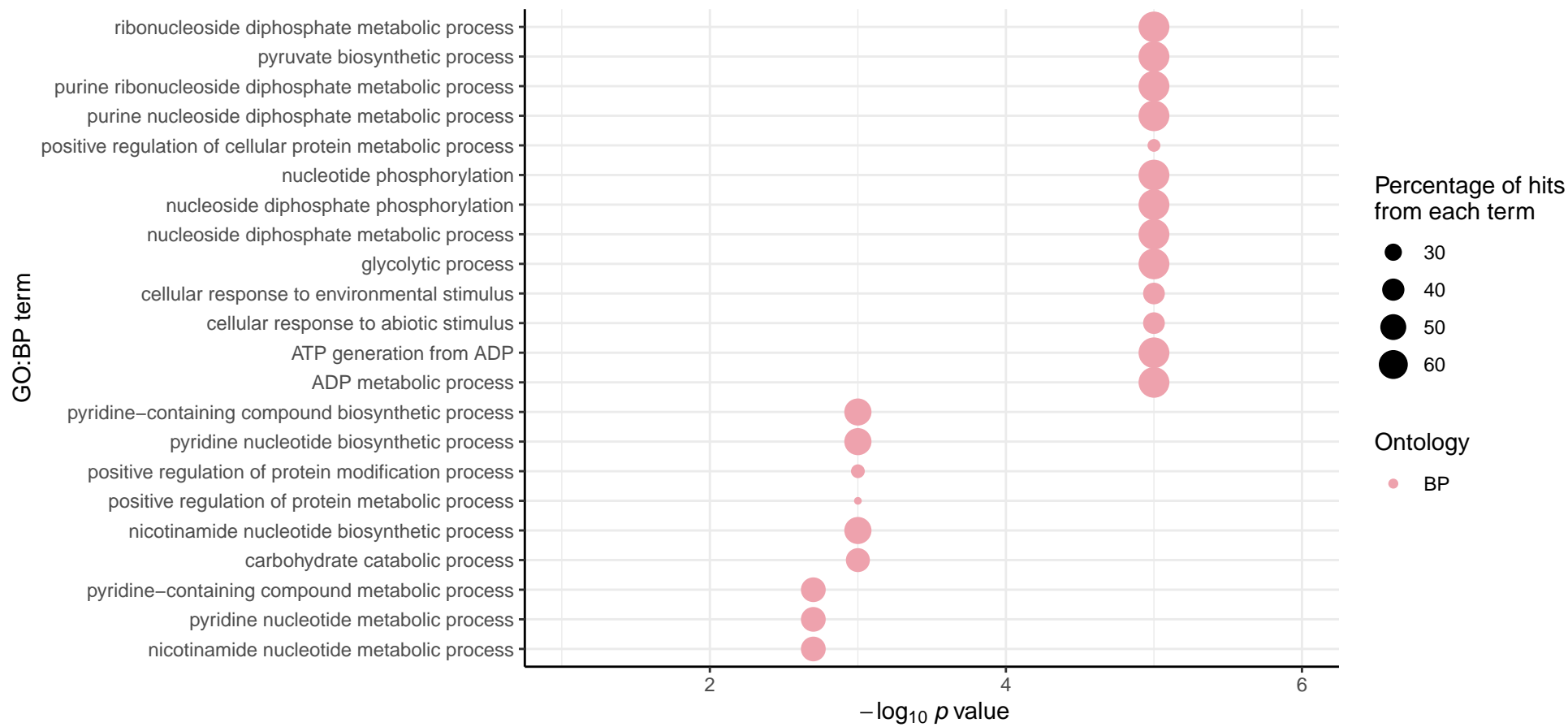


D

GO analysis on top 200 most time:layer-variable genes



E

GO analysis on top 200 most residuals-variable genes

