```
> cel-mir-62
GUGAGUUAGAUCUCAUAUCCUUCCGCAAAAUGGAAAUGAUAUGUAAUCUAGCUUACAG
> Unknown sequence 1
UUGAGCUAGAUUACAUAUCCGUAAGUGGGGCUUGAAUGAUAUGUAAUCUAGCUUACAG
Matching score: 189 (clustalw2)
> Unknown sequence 2
GUGAGUUAGAUCUCAUAUCCUUCCGCAAAAUGGAAAUCCGAUCUAAUCUACCUUACAG
Matching score: 249 (clustalw2)
```