



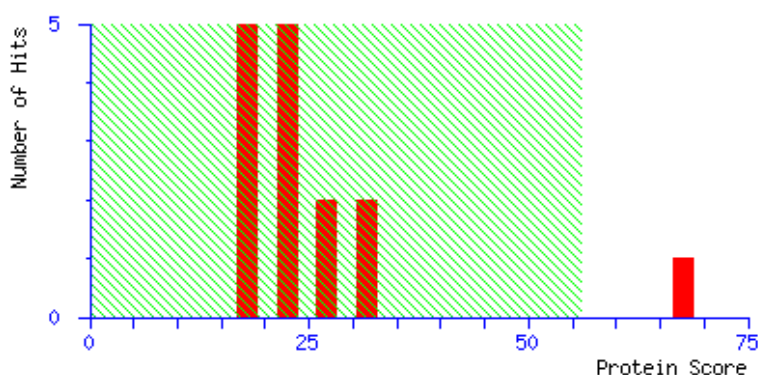
# Mascot Search Results

User : Tosifusas Toda  
 Email : ttoda@proteome.jp  
 Search title : TIG3-SSP1007  
 Database : SwissProt 2017\_11 (556196 sequences; 199341870 residues)  
 Taxonomy : Homo sapiens (human) (20244 sequences)  
 Timestamp : 24 Dec 2017 at 08:44:31 GMT  
 Top Score : 68 for **TCTP\_HUMAN**, Translationally-controlled tumor protein OS=Homo sapiens GN=TPT1 PE=1 SV=1

## Mascot Score Histogram

Protein score is  $-10 \cdot \log(P)$ , where P is the probability that the observed match is a random event.

Protein scores greater than 56 are significant ( $p < 0.05$ ).



## Concise Protein Summary Report

Format As

Concise Protein Summary

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 Significance threshold  $p <$   Max. number of hits 

 Preferred taxonomy 

Re-Search All

Search Unmatched

- [TCTP\\_HUMAN](#) Mass: 19583 Score: **68** Expect: 0.0035 Matches: 5  
 Translationally-controlled tumor protein OS=Homo sapiens GN=TPT1 PE=1 SV=1  
[CX042\\_HUMAN](#) Mass: 14154 Score: 23 Expect: 1e+02 Matches: 2  
 Putative uncharacterized protein CXorf42 OS=Homo sapiens GN=NKAPP1 PE=5 SV=3  
[TRA2B\\_HUMAN](#) Mass: 33646 Score: 22 Expect: 1.1e+02 Matches: 3  
 Transformer-2 protein homolog beta OS=Homo sapiens GN=TRA2B PE=1 SV=1  
[TCTP8\\_HUMAN](#) Mass: 15984 Score: 21 Expect: 1.5e+02 Matches: 2  
 Putative translationally-controlled tumor protein-like protein TPT1P8 OS=Homo sapiens GN=TPT1P8 PE=5 SV=2  
[UB2D3\\_HUMAN](#) Mass: 16676 Score: 21 Expect: 1.6e+02 Matches: 2  
 Ubiquitin-conjugating enzyme E2 D3 OS=Homo sapiens GN=UBE2D3 PE=1 SV=1  
[TEX43\\_HUMAN](#) Mass: 15442 Score: 21 Expect: 1.8e+02 Matches: 2  
 Testis-expressed protein 43 OS=Homo sapiens GN=TEX43 PE=2 SV=1  
[ECHP\\_HUMAN](#) Mass: 79445 Score: 20 Expect: 2e+02 Matches: 3