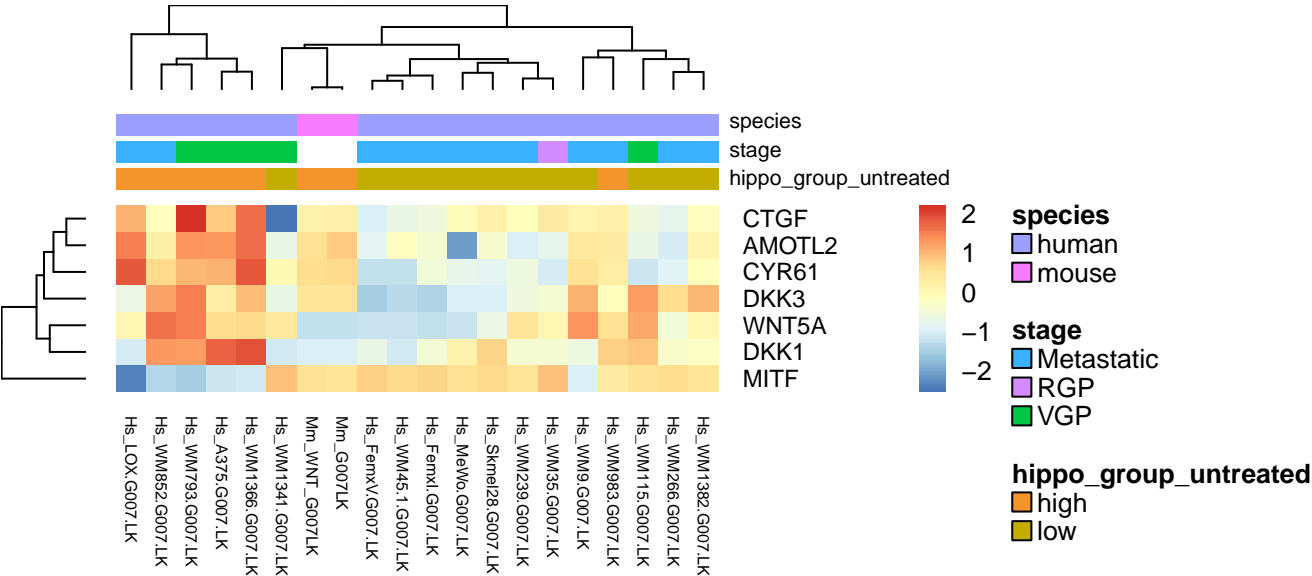


42_treated_v1, MITF_and_HIPPO_and_more, genes=7



samples: G007LK and also WNT_G007LK for mouse

genes: MITF and in comparison with the HIPPO genes and a few more

value: log2 of TPM

zscorecolors: row