Download the TCGA-LUAD dataset and the GEO dataset GSE68465 and collate them using 'merge.pl', 'moveMutFiles.pl', 'getClinical.pl' and 'probe2symbol.pl', TMB via 'TMb.pl'. Download the fatty acid metabolism gene "gene.txt" from MsigDB, <https://www.gsea-msigdb.org/gsea/msigdb/.> All subsequent analysis methods are presented using R files