

**Table** GSE59364\_DC\_all.csv ...

**File Path**

`${org.campagnelab.metaR.home}/data/GSE59364_DC_all.csv`

**Columns**

```
gene: string [ ID ]
mRNA len: numeric [ << ... >> ]
genomic span: numeric [ << ... >> ]
DC_normal: numeric [ << ... >> ]
DC_treated: numeric [ << ... >> ]
DC0904: numeric [ LPS=NO, counts ]
DC0907: numeric [ LPS=NO, counts ]
DCLPS0910: numeric [ LPS=NO, counts ]
DCLPS0913: numeric [ LPS=NO, counts ]
A_DC: numeric [ LPS=NO, counts ]
A_DC_LPS: numeric [ LPS=YES, counts ]
B_DC: numeric [ LPS=NO, counts ]
B_DC_LPS: numeric [ LPS=YES, counts ]
C_DC: numeric [ LPS=NO, counts ]
C_DC_LPS: numeric [ LPS=YES, counts ]
C2DC: numeric [ LPS=NO, counts ]
C2DCLPS: numeric [ LPS=YES, counts ]
C3DC: numeric [ LPS=NO, counts ]
C3DCLPS: numeric [ LPS=YES, counts ]
```