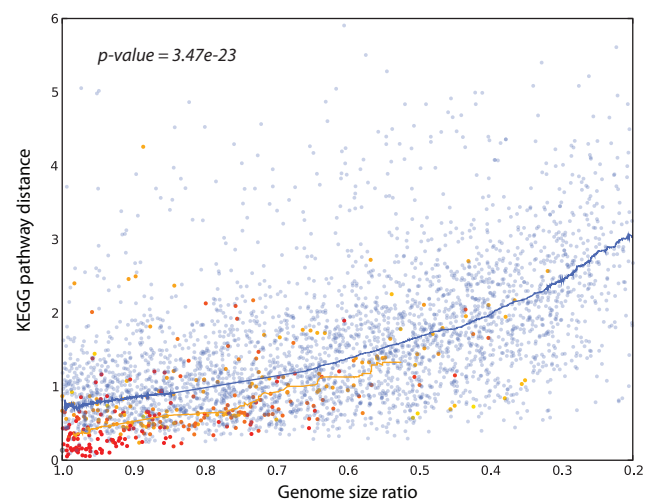
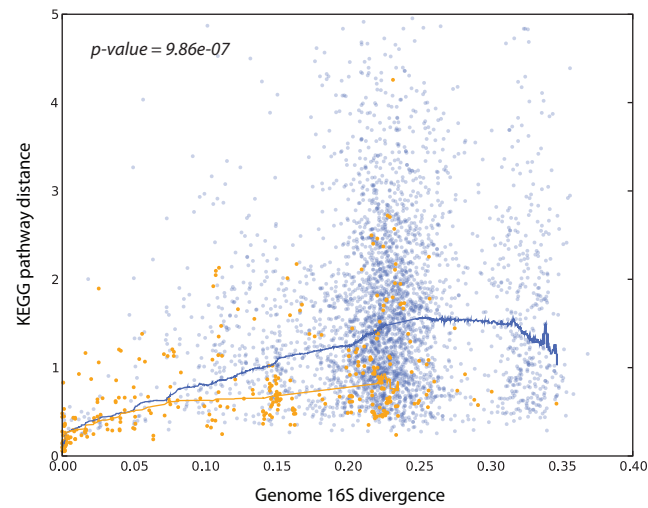
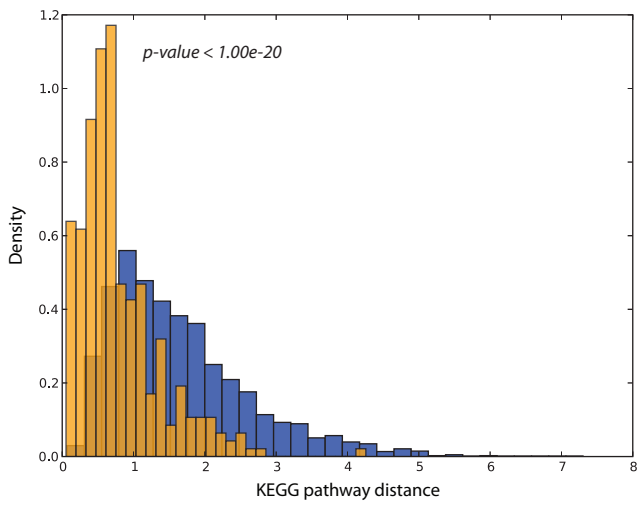
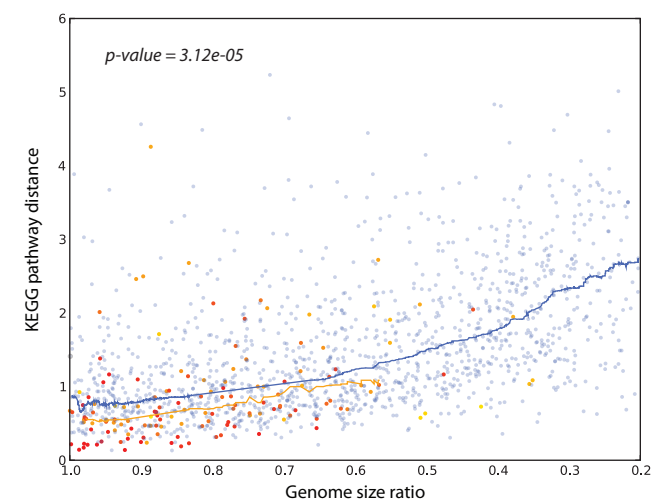
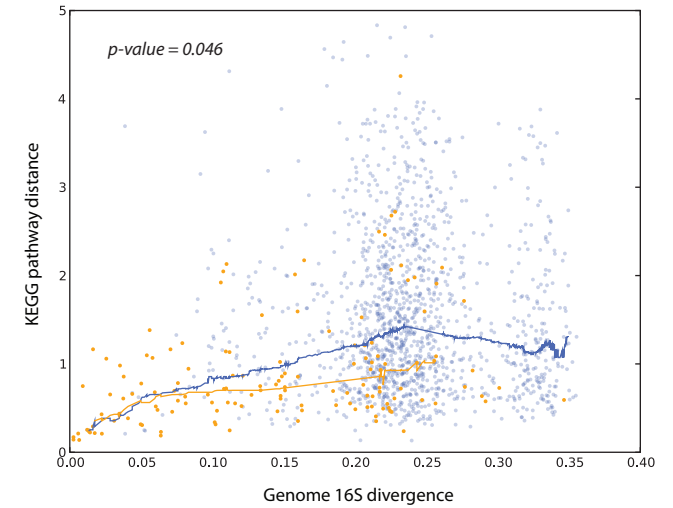
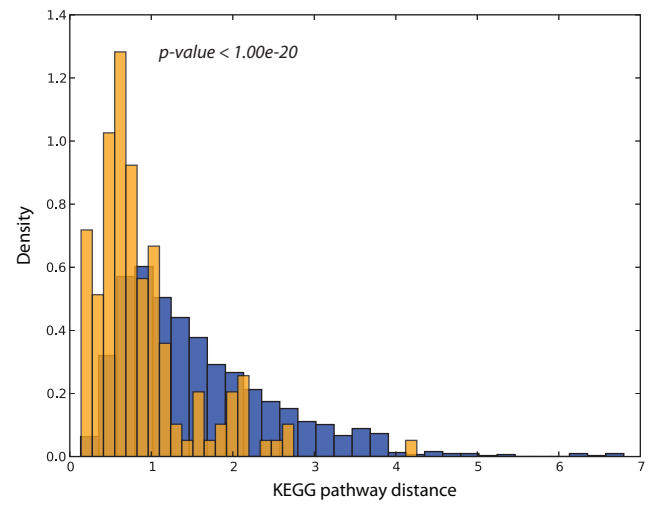




Strategy A:

considering all 16S genes in a genome,
mapping genomes to OTU by best identity hit
(min. length 700 nt, min. identity 97%)
599 genomes mapped



Strategy B:

considering all 16S genes in a genome,
best-reciprocal-hit mapping with OTUs
(based on identity, min. 05%; min. length 700 nt)
447 genomes mapped