SignalP

The SignalP page enables users to predict the presence and location of signal peptide cleavage sites in amino acid sequences from different organisms.

Usage:

1. Click the button in "Choose species" so that you can get the example gene IDs of corresponding versions.

2. Paste the gene list in the search box or upload the gene list file and click "Submit" to submit.

Results:

1. The result table of SignalP. Click the link above to download the result.

Version	ID	Prediction	Score	SP(Sec/SPI)	CS Position
MSU7	LOC_Os07g03288.1	SP	0.006804	0.993164	CS pos: 13-14. Pr: 0.9515
MSU7	LOC_Os09g26320.1	SP	0.000195	0.999856	CS pos: 13-14. Pr: 0.9889
MSU7	LOC_Os01g26390.1	SP	0.000224	0.999773	CS pos: 14-15. Pr: 0.9713

Totally 3 results have been found, this is page 1 of 1.

Begin << | > Next | >> END | Jump To page :