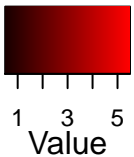
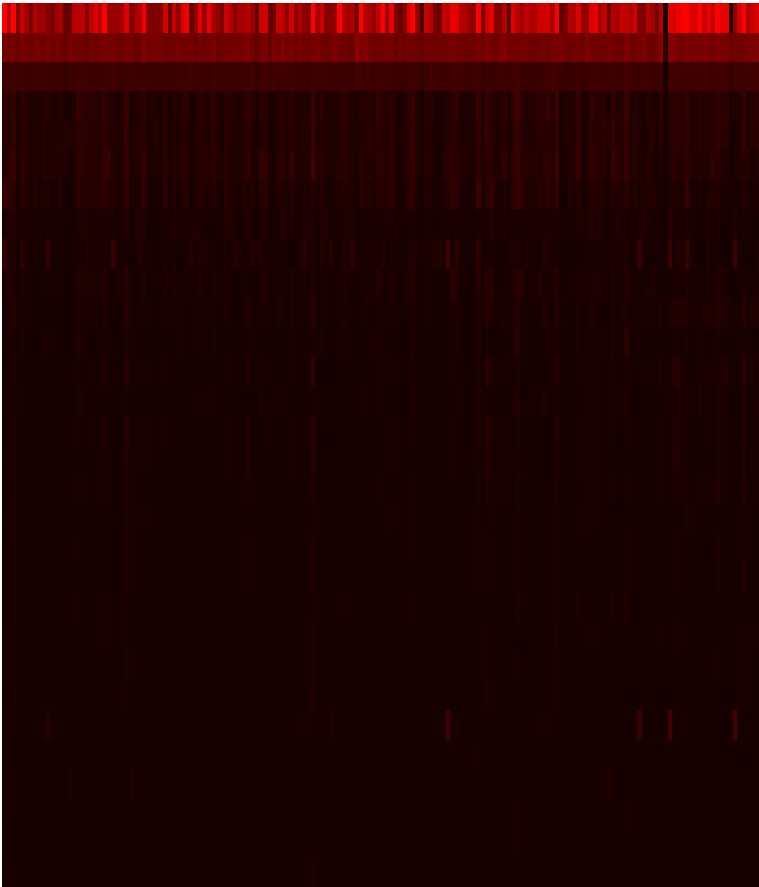


Color Key

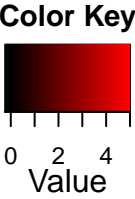


AGE Two-way test: $-\log_{10}(\text{p-value})$



510_120515
52_120416
689_120604
578_120521
160_120423
672_120530
316_120501
584_120522
708_120531
34_120416
705_120605
389_120503
584_120522
685_120604
286_120430
359_120502
494_120515
407_120503
280_120430
78_120417
537_120516
238_120426
255_120426
363_120502
435_120507
538_120516
676_120604
287_120430
637_120524
485_120515
545_120516
229_120425
406_120503
657_120529
323_120501
204_120424
472_120514
516_120515
846_130830
923_131031
936_131101
952_131107
956_131107
807_130701

hsa03010 Ribosome
hsa00190 Oxidative phosphorylation
hsa04260 Cardiac muscle contraction
hsa04971 Gastric acid secretion
hsa04970 Salivary secretion
hsa00010 Glycolysis / Gluconeogenesis
hsa04145 Phagosome
hsa00860 Porphyrin and chlorophyll metabolism
hsa04612 Antigen processing and presentation
hsa04540 Gap junction
hsa04720 Long-term potentiation and depression
hsa04976 Bile secretion
hsa04114 Oocyte meiosis
hsa04962 Vasopressin-regulated water transport
hsa04722 Neurotrophin signaling pathway
hsa04740 Olfactory transduction
hsa04912 GnRH signaling pathway
hsa04966 Collecting duct acid secretion
hsa04270 Vascular smooth muscle contraction
hsa04916 Melanogenesis
hsa04973 Carbohydrate digestion and absorption
hsa04130 SNARE interaction
hsa04020 Calcium signaling pathway
hsa04070 Phosphatidylinositol signaling system
hsa04141 Protein processing in endoplasmic reticulum
hsa04670 Leukocyte transmembrane protein signaling pathway
hsa00590 Arachidonic acid metabolism
hsa04972 Pancreatic secretion
hsa04810 Regulation of actin cytoskeleton
hsa04910 Insulin signaling pathway



GAGE test statistics

