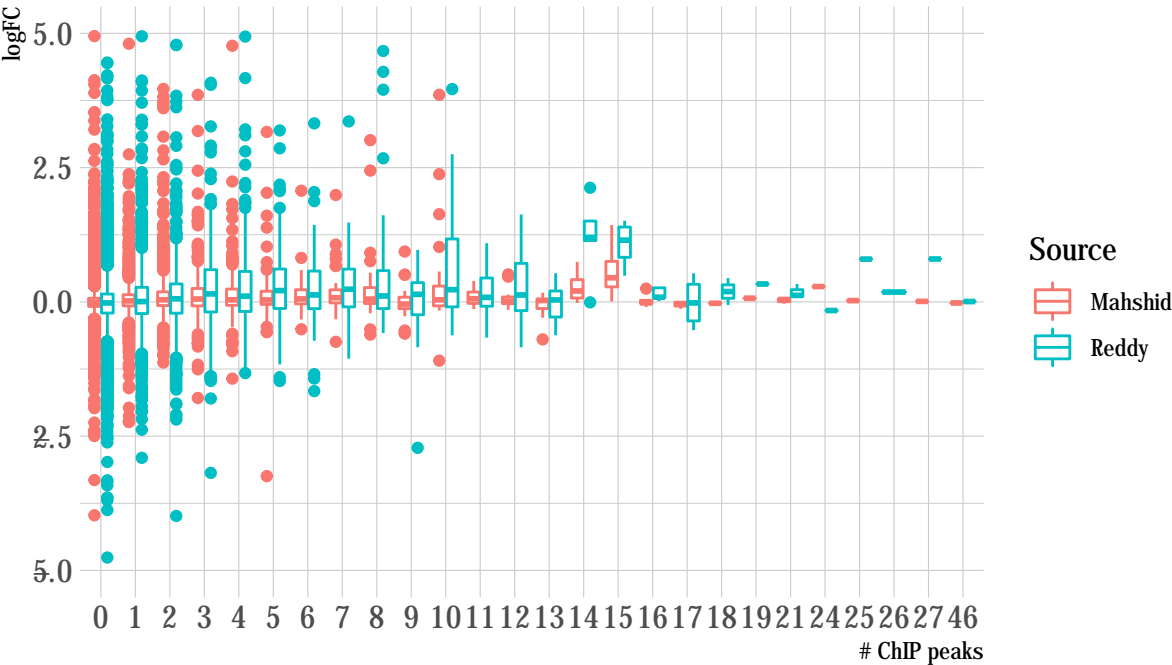
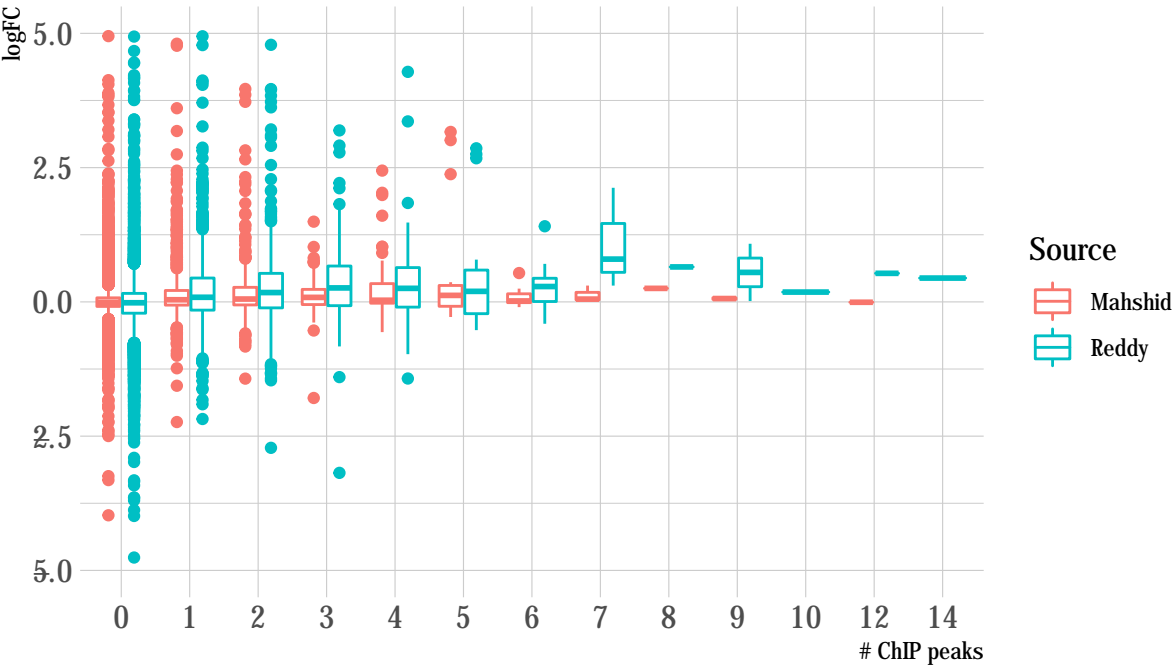


Distribution of logFC by number of nearby GRpeaks

Closest peak to gene no matter distance

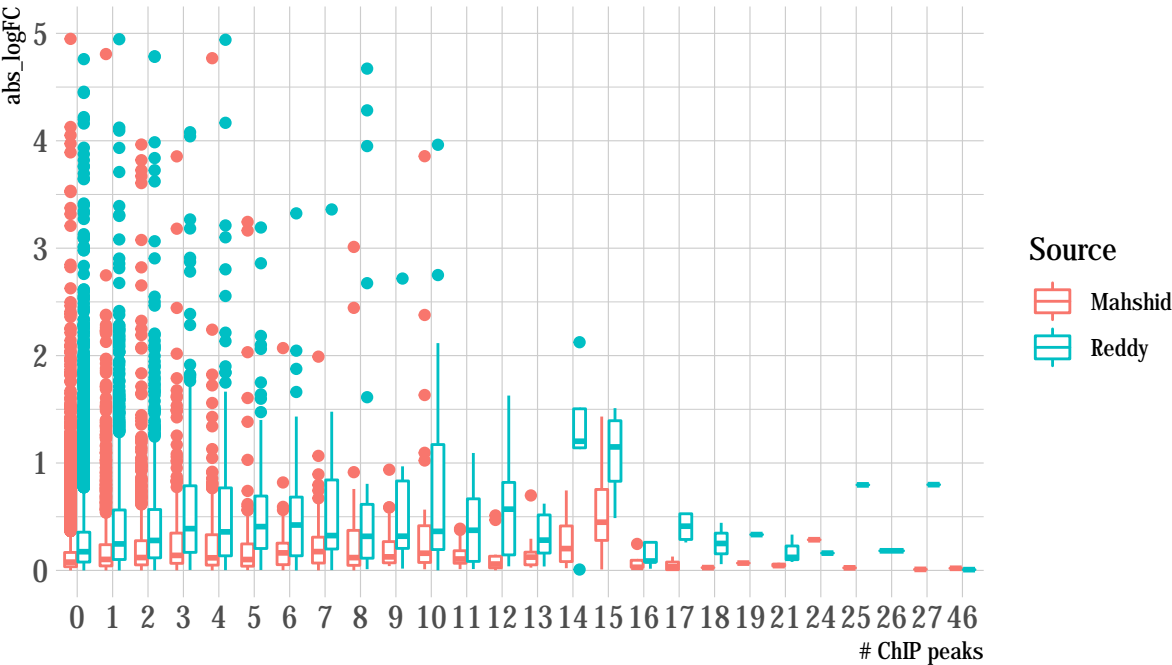


Peaks within 2500bp range



Distribution of ABSOLUTE logFCs by number of nearby GRpeaks

Closest peak to gene no matter distances



Peaks within 2500bp range

