

LLSSEKLIAGASTGGTEAIRH  
VLQPLPLSSPAVITQHMPPGF  
TRSFATTAEDG  
ERVTAARS  
GANVD  
VLFITGM  
GNDGALIAQN  
EASCVVFGMPREAINMGGVSEV  
VDLSQVSQQMLAKISAGQAIRI

Protein  
sequence

HHblits

Jackhmmer

PconsC3

Contact  
prediction

Protein  
structure

Pool of  
models

Model Quality  
Assessment

CONFOLD

Structure  
prediction

Contact  
maps

