**Conserved Divergent gene-CUT Pairs**

A. 

- **Average WT RNA-seq Coverage**
  - All Genes: N=4416
  - Conserved Divergent: N=375
  - p < 5x10^-6

- **Average rrp6Δ RNA-seq Coverage**
  - All Genes: N=4416
  - Conserved Divergent: N=375
  - p < 7x10^-6

- **Average Fold Change**
  - All Genes: N=4416
  - Conserved Divergent: N=375
  - p < 0.05

B. 

- **Average WT RNA-seq Coverage**
  - All Genes: N=4243
  - Conserved Divergent: N=366
  - p < 5x10^-6

- **Average rrp6Δ RNA-seq Coverage**
  - All Genes: N=4243
  - Conserved Divergent: N=366
  - p < 2x10^-7

- **Average Fold Change**
  - All Genes: N=4243
  - Conserved Divergent: N=366
  - p < 0.007

C. 

- **Average WT RNA-seq Coverage**
  - All Genes: N=4161
  - Conserved Divergent: N=365
  - p < 5x10^-6

- **Average rrp6Δ RNA-seq Coverage**
  - All Genes: N=4161
  - Conserved Divergent: N=365
  - p < 7x10^-5

- **Average Fold Change**
  - All Genes: N=4161
  - Conserved Divergent: N=365
  - p < 0.06

**Σ1278b**

**JAY291**

**S. paradoxus (N17)**