```
Columns (38):
C3DC.GSE59364 DC all.csv: numeric [ LPS=NO, counts ]
C3DC.another Table: numeric [ LPS=NO, counts ]
C2DC.another Table: numeric [ LPS=NO, counts ]
DC0904.another Table: numeric [ LPS=NO, counts ]
mRNA len.another Table: numeric [ << ... >> ]
B DC.GSE59364 DC all.csv: numeric [ LPS=NO, counts ]
mRNA len.GSE59364 DC all.csv: numeric [ << ... >> ]
B DC.another Table: numeric [ LPS=NO, counts ]
A DC.GSE59364 DC all.csv: numeric [ LPS=NO, counts ]
DCLPS0913.GSE59364 DC all.csv: numeric [ LPS=NO, counts ]
DC0904.GSE59364 DC all.csv: numeric [ LPS=NO, counts ]
C2DC.GSE59364 DC all.csv: numeric [ LPS=NO, counts ]
C DC LPS.GSE59364 DC all.csv: numeric [ LPS=YES, counts ]
DCLPS0910.another Table : numeric [ LPS=NO, counts ]
B DC LPS.another Table: numeric [ LPS=YES, counts ]
DC treated.another Table: numeric [ << ... >> ]
A DC LPS.another Table: numeric [ LPS=YES, counts ]
gene.GSE59364 DC all.csv: string [ ID ]
DC normal.another Table: numeric [ << ... >> ]
C3DCLPS.GSE59364_DC_all.csv: numeric [ LPS=YES, counts ]
C DC LPS.another Table: numeric [ LPS=YES, counts ]
B DC LPS.GSE59364 DC all.csv: numeric [ LPS=YES, counts ]
C2DCLPS.GSE59364_DC_all.csv: numeric [ LPS=YES, counts ]
DCLPS0910.GSE59364_DC_all.csv : numeric [ LPS=NO, counts ]
A DC LPS.GSE59364 DC all.csv: numeric [ LPS=YES, counts ]
DC_normal.GSE59364_DC_all.csv : numeric [ << ... >> ]
C3DCLPS.another Table: numeric [ LPS=YES, counts ]
C DC.another Table: numeric [ LPS=NO, counts ]
genomic span.GSE59364 DC all.csv : numeric [ << ... >> ]
C DC.GSE59364 DC all.csv: numeric [ LPS=NO, counts ]
C2DCLPS.another Table: numeric [ LPS=YES, counts ]
genomic span.another Table : numeric [ << ... >> ]
A DC.another Table: numeric [ LPS=NO, counts ]
DC treated.GSE59364 DC all.csv: numeric [ << ... >> 1
gene.another Table: string [ ID ]
DCLPS0913.another Table: numeric [ LPS=NO, counts ]
DC0907.another Table: numeric [ LPS=NO, counts ]
DC0907.GSE59364 DC all.csv: numeric [ LPS=NO, counts ]
```

path= /Users/fac2003/R RESULTS/table Results 0.tsv name: Results table.name Results groups= LPS=NO,counts,LPS=YES,ID