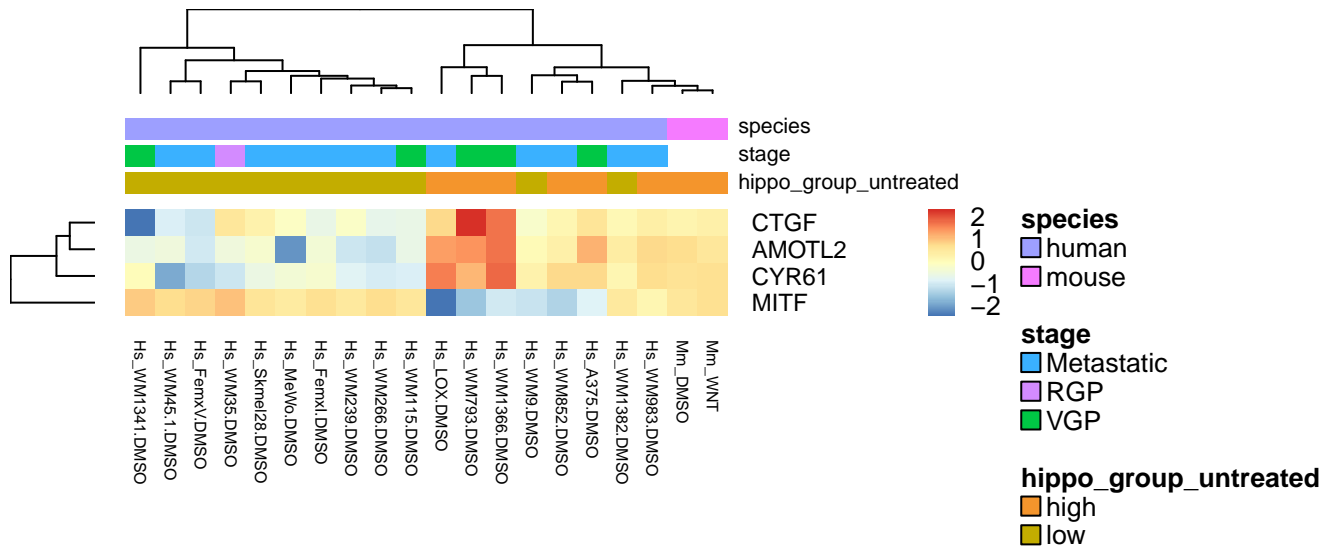


26_untreated_v2, MITF_and_HIPPO, genes=4



samples: DMSO and also WNT for mouse

genes: MITF in comparison with the HIPPO genes

value: log2 of TPM

zscorecolors: row