



Supplemental Figure S4: Tissue-specific expression patterns of methionine catabolic genes gamma-lyase (*MGL*) and S-adenosyl-methionine synthases (*SAM-1*, *SAM-2*, *SAM-3*, *SAM-4*): Expression analysis was performed by using publicly available data from the Genevestigator web site (www.genevestigator.ethz.ch) and plotted to compare gene expression among various tissues for all the genes. Error bars represent standard deviations.