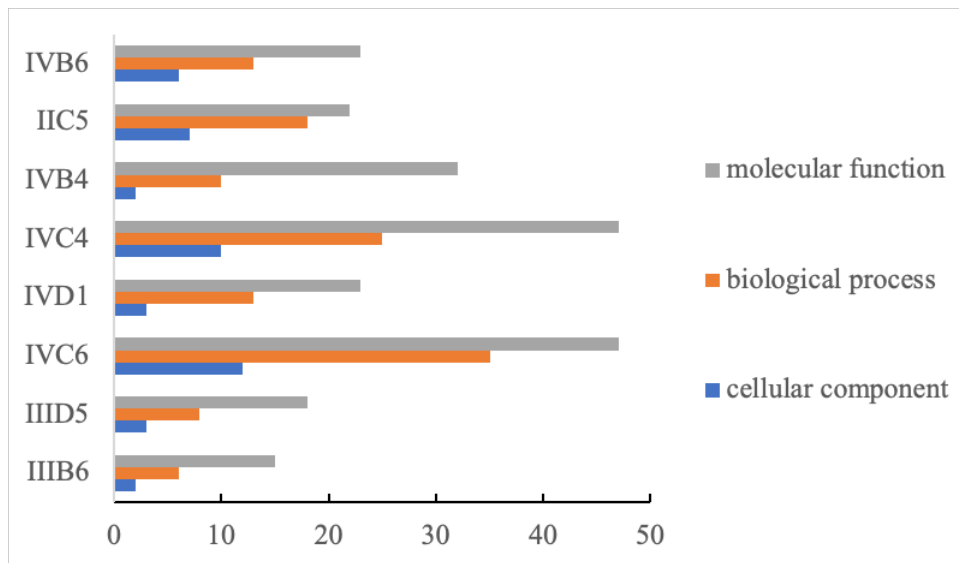




Supplementary Figure 2. GO counts for genes of each cell line. RNA-seq analysis identified 676 genes with annotated function. Of them, 36 genes were expressed commonly in all the eight lines while 640 genes were expressed preferentially and/or specifically in a certain line. Namely, 54 gene are specific to IVB6, 71 to IIC5, 61 to IVB4, 120 to IVC4, 53 to IVD1, 146 genes to IVC6, 38 to IIID5, and 31 to IIIB6. Those genes are listed up in Supplementary Table 1. Most cell lines show a similar ratio of GO categories.