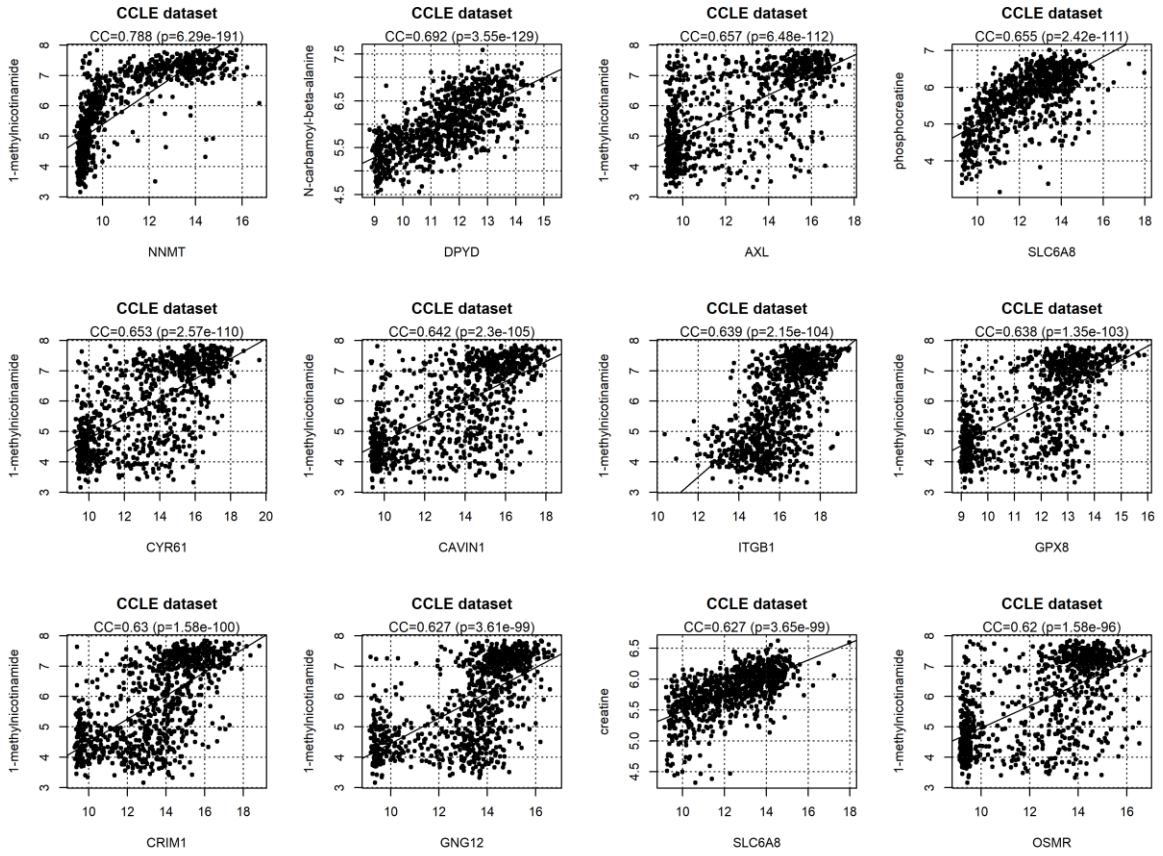


Figure S1: correlation between metabolite levels and gene expression in the CCLE dataset, showing the 12 pairs with the highest correlation coefficient.