

## Lnc RNAs: All

```
lncRNA group = IDRlessthan0_2 ::: data cols = bothPullDowns
```

PCA plot showing the relationship between log-transformed gene expression data (log(First Component) vs log(Second Component)). The plot displays two main clusters of data points, colored by the 'withinSubset' variable: red circles for 'rest' and teal triangles for 'true'.

Key labeled points include:

- 230937 (rest)
- 240647 (rest)
- 240647 (true)
- 231863 (rest)
- 231863 (true)
- 232389 (rest)
- 232389 (true)
- 245532 (true)
- 247844 (true)
- 254964 (rest)
- 237643 (rest)
- 249790 (rest)
- 234883 (rest)
- 234883 (true)
- 254964 (true)
- 237643 (true)
- 249790 (true)
- 1562 (true)

rest  
true