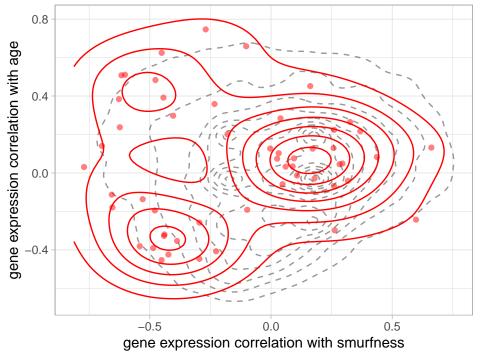
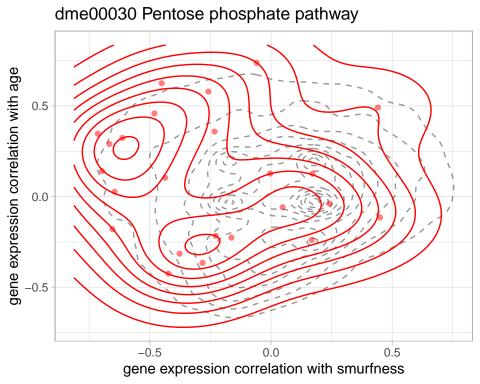
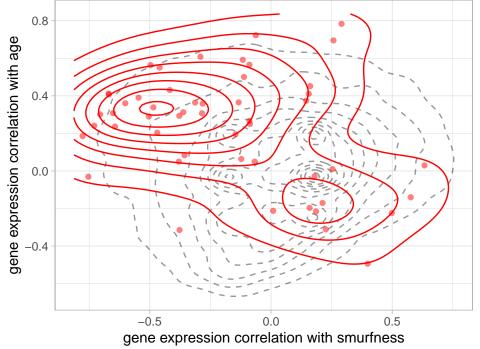
dme00010 Glycolysis / Gluconeogenesis

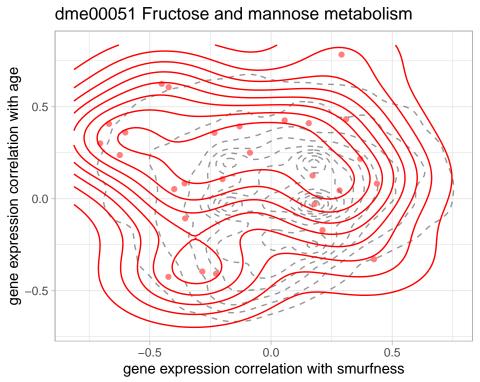


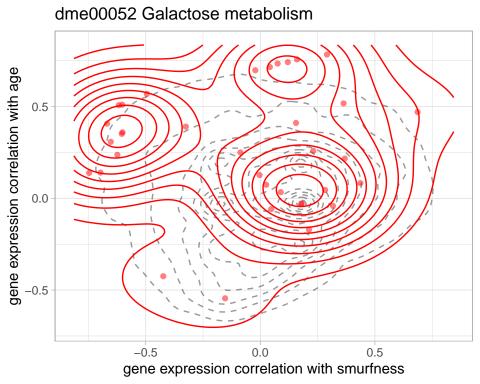
dme00020 Citrate cycle (TCA cycle) gene expression correlation with age 0.4 0.0 -0.4-0.50.0 0.5 gene expression correlation with smurfness



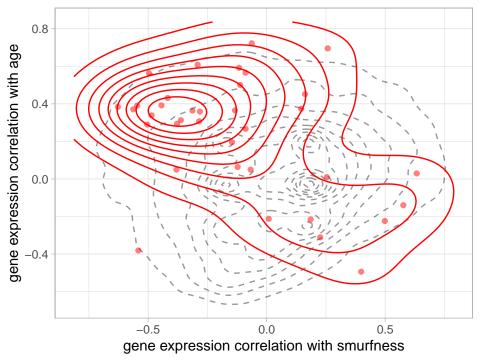
dme00040 Pentose and glucuronate interconversions

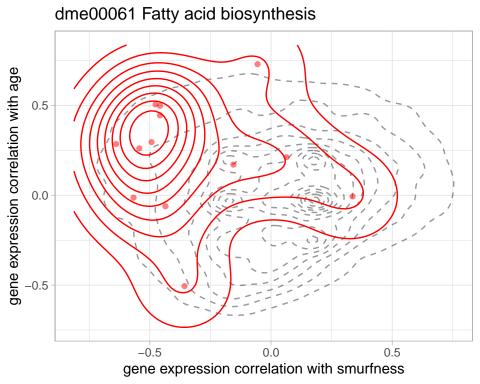


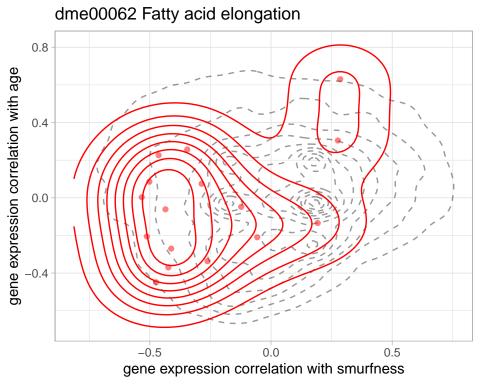




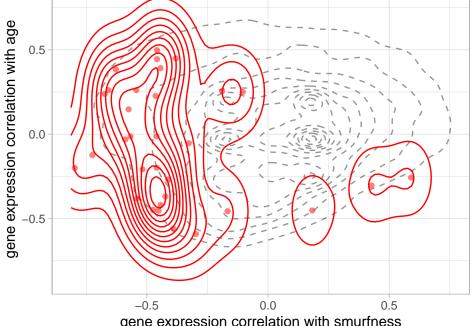
dme00053 Ascorbate and aldarate metabolism





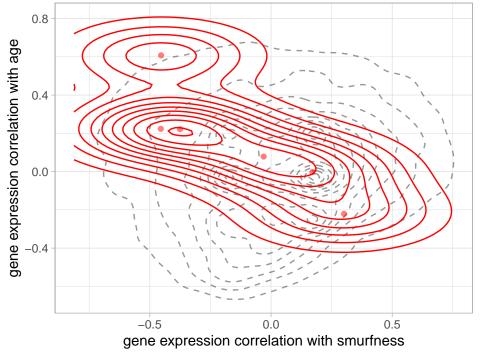


dme00071 Fatty acid degradation 0.5 0.0



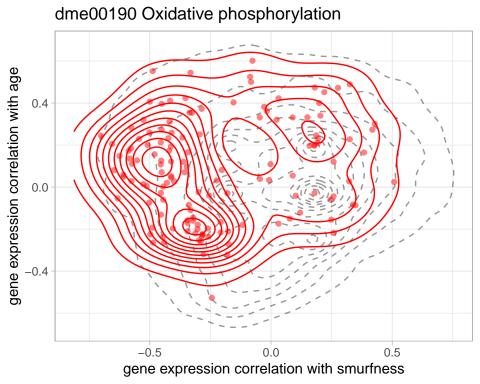
gene expression correlation with smurfness

dme00100 Steroid biosynthesis

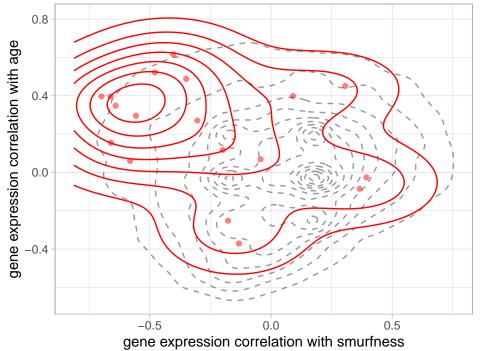


dme00130 Ubiquinone and other terpenoid-quinone bio 8.0 gene expression correlation with age 0.4 0.0 -0.40.0 -0.50.5

gene expression correlation with smurfness



dme00220 Arginine biosynthesis



dme00230 Purine metabolism 8.0 0.4 0.0 -0.4

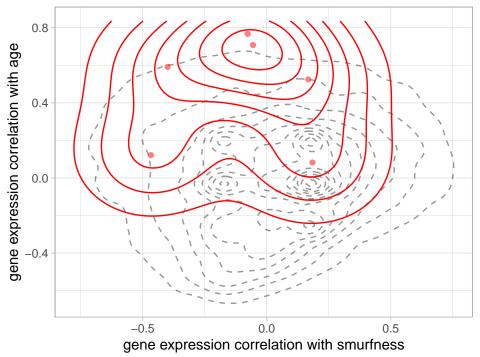
gene expression correlation with age

0.0 gene expression correlation with smurfness

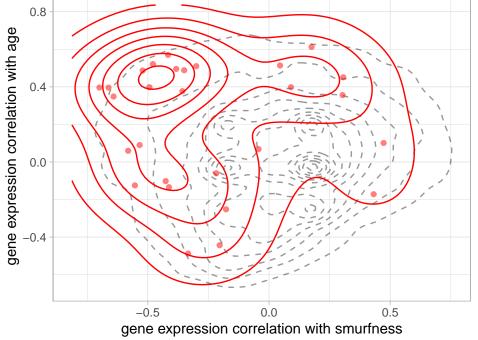
0.5

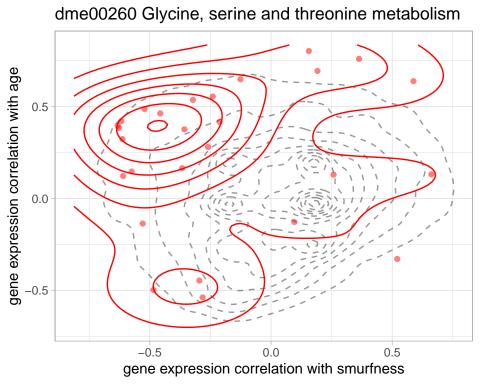
-0.5

dme00232 Caffeine metabolism

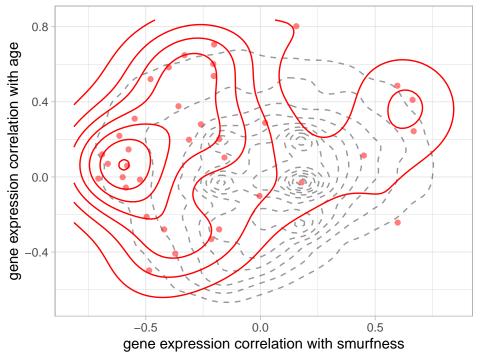


dme00240 Pyrimidine metabolism gene expression correlation with age 0.5 0.0 -0.5-0.50.0 0.5 gene expression correlation with smurfness dme00250 Alanine, aspartate and glutamate metabolism





dme00270 Cysteine and methionine metabolism

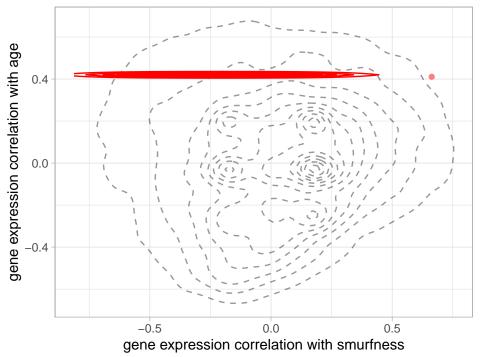


dme00280 Valine, leucine and isoleucine degradation 0.5 0.0 -0.50.0 -0.50.5

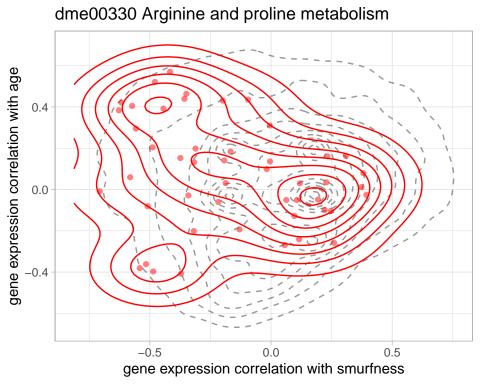
gene expression correlation with age

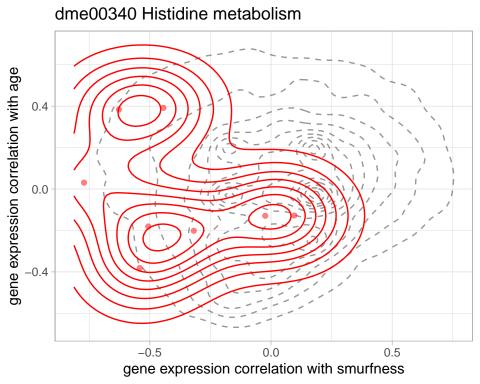
gene expression correlation with smurfness

dme00290 Valine, leucine and isoleucine biosynthesis

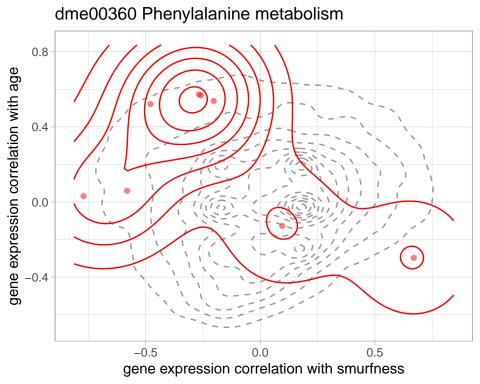


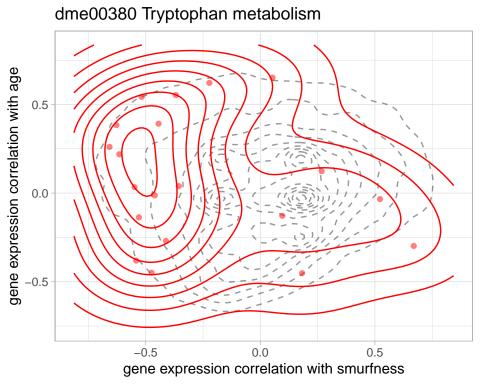
dme00310 Lysine degradation gene expression correlation with age 0.5 0.0 -0.50.0 0.5 gene expression correlation with smurfness

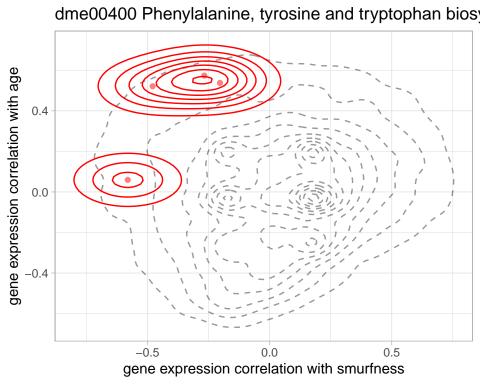


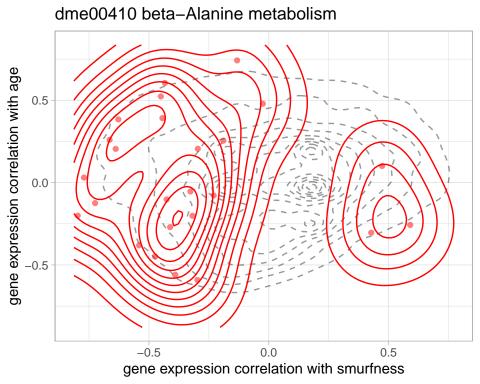


dme00350 Tyrosine metabolism 8.0 gene expression correlation with age 0.4 0.0 -0.4-0.50.0 0.5 gene expression correlation with smurfness







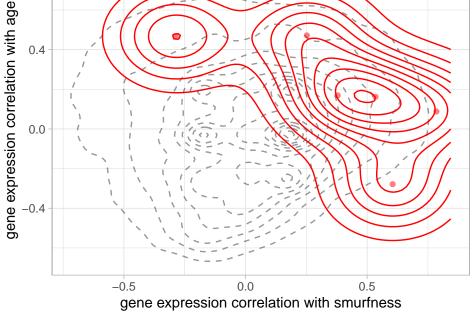


dme00430 Taurine and hypotaurine metabolism 0.5 0.0 -0.50.0 0.5

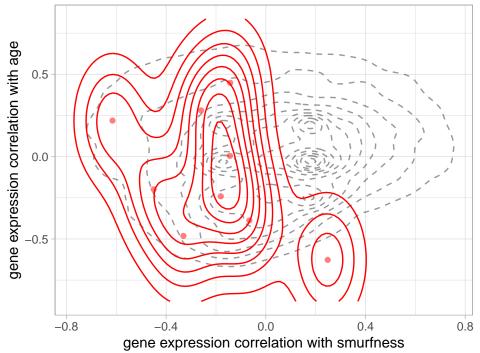
gene expression correlation with smurfness

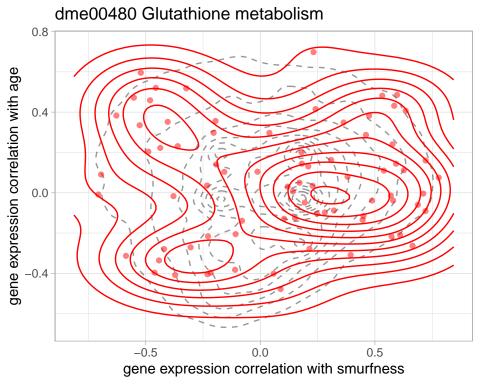
gene expression correlation with age

dme00440 Phosphonate and phosphinate metabolism 8.0 0.4 0.0

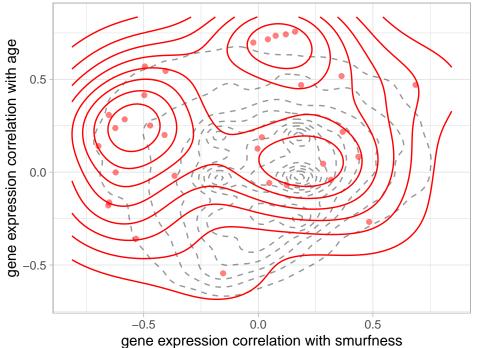


dme00450 Selenocompound metabolism

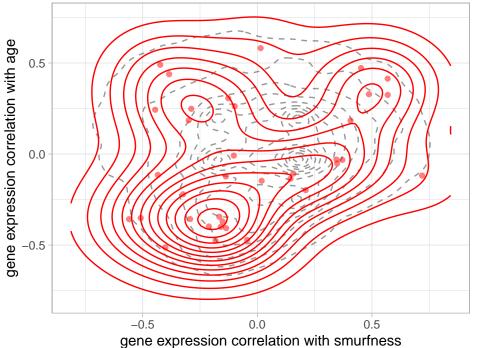


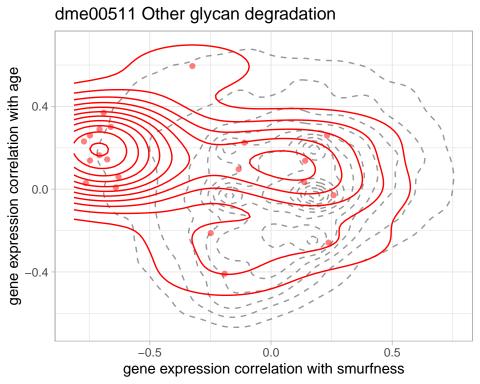


dme00500 Starch and sucrose metabolism

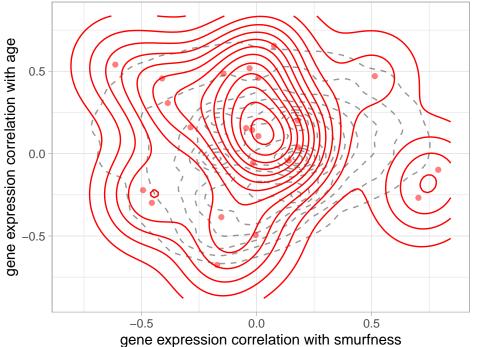


dme00510 N-Glycan biosynthesis

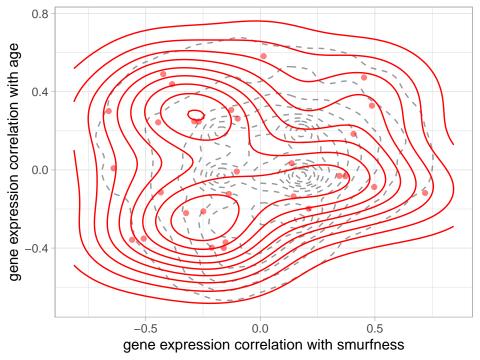




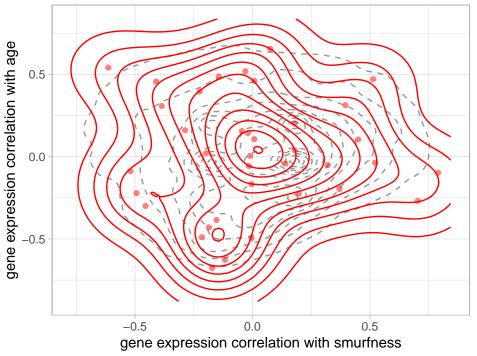
dme00512 Mucin type O-glycan biosynthesis



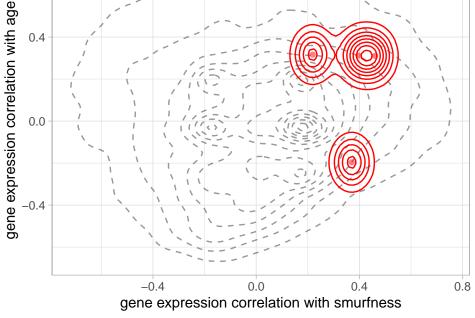
dme00513 Various types of N-glycan biosynthesis



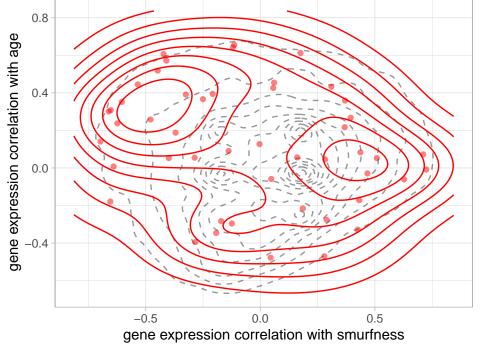
dme00514 Other types of O-glycan biosynthesis

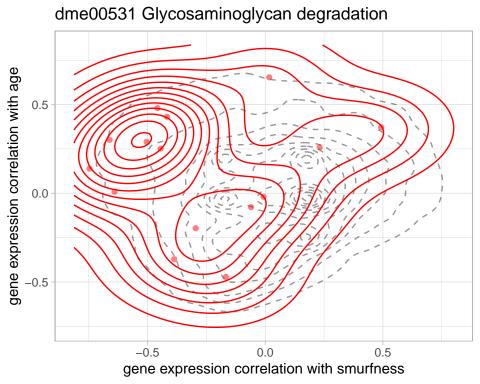


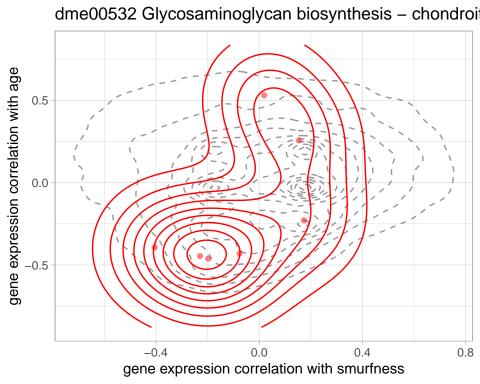
dme00515 Mannose type O-glycan biosynthesis 0.4 0.0



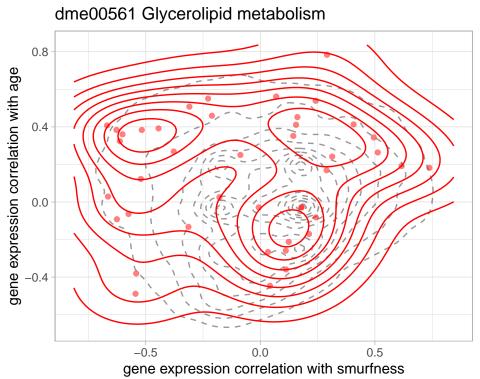
dme00520 Amino sugar and nucleotide sugar metabolis



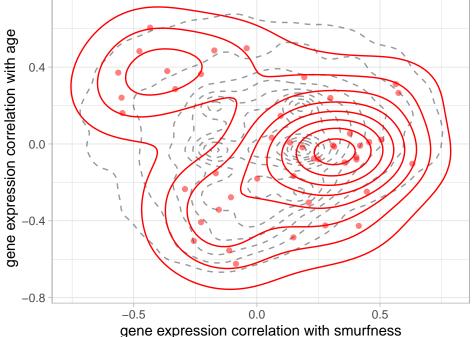




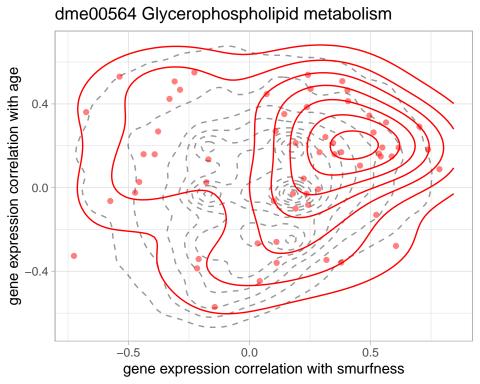
dme00534 Glycosaminoglycan biosynthesis – heparan gene expression correlation with age 0.4 0.0 -0.4 -0.8-0.50.0 0.5 gene expression correlation with smurfness



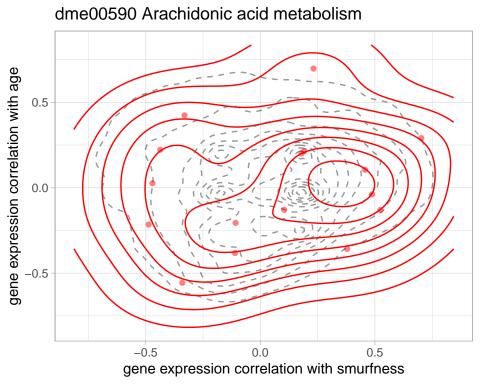
dme00562 Inositol phosphate metabolism



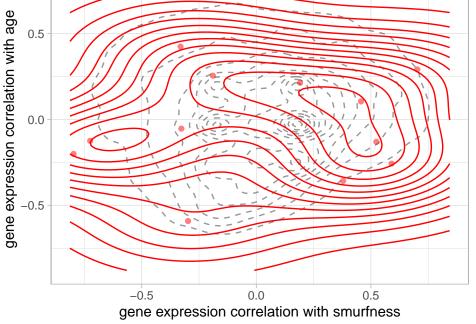
dme00563 Glycosylphosphatidylinositol (GPI)-anchor b gene expression correlation with age 0.4 0.0 -0.80.0 8.0 -0.40.4 gene expression correlation with smurfness

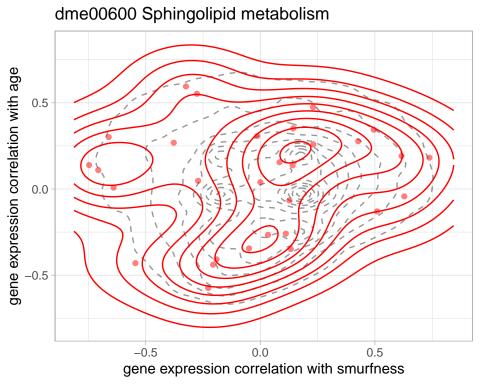


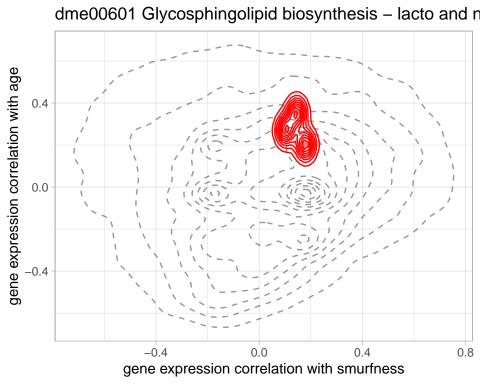
dme00565 Ether lipid metabolism gene expression correlation with age 0.4 0.0 -0.50.0 0.5 gene expression correlation with smurfness

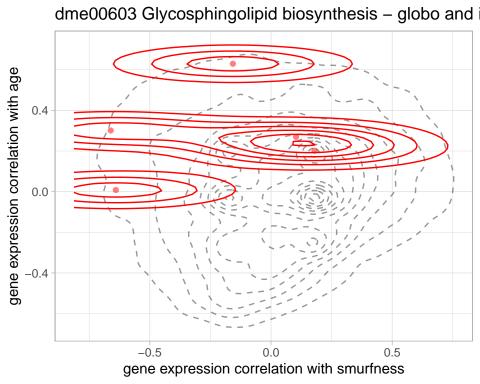


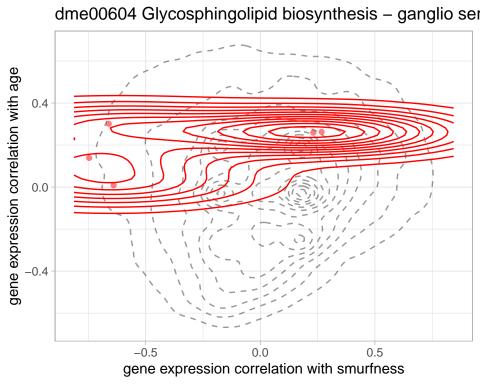
dme00592 alpha-Linolenic acid metabolism 0.5 0.0

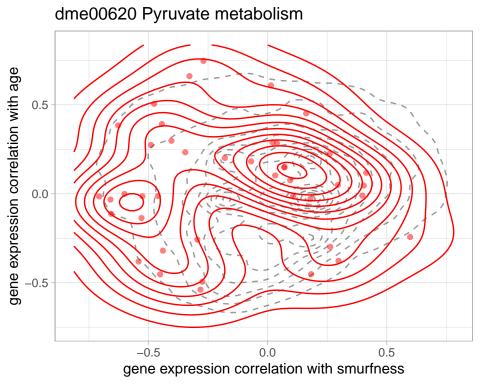


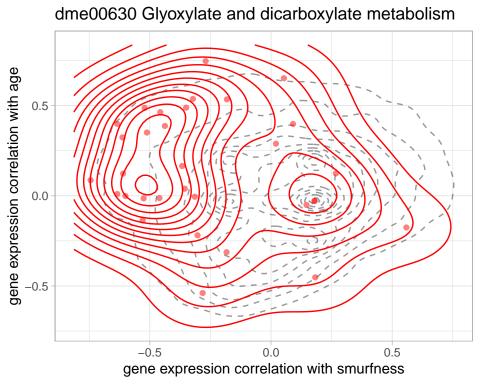




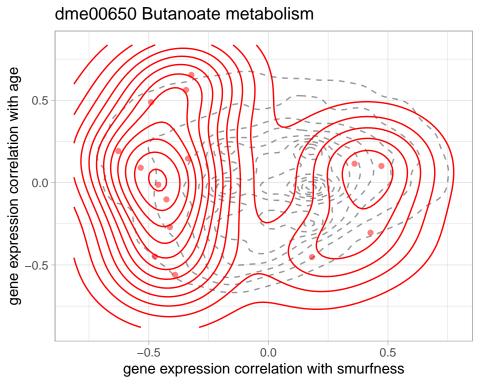








dme00640 Propanoate metabolism gene expression correlation with age 0.5 0.0 -0.50.0 0.5 gene expression correlation with smurfness

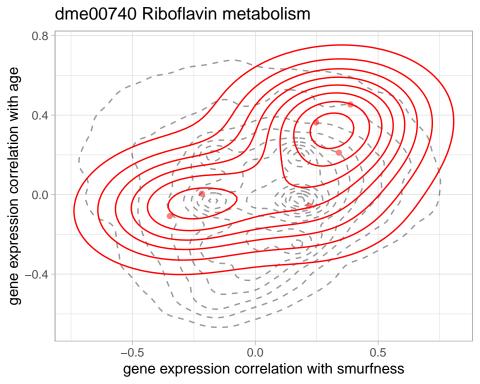


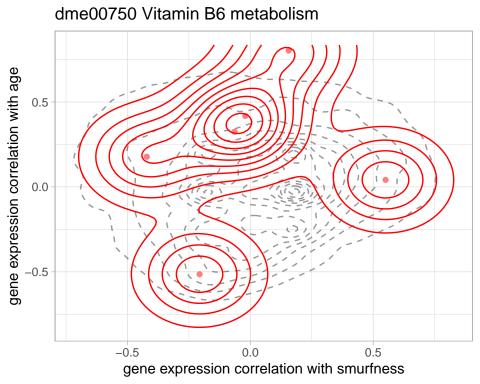
dme00670 One carbon pool by folate gene expression correlation with age 0.5 0.0 -0.5-0.50.0 0.5 gene expression correlation with smurfness

dme00730 Thiamine metabolism 0.4 0.0 -0.4-0.50.0 0.5

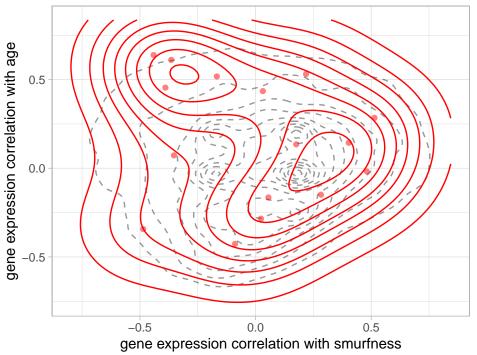
gene expression correlation with smurfness

gene expression correlation with age

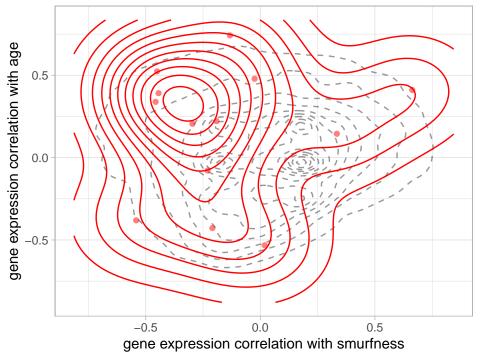




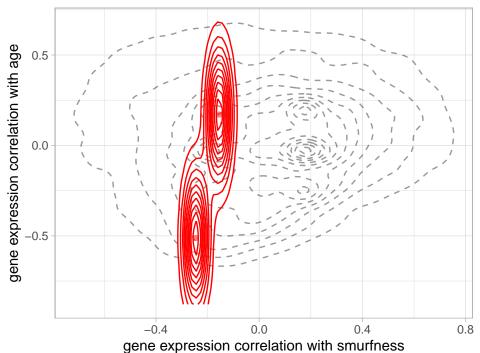
dme00760 Nicotinate and nicotinamide metabolism



dme00770 Pantothenate and CoA biosynthesis

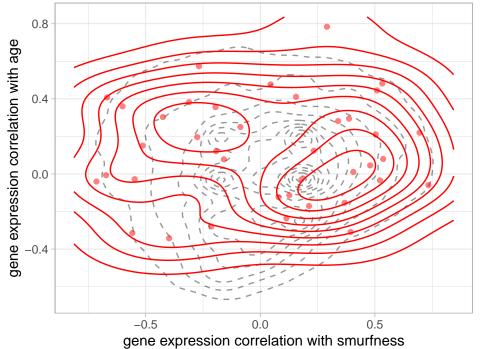


dme00780 Biotin metabolism

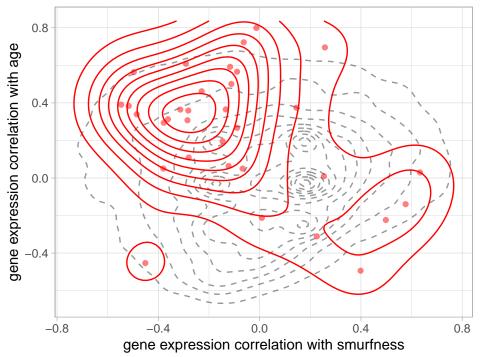


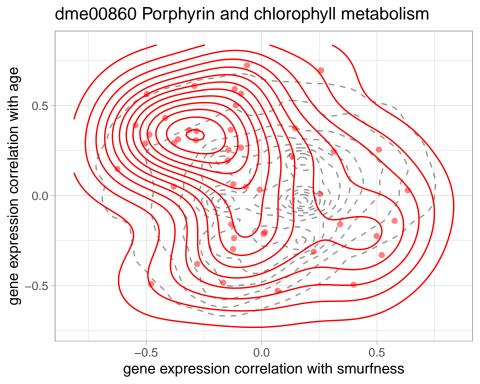
dme00785 Lipoic acid metabolism 8.0 gene expression correlation with age 0.4 0.0 -0.40.0 8.0 -0.40.4 gene expression correlation with smurfness

dme00790 Folate biosynthesis

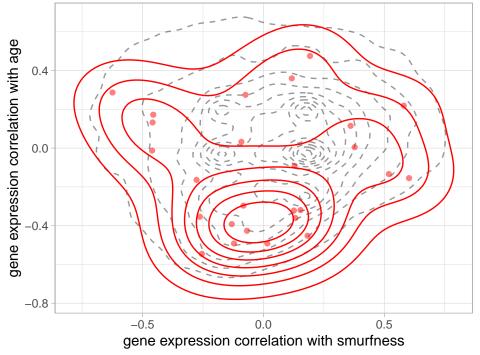


dme00830 Retinol metabolism

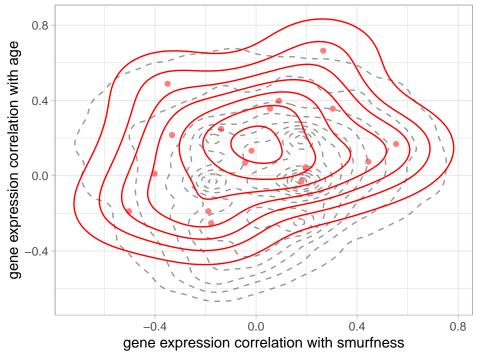


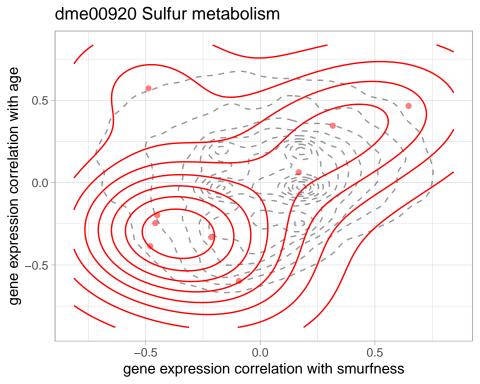


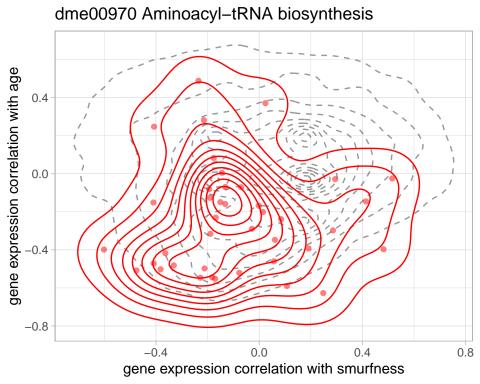
dme00900 Terpenoid backbone biosynthesis



dme00910 Nitrogen metabolism





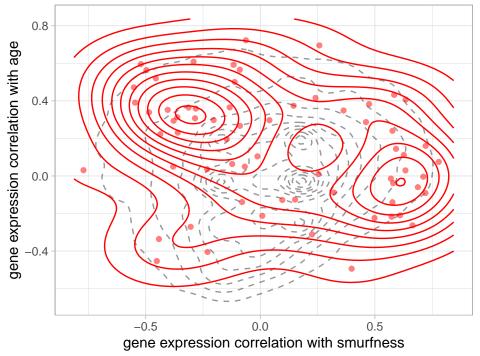


dme00980 Metabolism of xenobiotics by cytochrome P4 8.0 gene expression correlation with age 0.4 0.0 -0.4-0.50.0 0.5 gene expression correlation with smurfness

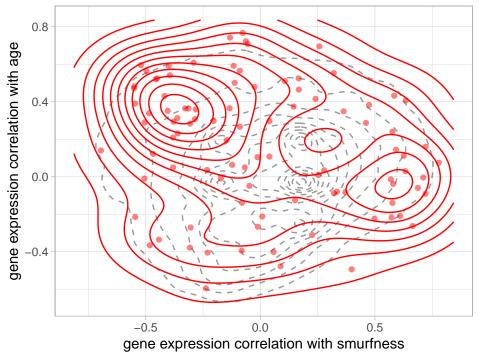
dme00981 Insect hormone biosynthesis 8.0 gene expression correlation with age 0.4 0.0 -0.4-0.50.0 0.5

gene expression correlation with smurfness

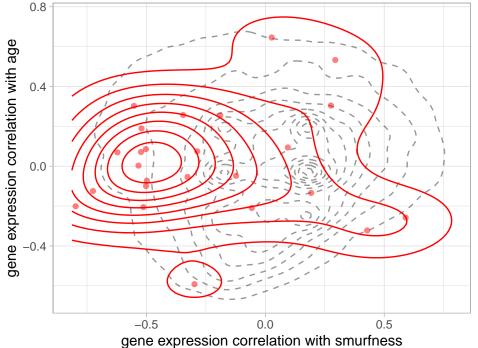
dme00982 Drug metabolism – cytochrome P450



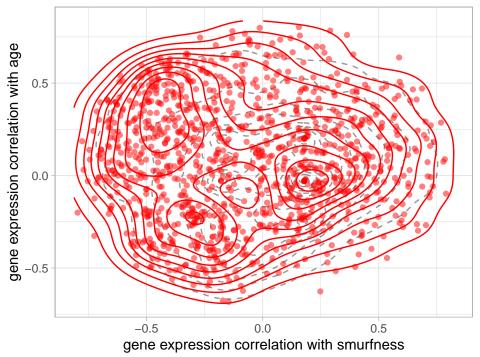
dme00983 Drug metabolism – other enzymes

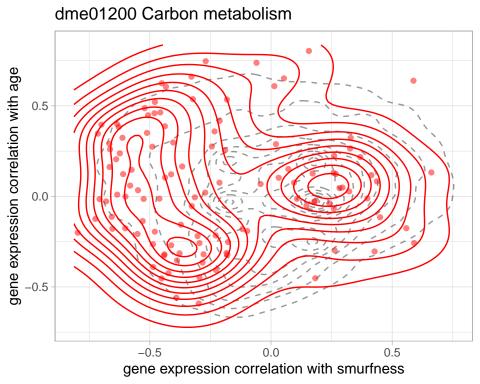


dme01040 Biosynthesis of unsaturated fatty acids

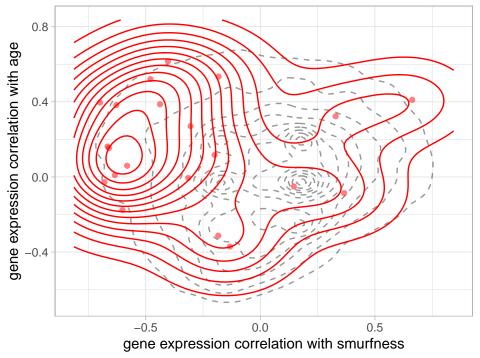


dme01100 Metabolic pathways





dme01210 2-Oxocarboxylic acid metabolism



dme01212 Fatty acid metabolism gene expression correlation with age 0.5 0.0 -0.5 -0.50.0 0.5

gene expression correlation with smurfness

dme01230 Biosynthesis of amino acids 0.5 0.0 -0.5

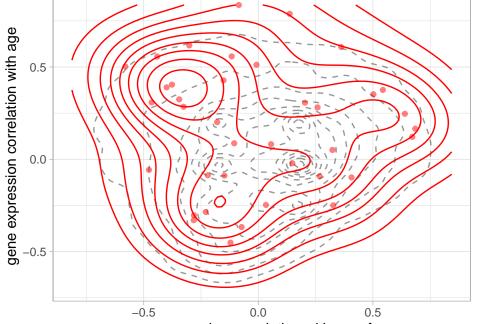
gene expression correlation with age

0.0 gene expression correlation with smurfness

0.5

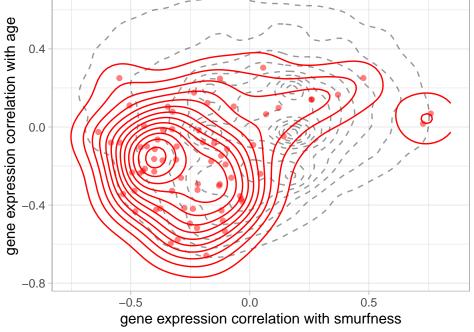
-0.5

dme02010 ABC transporters 0.5

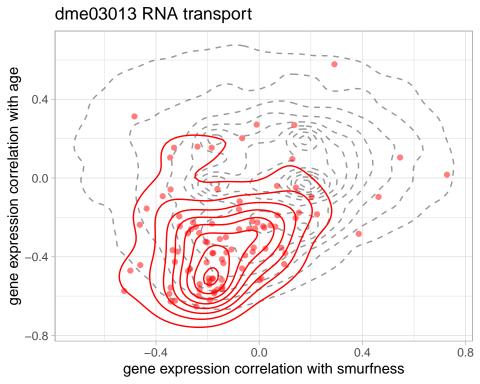


gene expression correlation with smurfness

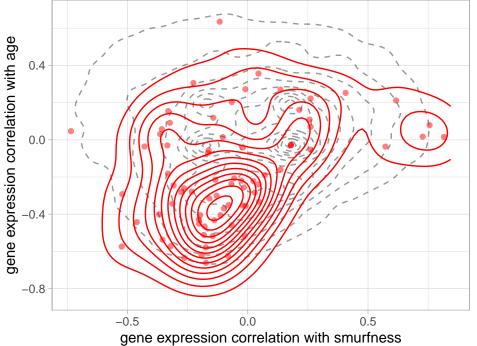
dme03008 Ribosome biogenesis in eukaryotes 0.4 0.0



dme03010 Ribosome gene expression correlation with age 0.4 0.0 -0.4-0.40.0 0.4 8.0 gene expression correlation with smurfness



dme03015 mRNA surveillance pathway

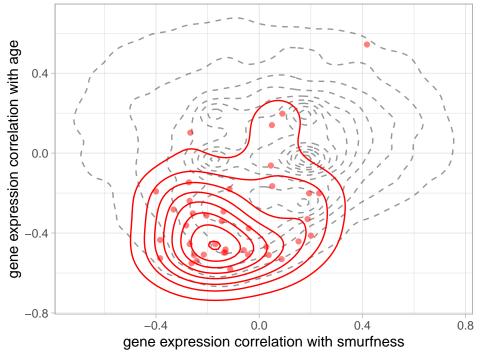


dme03018 RNA degradation gene expression correlation with age 0.4 0.0 -0.40.0 0.4 8.0 gene expression correlation with smurfness

dme03020 RNA polymerase gene expression correlation with age 0.4 0.0 -0.50.0 0.5

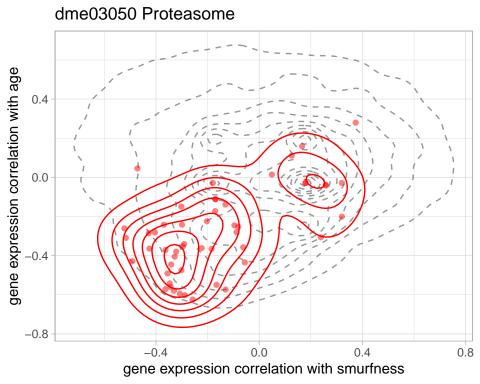
gene expression correlation with smurfness

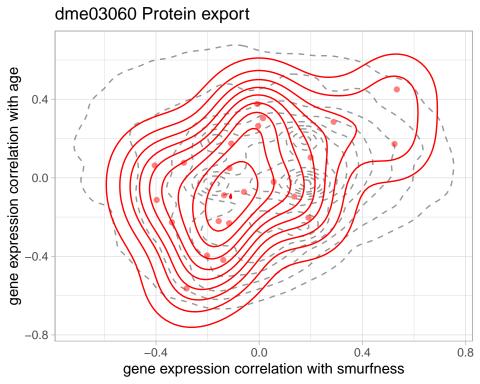
dme03022 Basal transcription factors



dme03030 DNA replication gene expression correlation with age 0.4 0.0 -0.8-0.40.0 0.4 8.0 gene expression correlation with smurfness

dme03040 Spliceosome gene expression correlation with age 0.4 0.0 -0.4 -0.8-0.40.0 0.4 8.0 gene expression correlation with smurfness

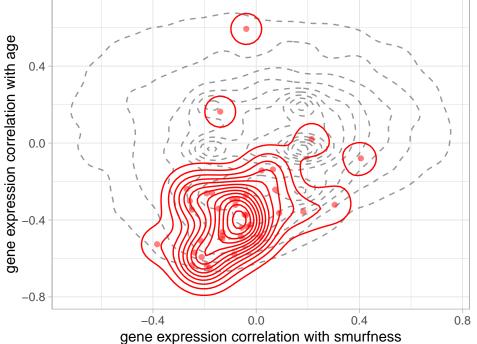




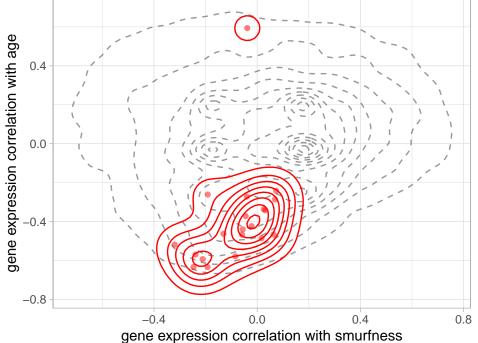
dme03410 Base excision repair gene expression correlation with age 0.5 0.0 -0.5-0.40.0 0.4 8.0

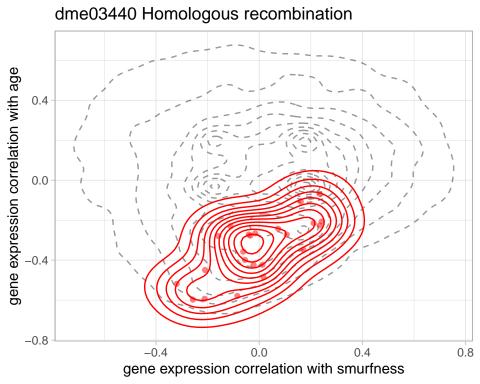
gene expression correlation with smurfness

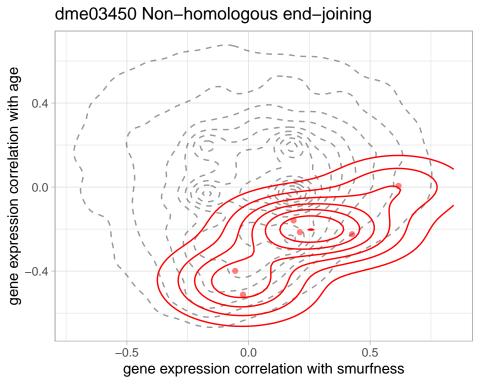
dme03420 Nucleotide excision repair



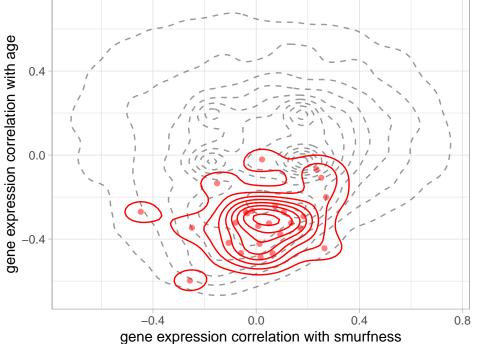
dme03430 Mismatch repair

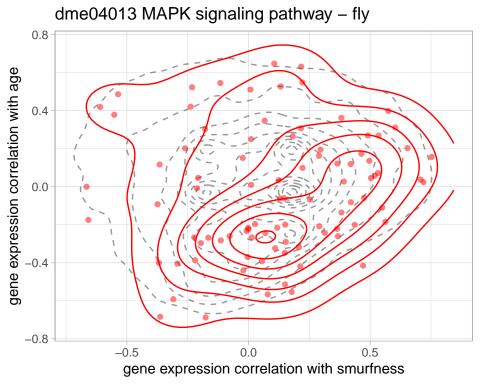


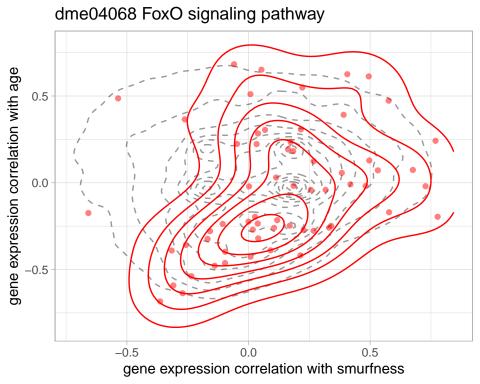


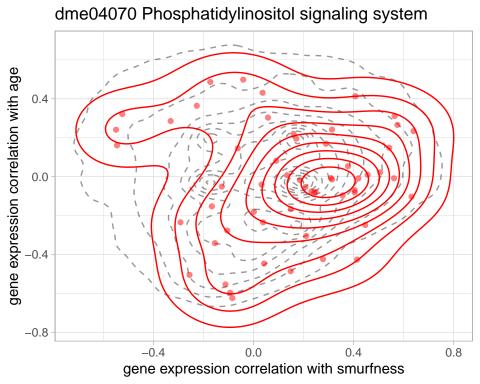


dme03460 Fanconi anemia pathway

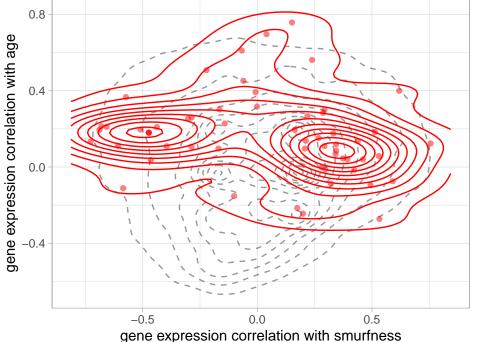




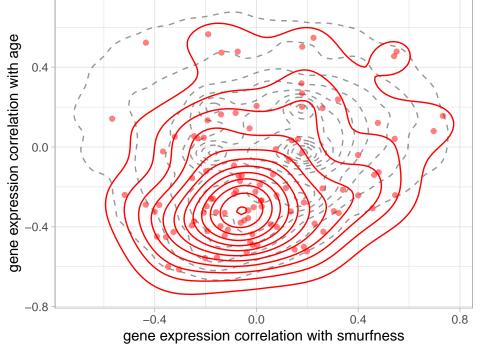




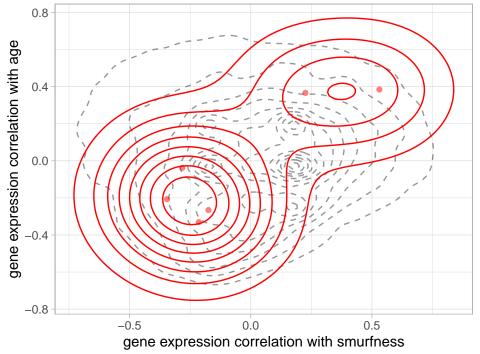
dme04080 Neuroactive ligand-receptor interaction



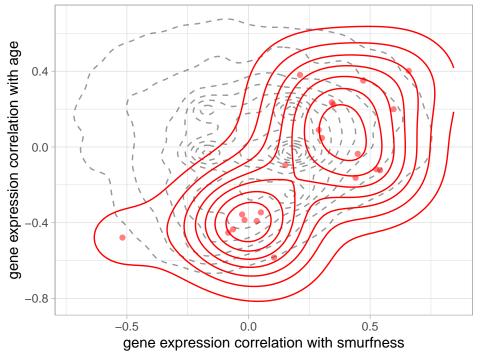
dme04120 Ubiquitin mediated proteolysis

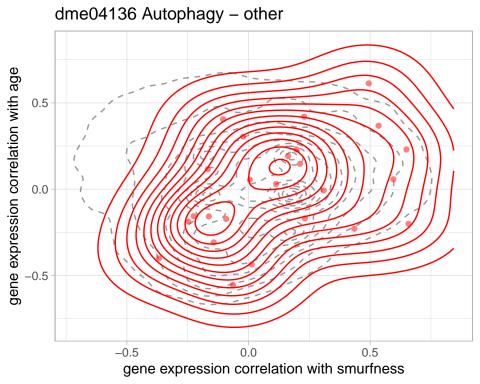


dme04122 Sulfur relay system

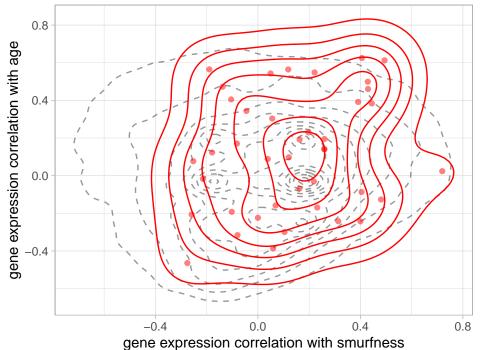


dme04130 SNARE interactions in vesicular transport

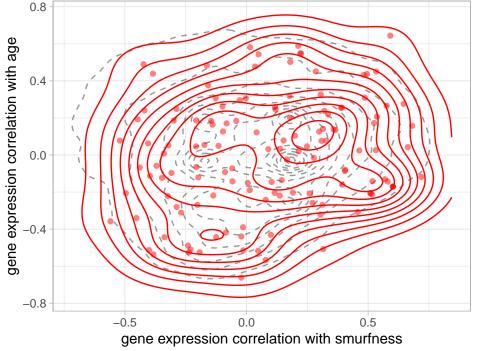


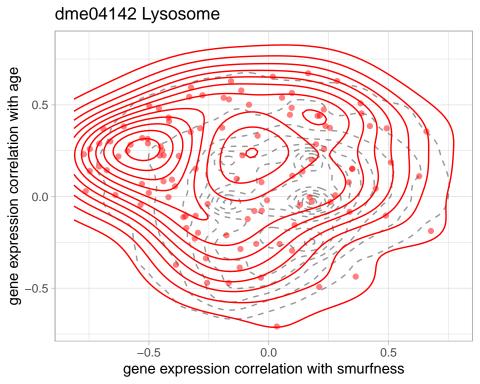


dme04137 Mitophagy - animal



dme04140 Autophagy - animal 8.0 gene expression correlation with age 0.4 0.0 -0.50.0 0.5 gene expression correlation with smurfness dme04141 Protein processing in endoplasmic reticulum





dme04144 Endocytosis gene expression correlation with age 0.5 0.0 -0.50.0 0.5 gene expression correlation with smurfness

dme04145 Phagosome 8.0 0.4 0.0 -0.4

gene expression correlation with age

0.0 gene expression correlation with smurfness

0.5

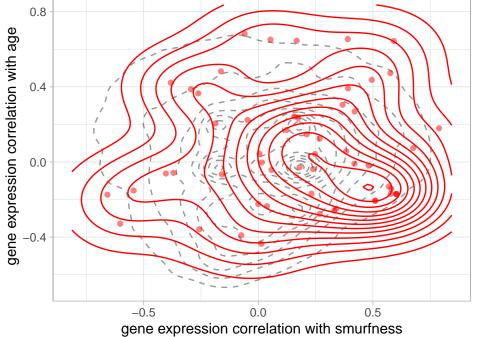
-0.5

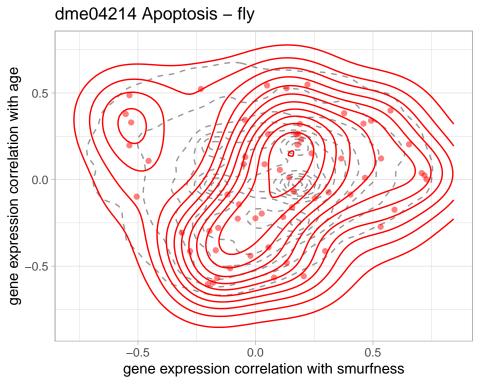
dme04146 Peroxisome gene expression correlation with age 0.5 0.0 -0.5-0.50.0 0.5

gene expression correlation with smurfness

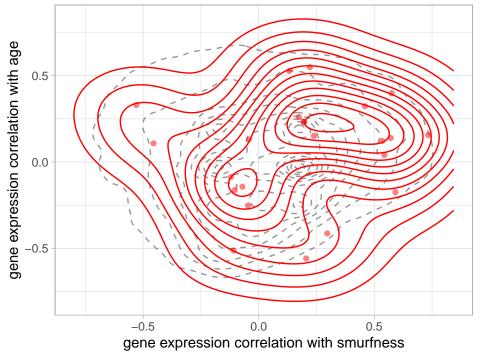
dme04150 mTOR signaling pathway gene expression correlation with age 0.4 0.0 -0.8-0.40.0 0.4 8.0 gene expression correlation with smurfness

dme04213 Longevity regulating pathway – multiple spec





dme04215 Apoptosis – multiple species



dme04310 Wnt signaling pathway gene expression correlation with age 0.4 0.0 -0.80.0 0.4 8.0 -0.4gene expression correlation with smurfness

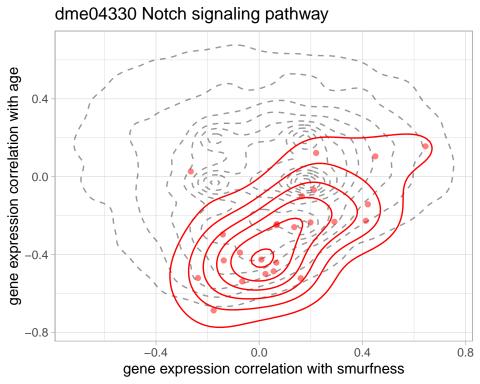
dme04320 Dorso-ventral axis formation 0.5 0.0 -0.5

gene expression correlation with age

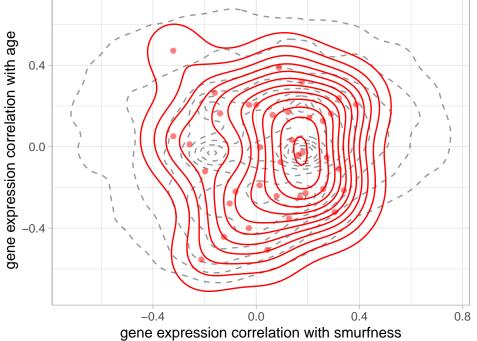
0.0 gene expression correlation with smurfness

0.5

-0.5



dme04341 Hedgehog signaling pathway – fly



dme04350 TGF-beta signaling pathway gene expression correlation with age 0.5 0.0 -0.5-0.40.0 0.4 8.0

gene expression correlation with smurfness

dme04391 Hippo signaling pathway - fly 0.5 0.0

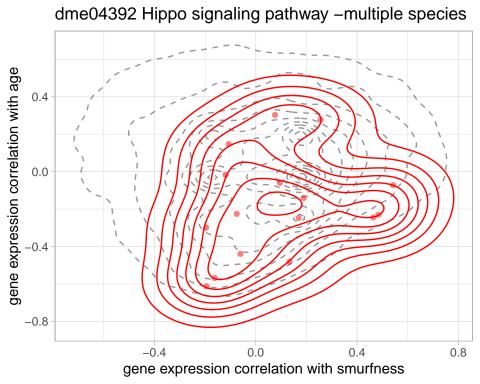
0.0

gene expression correlation with age

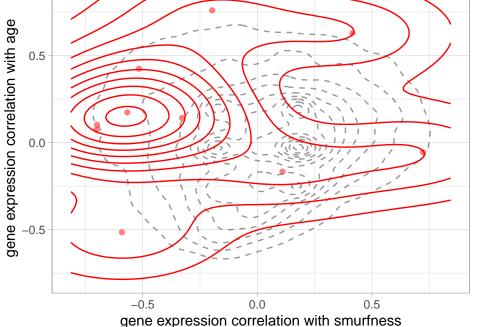
-0.5

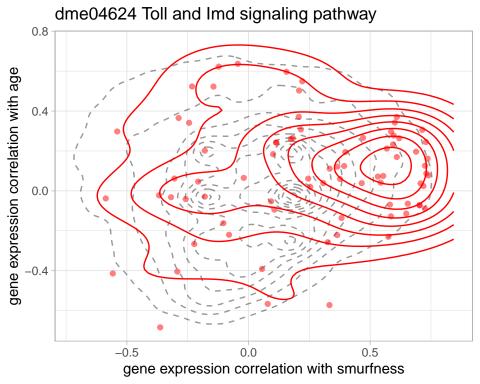
gene expression correlation with smurfness

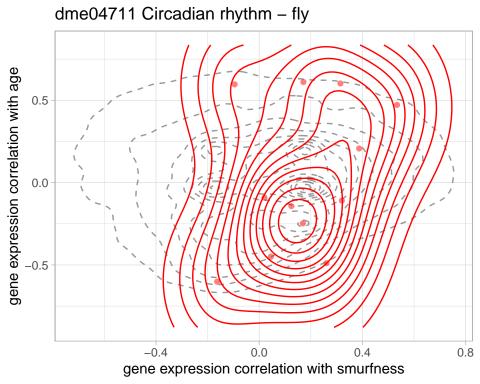
0.5



dme04512 ECM-receptor interaction 0.5 0.0







dme04745 Phototransduction - fly gene expression correlation with age 0.4 0.0

-0.4-0.40.0 0.4 8.0 gene expression correlation with smurfness