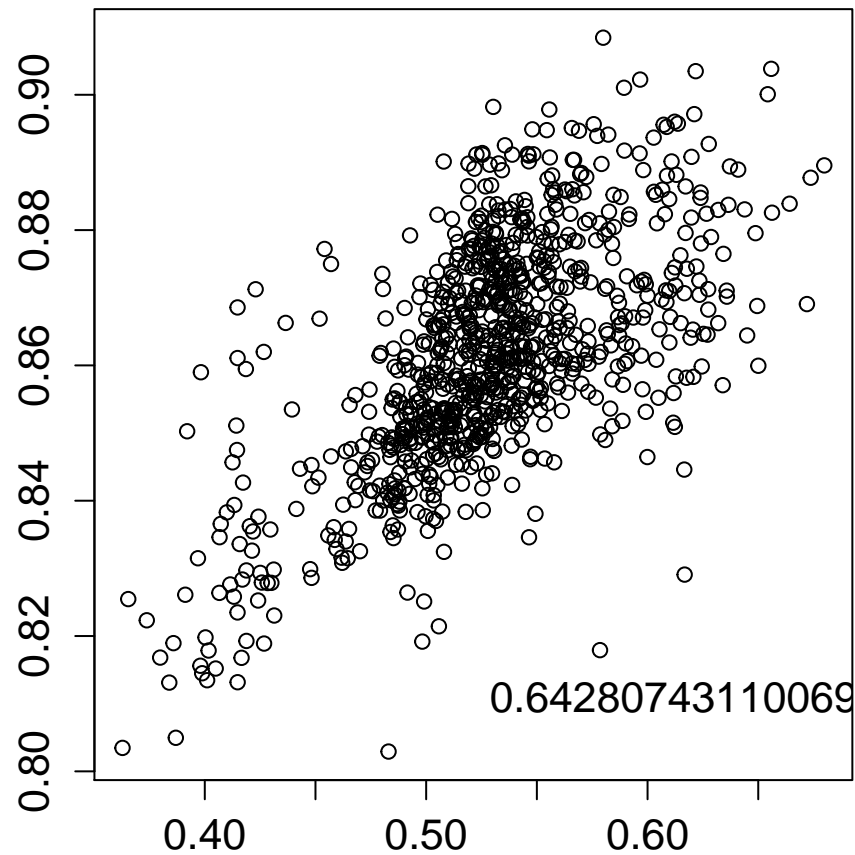
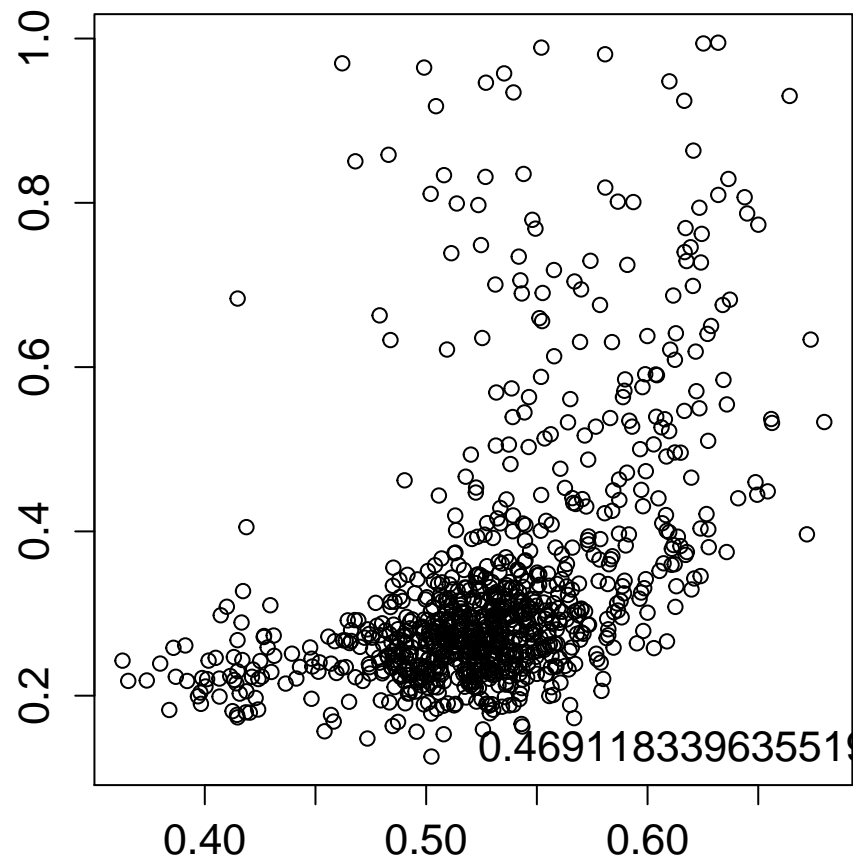


Accuracy of model reconstruction



N° rxns added for tasks



n gene expression between included vs excluded gem gene expression between included vs excluded gem