

# 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.

## SignalP:

The SignalP server predicts the presence and location of signal peptide cleavage sites in amino acid sequences from different organisms.

Choose species : **Click to get example gene IDs of corresponding version**

MSU7 RAPDB 9311 ZS97 MH63 R498 IR64

Enter The Gene ID: **Paste the gene list**

Data File :  未选择任何文件 **Upload the gene list file**

↑ **Click to submit**

Submit Reset

Results:

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

## The SignalP information of rice.

 Click here to download the information.

**← Click 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.**

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