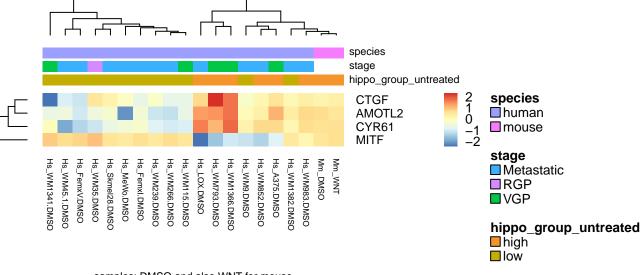
26_untreated_v2, MITF_and_HIPPO, genes=4



samples: DMSO and also WNT for mouse genes: MITF in comparison with the HIPPO genes value: log2 of TPM zscorecolors: row