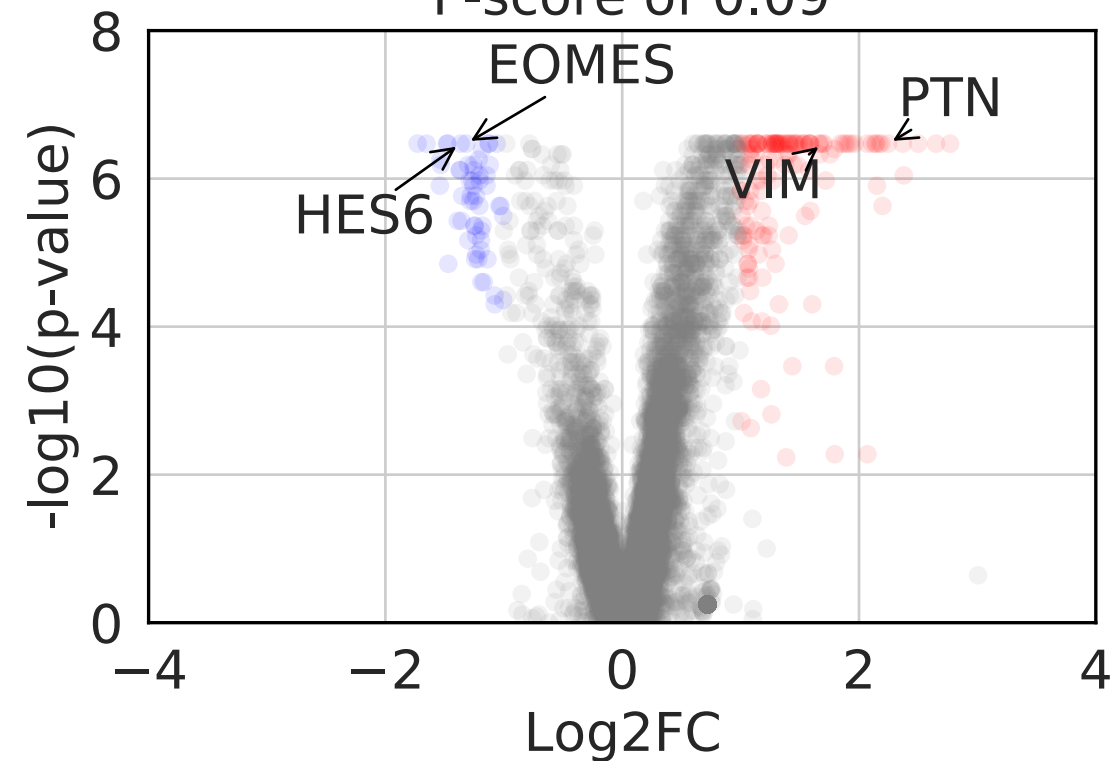
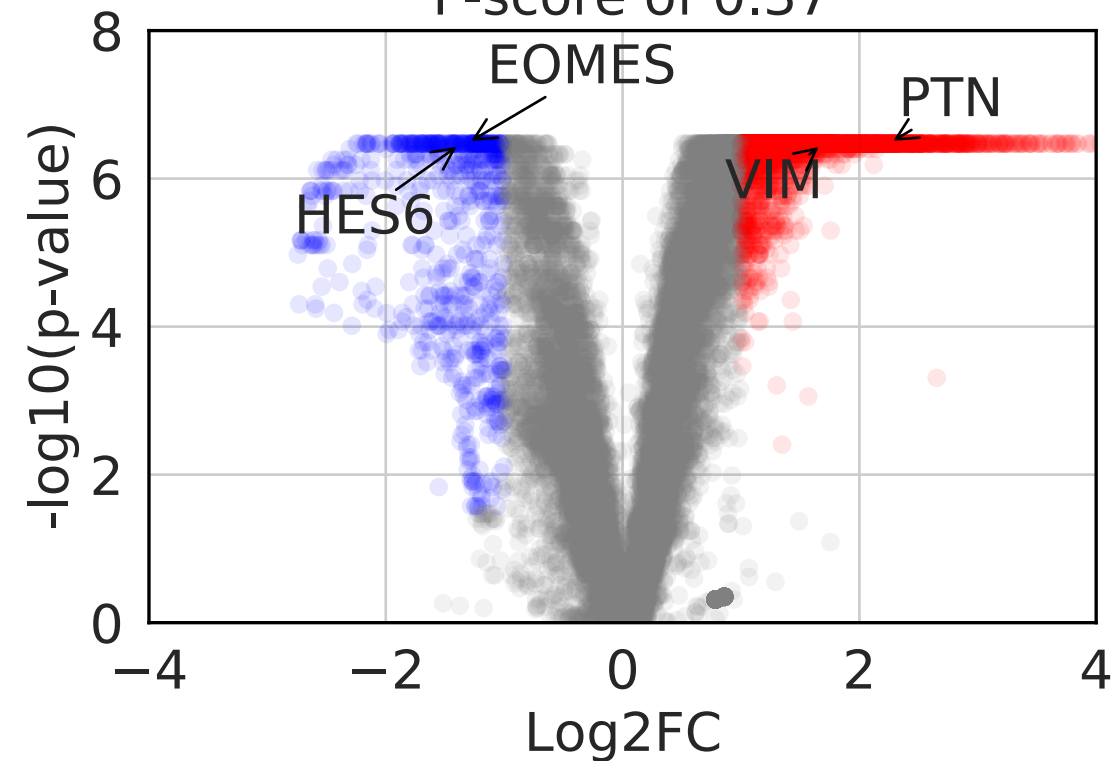


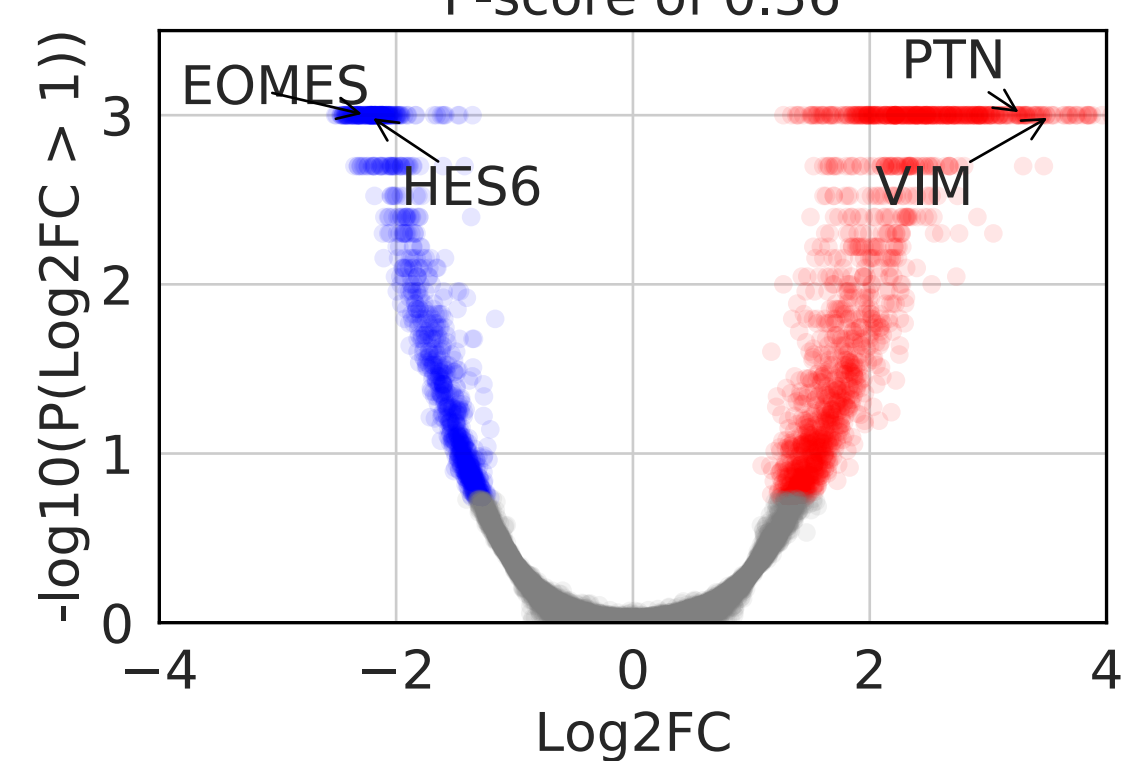
Wilcoxon Rank Sum Test on  
Uncorrected Counts (196 genes  
with  $\log_2FC > 1$  and  $FDR < 0.05$ )  
F-score of 0.09



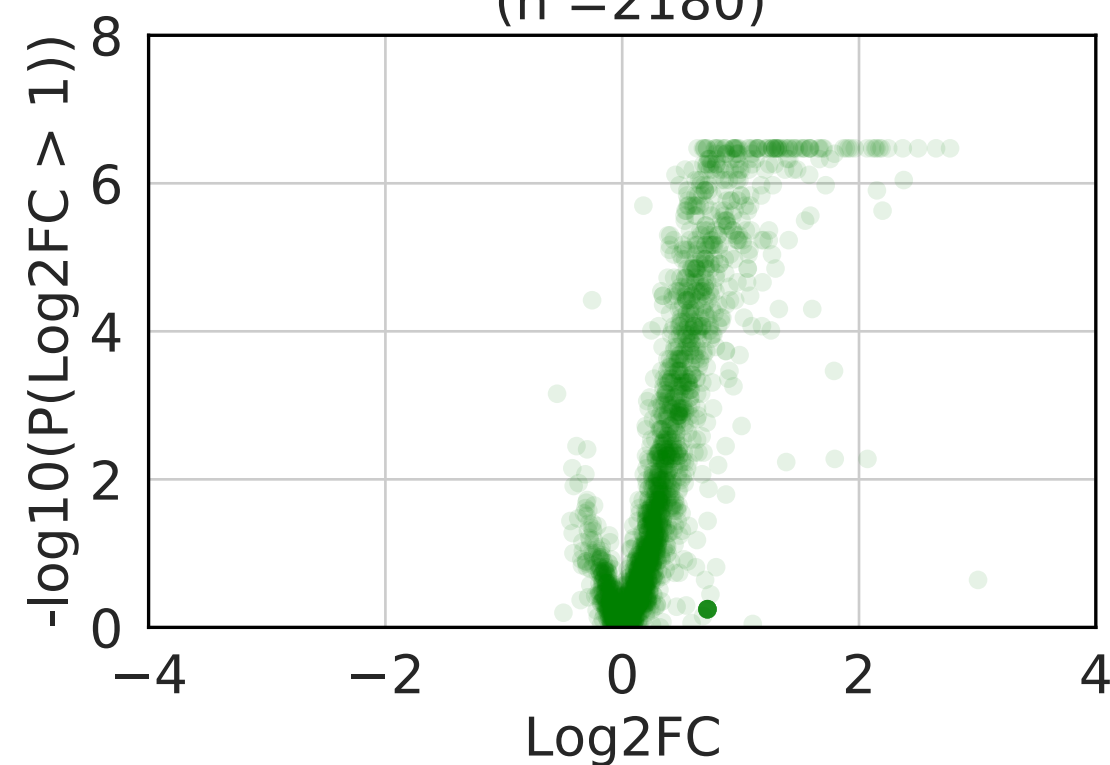
Wilcoxon Rank Sum Test on  
Corrected Counts (3334 genes  
with  $\log_2FC > 1$  and  $FDR < 0.05$ )  
F-score of 0.37



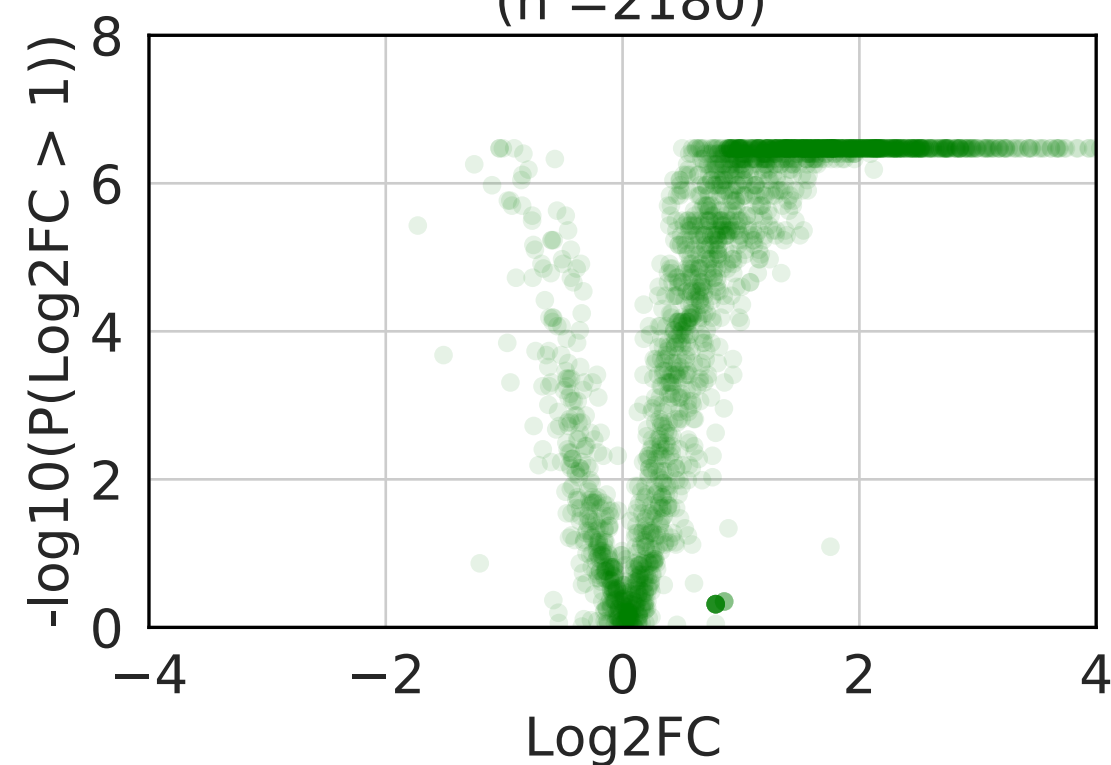
Probabilistic Generative Model  
(1334 genes with  $\log_2FC > 1$  and  $FDR < 0.05$ )  
F-score of 0.36



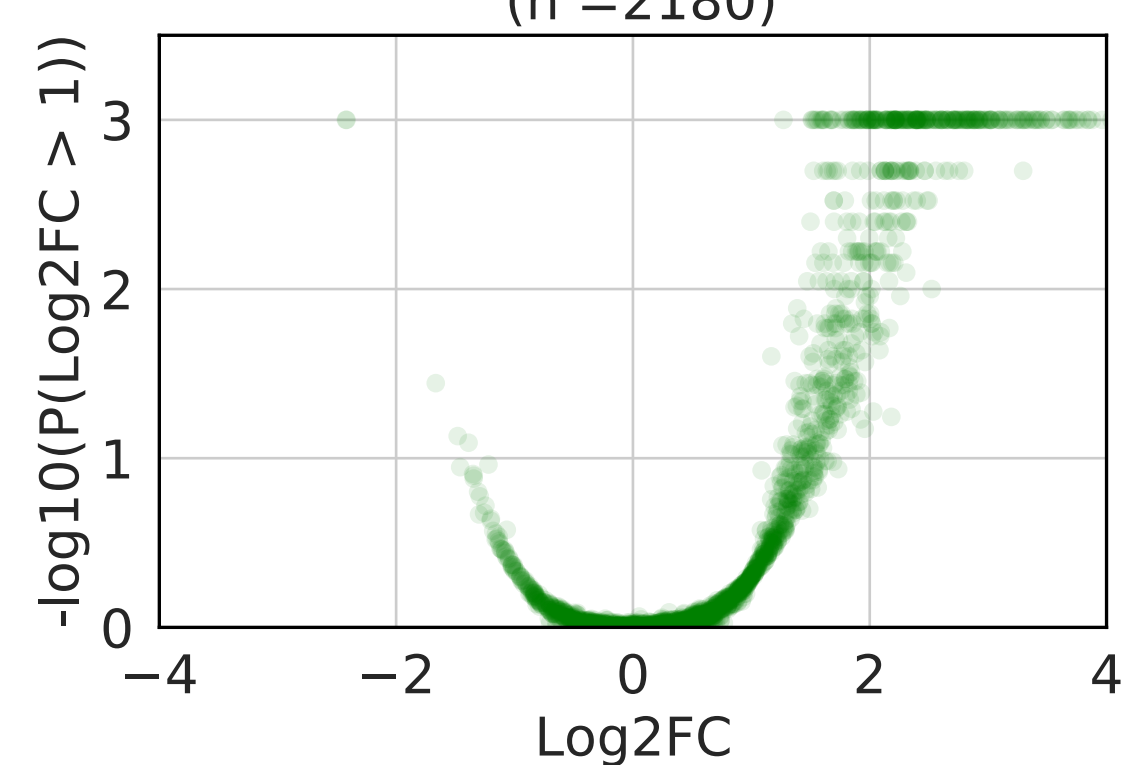
HOPX+ Reference Genes  
(n = 2180)



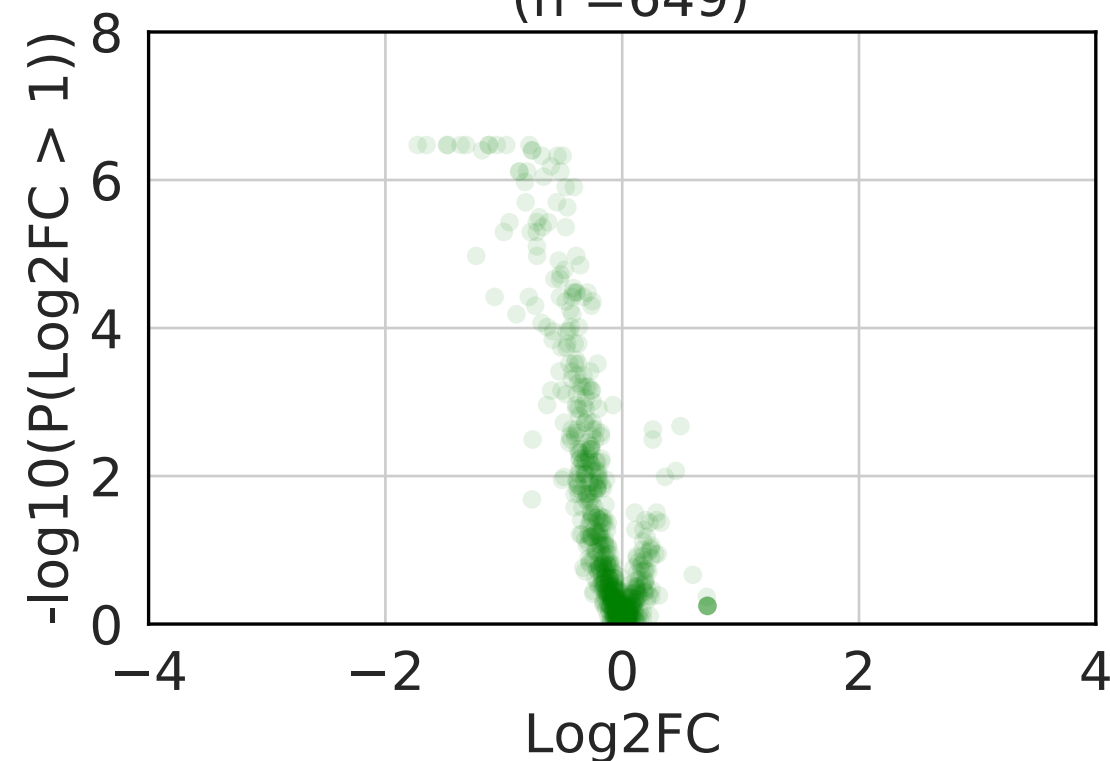
HOPX+ Reference Genes  
(n = 2180)



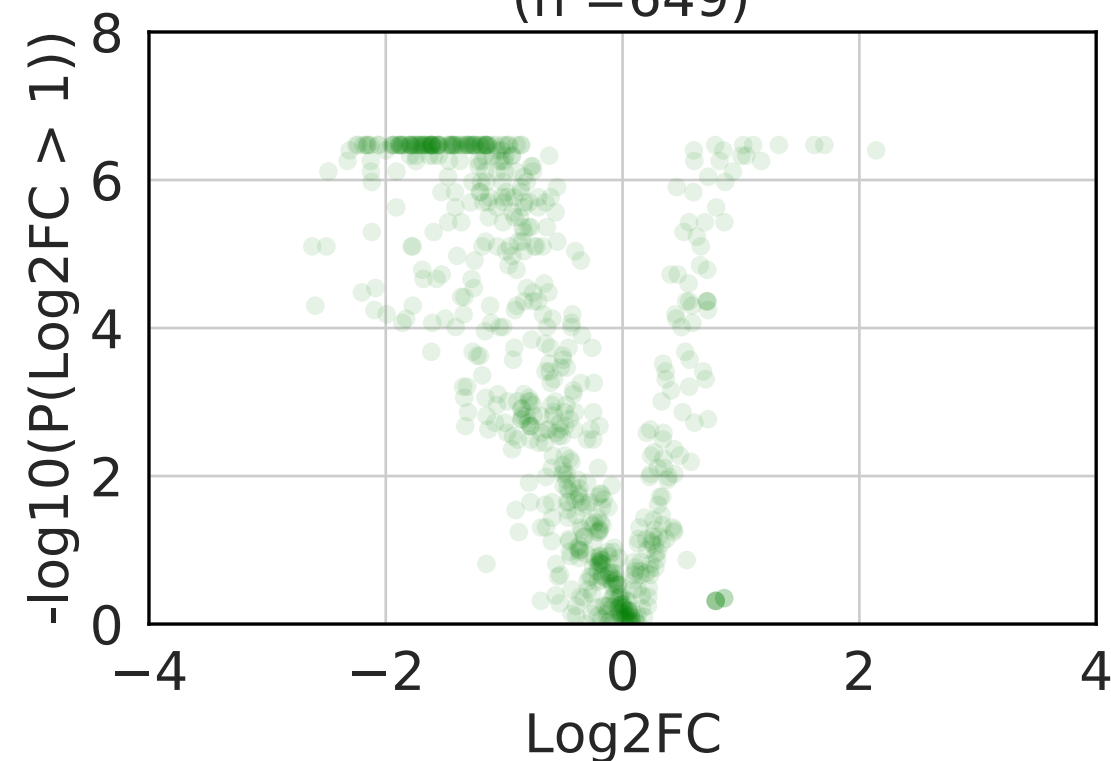
HOPX+ Reference Genes  
(n = 2180)



EOMES+ Reference Genes  
(n = 649)



EOMES+ Reference Genes  
(n = 649)



EOMES+ Reference Genes  
(n = 649)

