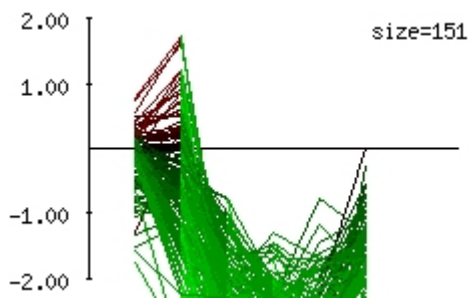
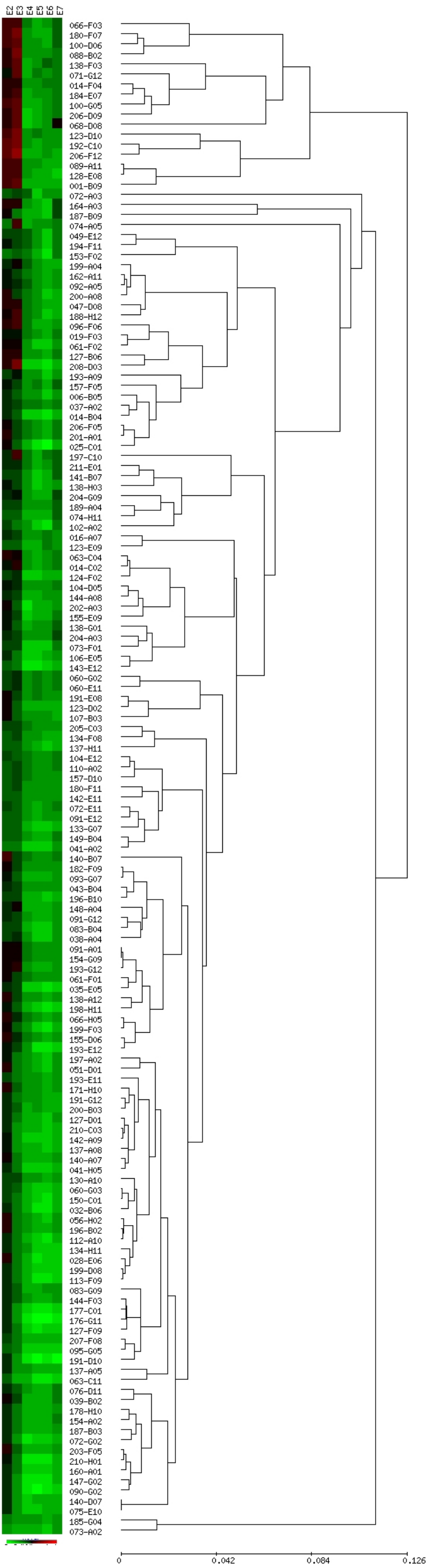


Subtree	Cluster size is 151, joined at the distance 0.125960				1:2			
Spot size:	Width:	Autom.	Height:	Autom.	Tree height:	Automatic	Show titles:	Redraw!
	Autom.							
Colors	Colors	NA color	Number of slots	Discretisation type				





066-F03 098-B11 Unknown

180-F07 107-H08 gi|13384684|ref|NP_079594.1| RIKEN cDNA 0610009E20 gene [Mus musculus]

>gi|13775220|ref|NP_112588.1| hypothetical protein MGC2963 [Homo sapiens]

>gi|14774367|ref|XP_040662.1| hypothetical protein MGC2963 [Homo sapiens]

>gi|12655115|gb|AAH01409.1|AAH01409 (BC001409) Similar to CG11781 gene product [Homo sapiens]

>gi|12833716|dbj|BAB22636.1| (AK003198) putative [Mus musculus] >gi|12835224|dbj|BAB23194.1|

(AK004149) putative [Mus musculus]

100-D06 100-D06 Unknown

088-B02 081-F02 Unknown

138-F03 138-F03 Unknown

071-G12 084-F07 Unknown

014-F04 163-A05 gi|7303345|gb|AAF58404.1| (AE003819) CG4643 gene product [Drosophila melanogaster]

184-E07 184-E07 Unknown

100-G05 100-G05 Unknown

206-D09 178-H07 Unknown

068-D08 068-D08 Unknown

123-D10 080-D06 Unknown

192-C10 192-C10 Unknown

206-F12 201-C03 Unknown

089-A11 061-F05 Unknown

128-E08 128-E08 Unknown

001-B09 001-B09 Unknown

072-A03 072-A03 Unknown

164-A03 164-A03 Unknown

187-B09 187-B09 Unknown

074-A05 074-A05 Unknown

049-E12 049-E12 gi|6841174|gb|AAF28940.1|AF161380_1 (AF161380) HSPC262 [Homo sapiens]

194-F11 079-H03 Unknown

153-F02 079-F09 gi|463046|gb|AAA49207.1| (L30111) cysteine proteinase [Cyprinus carpio]

199-A04 199-A04 Unknown

162-A11 162-A11 Unknown

092-A05 092-A05 Unknown

200-A08 200-A08 Unknown

047-D08 044-A09 Unknown

188-H12 188-H12 Unknown

096-F06 096-F06 Unknown

019-F03 019-F03 gi|13242156|gb|AAK16577.1| (AF331967) egg envelope protein ZP2 variant B [Danio rerio]

061-F02 061-F02 gi|2208895|dbj|BAA20516.1| (AB004740) beta-globin [Cyprinus carpio]

127-B06 127-B06 Unknown

208-D03 084-G05 Unknown

193-A09 158-D01 gi|8395614|ref|NP_059333.1| cytochrome c oxidase subunit I [Danio rerio]

>gi|8272688|gb|AAF74299.1|AC024175_3 (AC024175) cytochrome c oxidase subunit I [Danio rerio]

157-F05 157-F05 Unknown

006-B05 006-B05 Unknown

037-A02 132-C01 gi|15426528|gb|AAH13375.1|AAH13375 (BC013375) Unknown (protein for MGC:16436) [Homo sapiens]
014-B04 014-B04 Unknown
206-F05 206-F05 Unknown
201-A01 201-A01 Unknown
025-C01 025-C01 gi|12667276|gb|AAK01371.1| (AF315729) histone H2A [Carassius auratus]
197-C10 197-C10 Unknown
211-E01 211-E01 Unknown
141-B07 141-B07 Unknown
138-H03 170-B04 Unknown
204-G09 204-G09 Unknown
189-A04 189-A04 Unknown
074-H11 074-H11 Unknown
102-A02 102-A02 Unknown
016-A07 016-A07 Unknown
123-E09 151-C01 "gi|14744179|ref|XP_050764.1| COX15 (yeast) homolog, cytochrome c oxidase assembly protein [Homo sapiens] >gi|14744187|ref|XP_050763.1| COX15 (yeast) homolog, cytochrome c oxidase assembly protein [Homo sapiens] >gi|7619702|emb|CAB88198.1| (AL133353) bA483F11.2.1 (COX15 (yeast) homolog, cytochrome c oxidase assembly protein (isoform 1)) [Homo sapiens] >gi|12803157|gb|AAH02382.1|AAH02382 (BC002382) COX15 (yeast) homolog, cytochrome c oxidase assembly protein [Homo sapiens] >gi|15426569|gb|AAH13403.1|AAH13403 (BC013403) Unknown (protein for MGC:4234) [Homo sapiens]"
063-C04 063-C04 Unknown
014-C02 014-C02 Unknown
124-F02 124-F02 Unknown
104-D05 104-D05 Unknown
144-A08 144-A08 Unknown
202-A03 202-A03 Unknown
155-E09 155-E09 Unknown
138-G01 157-E11 Unknown
204-A03 204-A03 Unknown
073-F01 206-G08 Unknown
106-E05 106-E05 "gi|7512238|pir||JE0366 tight junction protein, ZO-2 - chicken >gi|3820580|gb|AAC95469.1| (AF085184) tight junction protein [Gallus gallus]"
143-E12 143-E12 Unknown
060-G02 060-G02 gi|13359451|dbj|BAB33421.1| (AB049723) putative senescence-associated protein [Pisum sativum]
060-E11 060-E11 gi|13435438|gb|AAH04587.1|AAH04587 (BC004587) Unknown (protein for IMAGE:3587950) [Mus musculus]
191-E08 191-E08 Unknown
123-D02 123-D02 Unknown
107-B03 059-C02 gi|113726|sp|P00811|AMPC_ECOLI BETA-LACTAMASE PRECURSOR (CEPHALOSPORINASE) >gi|67775|pir||QKEC beta-lactamase (EC 3.5.2.6) precursor - Escherichia coli >gi|145267|gb|AAA23441.1| (J01611) beta-lactamase [Escherichia coli] >gi|536994|gb|AAA97049.1| (U14003) beta-lactamase [Escherichia coli] >gi|1790593|gb|AAC77110.1| (AE000487) beta-lactamase; penicillin resistance [Escherichia coli K12]

205-C03 205-C03 Unknown
134-F08 134-F08 Unknown
137-H11 137-H11 Unknown
104-E12 104-E12 gi|135410|sp|P06604|TBA2_DROME TUBULIN ALPHA-2 CHAIN >gi|85173|pir||B26488
tubulin alpha-2 chain - fruit fly (*Drosophila melanogaster*) >gi|158733|gb|AAA28986.1|
(M14644) alpha-tubulin 2 [*Drosophila melanogaster*] >gi|7299237|gb|AAF54433.1| (AE003684)
alphaTub85E gene product [*Drosophila melanogaster*]
110-A02 110-A02 Unknown
157-D10 157-D10 Unknown
180-F11 180-F11 Unknown
142-E11 142-E11 Unknown
072-E11 072-E11 Unknown
091-E12 058-D01 Unknown
133-G07 104-H10 Unknown
149-B04 014-G02 Unknown
041-A02 011-D11 Unknown
140-B07 140-B07 Unknown
182-F09 182-F09 Unknown
093-G07 093-G07 Unknown
043-B04 102-C10 gi|13650899|ref|XP_017654.1| x 009 protein [*Homo sapiens*]
196-B10 196-B10 Unknown
148-A04 148-A04 Unknown
091-G12 092-F11 Unknown
083-B04 083-B04 Unknown
038-A04 087-H08 Unknown
091-A01 091-A01 Unknown
154-G09 154-G09 Unknown
193-G12 192-G12 Unknown
061-F01 061-F01 Unknown
035-E05 021-B02 Unknown
138-A12 138-A12 Unknown
198-H11 007-C10 Unknown
066-H05 163-B09 gi|15801165|ref|NP_287182.1| periplasmic glucans biosynthesis protein
[*Escherichia coli* 0157:H7 EDL933] >gi|15830680|ref|NP_309453.1| periplasmic glucans
biosynthesis protein [*Escherichia coli* 0157:H7] >gi|12514580|gb|AAG55794.1|AE005315_8
(AE005315) periplasmic glucans biosynthesis protein [*Escherichia coli* 0157:H7 EDL933]
>gi|13360890|dbj|BAB34849.1| (AP002555) periplasmic glucans biosynthesis protein
[*Escherichia coli* 0157:H7]
199-F03 199-F03 Unknown
155-D06 155-D06 Unknown
193-E12 024-G12 Unknown
197-A02 197-A02 Unknown
051-D01 051-D01 Unknown
193-E11 154-F11 Unknown
171-H10 119-F12 Unknown
191-G12 191-G12 Unknown
200-B03 200-B03 Unknown

127-D01 176-G02 Unknown
 210-C03 210-C03 Unknown
 142-A09 147-E05 gi|12848186|dbj|BAB27861.1| (AK011820) putative [Mus musculus]
 137-A08 154-G05 Unknown
 140-A07 140-A07 Unknown
 041-H05 041-H05 Unknown
 130-A10 181-A04 Unknown
 060-G03 060-G03 Unknown
 150-C01 150-C01 Unknown
 032-B06 032-B06 Unknown
 056-H02 117-D06 gi|773414|gb|AAB05989.1| (U23751) beta galactosidase [Cloning vector pBBR1MCS-2]
 >gi|833819|gb|AAB06689.1| (U25059) LacZ alpha peptide [Cloning vector pBBR1MCS-3]
 >gi|833823|gb|AAB06692.1| (U25060) LacZ alpha peptide [Cloning vector pBBR1MCS-4]
 >gi|833827|gb|AAB06695.1| (U25061) LacZ alpha peptide [Cloning vector pBBR1MCS-5]
 196-B02 196-B02 Unknown
 112-A10 112-A10 Unknown
 134-H11 134-H11 Unknown
 028-E06 181-D04 Unknown
 199-D08 199-D08 Unknown
 113-F09 113-F09 Unknown
 083-G09 080-F05 Unknown
 144-F03 144-F03 Unknown
 177-C01 177-C01 Unknown
 176-G11 188-A04 Unknown
 127-F09 151-F02 gi|6685321|sp|Q9YH91|CLDY_BRARE CLAUDIN-LIKE PROTEIN ZF-A89
 >gi|4127541|emb|CAA09777.1| (AJ011789) claudin-like [Danio rerio]
 207-F08 002-G05 gi|6685320|sp|Q9YH90|CLDZ_BRARE CLAUDIN-LIKE PROTEIN ZF-A9
 >gi|4127543|emb|CAA09778.1| (AJ011790) claudin-like protein [Danio rerio]
 095-G05 095-G05 Unknown
 191-D10 076-G04 Unknown
 137-A05 137-A05 Unknown
 063-C11 063-C11 Unknown
 076-D11 174-C09 Unknown
 039-B02 039-B02 Unknown
 178-H10 056-D07 Unknown
 154-A02 154-A02 Unknown
 187-B03 187-B03 Unknown
 072-G02 072-G02 Unknown
 203-F05 203-F05 Unknown
 210-H01 210-H01 Unknown
 160-A01 174-G03 Unknown
 147-G02 147-G02 Unknown
 090-G02 094-D06 Unknown
 140-D07 140-D07 Unknown
 075-E10 075-E10 Unknown
 185-G04 185-G04 Unknown
 073-A02 073-A02 Unknown

