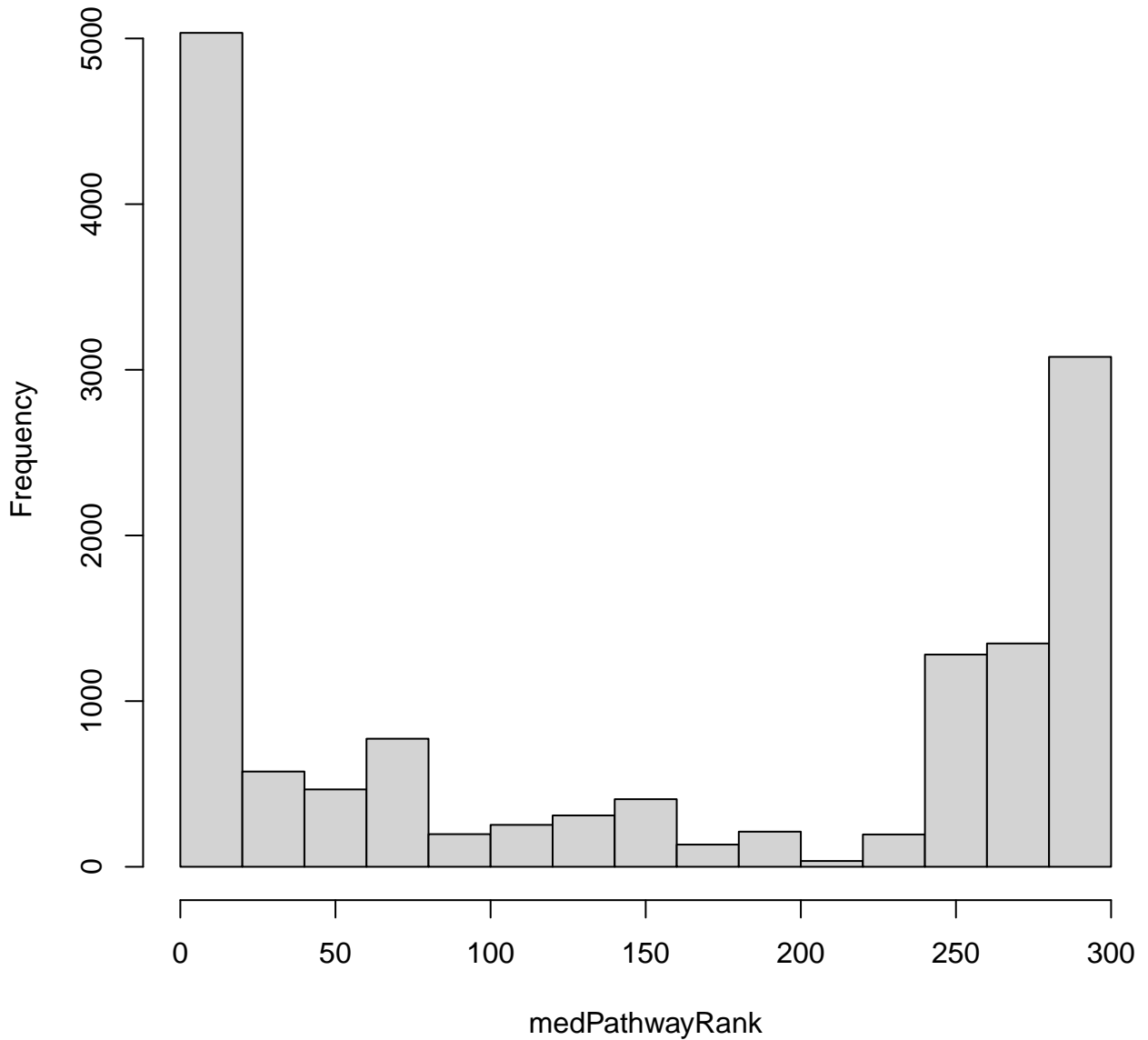
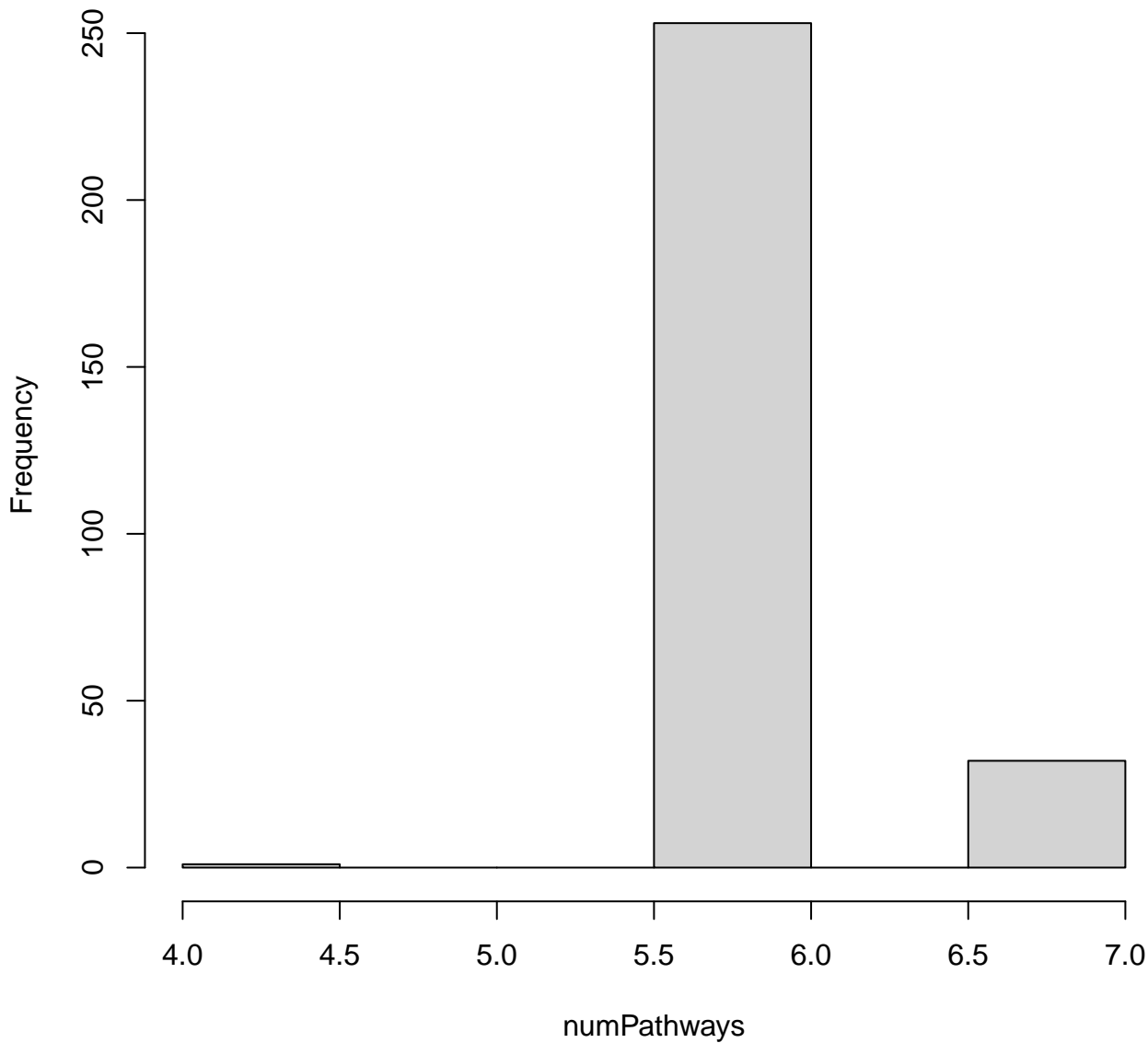


## Distribution of all P Values



## Number of Pathways overexpressed



## Distribution of the fold change applied to the cells

