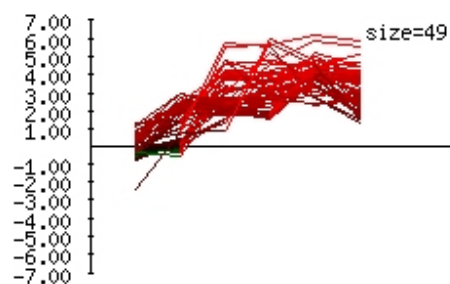
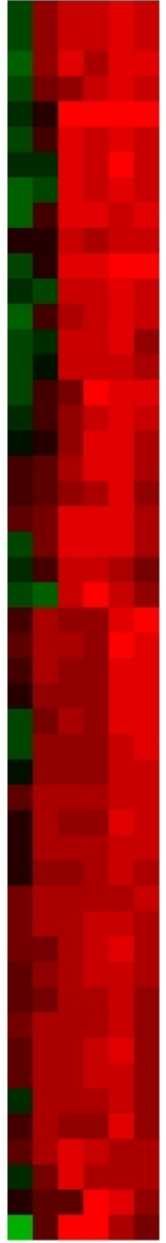


Subtree	Cluster size is 49, joined at the distance 0.203098				1:23
Spot size:	Width:	Autom.	Height:	Autom.	Tree height: Automatic Show titles:
		Autom.			
Colors	Colors	NA color	Number of slots	Discretisation type	Redraw!

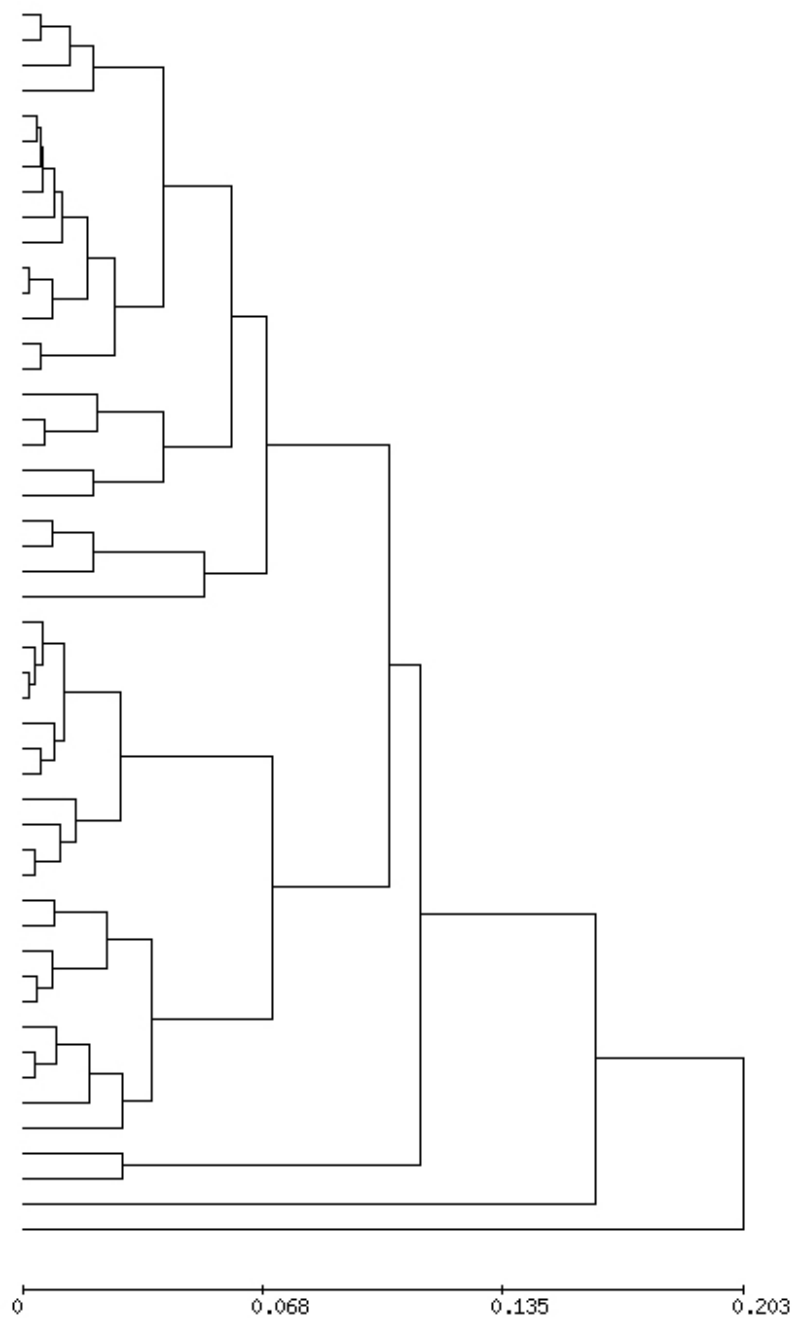


3sh3j.corr.dist.ave.cluster

23 24 25 26 27



088-G11
052-B06
189-C04
051-H05
104-B10
025-A07
091-H03
058-D03
204-B03
052-C05
200-F09
062-F07
008-A04
011-A01
213-D09
085-A01
002-A09
123-A02
209-F02
101-E03
074-D10
184-F07
142-F03
126-C08
193-B10
200-B07
051-E06
156-H10
155-E05
060-B06
162-H11
113-A05
113-C05
037-B10
123-D09
039-D08
128-H09
049-D01
168-B08
081-C07
064-E12
199-B07
095-E12
208-B11
197-D09
194-F09
060-D05
080-D01
092-E10



088-G11 174-A12 Unknown
052-B06 052-B06 gi|10242166|gb|AAG15318.1|AF255556_1 (AF255556) alpha tubulin [Notothenia coriiceps]
189-C04 189-C04 Unknown
051-H05 051-H05 gi|15293869|gb|AAK95127.1|AF401555_1 (AF401555) ribosomal protein L4 [Ictalurus punctatus]
104-B10 135-C12 Unknown
025-A07 025-A07 gi|5813855|gb|AAD52042.1|AF084461_1 (AF084461) type I cytokeratin; cyt1 [Danio rerio]
091-H03 008-B01 Unknown
058-D03 151-F06 gi|14017947|dbj|BAB47494.1| (AB058768) KIAA1865 protein [Homo sapiens]
204-B03 204-B03 Unknown
052-C05 142-H12 gi|6180205|gb|AAF05846.1|AF197880_1 (AF197880) type I cytokeratin [Danio rerio]
200-F09 200-F09 Unknown
062-F07 058-E03 Unknown
008-A04 004-D09 gi|15294013|gb|AAK95183.1|AF402809_1 (AF402809) 40S ribosomal protein S2 [Ictalurus punctatus]
011-A01 081-E10 Unknown
213-D09 213-D09 Unknown
085-A01 165-A01 gi|3913050|sp|042363|APA1_BRARE APOLIPOPROTEIN A-I PRECURSOR (APO-AI) >gi|2440055|emb|CAA74004.1| (Y13653) apolipoprotein A-I precursor protein [Danio rerio]
002-A09 002-A09 gi|7709990|ref|NP_057877.1| betaine-homocysteine methyltransferase [Mus musculus] >gi|5915784|sp|035490|BHMT_MOUSE BETAIN--HOMOCYSTEINE S-METHYLTRANSFERASE >gi|2645804|gb|AAB87501.1| (AF033381) betaine homocysteine methyl transferase [Mus musculus] >gi|9
123-A02 123-A02 gi|1752664|emb|CAA69623.1| (Y08321) cathepsin L [Danio rerio]
209-F02 170-B02 Unknown
101-E03 159-D06 Unknown
074-D10 202-A09 Unknown
184-F07 184-F07 Unknown
142-F03 142-F03 Unknown
126-C08 126-C08 Unknown
193-B10 193-B10 gi|8096950|gb|AAF61951.2|AF237950_1 (AF237950) arginine:glycine amidinotransferase [Gallus gallus]
200-B07 200-B07 Unknown
051-E06 051-E06 Unknown
156-H10 156-H10 Unknown
155-E05 092-G12 gi|8698673|gb|AAF78470.1|AF180887_1 (AF180887) skeletal alpha1 actin [Danio rerio]
060-B06 060-B06 Unknown
162-H11 211-G07 Unknown
113-A05 013-F11 Unknown
113-C05 113-C05 gi|6653228|gb|AAF22646.1| (AF190473) skeletal alpha-actin [Sparus aurata]
037-B10 189-E05 Unknown
123-D09 180-H03 Unknown
039-D08 039-D08 Unknown

128-H09 005-C08 gi|9988777|gb|AAG10787.1|AF292096_1 (AF292096) protein kinase AIRK2 [Xenopus laevis]
049-D01 049-D01 Unknown
168-B08 162-C10 gi|4929579|gb|AAD34050.1|AF151813_1 (AF151813) CGI-55 protein [Homo sapiens]
081-C07 081-C07 Unknown
064-E12 066-A01 gi|15489106|gb|AAH13665.1|AAH13665 (BC013665) Similar to RIKEN cDNA 1200009K13 gene [Mus musculus]
199-B07 028-G01 Unknown
095-E12 179-D11 Unknown
208-B11 194-C09 gi|13385280|ref|NP_080090.1| RIKEN cDNA 1200009K13 gene [Mus musculus]
 >gi|12836024|dbj|BAB23466.1| (AK004678) putative [Mus musculus]
197-D09 148-H04 Unknown
194-F09 194-F09 Unknown
060-D05 060-D05 Unknown
080-D01 147-F04 gi|14726331|ref|XP_003920.4| betaine-homocysteine methyltransferase [Homo sapiens] >gi|4530461|gb|AAD22043.1| (AF118378) betaine-homocysteine methyltransferase [Homo sapiens] >gi|15214972|gb|AAH12616.1|AAH12616 (BC012616) Unknown (protein for MGC:13645) [
092-E10 188-E11 "gi|2808622|emb|CAA74664.1| (Y14289) K18, simple type I keratin [Oncorhynchus mykiss]"