

is decoy

false

Peptide

start

164

end

174

MS:1001331

MS:1001330

peptide_prophet_probability 0.99962

interprophet_probability 0.994339

pre

post

VDCDQVSAVLK

UNIMOD:4 [Carbamidomethyl] C 3 - avg mass delta: 57,0215

SPECTRUM IDENTIFICATION ITEM

*MS/MS spectrum is shown if <Fragmentation>

element is present in the mzid file

xtandem:hyperscore

xtandem:expect

Spectrum Identification Item PSI-MS CV Parameters

Spectrum Identification Item User Parameters

Protein Ambiguity Groups Protein Group ID

Gene Name(s) CYC1 MP65

Protein Detection Hypothesis/Hypotheses

CGD ID pass thrshld name

Protein Description PDH 52 1

n indist proteins

probability

Peptide Spectrum Matches

Peptide Sequence

ESQIASEIAQLSGFDVIR

ITYSPYSDNGGCK

VDCDQVSAVLK

MP65 CGDID:CAL0000936... Verified ORF; Cell surface mannoprotein;

Accession(s) / Id(s)

orf19.1770

orf19.1779

PD 1 Mzid 14 PDL 1

PDP 1

X!Tandem

orf19.1779 CGD

Protein Detection Hypothesis PSI-MS CV Parameters MS:1001093

322

322

0.024

DB sequence

orf19.1779

PDH ID

xtandem:hyperscore **Protein Detection Hypothesis User Parameters**

orf19.1779 MLFKSFVTFTVLANALAAPLAHQHHQHKEEKRAVHVVTTTNVVVVTIGNGDQTTTFAAPSVAAESSVSVSVNTEPPQNHPTTTQDVASAST

YPSSTDGSAASSSAAASSSSQAGSEPSGGVGSGGAKG ITYSPYSDNGGCKSESQIASEIAQLSGFDVIR LYG VDCDQVSAVLK AKTSSOKIFAGIFDVSNIASGIESLAEAVEACGSWDDIYTVSIGNELVNAGSATPSOIKAYVDEGRKALKAAGYTGPVVSVDTF...

Spectrum Identification Item(s)

SIL Candida4ug SII 2757 1

SIL Candida2ug SII 60 1

SIL Candida4ug SII 4141 1

PSM count

1

1

1000