# STAT1 SCEPTRE vs Seurat With Pvalue QC Summary

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## Introduction

In this writeup I will change the method of QC for the CHIP-seq data to be such that peaks are only called if they are in the top $\alpha$ percentile of peaks with respect to the pvalue reported. I will return the odds ratios that SCEPTRE and Seurat give with respect to the post-QC chipseq data. For more information such as the actual number of discoveries, see "STAT1-analysis-pval-QC-individual".

Table 1: Enrichment odds ratios, comparing to ChIP-seq target assignments with changing quantile threshold

| Ground truth \ Method | SCEPTRE | Seurat | SCEPTRE-pval | Seurat-pval |
|---|---|---|---|---|
| 0.1 | 1.36 | 1.28 | $6.67e-12$ | $8.69e-09$ |
| 0.25 | 1.30 | 1.23 | $4.47e-10$ | $4.37e-07$ |
| 0.5 | 1.25 | 1.16 | $3.61e-09$ | $5.23e-05$ |
| 0.75 | 1.31 | 1.24 | $3.36e-12$ | $9.97e-09$ |