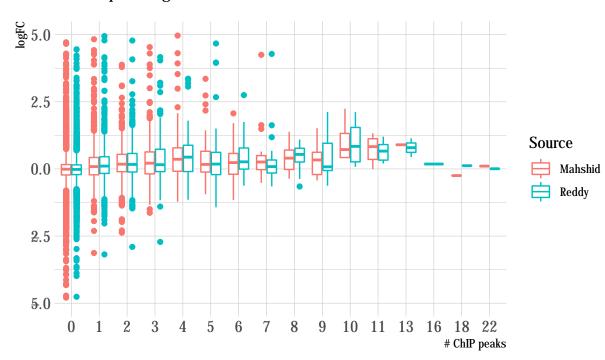
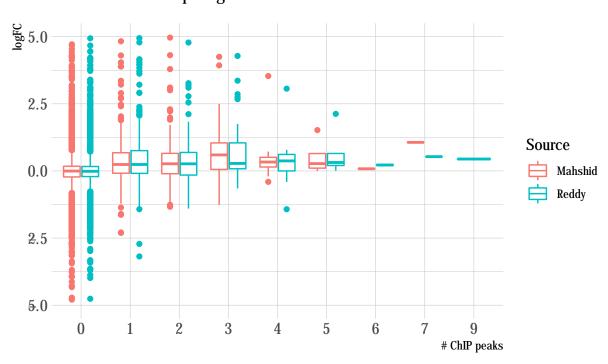
## Distribution of logFC by number of nearby GRpeaks

Closest peak to gene no matter distance

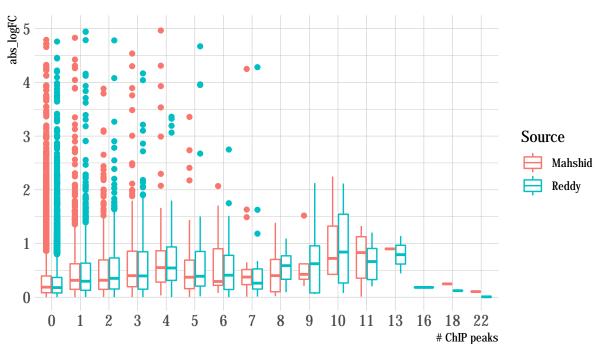


## Peaks within 2500bp range



## Distribution of ABSOLUTE logFCs by number of nearby GRpeaks

Closest peak to gene no matter distances



## Peaks within 2500bp range

