

1. Input a sequence or upload a sequence file(.fasta) to Input area.

**1.Sequence Input**  
*Predict GO terms for a protein sequence. Please paste your sequence in fasta format.*

Paste your sequence

**ADD SAMPLE** Or **CHOOSE FILE**

2. Select predict type

**2.Predict Type**  
*Select which category you want to predict*

Eukaryotes 真核生物 ▼

3. Enter your email(optional), and we will send you an email when the prediction is finished.

**3.Your Email(Optional)**  
*Specify your email address if your want to be notified on prediction completion (optional)*

example@gmail.com(optional)

4. Click submit