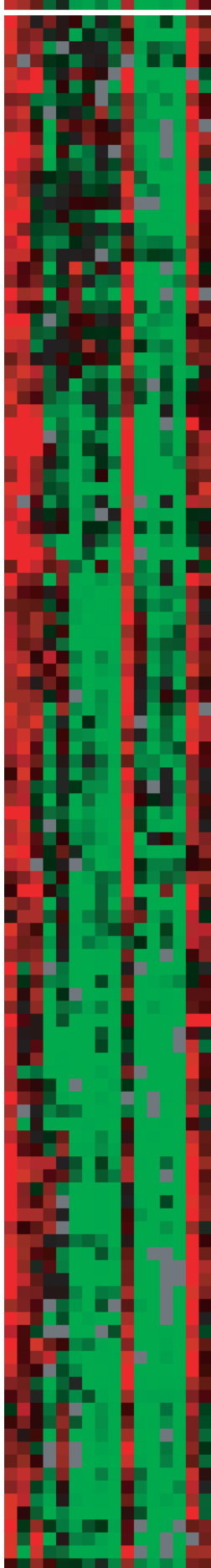


66 E11 head  
30 E12 head  
31 E13 head  
40 E15 head  
19 E19 head  
39 E19 head  
xx N6 head  
xx N8 head  
xx N10 head  
66 N10 cerebellum  
15 cerebellum  
07 brain  
75 cerebellum  
63 medulla oblongata  
64 olfactory brain



2610529E15 Tera (Tera) mRNA  
2610529P06 EST  
2810406G03 homolog to HYPOTHETICAL PROTEIN KIAA0101.[fantom]  
1110030E03 unclassifiabletranscript[fantom]  
2510004L17 ESTs  
1200001C16 GLN retrotransposon family 3-1  
1600019P09 U2-snRNP b" (pRNP11)  
1500006J04 DNA primase, p49 subunit  
2700078I24 ESTs  
1700008B16 karyopherin (importin) alpha 2  
2610005B21 homolog to FLS353.[fantom]  
1700026B03 RAS-like, family 2, locus 9[fantom]  
1110036C23 DNA methyltransferase (cytosine-5)  
1190024M07 mini chromosome maintenance deficient 4[fantom]  
1700028H20 serine/threonine kinase 18[fantom]  
1110061N19 nuclear protein 95[fantom]  
1700010N24 ESTs, similar to ACTIVATOR 1 37 KD SUBUNIT [Homo sapiens]  
1700029M11 ADP-ribosyltransferase (NAD+; poly (ADP-ribose) polymerase) 1  
1700029M02 actin-like 6[fantom]  
2610024A10 MAD2 (mitotic arrest deficient, homolog)[fantom]  
3110001D19 homolog to MITOTIC KINASE-LIKE PROTEIN-1.[fantom]  
2010204O16 high mobility group protein 2[fantom]  
2610009D01 ESTs, similar to unknown [C.elegans]  
2400001O07 EST  
1810016H06 ADP-ribosyltransferase (NAD+; poly (ADP-ribose) polymerase) 1  
1700001K22 similar to REPLICATION FACTOR C, 40 KDA SUBUNIT.[fantom]  
1110007D06 ring finger protein 2[fantom]  
1700029D20 small EDRK-rich factor 1[fantom]  
2700088O10 hyaluronidase 2  
1810011I18 homolog to U5 SNRNP-SPECIFIC 40 KDA PROTEIN.[fantom]  
2510001I21 proliferation-associated protein 1  
2410002O03 ADP-ribosylarginine hydrolase  
2410004F21 cyclin B1, related sequence 1  
2310032K12 mini chromosome maintenance deficient 7 (S. cerevisiae)  
2510004H09 poly A binding protein, cytoplasmic 1  
2410004M23 amelogenin  
2400002I05 nucleophosmin 1  
1810019E15 DEK oncogene (DNA binding)[fantom]  
2810421N01 LIPOATE-PROTEIN LIGASE B (LIPOATE BIOSYNTHESIS PROTEIN B)  
1810026F04 homolog to HSPCO25.[fantom]  
2510002A06 transcriptional intermediary factor 1, beta  
2810002B04 hemoglobin X, alpha-like embryonic chain in Hba complex  
2700038P03 tubulin, beta 5[fantom]  
2210420E09 high mobility group protein 1  
2810402H01 splicing factor, arginine/serine-rich 3 (SRp20)  
1700016F14 MDJ6.[fantom]  
2610034C21 splicing factor, arginine/serine-rich 3[fantom]  
2310043F15 ESTs, similar to heterogenous nuclear ribonucleoprotein A2/B1 [M.musculus]  
2410006F02 syntrophin, acidic 1  
2810017D11 splicing factor, arginine/serine-rich 3[fantom]  
2610027C05 ESTs, similar to PRE-MRNA SPLICING FACTOR SRP75 [Homo sapiens]  
2610025E13 ESTs  
3300002D11 Ewing sarcoma homolog[fantom]  
2600017F24 ESTs  
1700011N22 high mobility group protein 1  
1500005O20 ESTs  
1600012G22 splicing factor, arginine/serine-rich 3 (SRp20)  
1500015E08 ESTs  
0610016P10 ESTs, similar to nucleoporin Nup84 [R.norvegicus]  
1700008B17 Mus musculus 26S proteasome ATPase (CIP21) mRNA, complete cds  
1500005D11 hexosaminidase  
1700028K07 RAN, member RAS oncogene family[fantom]  
1190032N16 glycosyl-phosphatidyl-inositol-anchored protein homolog  
1500034K21 NUCLEOSOME ASSEMBLY PROTEIN 1-LIKE 1  
2810003H07 UDP-glucuronosyltransferase 1 family, me[fantom]  
2810003C07 RNA binding motif protein, X chromosome retrogene  
2600001I05 nucleolin  
2400002A21 prothymosin alpha  
2500002K24 similar to PROLIFERATING CELL NUCLEAR ANTIGEN (PCNA) (CYCLIN).[fantom]  
1110001C01 ESTs, similar to putative deubiquitinating enzyme UBPY [M.musculus]  
2610524I02 ESTs  
2810403C15 splicing factor, arginine/serine-rich 3 (SRp20)  
1190033O17 cytokine inducible SH2-containing protein 7  
2610044P12 inosine 5'-phosphate dehydrogenase 2[fantom]  
2410003J23 ribosomal protein S3  
2010016F17 RNA binding protein 1  
2810417B20 similar to BAF57[fantom]  
2700082L14 EST  
2410003G18 hypothetical protein[fantom]  
2410003A12 C9[fantom]  
2700047H10 high mobility group protein 17[fantom]  
2510042E23 EST  
2410006I22 solute carrier family 10 (sodium/bile acid cotransporter family), member 1  
2810004G22 homolog to UBIQUITIN-CONJUGATING ENZYME E2 H10 (EC 6.3.2.19) (UBIQUITIN-PROTEIN LIGASE) (UBIQUITIN CARRIER PROTEIN).[fantom]  
1110014E11 centromere autoantigen A  
2600017H22 Rad51 homolog (S. cerevisiae)[fantom]  
2700088A02 ESTs  
2610017G21 thymidylate synthase[fantom]  
2810406C15 related to 13S condensin XCAP-D2 subunit[fantom]  
2610033C07 homolog to HYPOTHETICAL PROTEIN KIAA0101.[fantom]  
2700018L05 unclassifiable transcript[fantom]  
2410044B12 karyopherin (importin) alpha 2[fantom]  
2610021G09 SM-20 (factor-responsive smooth muscle protein ) [fantom]  
3110002M17 EST, similar to UBIQUITIN-CONJUGATING ENZYME  
0610017N24 ESTs  
2610019J24 high mobility group protein 2[fantom]  
2700085B09 high mobility group protein 2[fantom]  
2610021J01 homolog to high-mobility group (nonhistone chromosomal) protein 1[fantom]  
2610510C05 cyclin B1, related sequence 1[fantom]  
2610042I15 mini chromosome maintenance deficient 4[fantom]  
1700018M23 high mobility group protein 2  
2010013J11 CYCLIN-DEPENDENT KINASES REGULATORY SUBUNIT 2[fantom]  
3000002M17 Cyclin B1, related sequence 1  
2010107E01 nuclear protein 95[fantom]  
3110001H06 ESTs  
2810403E20 high mobility group protein 1  
1500011C13 ESTs, similar to (define not available 3668141) [H.sapiens]  
2700078C08 major histocompatibility protein class II alpha  
2610034C13 ribosomal protein S3[fantom]  
2310046J12 homolog to RIBOSOMAL PROTEIN L41 (FRAGMENT).[fantom]  
2700043E10 ring finger protein 2[fantom]  
2510027A07 coproporphyrinogen oxidase  
2610014N04 ESTs  
1110037B17 ESTs  
1700027H19 oxidative stress induced  
1110056L21 proliferating cell nuclear antigen  
2500004E18 centromere autoantigen A  
2510004J15 CD24a antigen  
2810004I20 cyclin A2  
2210404L05 ESTs