Skyline-daily

File Edit View Settings Tools Help

Targets

- sp|O43670|ZN207_HUMAN
  - K.ETIDAVPNAIPGR.T [59, 71]
- sp|Q96BR1|SGK3_HUMAN
  - K.ESCPSVSIPSSDEHR.E [10, 24]
  - K.VLVSVGR.S [35, 41]

Peptide Settings

Digestion | Prediction | Filter | Library | Modifications | Quantification

Enzyme: Trypsin [KR | P]

Max missed cleavages: 0

Background proteome: HUMAN_9606_Uniprot

Enforce peptide uniqueness by: None

OK Cancel

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sp|Q96BR1|SGK3_HUMAN Peptides

- -.MQR.D [0, 2]
- R.DHTMDYK.E [3, 9]
- K.ESCPSVSIPSSDEHR.E [10, 24]
- K.VLVSVGR.S [35, 41]
- R.YPELYNHPDVR.A [103, 113]
- R.AFLQMDSPK.H [114, 122]
- K.HQSDPSEDEDER.S [123, 134]
- R.SSQK.L [135, 138]
- K.LHSTSQNINLGPSGNPHAKPTDFDFLK.V [139, 165]
- K.VIGK.G [166, 169]
- K.GSFGK.V [170, 174]
- K.FYAVK.V [186, 190]

sp|Q96BR1|SGK3_HUMAN

Name	sp|Q96BR1|SGK3_HUMAN
Accession	Q96BR1
Preferred Name	SGK3_HUMAN
Gene	SGK3
Species	Homo sapiens
Description	Serine/threonine-protein kinase Sgk3 OS=Homo sapiens GN=SGK3 PE=1 SV=1

MQRDHTMDYKESCPSVSIPSSDEHREKKKRFTVYKVLVSVGRSEWFVFRRYAEFDKLYNTLKKQFPAMALKIPAKNFGDNFDP
DFIKQRRAGLNEFIQNLVRYPELYNHPDVRAFLQMDSPKHQSDPSEDEDERSSQKLHSTSQNINLGPSGNPHAKPTDFDFLKVIG
KGSFGKVLLAKRKLDGKFYAVKVLQKKIVLNRKEQKHIMAERNVLLKNVKHPFLVGLHYSFQTTEKLYFVLDFVNGGELFFHLQRE
RSFPEHRARFYAAEIASALGYLHSIKIVYRDLKPENILLDSVGHVVLTDFGLCKEGIAISDTTTTFCGTPEYLAPEVIRKQPYDNTVD
WWCLGAVLYEMLYGLPPFYCRDVAEMYDNILHKPLSLRPGVSLTAWSILEELLEKDRQNRLGAKEDFLEIQNHPFFESLSWADLV
QKKIPPPFNPNVAGPDDIRNFDTAFTEETVPYSVCVSSDYSIVNASVLEADDAFVGFSYAPPSEDLFL