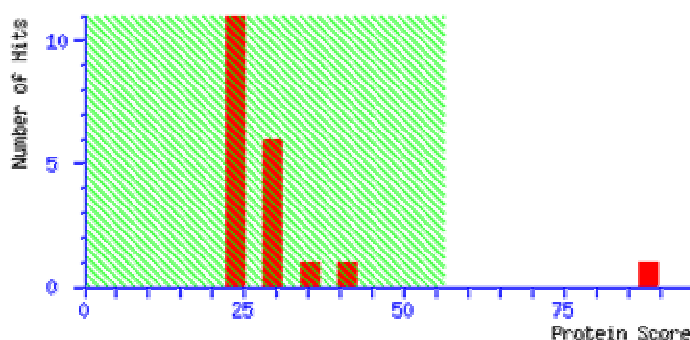


MATRIX SCIENCE Mascot Search Results

User : Tosifusa Toda
 Email : ttoda@proteome.jp
 Search title : TIG3-ssp2417
 Database : SwissProt 2017_12 (556388 sequences; 199430408 residues)
 Taxonomy : Homo sapiens (human) (20245 sequences)
 Timestamp : 26 Dec 2017 at 01:51:17 GMT
 Top Score : 88 for **VIME_HUMAN**, Vimentin OS=Homo sapiens GN=VIM PE=1 SV=4

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	Help
Significance threshold p<	0.05	Max. number of hits AUTO
Preferred taxonomy	All entries	

[Re-Search All](#)
[Search Unmatched](#)

- VIME_HUMAN** **Mass:** 53676 **Score:** 88 **Expect:** 3.1e-05 **Matches:** 8
 Vimentin OS=Homo sapiens GN=VIM PE=1 SV=4
FEZF2_HUMAN **Mass:** 50034 **Score:** 34 **Expect:** 7.4 **Matches:** 4
 Fez family zinc finger protein 2 OS=Homo sapiens GN=FEZF2 PE=2 SV=2
CX7B2_HUMAN **Mass:** 9129 **Score:** 27 **Expect:** 39 **Matches:** 2
 Cytochrome c oxidase subunit 7B2, mitochondrial OS=Homo sapiens GN=COX7B2 PE=3 SV=2
YC023_HUMAN **Mass:** 10927 **Score:** 26 **Expect:** 54 **Matches:** 2
 Putative uncharacterized protein LOC152225 OS=Homo sapiens PE=5 SV=1
BCAS3_HUMAN **Mass:** 102484 **Score:** 25 **Expect:** 63 **Matches:** 4
 Breast carcinoma-amplified sequence 3 OS=Homo sapiens GN=BCAS3 PE=1 SV=3
RNF25_HUMAN **Mass:** 51814 **Score:** 24 **Expect:** 82 **Matches:** 3
 E3 ubiquitin-protein ligase RNF25 OS=Homo sapiens GN=RNF25 PE=1 SV=1
F90A5_HUMAN **Mass:** 50379 **Score:** 24 **Expect:** 84 **Matches:** 3
 Putative protein FAM90A5P OS=Homo sapiens GN=FAM90A5P PE=5 SV=1
F90A9_HUMAN **Mass:** 50277 **Score:** 24 **Expect:** 86 **Matches:** 3
 Putative protein FAM90A9P OS=Homo sapiens GN=FAM90A9P PE=5 SV=1
F90AI_HUMAN **Mass:** 50219 **Score:** 24 **Expect:** 86 **Matches:** 3
 Putative protein FAM90A18P/FAM90A19P OS=Homo sapiens GN=FAM90A18P PE=5 SV=1
F90AE_HUMAN **Mass:** 50492 **Score:** 24 **Expect:** 90 **Matches:** 3
 Putative protein FAM90A14P OS=Homo sapiens GN=FAM90A14P PE=5 SV=1
F90AJ_HUMAN **Mass:** 50434 **Score:** 24 **Expect:** 90 **Matches:** 3
 Putative protein FAM90A15P OS=Homo sapiens GN=FAM90A15P PE=5 SV=1