

BASE DE DATOS DE SECUENCIAS

parámetros de búsqueda:

- péptidos tripticos
- masa péptido precursor
- rango de tolerancia de masas ion precursor

```
>orf19.6109 TUP1 CGDID:CAL0005939 Verified ORF  
MYPQRTQHQR ...K EIPPFLQDLIAK ... WKWTKK*
```

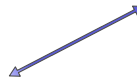
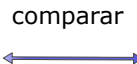
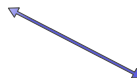
```
> .  
> .  
> .
```

```
>orf19.3001 TEM1 CGDID:CAL0001322 Verified ORF  
MEEDNQDQNNQVALK ...K AILFMFDLTR ... LNLDEIVNVGEPILYK*
```

```
> .  
> .  
> .
```

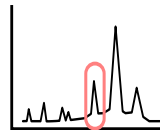
```
>DEC0Y_orf19.802 Decoy sequence  
MSSKSPÉPSYVR ...R LTILDQPFILK ... LGYS*
```

Espectros teóricos



comparar

MS/MS



Espectro adquirido

