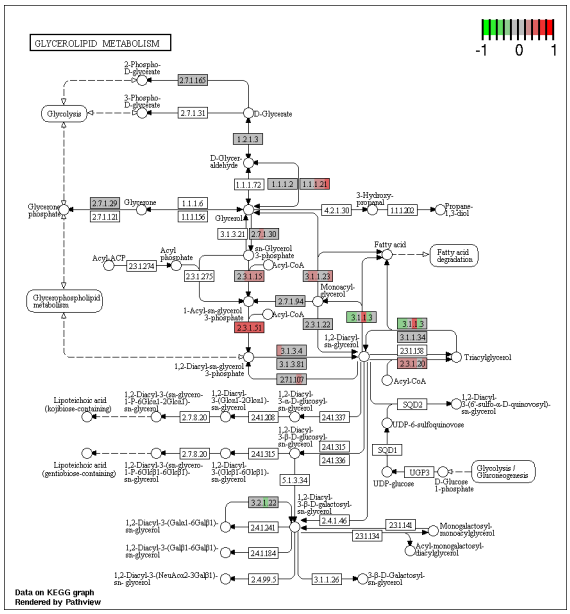


KEGG view representation of Glycerolipid metabolism (mmu00561). Box of expressed genes are colored according to the scaled log2FC between epi and ing expression levels. Each box is partioned into six parts, each representing one time point (-2,0,2,4,6,8), so that the whole time course for the gene is shown. The log2FC is scaled between -1 to 1. When the gene is higher expressed in ing the color is in a shade of green, while it is red, when the gene is higher expressed in epi. Grey color indicates a similar expression intensity in both adipose tissues, and no filling means no expression was measured.