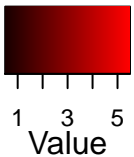
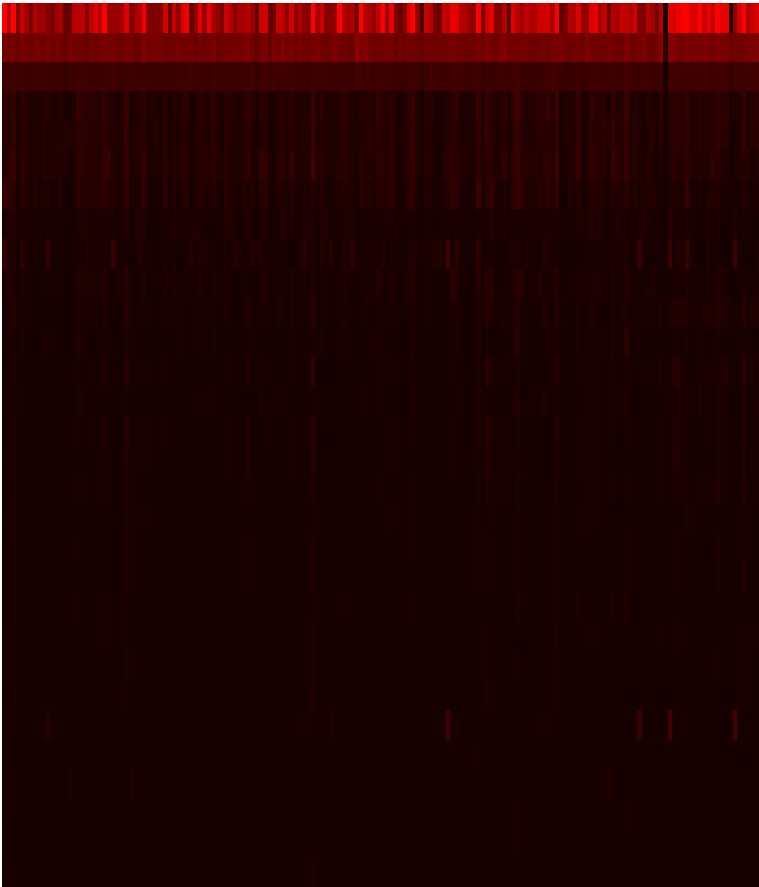


Color Key

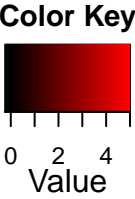


# AGE Two-way test: $-\log_{10}(\text{p-value})$



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

