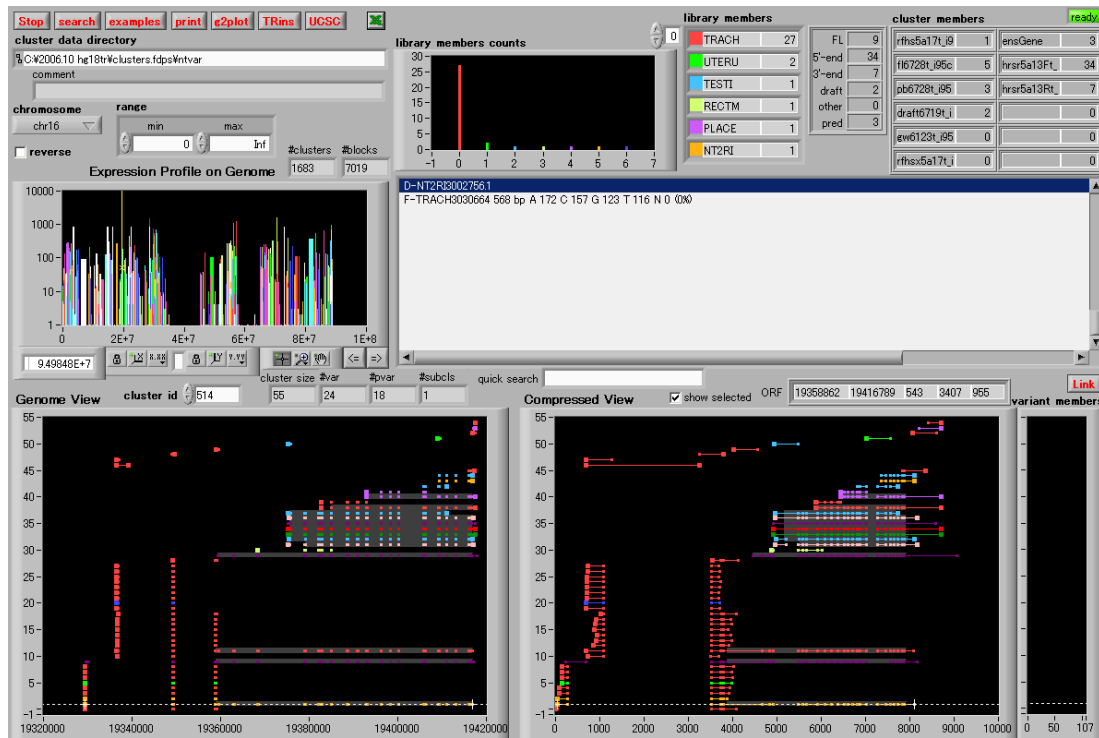


FLJ55256



NM_012252.2

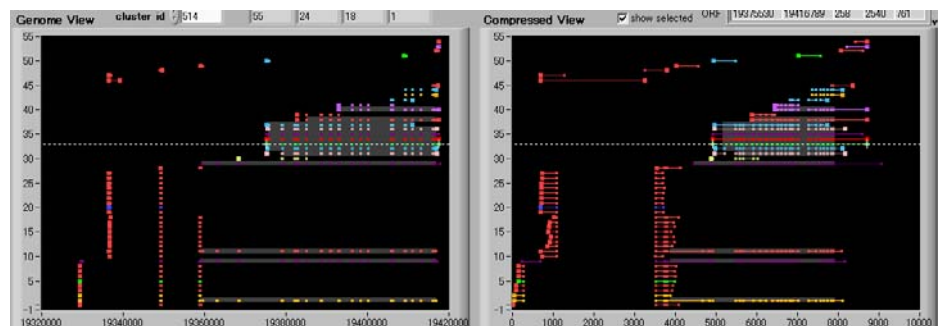
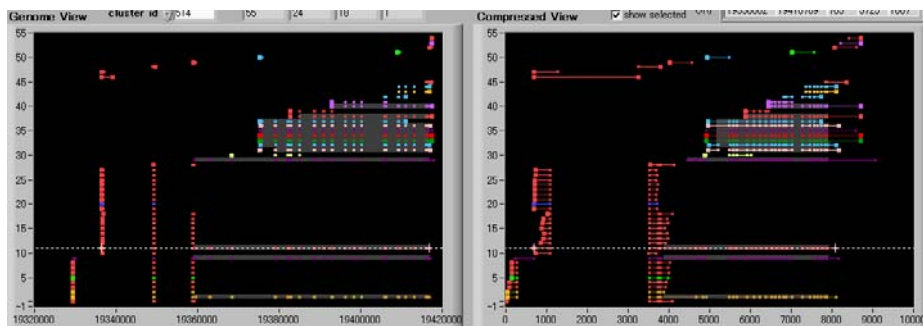
TFEC_FLJ55256,
NM_012252.2



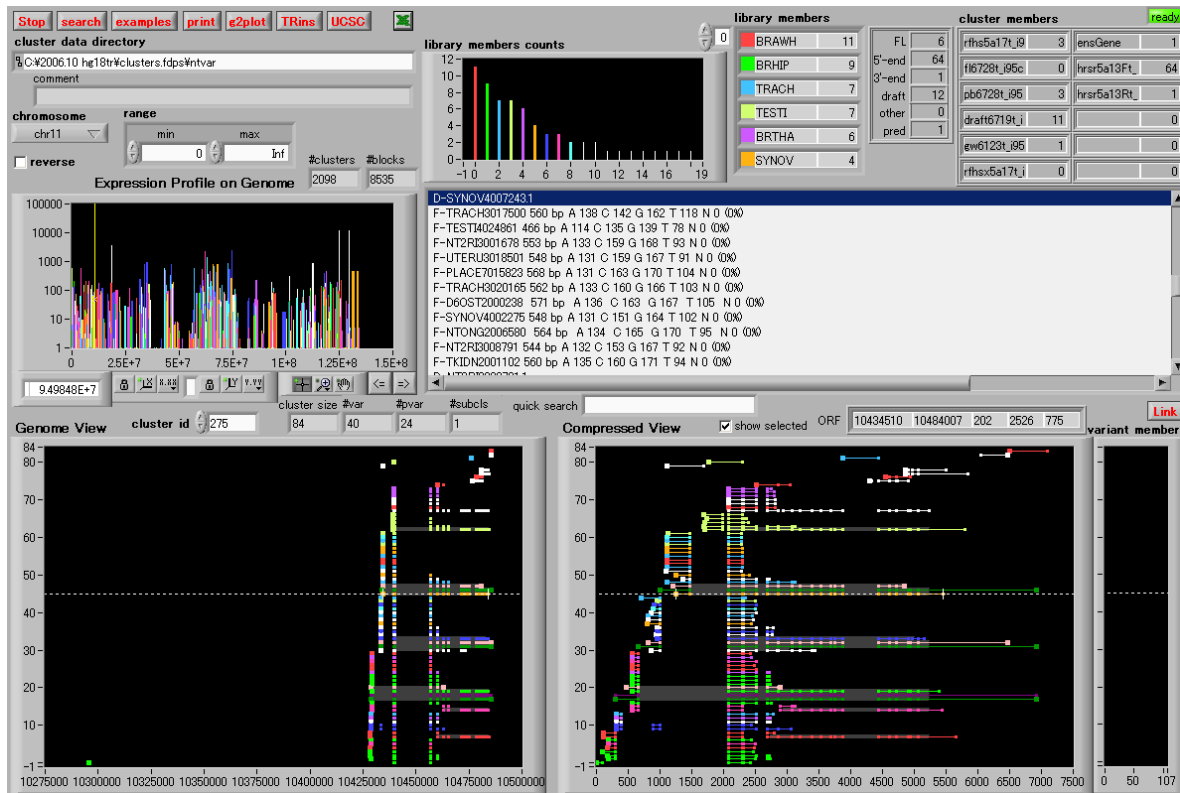
FLJ54454

FLJ54906

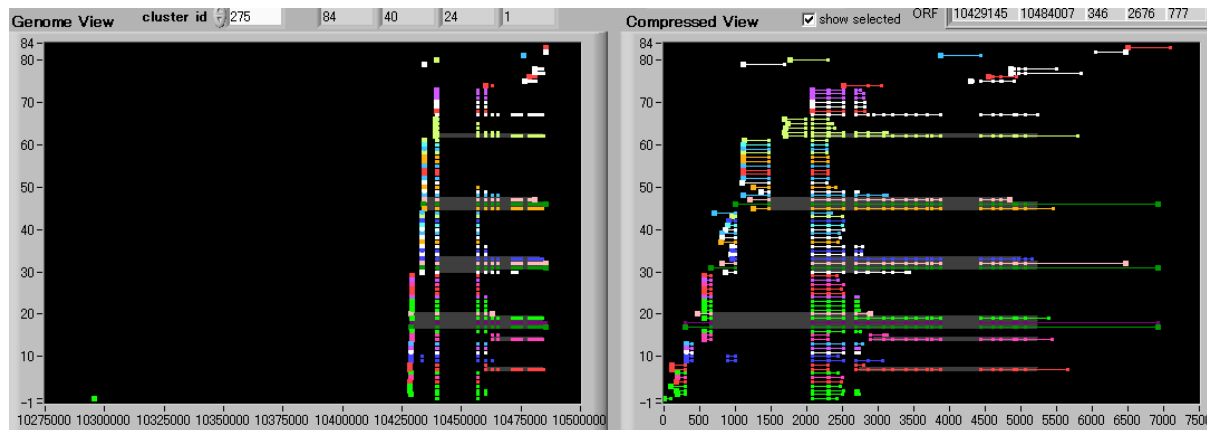
NM_024780.3



TMC5_FLJ54454,
FLJ54906,
NM_024780.3

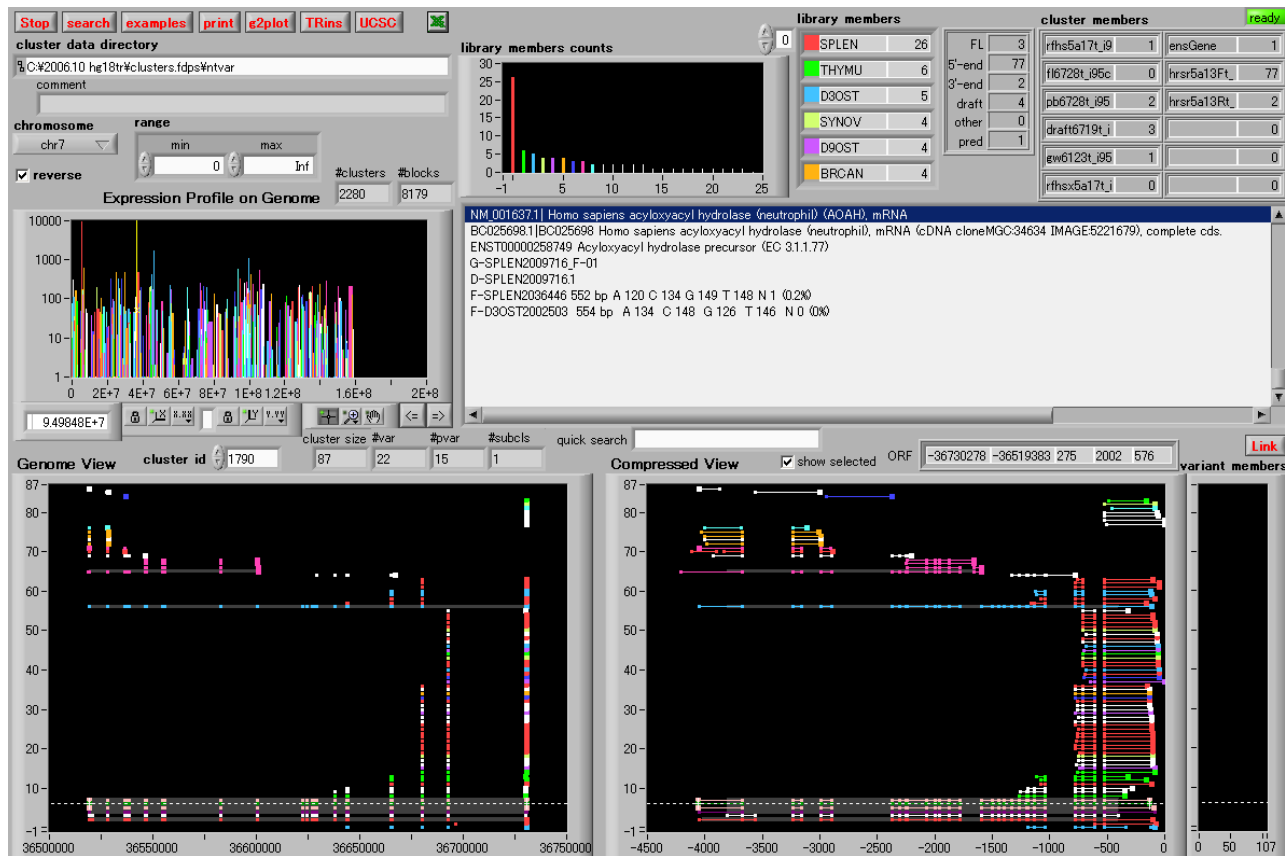


FLJ51124

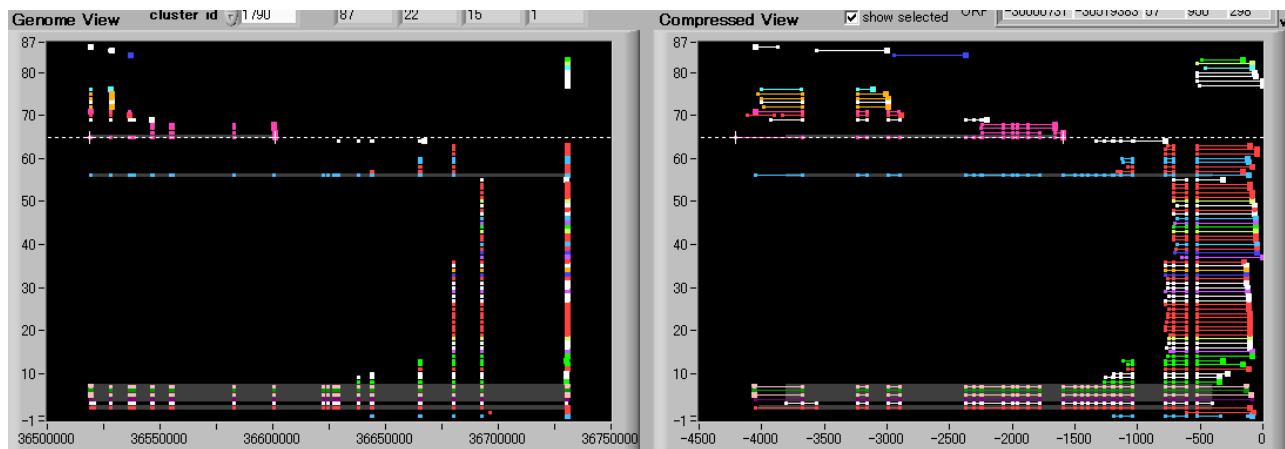


NM_000480.2

AMPD3_FLJ51124,
NM_000480.2

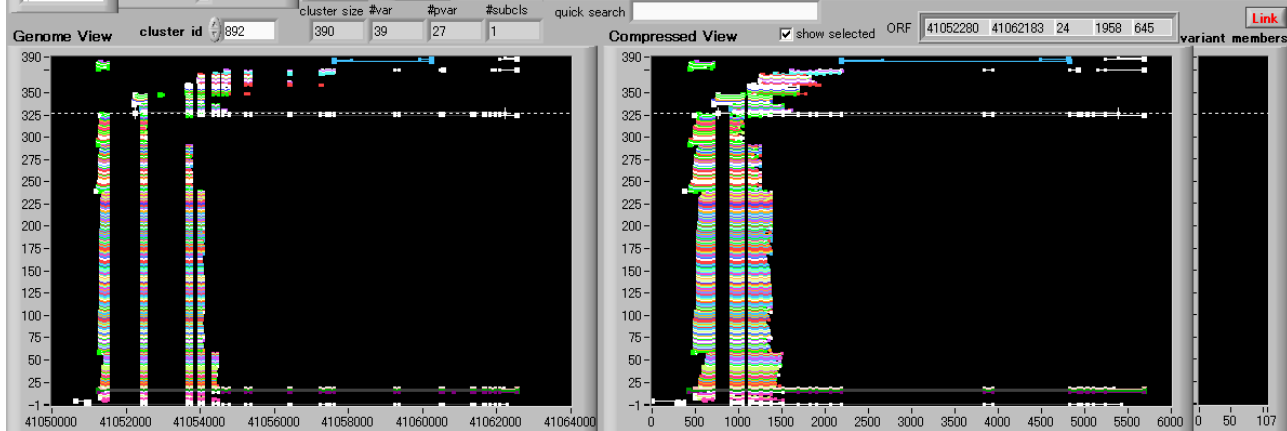
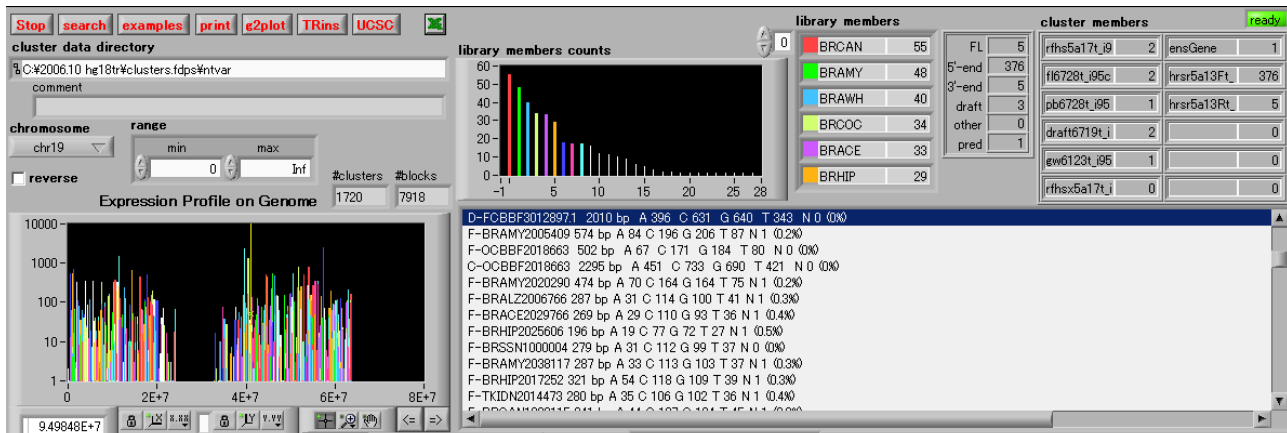


NM_001637.1

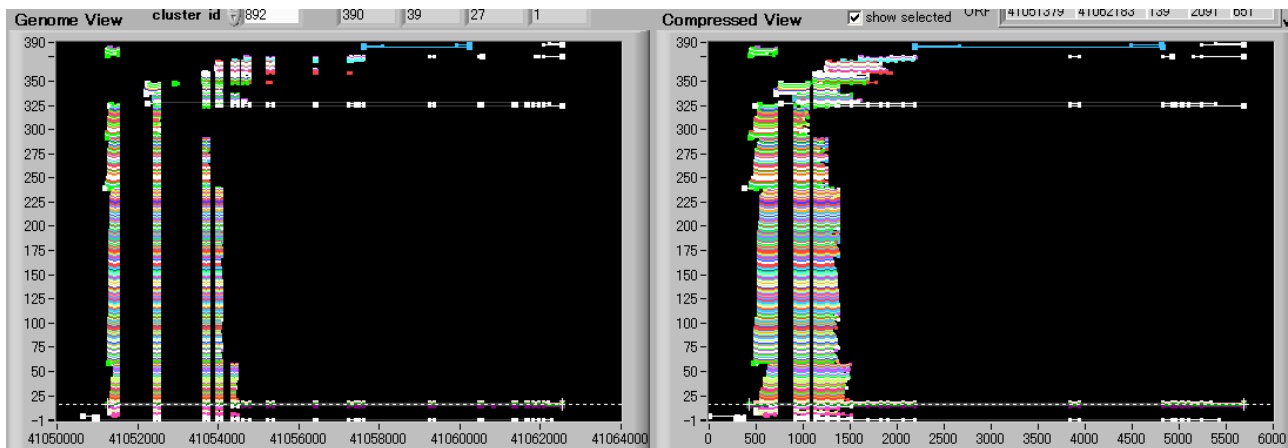


FLJ51934

AOAH_NM_001637.1,
 FLJ51934

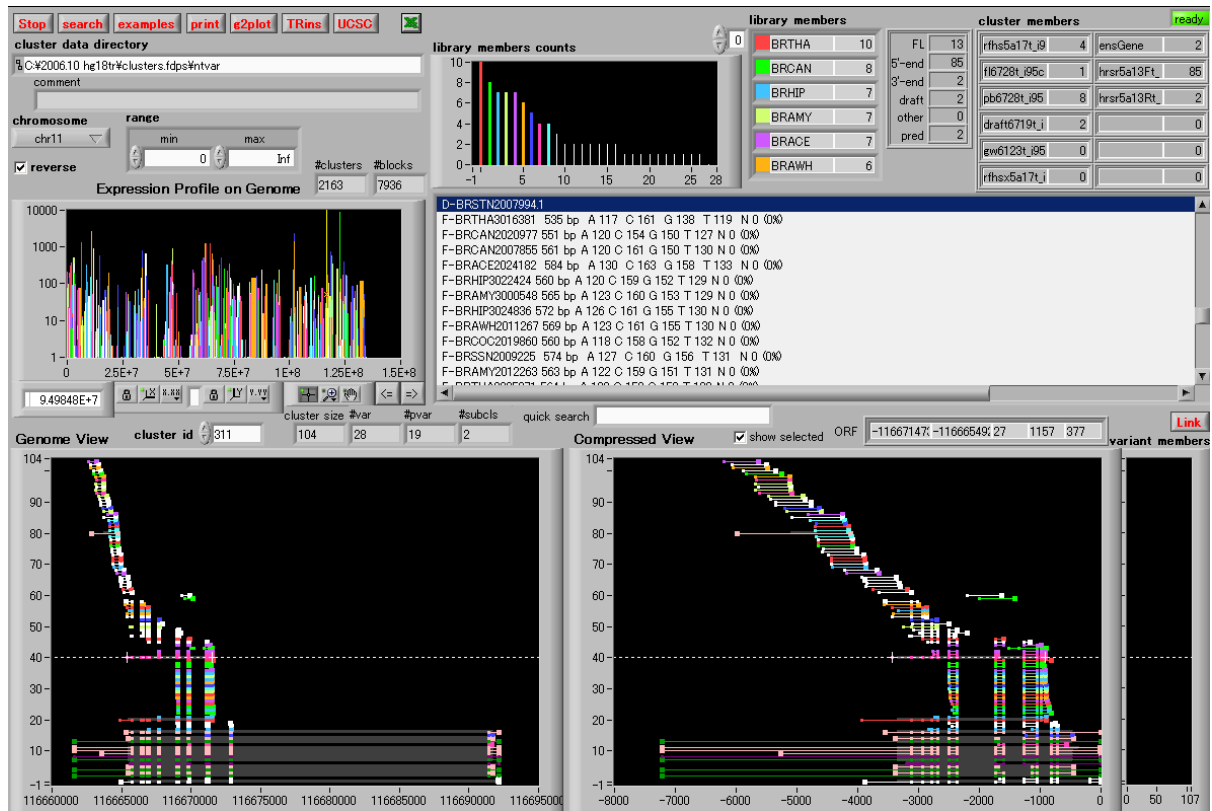


FLJ56046

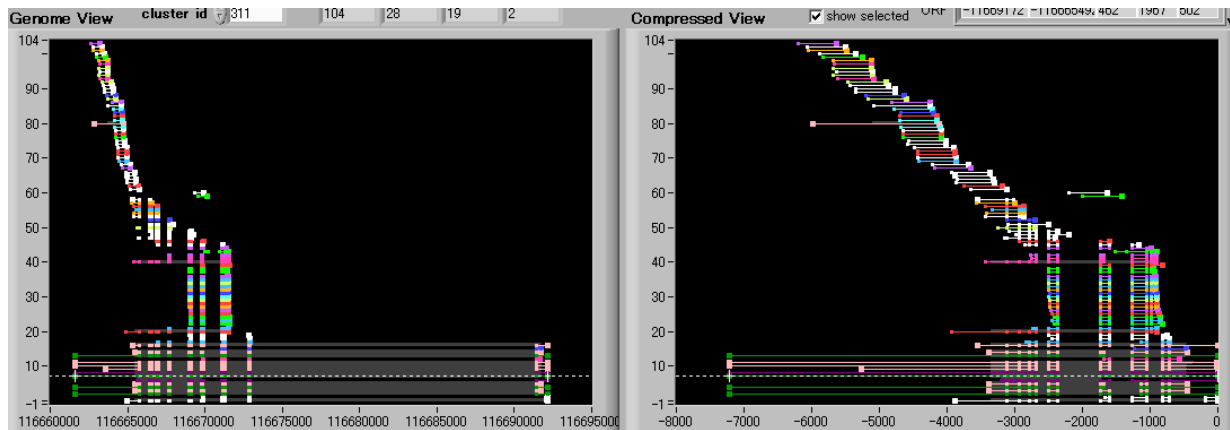


NM_005166.3

APLP1_FLJ56046,
 NM_005166.3

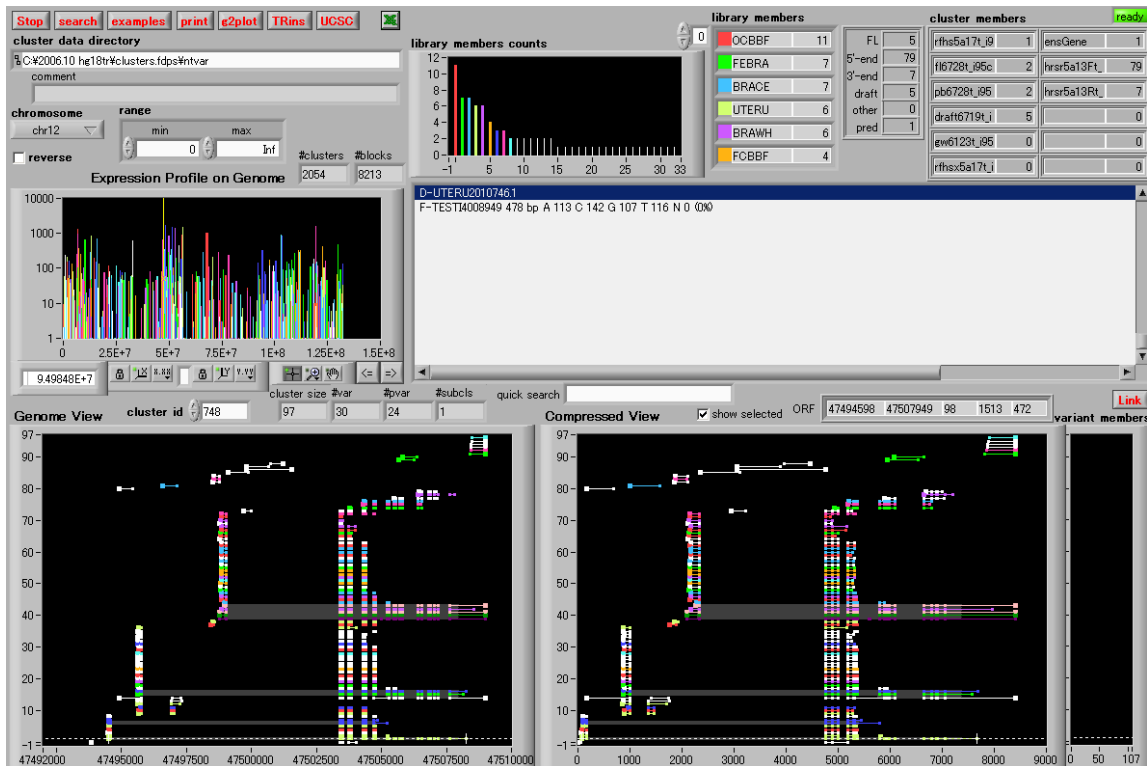


FLJ54690



NM_012104.3

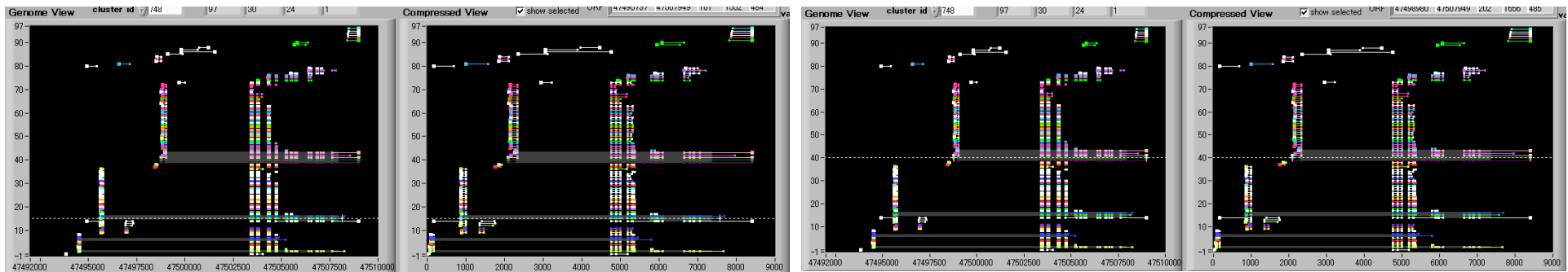
BACE1_FLJ54690,
NM_012104.3



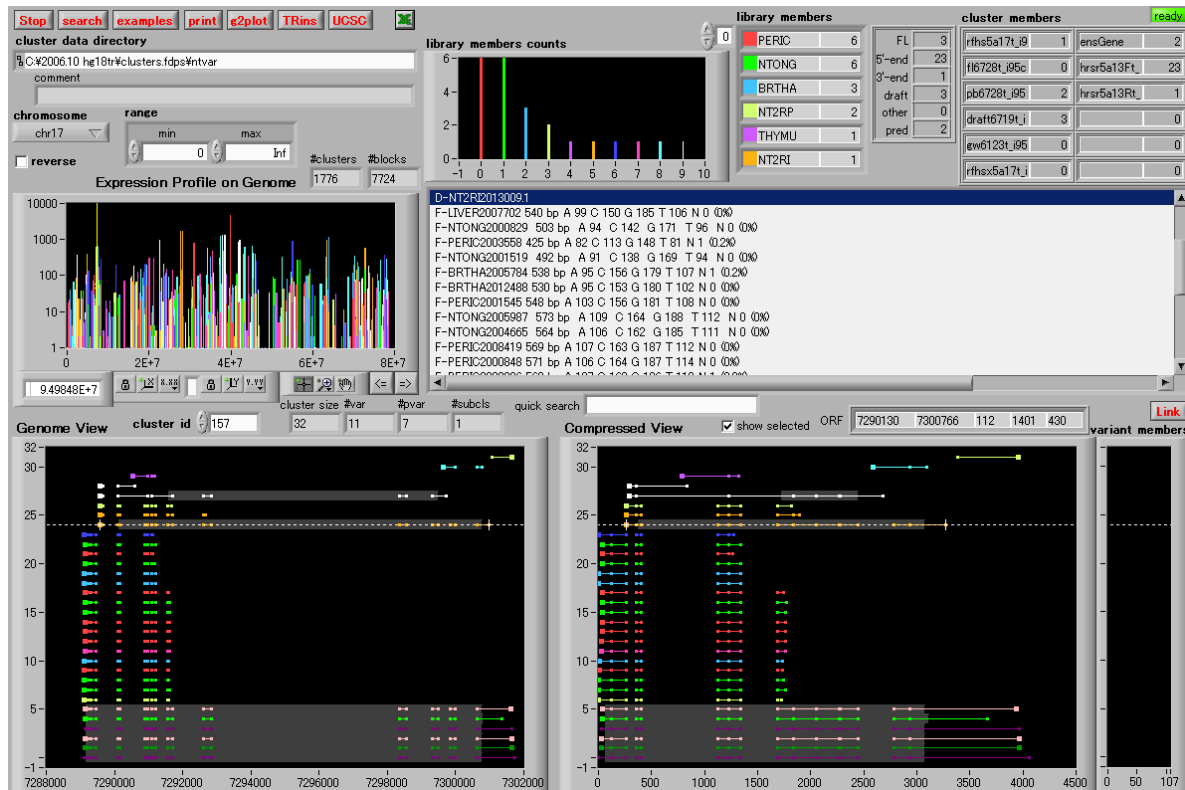
FLJ58411

FLJ57401

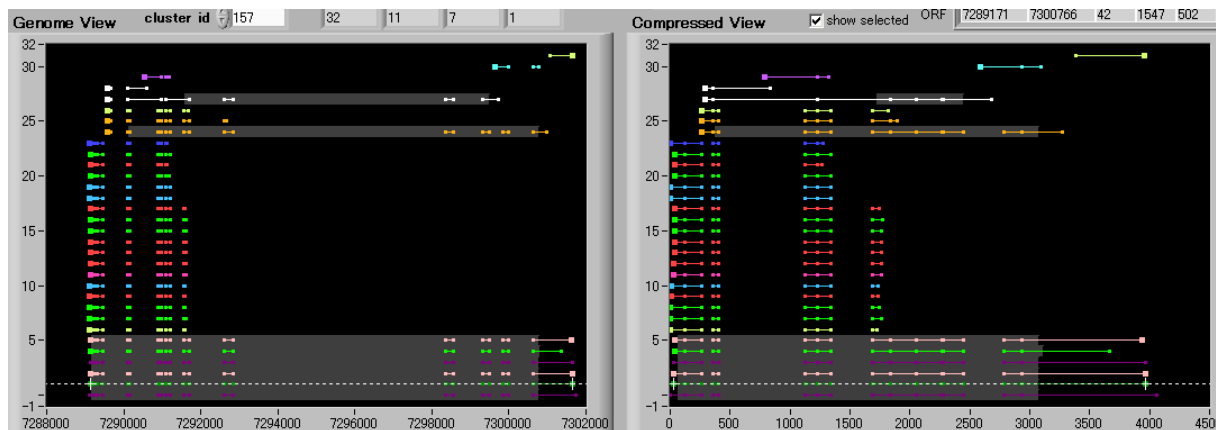
NM_000725.2



CACNB3_FLJ58411,
FLJ57401,
NM_000725.2

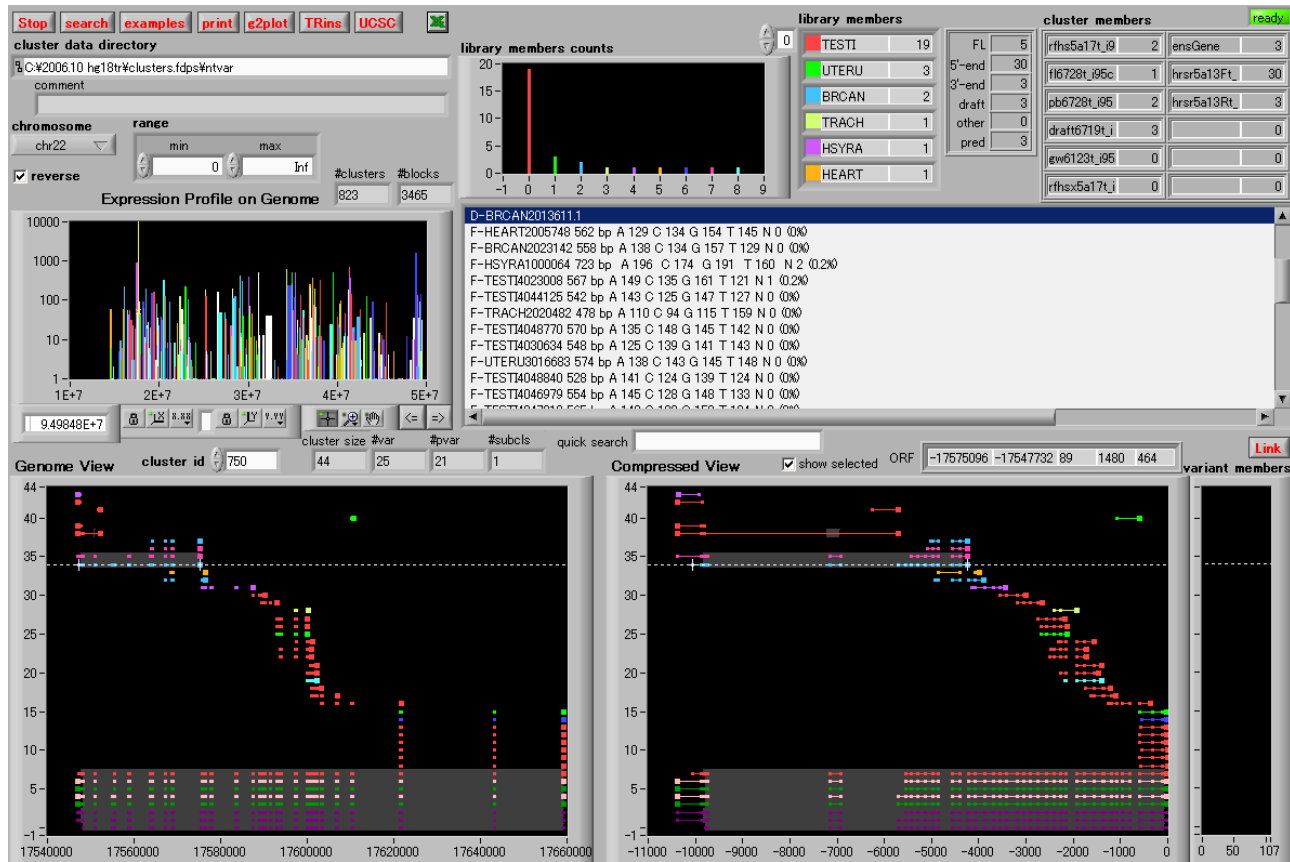


FLJ52354

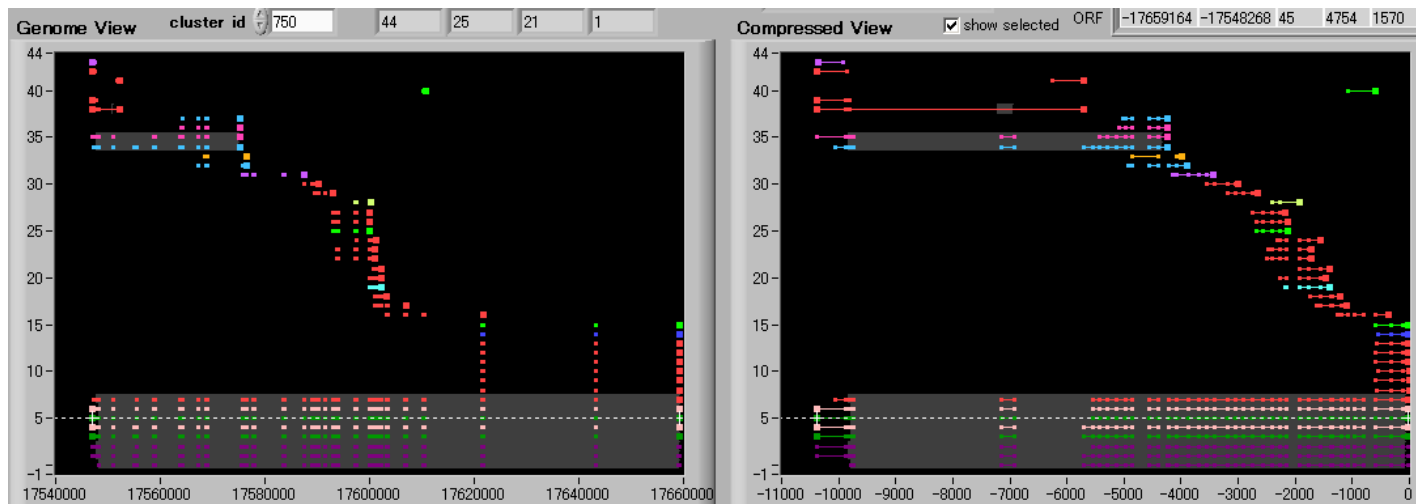


NM_000747.2

CHRN1_FLJ52354,
NM_000747.2



FLJ56961



NM_001835.1

CLTCL1_FLJ56961,
NM_001835.1

[Stop](#) [search](#) [examples](#) [print](#) [e2plot](#) [TRIns](#) [UCSC](#)

cluster data directory
 C:\2006.10 hg18tr\clusters.fdfs\ntvar
 comment

chromosome chr10 **range** min max
 reverse #clusters: 1961 #blocks: 7027

Expression Profile on Genome

library members counts

Library Member	Count
TESTI	15
CTONG	7
ASTRO	5
NT2RI	4
BRHIP	4
TRACH	3

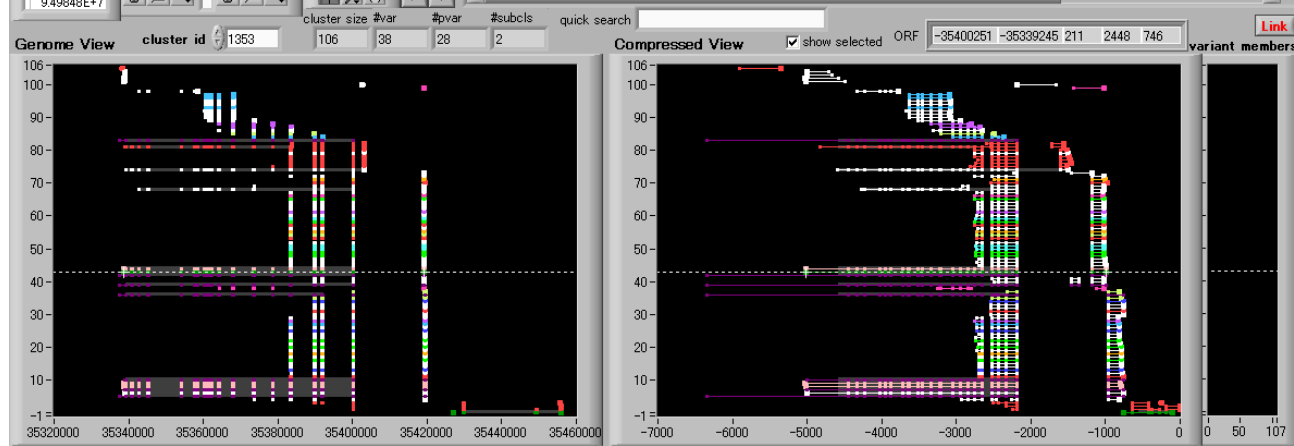
Category	Count
FL	6
5'-end	85
3'-end	6
draft	3
other	0
pred	7

Cluster Member	Count
rths5a17L_i9	1
ensGene	7
fl6728t_i95c	1
hrrs5a13Ft	85
pb6728t_i95	3
hrrs5a13Rt	5
draft6719t_i	3
ew6123t_i95	0
rthsx5a17L_i	1
	0

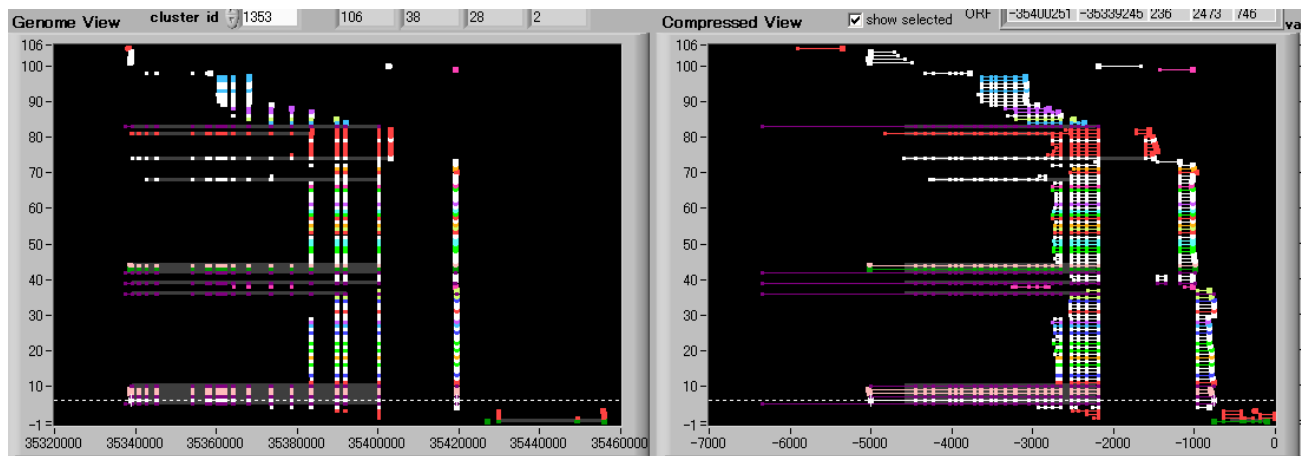
library members
 TESTI: 15, CTONG: 7, ASTRO: 5, NT2RI: 4, BRHIP: 4, TRACH: 3

cluster members ready

NM_003591.2 Homo sapiens cullin 2 (CUL2), mRNA
 ENST00000374749 Cullin-2 (CUL2-2).
 F-BRAMY3013213 600 bp A 146 C 166 G 143 T 145 N 0 (0%)
 F-HCASM2002340 559 bp A 132 C 160 G 130 T 137 N 0 (0%)
 ENST00000374748 Cullin-2 (CUL2-2).
 F-FCBBF3000883 554 bp A 179 C 121 G 113 T 141 N 0 (0%)
 F-NT2R2011307 295 bp A 69 C 79 G 78 T 68 N 1 (0.3%)
 ENST00000374754 Cullin-2 (CUL2-2).
 F-NT2R2024160 537 bp A 142 C 114 G 141 T 140 N 0 (0%)
 F-FEBRA2004209 562 bp A 145 C 122 G 151 T 144 N 0 (0%)
 F-BRACE3014507 562 bp A 145 C 122 G 151 T 144 N 0 (0%)
 F-TKIDN2015572 569 bp A 148 C 123 G 152 T 146 N 0 (0%)

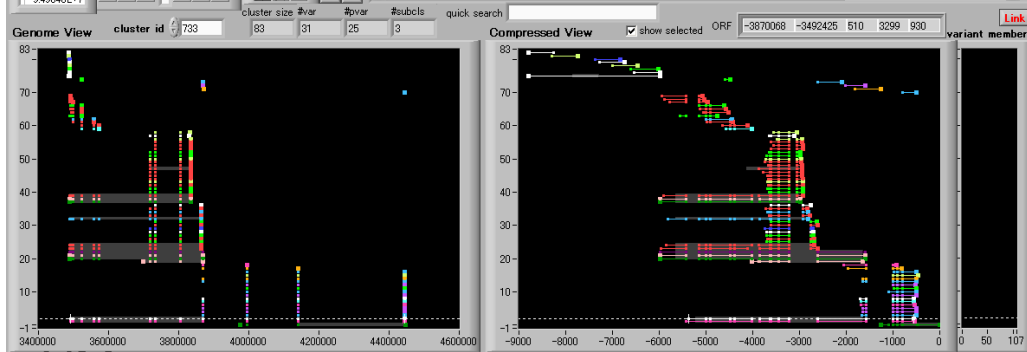
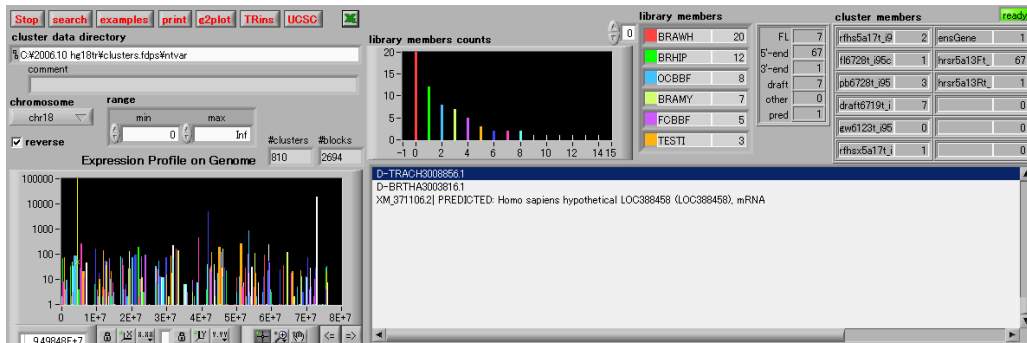


NM_003591.2



FLJ37898

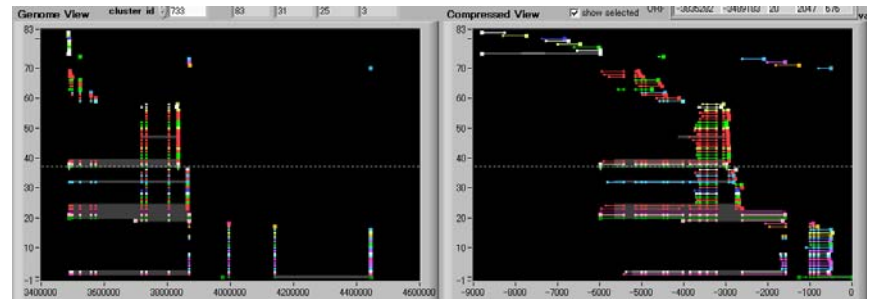
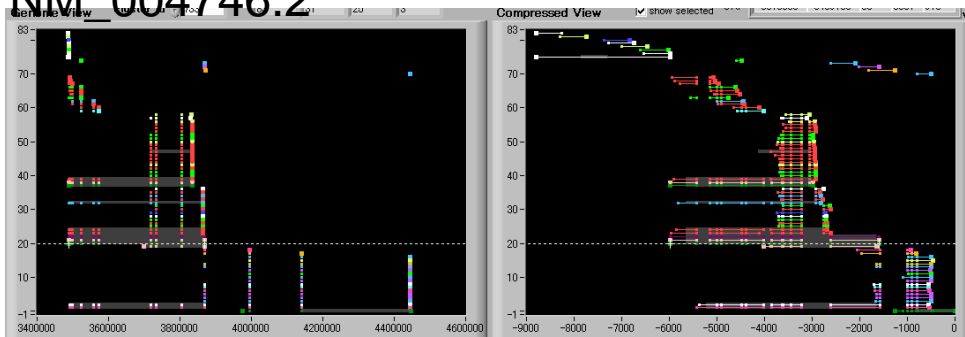
CUL2_NM_003591.2,
FLJ37898



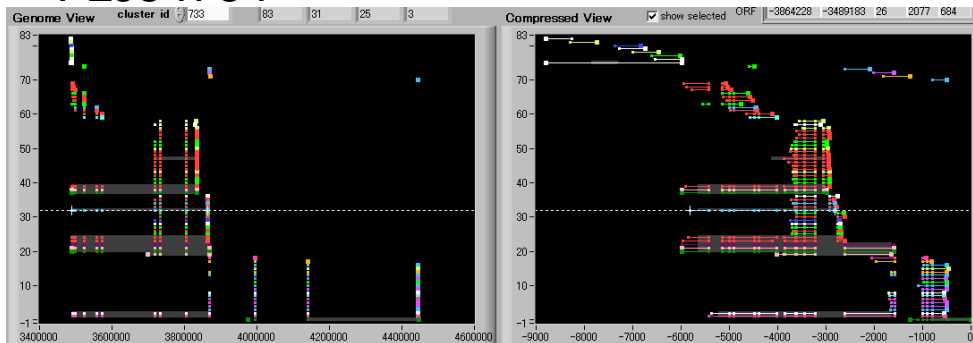
FLJ56525

NM_004746.2

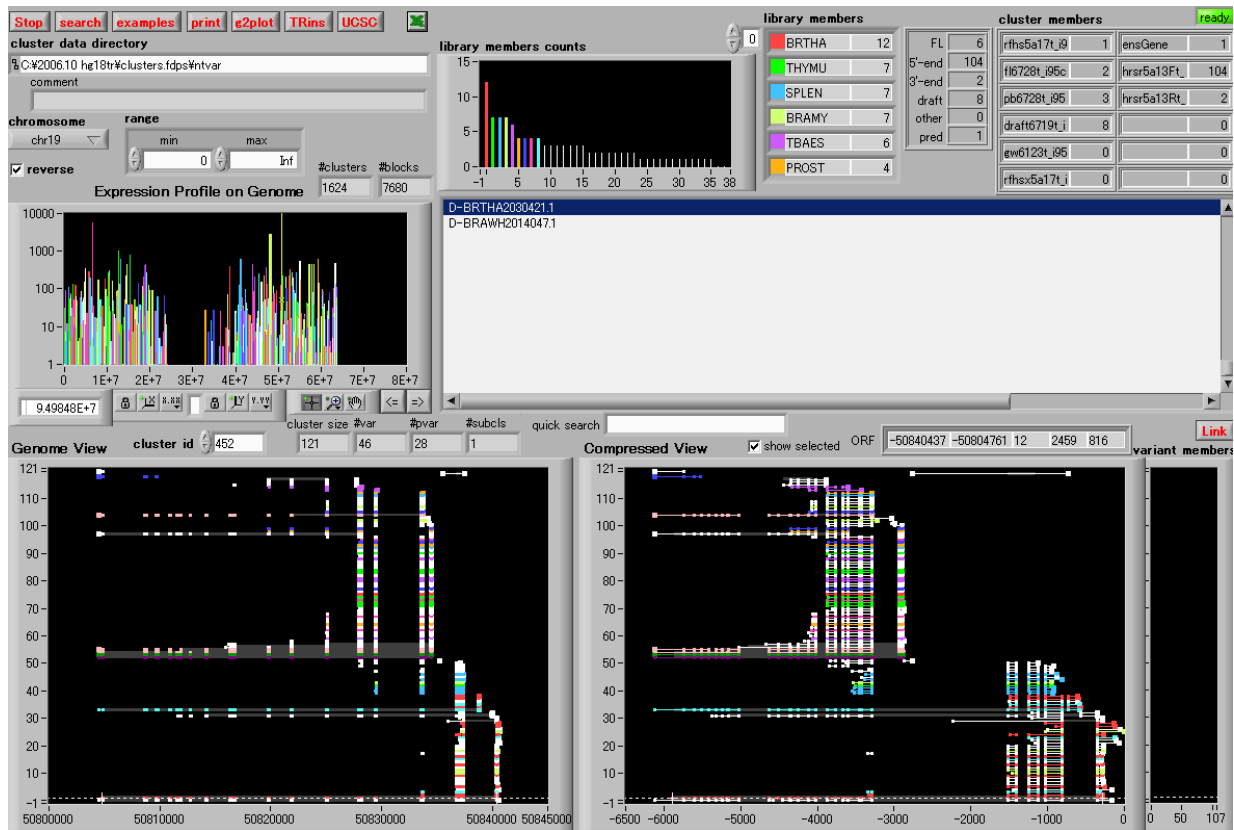
NM_001003809.1



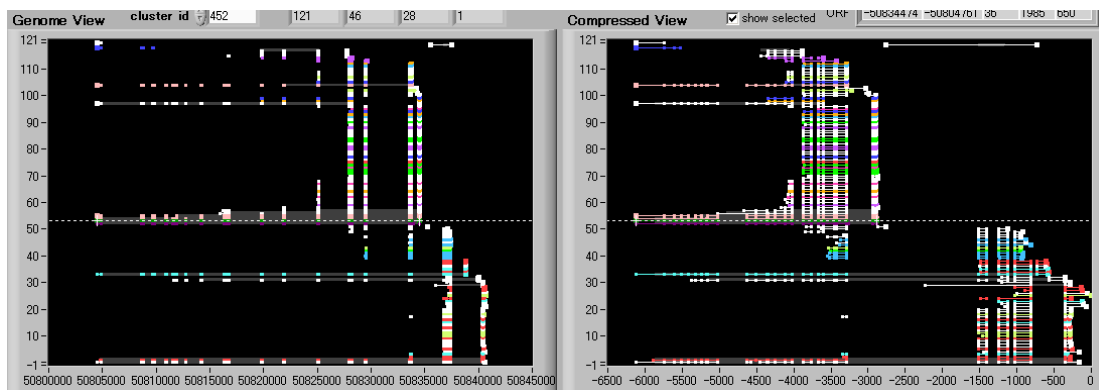
FLJ54734



DLGAP1_FLJ56525,
NM_004746.2,
FLJ54734,
NM_001003809.1

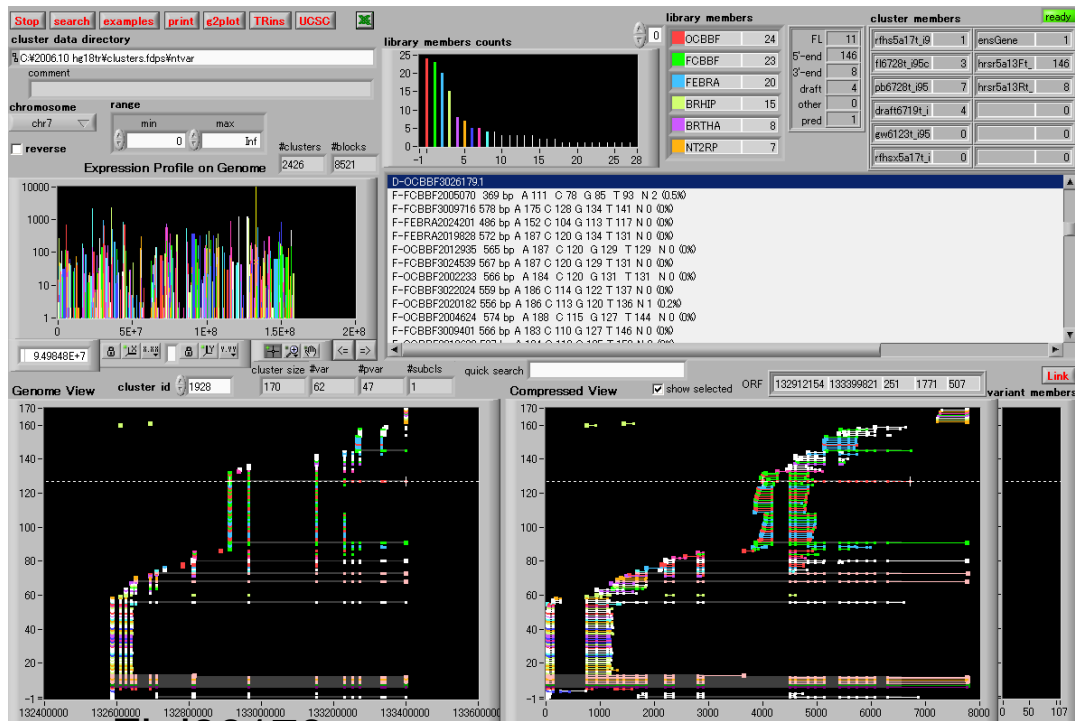


FLJ56452



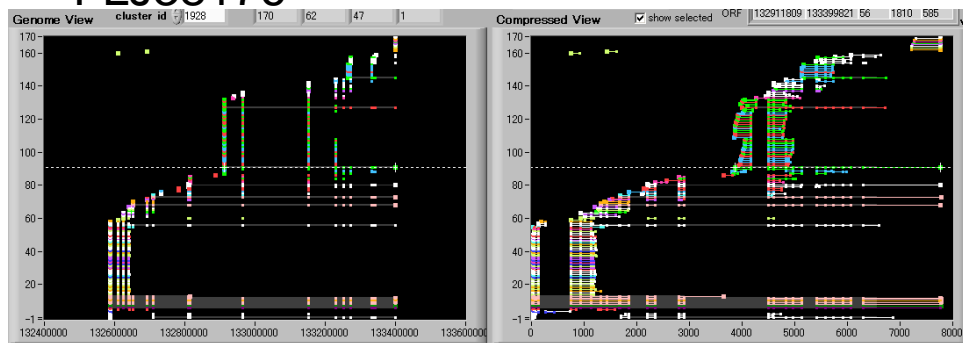
NM_012155.1

EML2_FLJ56452,
 NM_012155.1

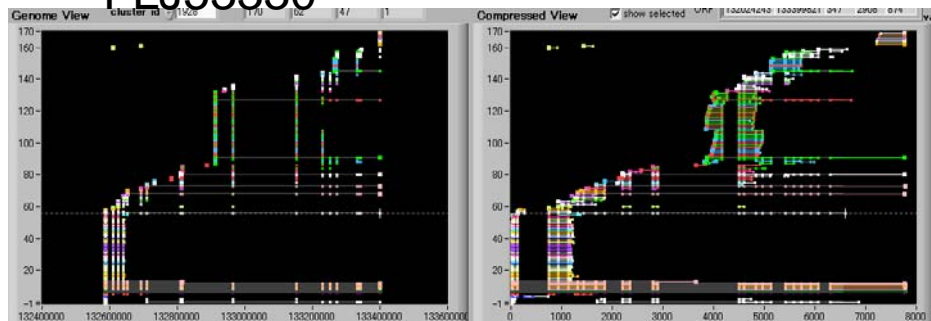
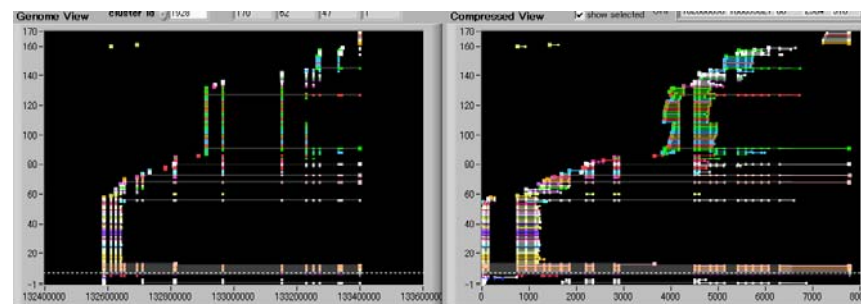


FLJ54541

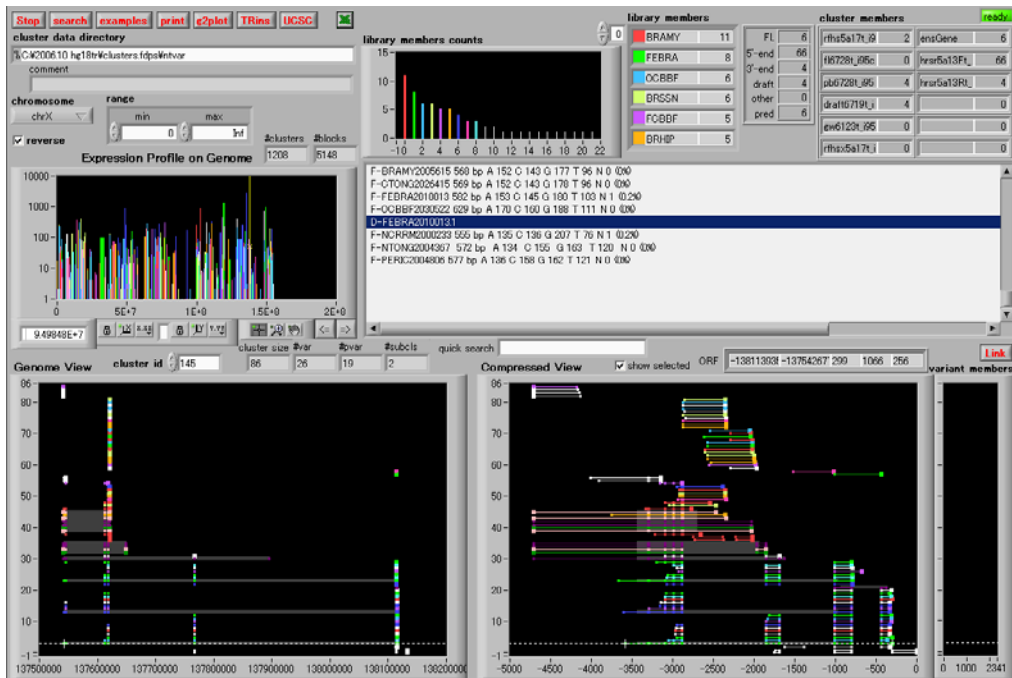
NM_021807.2



FLJ53330

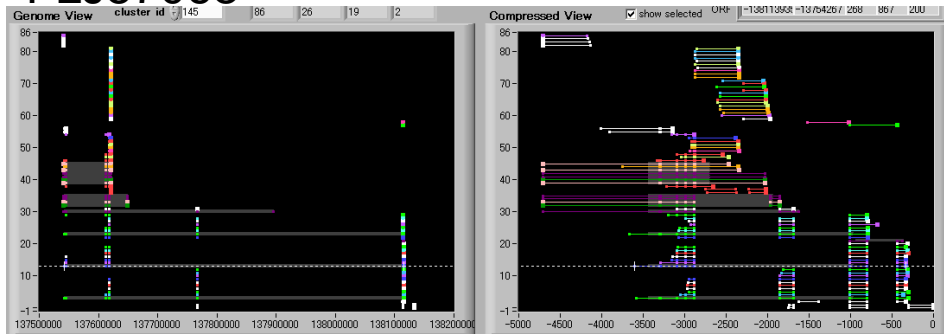


EXOC4_FLJ54541,
FLJ38176, FLJ53330,
NM_021807.2

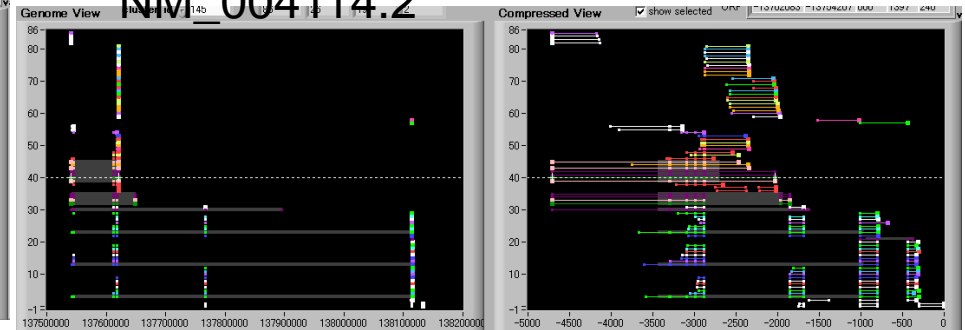


FLJ57884

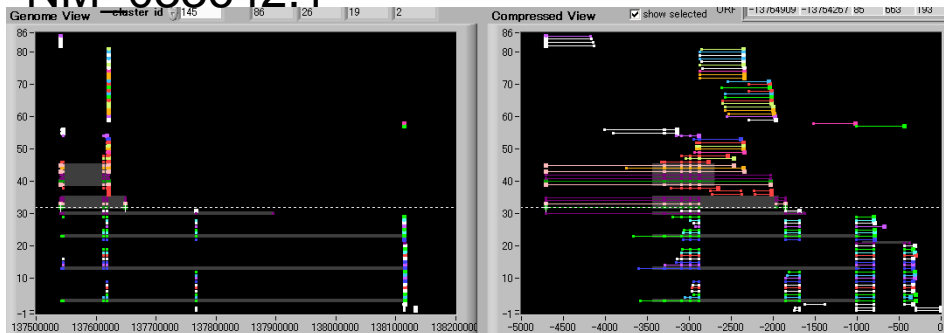
FLJ57068



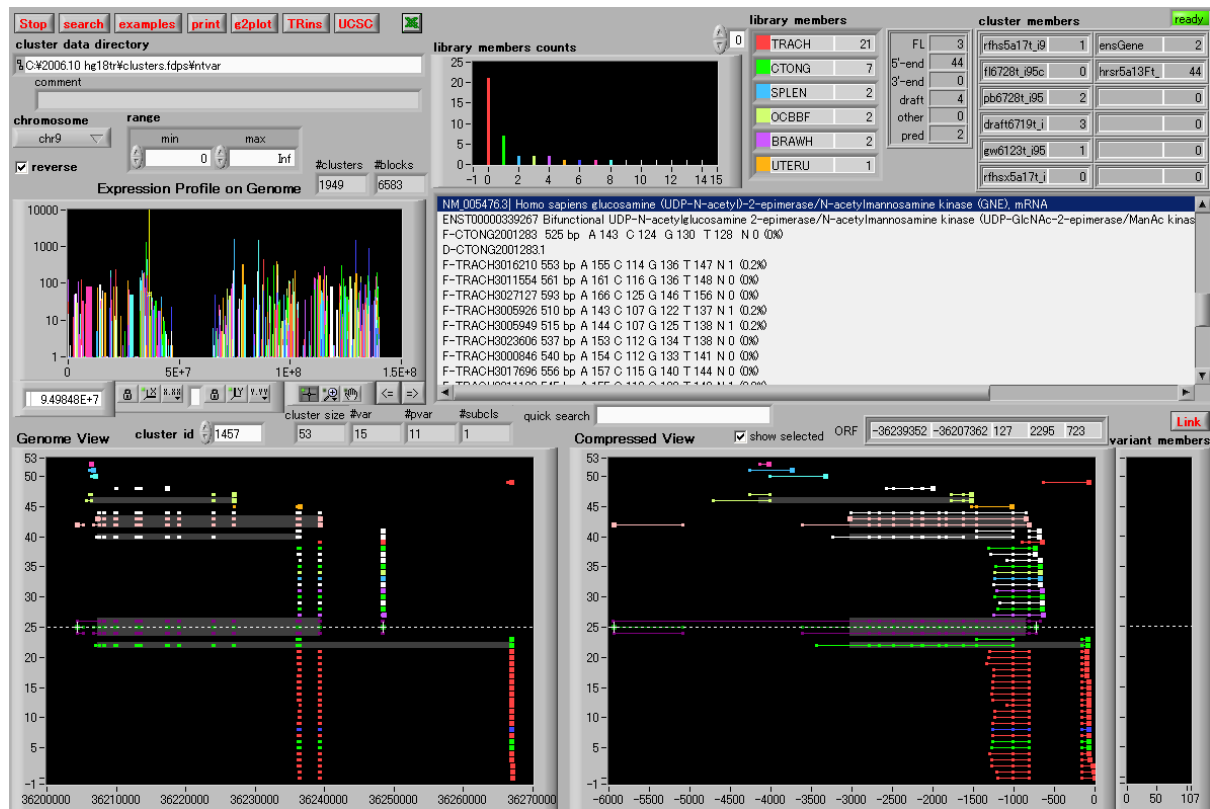
NM_004114.2



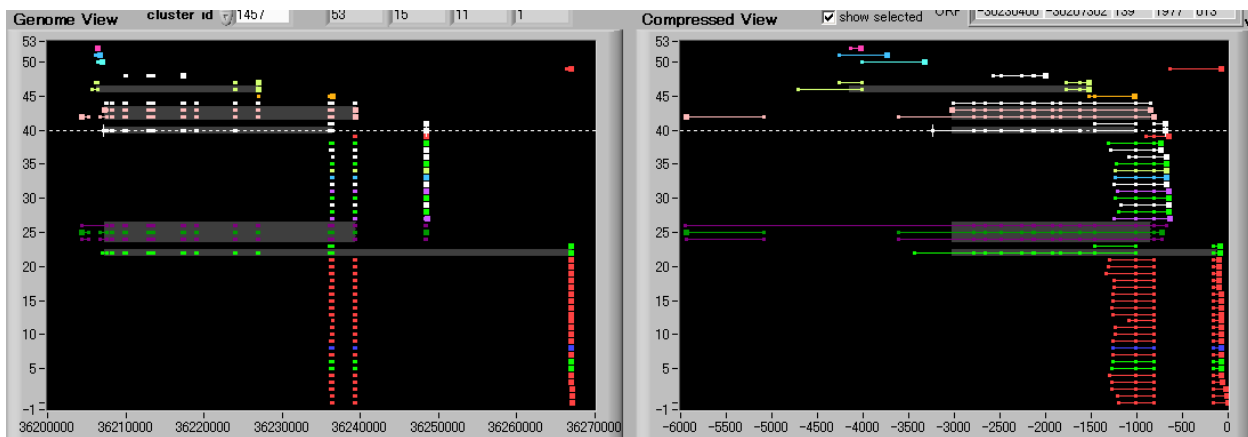
NM_033642.1



FGF13_FLJ57884,
FLJ57068,
NM_033642.1,
NM_004114.2

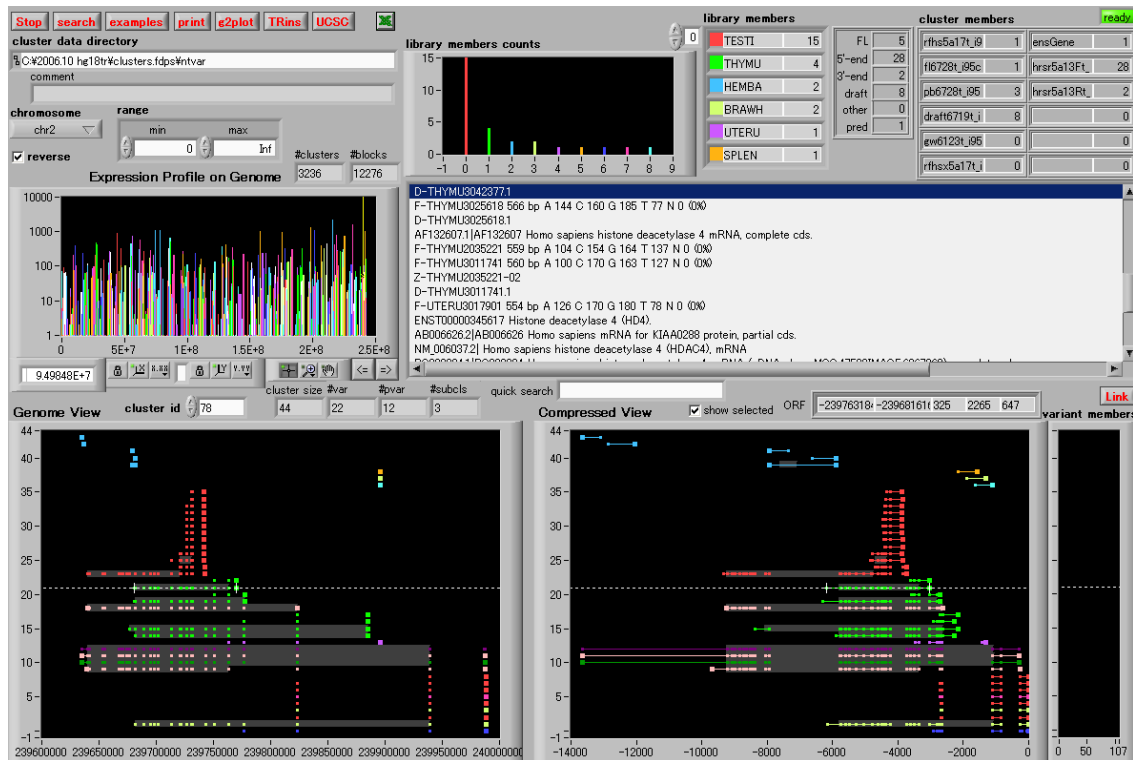


NM_005476.3



FLJ51479

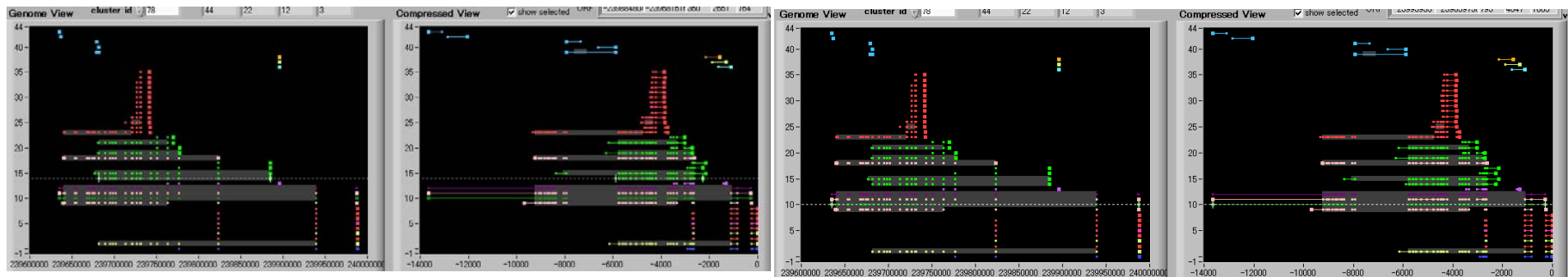
GNE_NM_005476.3,
FLJ51479



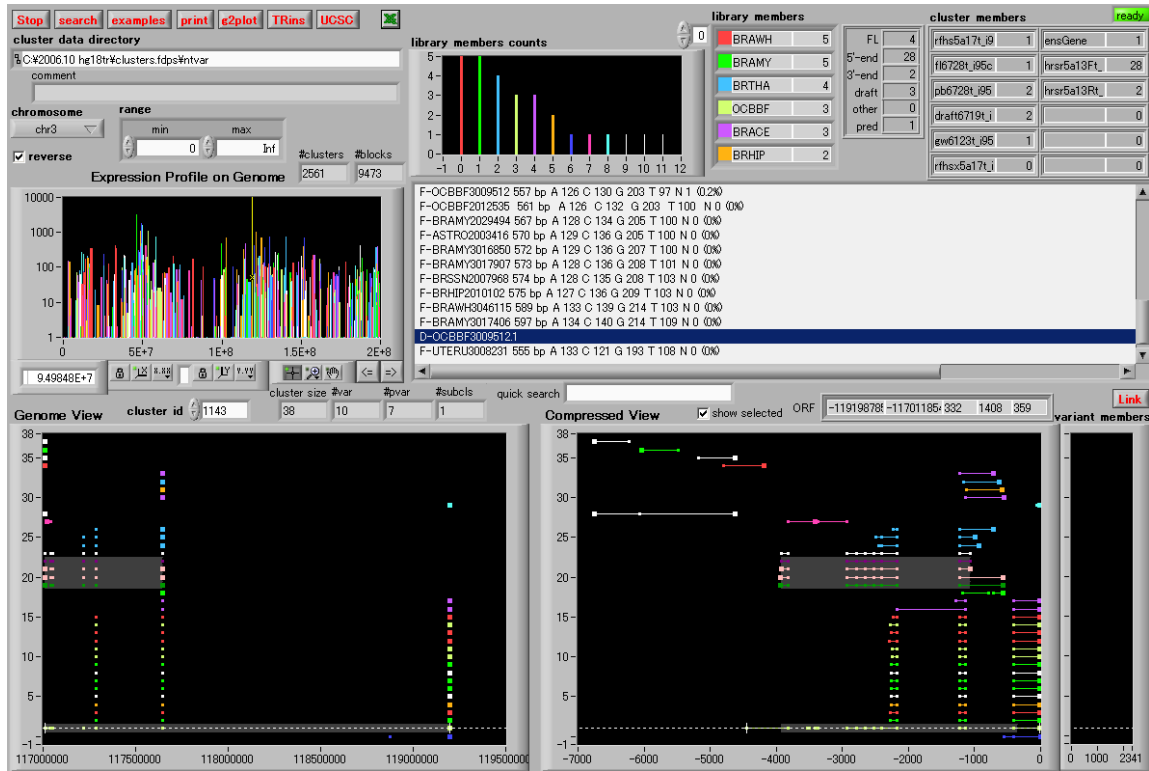
FLJ51177

FLJ51174

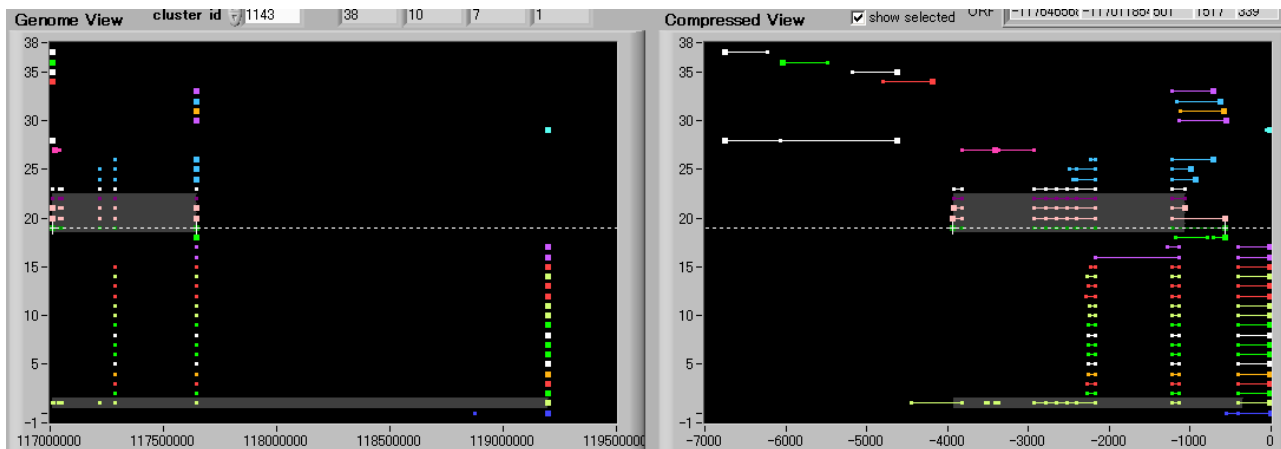
NM_006037.2



HDAC4_FLJ51177,
FLJ51174,
NM_006037.2

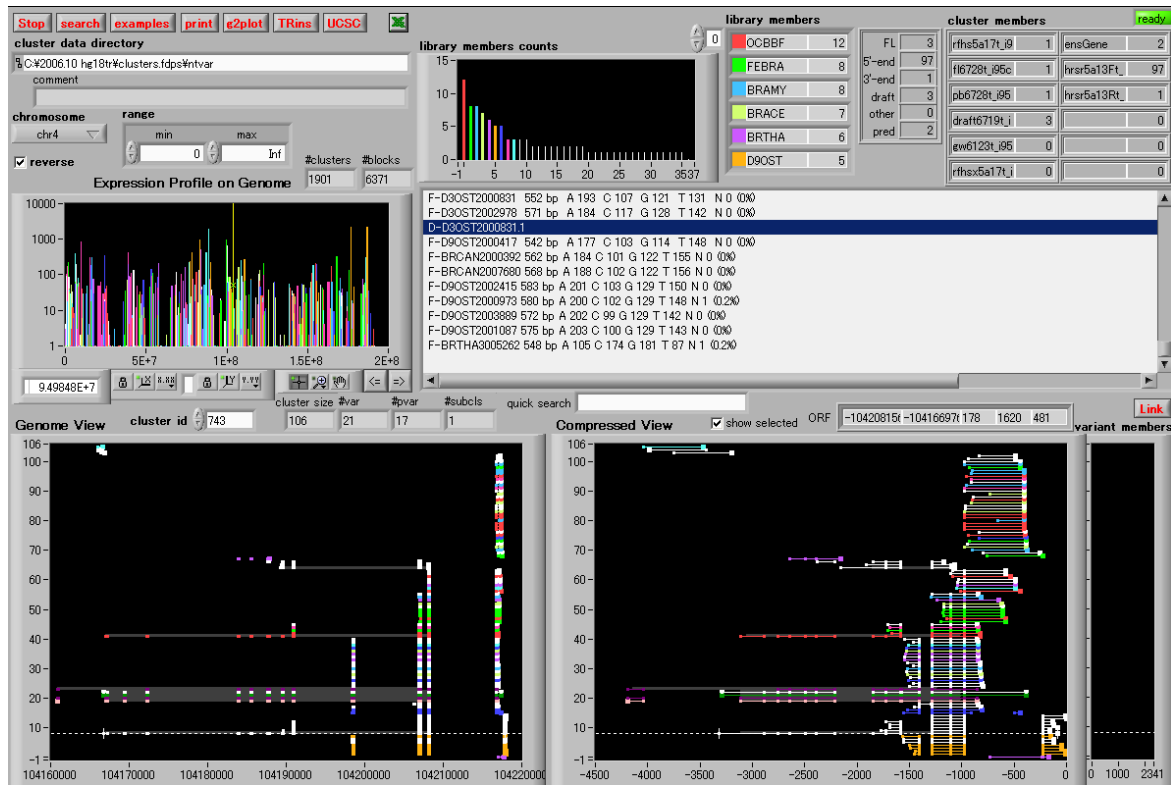


FLJ54658

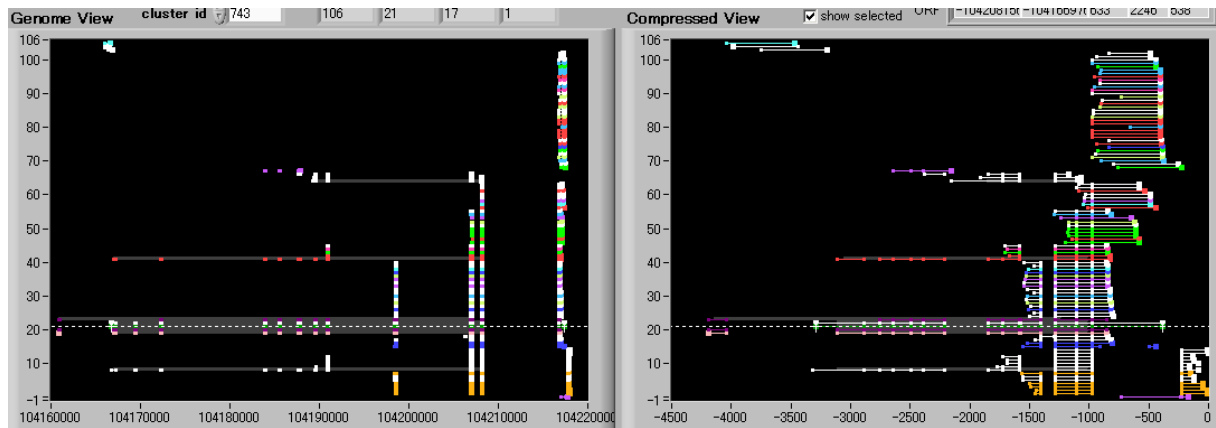


NM_002338.2

LSAMP_FLJ54658,
 NM_002338.2

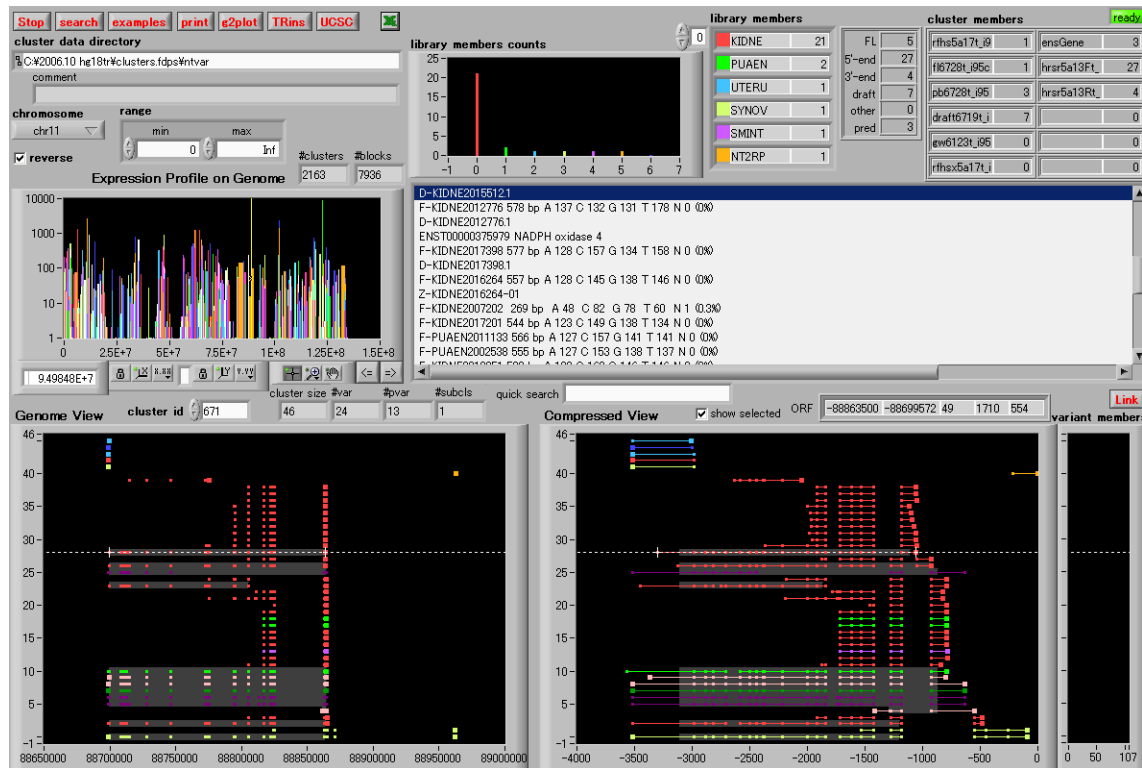


FLJ54331



NM_178833.3

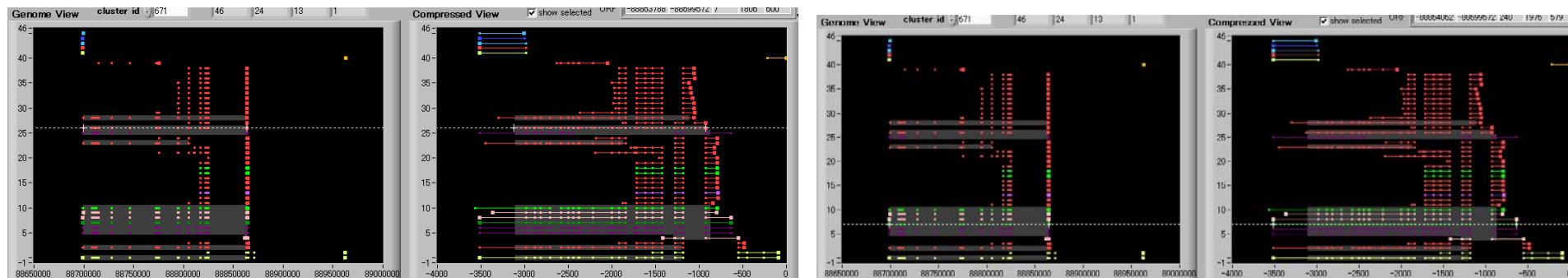
NHEDC2_FLJ54331,
NM_178833.3



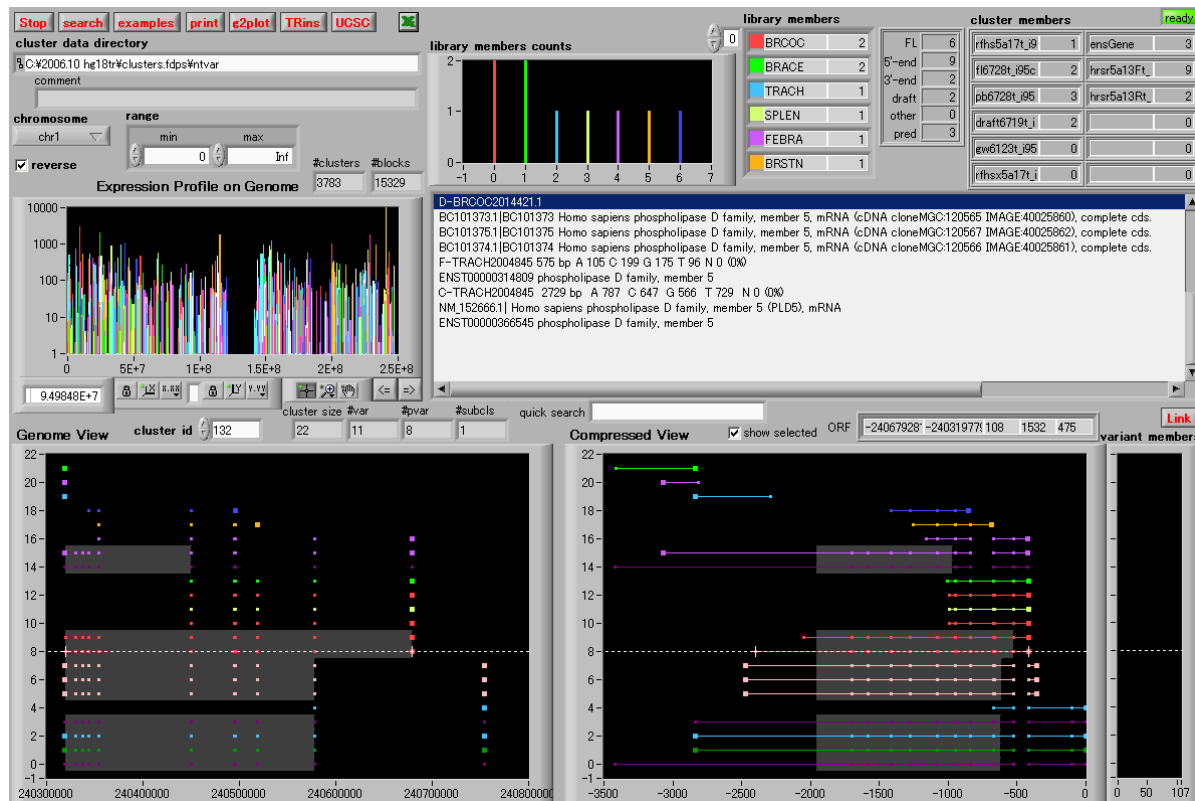
FLJ51027

FLJ51025

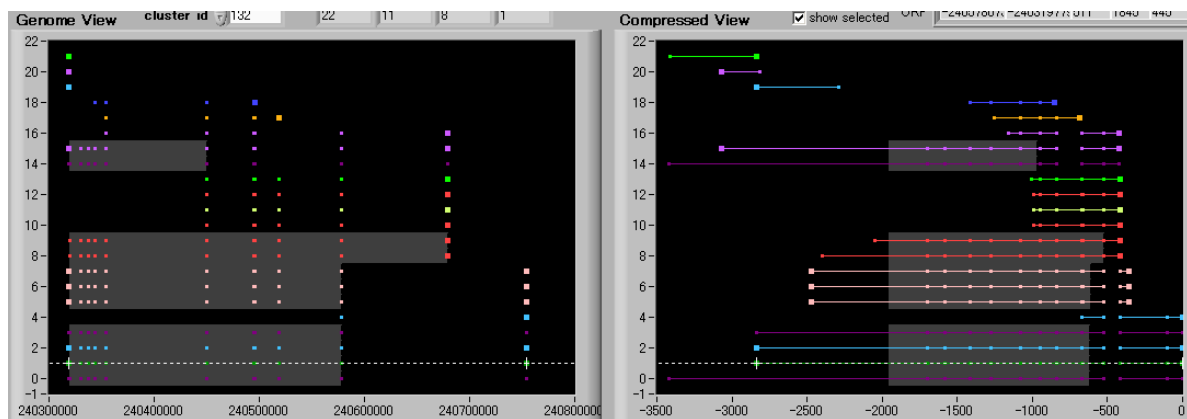
NM_016931.2



NOX4_FLJ51027,
FLJ51025,
NM_016931.2

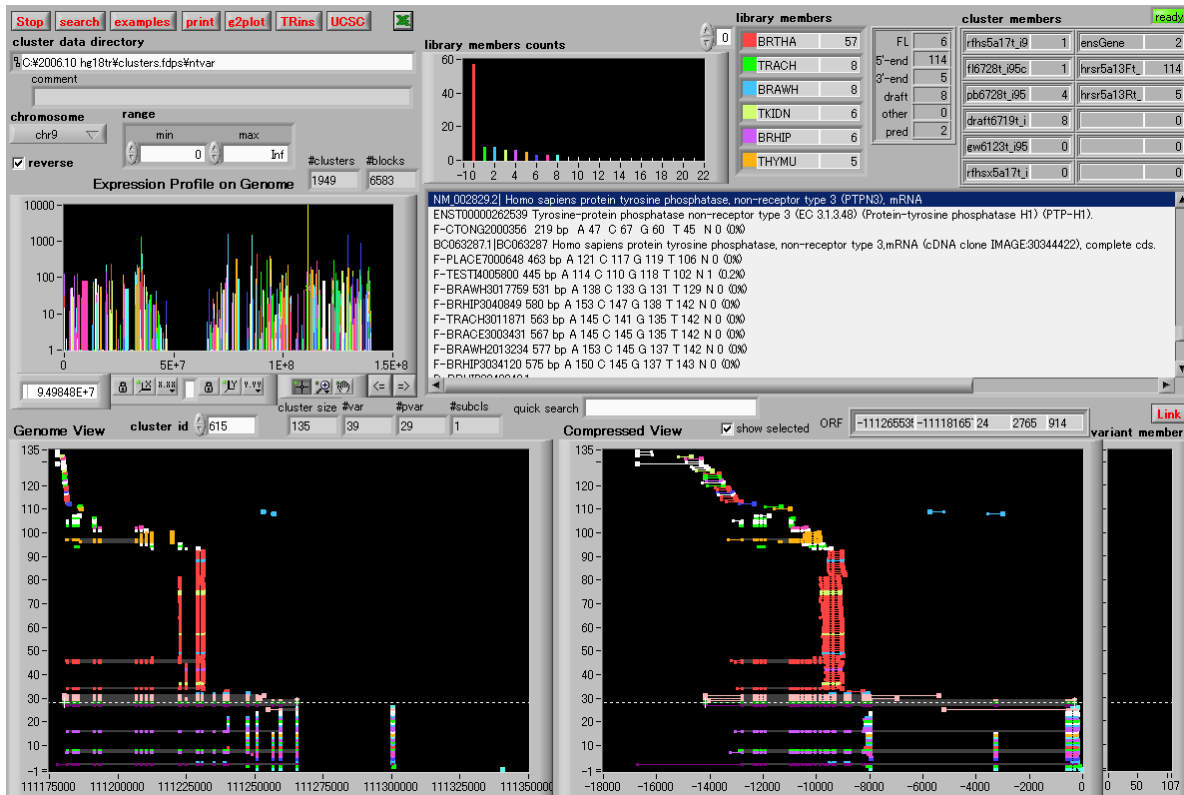


FLJ57051

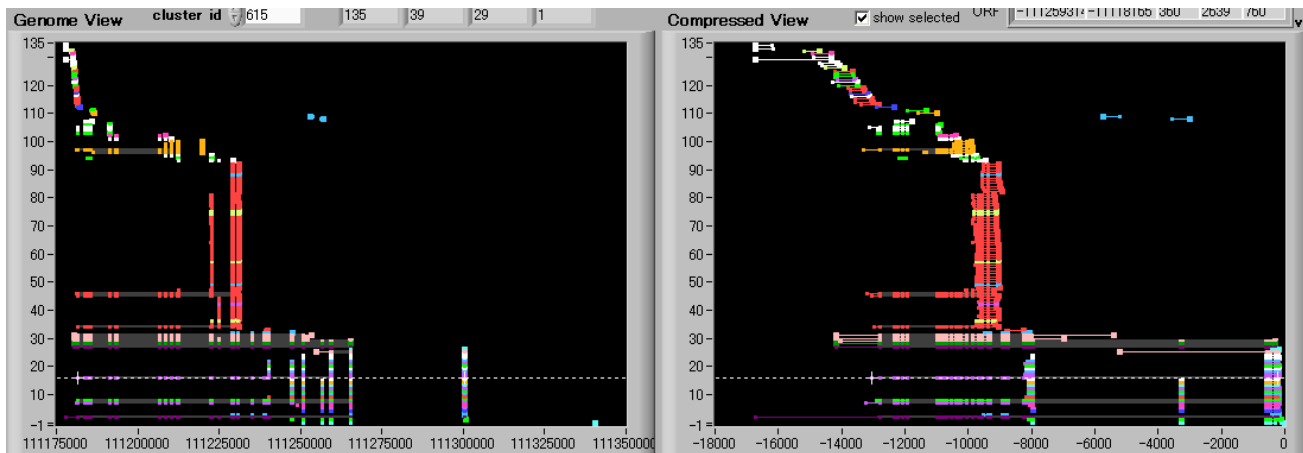


NM_152666.1

PLD5_FLJ57051,
NM_152666.1

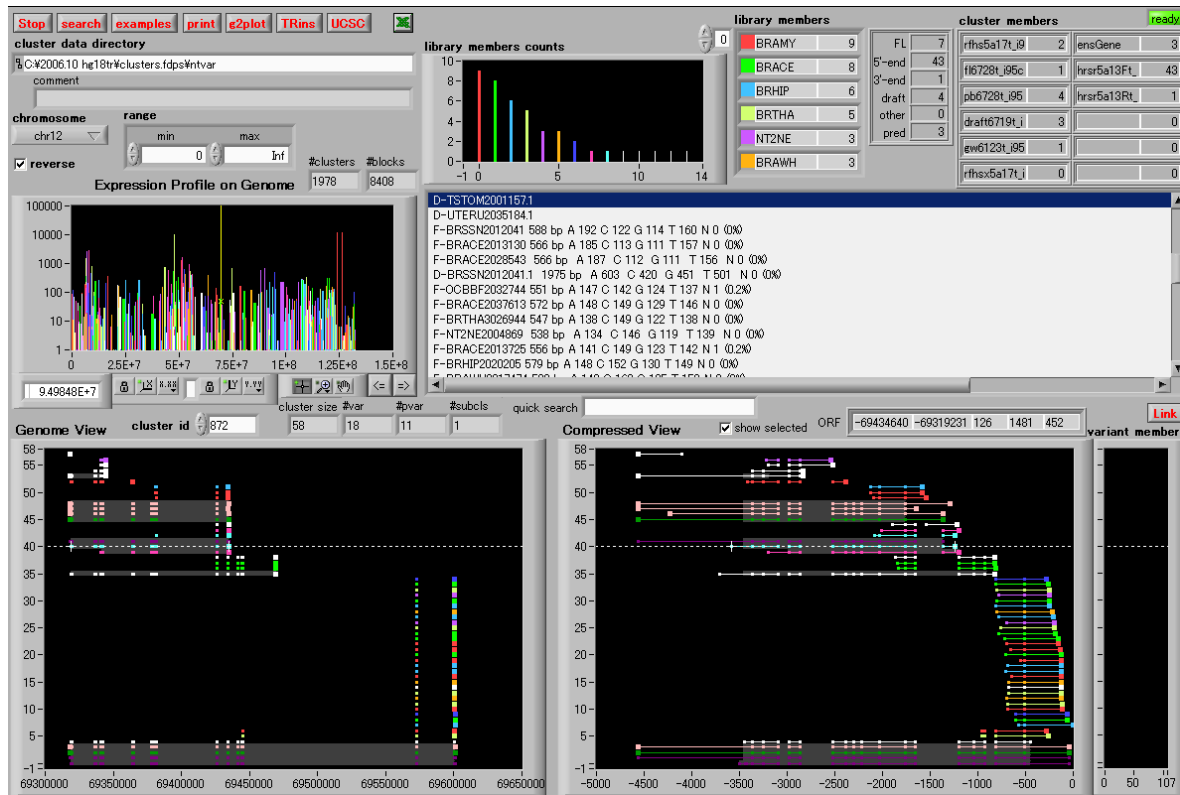


NM_002829.2

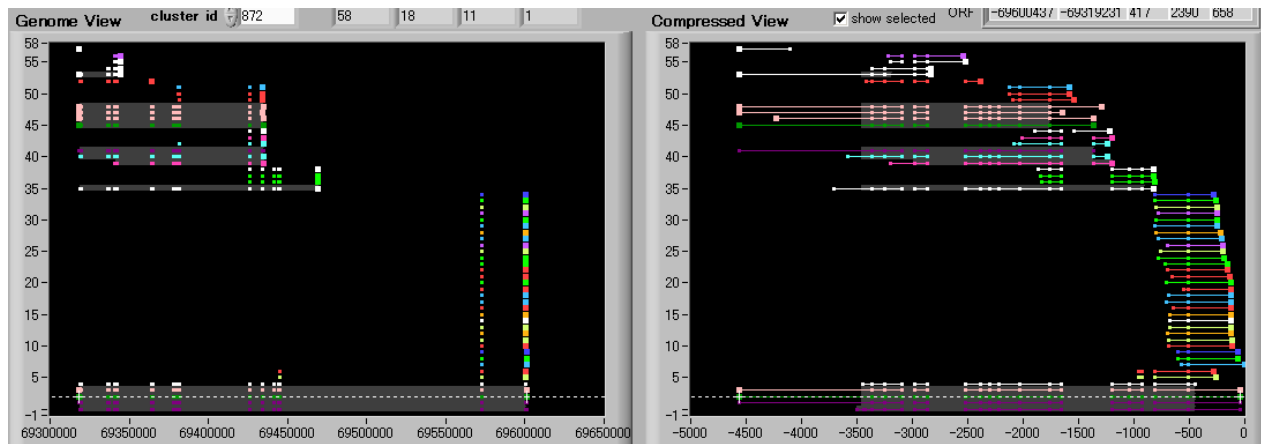


FLJ50545

PTPN3_NM_002829.2,
FLJ50545

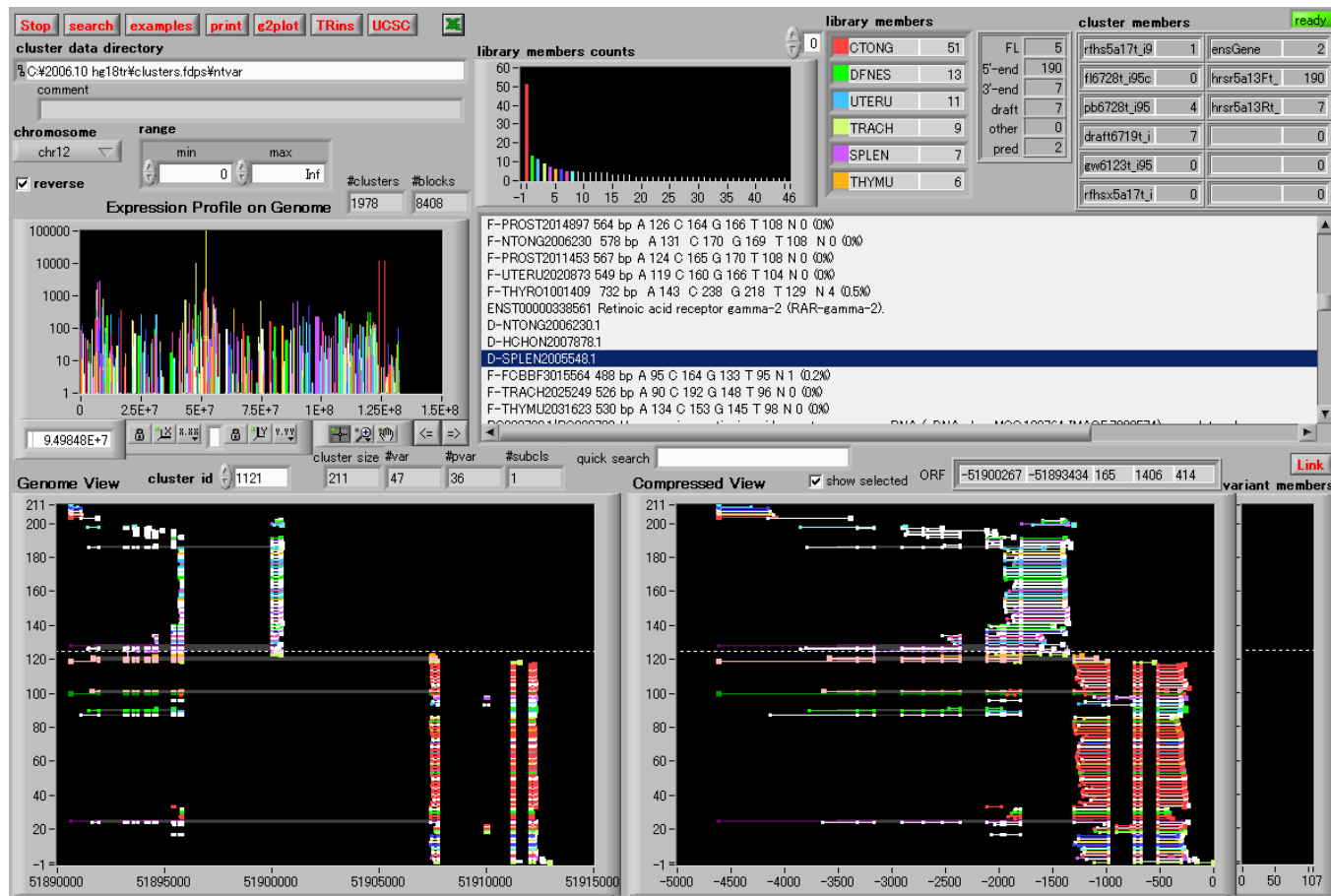


FLJ55167



NM_002849.2

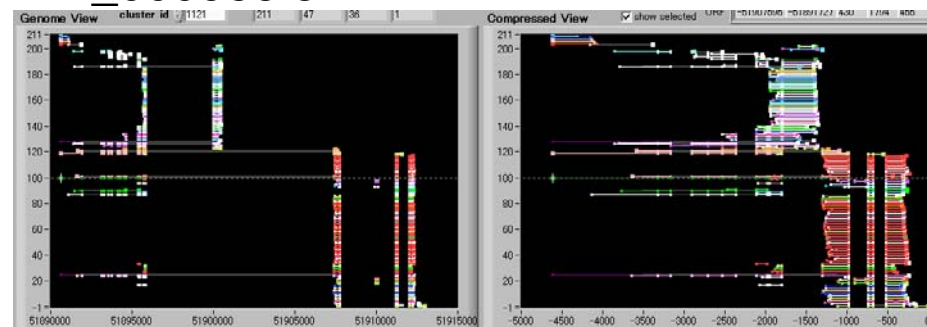
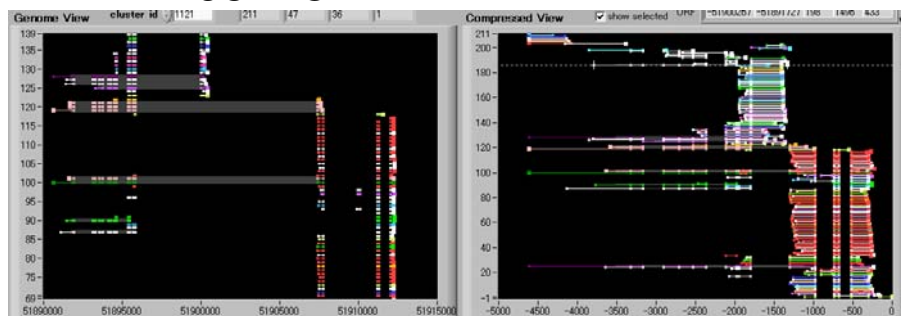
PTPRR_FLJ55167,
NM_002849.2



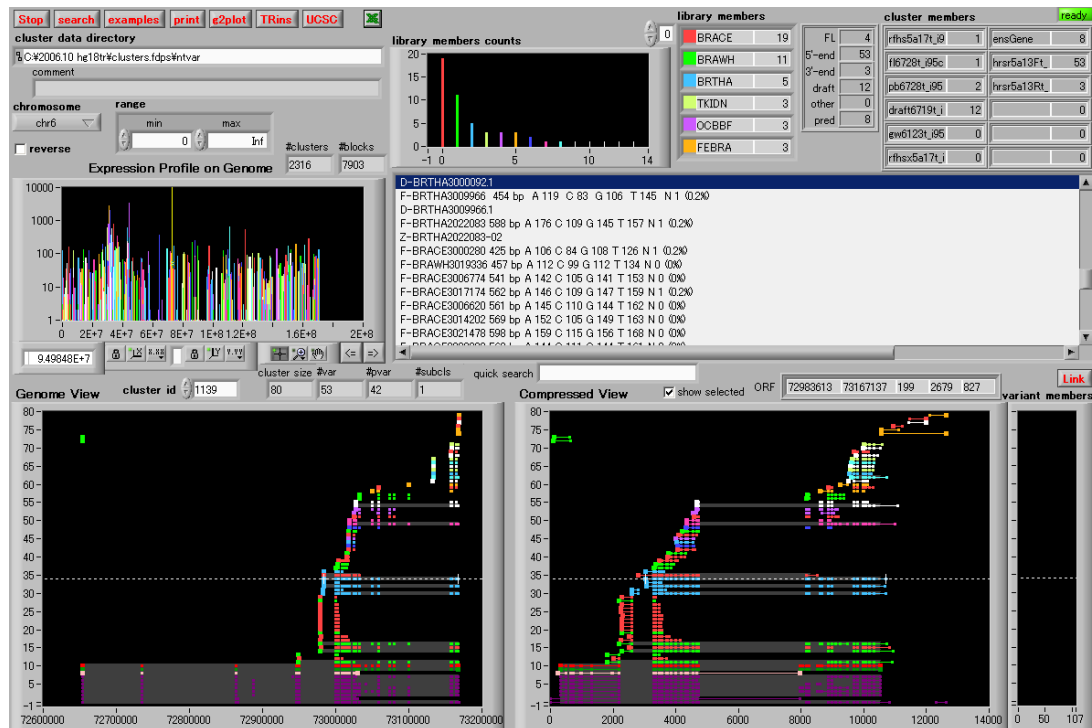
FLJ54463

FLJ54482

NM_000966.3



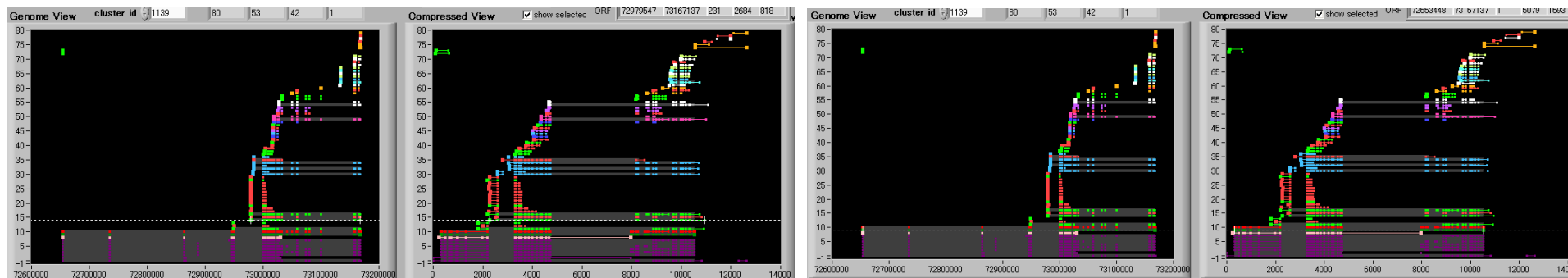
RARG_FLJ54463,
FLJ54482,
NM_000966.3



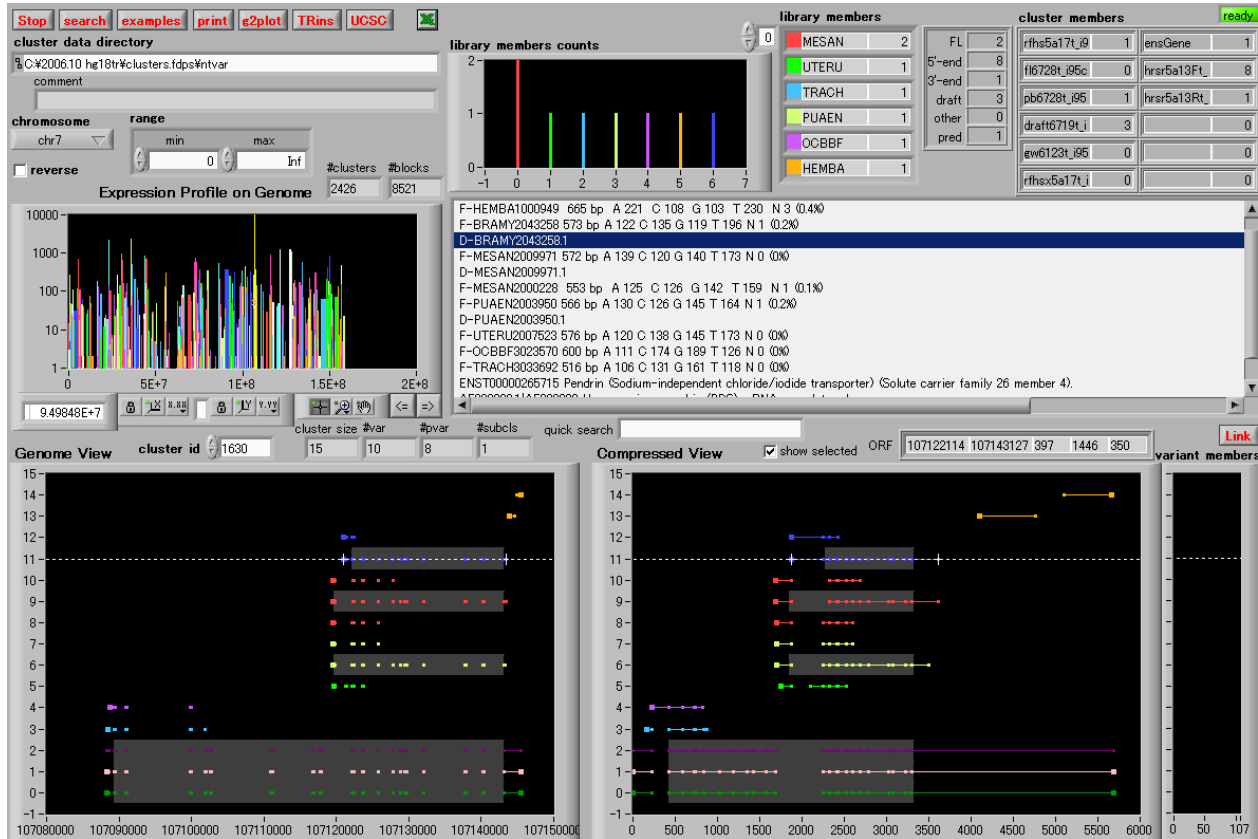
FLJ52438

FLJ53578

NM_014989.2



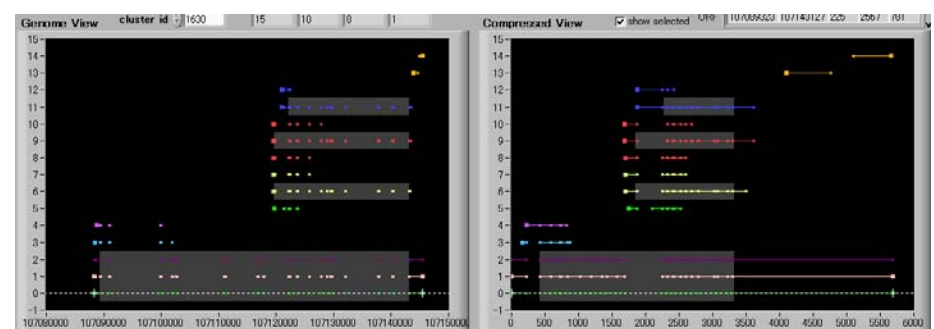
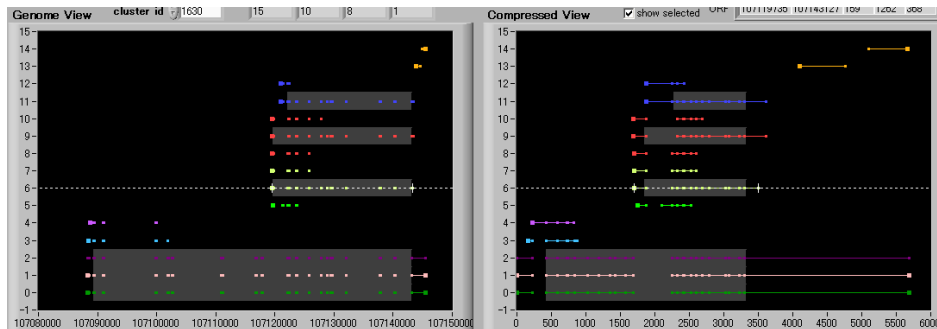
RIMS1_FLJ52438,
FLJ53578,
NM_014989.2



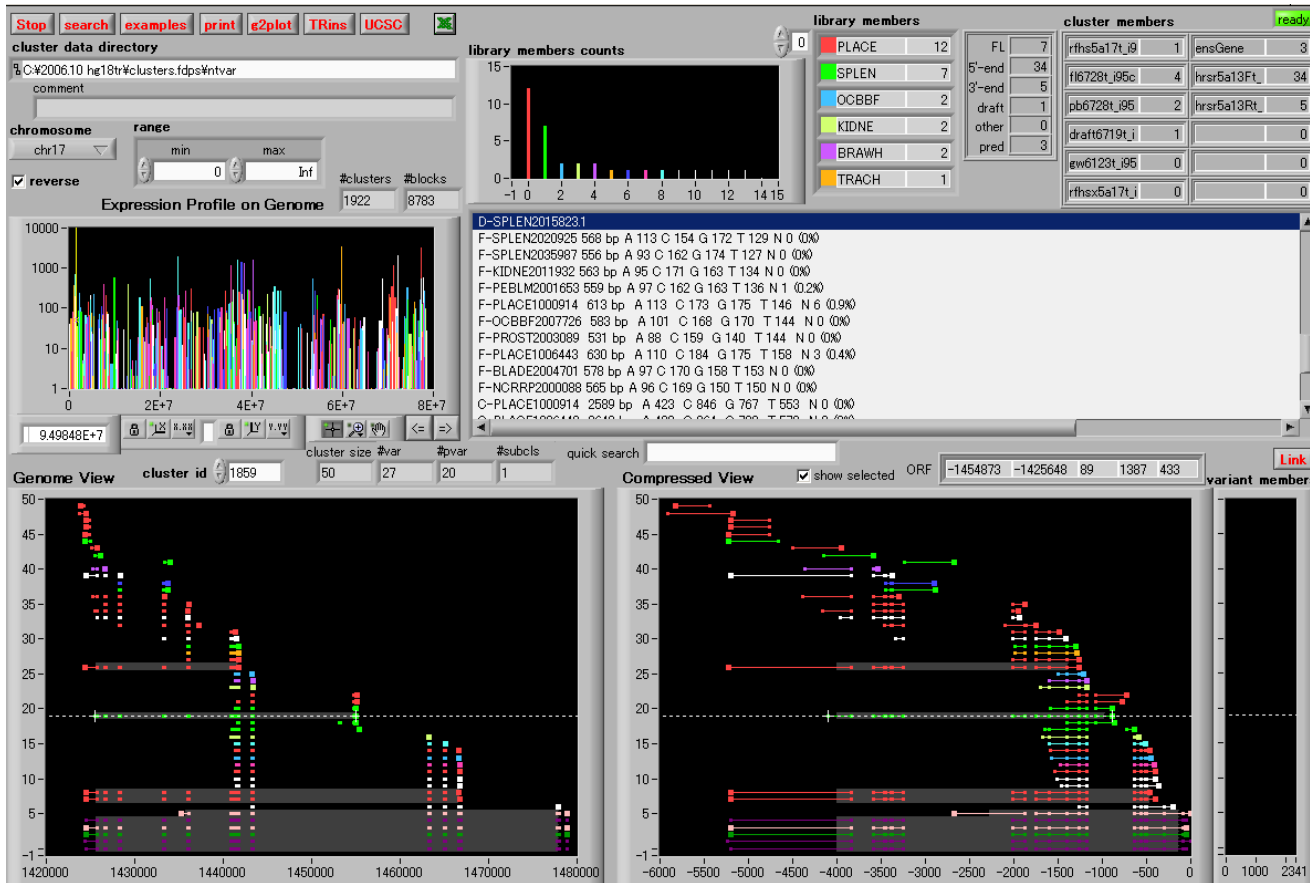
FLJ50484

FLJ50684

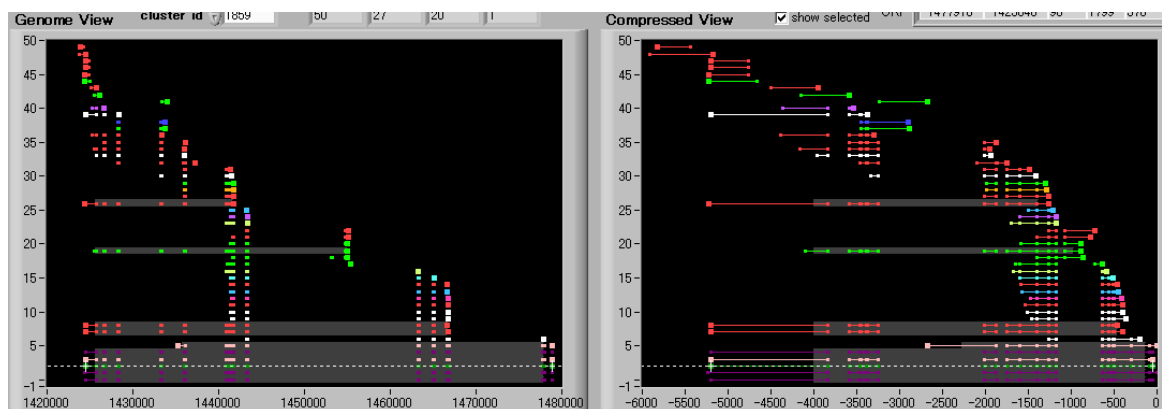
NM_000441.1



SLC26A4_FLJ50484,
FLJ50684,
NM_000441.1

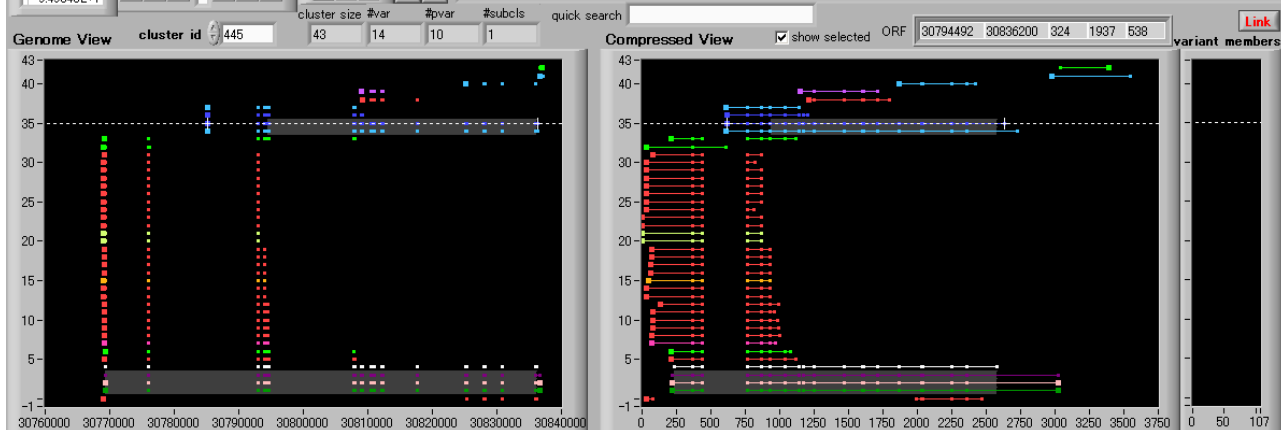
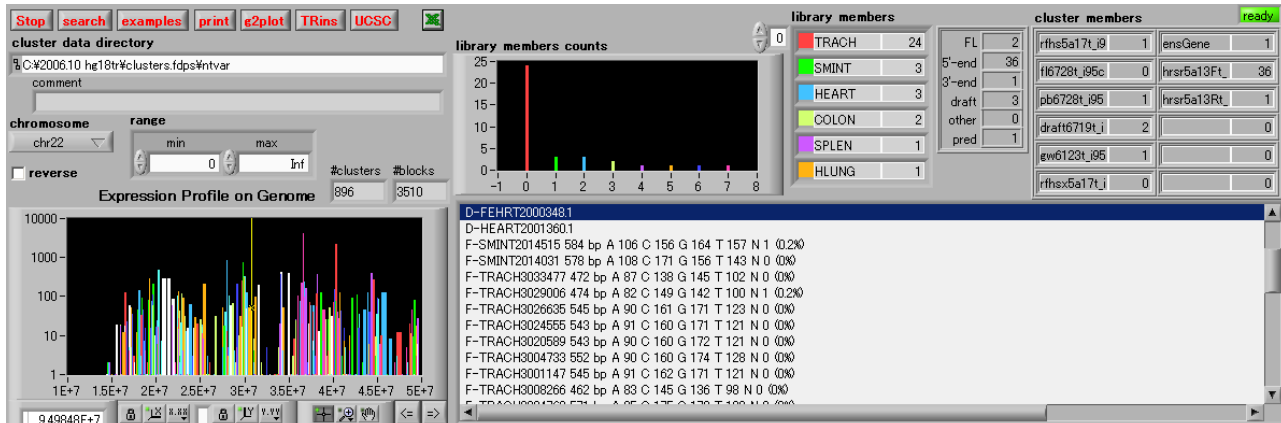


FLJ55865

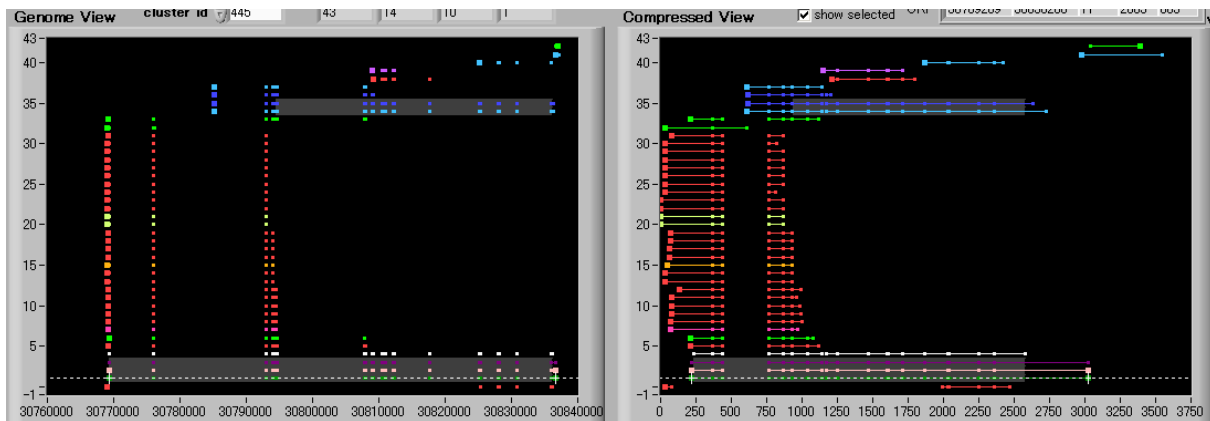


NM_152346.1

SLC43A2_FLJ55865,
 NM_152346.1



FLJ55281



NM_000343.1

SLC5A1_FLJ55281,
 NM_000343.1