



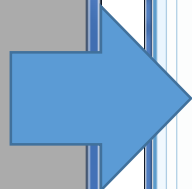
Skyline-daily - MRMer.sky

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Targets

- YAL038W
  - R.LTSLNVVAGSDLR
  - K.TNNPETLVALR.K [3]
  - K.GVNLPGTDVDLPAL
  - R.TANDVLTIR.E [216]
  - R.GDLGIEIPAEVLAV
  - K.EPVSDDWTDDEAR.R [446, 456]
- YCR012W
  - K.LSVQDLDLK.D [6, 14]
  - R.VDFNVPLDGK.K [22, 31]
  - K.ASAPGSVILLENLR.Y [108, 121]
  - R.YHIEEESR.K [122, 130]
  - K.ELPGVAFLSEK.K [404, 414]
- YOR332W
  - K.LLSEEALPAIR.L [208, 218]
- YLR340W
  - R.GFLSDLPDFEK.L [64, 74]
  - K.GNVGFVFTNEPLTEIK.N [81, 96]
  - R.AGAVAPEDIWV.R.A [110, 121]
- S rerdcj
  - K.LVEDPQVIAPFLGK.L [275, 288]
  - K.SNFATIADPEAR.E [295, 306]
  - K.FPEPGYLEGVK.T [649, 659]
- YMR116C
  - R.LWDVATGETYQR.F [90, 101]
- YDL055C

Ready 1/24 prot 1/44 pep 1/88 prec 1/296 tran



Protter - customizable proteo x

wlab.ethz.ch/protter/?file=95b2c78e-f64a-4b29-8780-1db464315f44.csv#up=YAL038W&peptides=LTSLNVVAGSDLR,TNNPETLVALR,GVNLPGTDVDLPALSEK,TANDVLTIR,GD

PROTTER

1 protein 2 topology 3 styles 4 misc. ? help feedback to protter@gmx.ch ETH zürich

24 proteins:

- YAL038W 6 peptides
  - LTSLNVVAGSDLR
  - TNNPETLVALR
  - GVNLPGTDVDLPALSEK
  - TANDVLTIR
  - GDLGIEIPAEVLAVQK
  - EPVSDDWTDDEAR
- YCR012W 5 peptides
- YOR332W 1 peptide
- YLR340W 3 peptides
- S rerdcj 3 peptides
- YMR116C 1 peptide
- YDL055C 2 peptides
- YHR174W 2 peptides
- YKL152C 4 peptides
- YOL086C 2 peptides
- YLR044C 1 peptide
- YOL127W 1 peptide

by accession please enter a UniProt protein accession:  
YAL038W submit load example

by sequence

refresh export email open in UniProt

Legend:   
● exp. pept.   
● signal pept.   
● disulfide bonds   
● variants   
■ PTMs   
● exp. pept.   
● N-term. Phosph.   
● TM/MS. UniProt