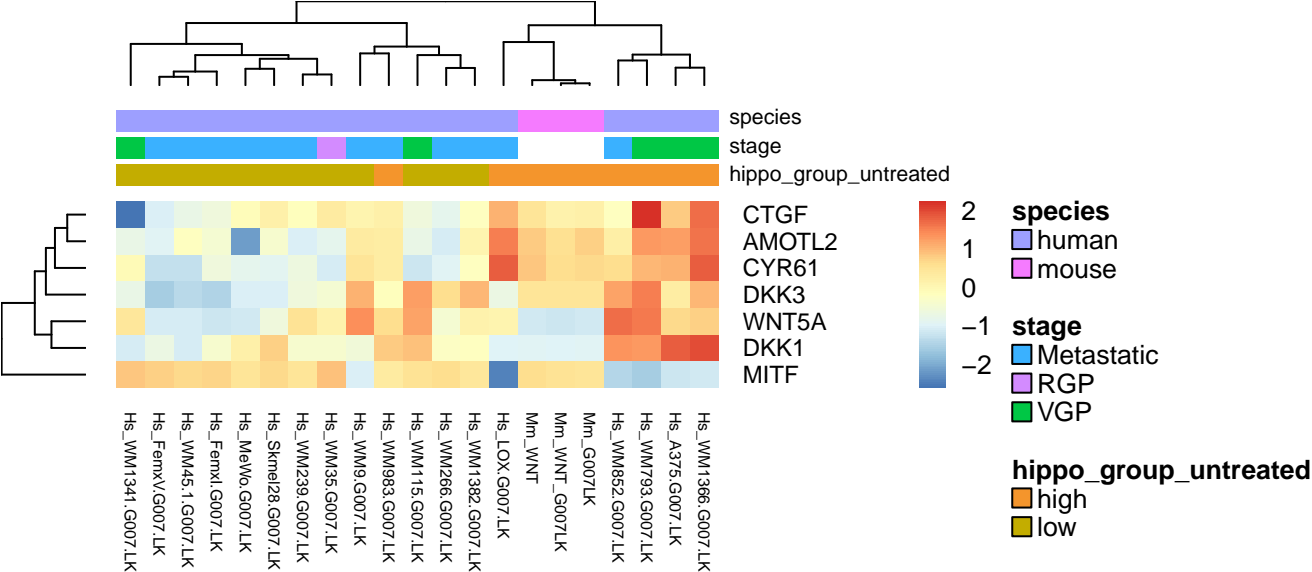


57_treated_v2, MITF_and_HIPPO_and_more, genes=7



samples: G007LK and also WNT_G007LK and WNT for mouse
genes: MITF and in comparison with the HIPPO genes and a few more
value: log2 of TPM
zscorecolors: row