X11 Applications Edit Window (100%) 5:49 PM (月)

xced version 8a.3.93 - Protein

File Mode Edit View Distance Align Name Help

Foreward | Backward | Sort | Score | JTT 200PAM | AO = +0.06 | OGC = 2.4 | EGC = 0.0 | 9 | 478

Progressive alignment

Algorithm

A A+ C/A+

FFT

Terminal gaps (modified at 97/6/15)

Penalize

OK Cancel

9 9=BCT 327 MPU00089 U00089 9611 Mycoplasma pneumoniae comp

NCBI Data

```
Sequence Check .. OK (ch5)
Backup.  Mon Jan 13 17:25:27 JST 2003
Sequence Check .. OK (ch5)
Backup.  Mon Jan 13 17:40:27 JST 2003
```

NCBI Blast

Options for advanced blasting

Internet zone