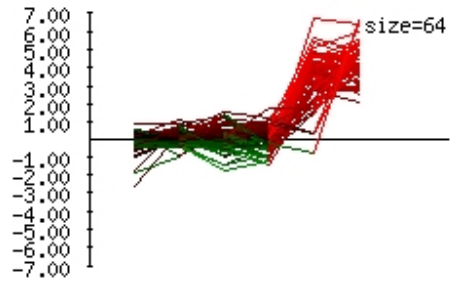
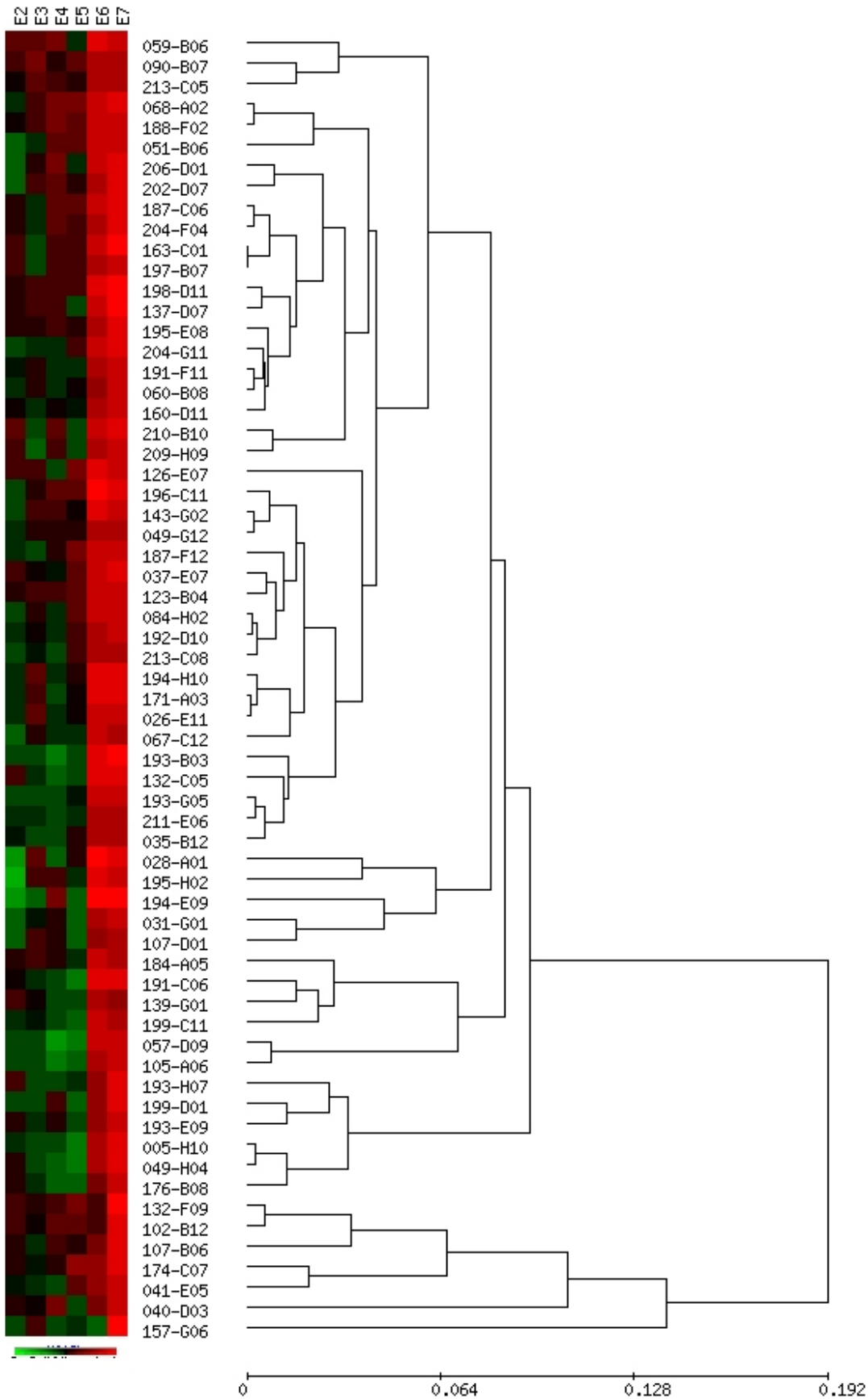


Subtree	Cluster size is 64, joined at the distance 0.191553				1:7			
Spot size:	Width:	Autom.	Height:	Autom.	Tree height:	Automatic	Show titles:	Redraw!
	Autom.							
Colors	Colors	NA color	Number of slots	Discretisation type				



Ex102.corr.dist.ave.cluster



CLUSTER_0001: 059-B06 090-B07 213-C05 068-A02 188-F02 051-B06 206-D01 202-D07 187-C06 204-F04 163-C01 197-B07 198-D11 137-D07 195-E08 204-G11 191-F11 060-B08 160-D11 210-B10 209-H09 126-E07 196-C11 143-G02 049-G12 187-F12 037-E07 123-B04 084-H02 192-D10 213-C08 194-H10 171-A03 026-E11 067-C12 193-B03 132-C05 193-G05 211-E06 035-B12 028-A01 195-H02 194-E09 031-G01 107-D01 184-A05 191-C06 139-G01 199-C11 057-D09 105-A06 193-H07 199-D01 193-E09 005-H10 049-H04 176-B08 132-F09 102-B12 107-B06 174-C07 041-E05 040-D03 157-G06
059-B06 059-B06 Unknown
090-B07 150-A08 Unknown
213-C05 190-C01 gi|8698673|gb|AAF78470.1|AF180887_1 (AF180887) skeletal alpha1 actin [Danio rerio]
068-A02 068-A02 Unknown
188-F02 188-F02 Unknown
051-B06 051-B06 gi|131096|sp|P09227|PRVA_CYPCA PARVALBUMIN ALPHA (A1)
206-D01 206-D01 Unknown
202-D07 202-D07 Unknown
187-C06 092-G12 gi|387082|gb|AAA37143.1| (M12233) skeletal muscle alpha-actin [Mus musculus]
204-F04 204-F04 gi|9294786|gb|AAF86670.1|AF178449_1 (AF178449) beta-galactosidase alpha peptide [Integration vector pCD11PKS] >gi|9294795|gb|AAF86676.1|AF178452_1 (AF178452) beta-galactosidase alpha peptide [Integration vector pCD13PKS]
163-C01 171-C06 gi|3676660|gb|AAC62069.1| (AF082662) embryonic 1 beta-globin [Danio rerio]
197-B07 197-B07 Unknown
198-D11 192-C12 Unknown
137-D07 137-D07 Unknown
195-E08 195-E08 Unknown
204-G11 204-G11 Unknown
191-F11 191-F11 Unknown
060-B08 026-B02 Unknown
160-D11 160-D11 Unknown
210-B10 159-B10 "gi|136216|sp|P02644|TRIF_CHICK TROPONIN I, FAST SKELETAL MUSCLE (TROPONIN I, FAST-TWITCH ISOFORM) >gi|86124|pir|A23569 troponin I, fast skeletal muscle - quail >gi|625311|pir|TPCHIS troponin I, fast skeletal muscle - chicken >gi|213644|gb|AAB00122.1| (
209-H09 209-H09 Unknown
126-E07 158-B04 gi|6009664|dbj|BAA85018.1| (AB026052) embryonic alpha-type globin [Oryzias latipes]
196-C11 196-C11 Unknown
143-G02 143-G02 Unknown
049-G12 049-G12 gi|8698675|gb|AAF78471.1|AF180888_1 (AF180888) parvalbumin [Danio rerio]
187-F12 187-F12 Unknown
037-E07 063-B05 gi|8698679|gb|AAF78473.1|AF180890_1 (AF180890) fast skeletal muscle troponin C [Danio rerio]
123-B04 007-G01 Unknown
084-H02 081-B06 Unknown
192-D10 013-F09 Unknown
213-C08 213-C08 Unknown
194-H10 194-H10 Unknown
171-A03 103-E04 Unknown

026-E11 144-H03 Unknown
067-C12 137-G05 gi|3986184|dbj|BAA34949.1| (AB015449) embryonic alpha-type globin2
 [Oncorhynchus mykiss]
193-B03 192-B03 Unknown
132-C05 132-C05 Unknown
193-G05 192-G05 Unknown
211-E06 211-E06 Unknown
035-B12 035-B12 Unknown
028-A01 010-C02 gi|2565196|gb|AAB81938.1| (AF000381) non-functional folate binding protein
 [Homo sapiens]
195-H02 195-H02 Unknown
194-E09 194-E09 Unknown
031-G01 031-G01 Unknown
107-D01 102-E07 "gi|1170206|sp|P43091|HEM1_OPSTA 5-AMINOLEVULINIC ACID SYNTHASE, NONSPECIFIC,
 MITOCHONDRIAL PRECURSOR (DELTA-AMINOLEVULINATE SYNTHASE) (DELTA-ALA SYNTHETASE) (ALAS-H)
 >gi|532630|gb|AAA49435.1| (L35915) 5-aminolevulinate synthase [Opsanus tau]"
184-A05 181-F03 Unknown
191-C06 191-C06 Unknown
139-G01 139-G01 Unknown
199-C11 199-C11 Unknown
057-D09 057-D09 Unknown
105-A06 001-F03 "gi|136083|sp|P13104|TPMA_BRARE TROPOMYOSIN ALPHA CHAIN, SKELETAL MUSCLE
 >gi|2133835|pir||I51731 alpha-tropomyosin - zebra fish >gi|214974|gb|AAA50021.1| (M24635)
 alpha-tropomyosin [Danio rerio] >gi|8698683|gb|AAF78475.1|AF180892_1 (AF180892) alpha tropom
193-H07 192-H07 Unknown
199-D01 199-D01 gi|8698677|gb|AAF78472.1|AF180889_1 (AF180889) fast skeletal muscle troponin
 T [Danio rerio]
193-E09 191-H06 "gi|4502699|ref|NP_003662.1| CDC14 homolog B, isoform 1 [Homo sapiens]
 >gi|14743349|ref|XP_011585.4| S. cerevisiae CDC14 homolog, gene B [Homo sapiens]
 >gi|2662463|gb|AAB88293.1| (AF023158) tyrosine phosphatase [Homo sapiens]"
005-H10 142-E08 gi|15921448|ref|NP_377117.1| 109aa long hypothetical protein [Sulfolobus
 tokodaii] >gi|15622234|dbj|BAB66226.1| (AP000985) 109aa long hypothetical protein
 [Sulfolobus tokodaii]
049-H04 049-H04 Unknown
176-B08 176-B08 Unknown
132-F09 132-F09 gi|117451|sp|P10044|CRG2_CYPKA GAMMA CRYSTALLIN M2 (GAMMA-M2)
 >gi|71519|pir||CYCAG2 gamma-crystallin m2 precursor - common carp >gi|62604|emb|CAA31388.1|
 (X12903) gamma-crystallin gamma-m2 (AA 1 - 173) [Cyprinus carpio]
102-B12 102-B05 gi|3695368|gb|AAC62774.1| (AF093129) zfY1 [Danio rerio]
107-B06 009-B02 gi|3421386|gb|AAC32193.1| (AF081462) myosin light chain 2 [Danio rerio]
174-C07 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) [
041-E05 037-D08 gi|15294013|gb|AAK95183.1|AF402809_1 (AF402809) 40S ribosomal protein S2
 [Ictalurus punctatus]
040-D03 040-D03 "gi|14773188|ref|XP_006406.4| troponin I, skeletal, fast [Homo sapiens]"

157-G06 173-F09 gi|11693032|gb|AAG38940.1| (AY013297) calcineurin-binding protein calsarcin-2
[Homo sapiens]