

V I L K K Q Q A A L E R T - - V - - - H P N S P G I P Y R T L E



- ☒ sp|Q9UKV0|HDAC9_HUMAN
- ☒ sp|Q9UQL6|HDAC5_HUMAN
- ☒ sp|P56524|HDAC4_HUMAN

Q8WUI4:Sequence conflict

- ☐ sp|Q9BY41|HDAC8_HUMAN
- ☐ sp|O15379|HDAC3_HUMAN
- ☐ sp|Q92769|HDAC2_HUMAN
- ☐ sp|Q13547|HDAC1_HUMAN
- ☐ sp|Q9UBN7|HDAC6_HUMAN
- ☒ sp|Q8WUI4|HDAC7_HUMAN
- ☒ sp|Q9UKV0|HDAC9_HUMAN
- ☒ sp|Q9UQL6|HDAC5_HUMAN
- ☒ sp|P56524|HDAC4_HUMAN

Q8WUI4:Sequence conflict

- ☐ sp|Q9BY41|HDAC8_HUMAN
- ☐ sp|O15379|HDAC3_HUMAN
- ☐ sp|Q92769|HDAC2_HUMAN
- ☐ sp|Q13547|HDAC1_HUMAN
- ☐ sp|Q9UBN7|HDAC6_HUMAN
- ☒ sp|Q8WUI4|HDAC7_HUMAN
- ☒ sp|Q9UKV0|HDAC9_HUMAN
- ☒ sp|Q9UQL6|HDAC5_HUMAN
- ☒ sp|P56524|HDAC4_HUMAN

Q8WUI4:Sequence conflict

- ☐ sp|Q9BY41|HDAC8_HUMAN
- ☐ sp|O15379|HDAC3_HUMAN
- ☐ sp|Q92769|HDAC2_HUMAN
- ☐ sp|Q13547|HDAC1_HUMAN
- ☐ sp|Q9UBN7|HDAC6_HUMAN
- ☒ sp|Q8WUI4|HDAC7_HUMAN
- ☒ sp|Q9UKV0|HDAC9_HUMAN
- ☒ sp|Q9UQL6|HDAC5_HUMAN
- ☒ sp|P56524|HDAC4_HUMAN

Q8WUI4:Sequence conflict

- ☐ sp|Q9BY41|HDAC8_HUMAN
- ☐ sp|O15379|HDAC3_HUMAN
- ☐ sp|Q92769|HDAC2_HUMAN
- ☐ sp|Q13547|HDAC1_HUMAN
- ☐ sp|Q9UBN7|HDAC6_HUMAN
- ☒ sp|Q8WUI4|HDAC7_HUMAN
- ☒ sp|Q9UKV0|HDAC9_HUMAN
- ☒ sp|Q9UQL6|HDAC5_HUMAN
- ☒ sp|P56524|HDAC4_HUMAN

Q8WUI4:Sequence conflict




FLLSKSK - - EPTPGGLNHS L P Q H P K C W G A - - F
FVLNKKK - - ALAHRNLNHC I SSDPRYWYGKTC

PEHFFPLRKT VSEPNLKLRYKPKKS - LERRKNF
KDDFPLRKTASEPNLKVRSRLKQKVAERRSSF
RDDFPLRKTASEPNLKVRSRLKQKVAERRSSF
KDDFPLRKTASEPNLKLRSRLKQKVAERRSSF

SSSTPASGCSSPNDSEHGP - - - - NP
SSSSPGSGPSSPNNGPTGSVTENETSVLPPTF
CNSAPGSGPSSPNNSSH - STIAENGFTGSVPN
CSSAPGSGPSSPNNSSGGSVSAENGIAPAVPS

PERLHAIKEQLIQEGLLDRCVSFQARFAEKEE
TLGLPAPARADS - - - -
TLGLPAVPSQLNA - - - - SNSLKEKQ
SLGLQATVTVTNS - - - - HLTASPKLSTQQE
TLGLPATGPSAGT - - - - AGQQ

SVYLHPNSYSCACLASGSVLRLVDAVLGAEIF
- - - H - - - - TPLFLP - HGLEPEA - GGTLP SRL
GSIP - - - - ASSSHPHVTLEGKPPNSSHQALL
STSS - - - - IPGCLLGVALEGDGSPHGHASLL
PGTH - - - - LTPYLSSTSPLERDG - GAAHSPLL

- ☐  sp|Q9BY41|HDAC8_HUMAN
- ☐  sp|O15379|HDAC3_HUMAN
- ☐  sp|Q92769|HDAC2_HUMAN
- ☐  sp|Q13547|HDAC1_HUMAN
- ☐  sp|Q9UBN7|HDAC6_HUMAN
- ☒  sp|Q8WUI4|HDAC7_HUMAN
- ☒  sp|Q9UKV0|HDAC9_HUMAN
- ☒  sp|Q9UQL6|HDAC5_HUMAN
- ☒  sp|P56524|HDAC4_HUMAN

- - - - -0

- - - - -0

- - - - -0

- - - - -0

Y A Q Q K H R I R R V L I V D W D V H H G Q G T Q F T F D Q D F300

- - - - - P F H F A Q S L M T T E R - L367

- - - - - P L H P Q S P L A T K E R - I458

- - - - - P L H G Q S P L V T G E R - V507

- - - - - P L H A Q S L - V G A D R - V476

Q8WUI4:Sequence conflict

- ☐  sp|Q9BY41|HDAC8_HUMAN
- ☐  sp|O15379|HDAC3_HUMAN
- ☐  sp|Q92769|HDAC2_HUMAN
- ☐  sp|Q13547|HDAC1_HUMAN
- ☐  sp|Q9UBN7|HDAC6_HUMAN
- ☒  sp|Q8WUI4|HDAC7_HUMAN
- ☒  sp|Q9UKV0|HDAC9_HUMAN
- ☒  sp|Q9UQL6|HDAC5_HUMAN
- ☒  sp|P56524|HDAC4_HUMAN

- - - - -0

- - - - -0

- - - - -0

- - - - -0

- Q G Y T I N V P W N Q V G M R D A D Y I - A A F L H - - V L L344

P P P P G P M Q P R L E Q L K T H - - - - - V Q V I K R S411

L A Q L V I Q Q Q H Q Q F L E K Q K Q Y - - Q Q Q I H M N K L L520

L Q Q L V M Q Q Q H Q Q F L E K Q K Q - - - - Q Q L Q L G K I L567

L Q H L V I Q Q Q H Q Q F L E K H K Q Q F Q Q Q Q L Q M N K I I540

Q8WUI4:Sequence conflict

- ☐  sp|Q9BY41|HDAC8_HUMAN
- ☐  sp|O15379|HDAC3_HUMAN
- ☐  sp|Q92769|HDAC2_HUMAN
- ☐  sp|Q13547|HDAC1_HUMAN
- ☐  sp|Q9UBN7|HDAC6_HUMAN
- ☒  sp|Q8WUI4|HDAC7_HUMAN
- ☒  sp|Q9UKV0|HDAC9_HUMAN
- ☒  sp|Q9UQL6|HDAC5_HUMAN
- ☒  sp|P56524|HDAC4_HUMAN

- - - - -0

- - - - -0

- - - - -0

- - - - -0

F D A L Q G D P K G E M A A - - - - T P A G F A Q L T H L L M G404

- - - - - D G G G - - - - P G Q V V D D G L - - - - -430

- - - - P S S G N S T R S D S S A C V D D T L G Q V G A V - - -559

A L T M P R E G S T E - - - - - S E S T Q E - - -608

L D R L P G Q K E A H - - - - - A Q A G V - - - -569

Q8WUI4:Sequence conflict

- ☐  sp|Q9BY41|HDAC8_HUMAN
- ☐  sp|O15379|HDAC3_HUMAN
- ☐  sp|Q92769|HDAC2_HUMAN
- ☐  sp|Q13547|HDAC1_HUMAN
- ☐  sp|Q9UBN7|HDAC6_HUMAN
- ☒  sp|Q8WUI4|HDAC7_HUMAN
- ☒  sp|Q9UKV0|HDAC9_HUMAN
- ☒  sp|Q9UQL6|HDAC5_HUMAN
- ☒  sp|P56524|HDAC4_HUMAN

- - - - -0

- - - - -0

- - - - -0

- - - - -0

D P C P M L E S P G A P C R S A Q A S V S C A L E A L E P F W E459






H G Q P E A R G P A P - - - - - L Q Q H P Q V L L W E484

- - - - - I Q E - - - M E S G E Q A A F M Q Q P F L E609

G E S G A E E G P D L E E - - - P - - - - G A G Y K K L F S659

A E P P R E V E P G Q R Q - - - P S E Q E L L F R Q Q A L L L E630

Q8WUI4:Sequence conflict

- ☐  sp|Q9BY41|HDAC8_HUMAN
- ☐  sp|O15379|HDAC3_HUMAN
- ☐  sp|Q92769|HDAC2_HUMAN
- ☐  sp|Q13547|HDAC1_HUMAN
- ☐  sp|Q9UBN7|HDAC6_HUMAN

- - - - - M E E P E E P A D S -31

- - - - - -22

- - - - - M A Y S Q G -29

- - - - - M A Q T Q -28

E - - - - - E G P W E P P V L F500

O C C D A A D A C L S A D E R A C O A R V L S C C E T R A D

<input checked="" type="checkbox"/>	sp Q8WUI4 HDAC7_HUMAN	QSSSPAASVLPHPA - - - - - MDRPL -	541
<input checked="" type="checkbox"/>	sp Q9UKV0 HDAC9_HUMAN	QSSSPAAPGGMKSP - - - - - PDQPV -	654
<input checked="" type="checkbox"/>	sp Q9UQL6 HDAC5_HUMAN	QSSPASATFPVSV - - - - - QEPPT -	704
<input checked="" type="checkbox"/>	sp P56524 HDAC4_HUMAN		675

Q8WUI4:Sequence conflict

<input type="checkbox"/>	sp Q9BY41 HDAC8_HUMAN	AKIPKRASMVHSLIEAYALHKQMRIVKPKVA	592
<input type="checkbox"/>	sp O15379 HDAC3_HUMAN	PMKPHRLALTHSLVLHYGLYKKMIVFKPYQAS	84
<input type="checkbox"/>	sp Q92769 HDAC2_HUMAN	PMKPHRIRMTHNLLLNLYGLYRKMEIYRPHKAT	91
<input type="checkbox"/>	sp Q13547 HDAC1_HUMAN	PMKPHRIRMTHNLLLNLYGLYRKMEIYRPHKAN	90
<input type="checkbox"/>	sp Q9UBN7 HDAC6_HUMAN	PEVPQRI LRIMCRLEELGLAGRCLTLTPRPA	559
<input checked="" type="checkbox"/>	sp Q8WUI4 HDAC7_HUMAN	PEHAGRIQS IWSRLQERGLR	604
<input checked="" type="checkbox"/>	sp Q9UKV0 HDAC9_HUMAN	PEHAGRIQS IWSRLQETGL LNKCEIRIQRKA	717
<input checked="" type="checkbox"/>	sp Q9UQL6 HDAC5_HUMAN	PEHAGRIQS IWSRLQETGL LSKCEIRIRGRKA	767
<input checked="" type="checkbox"/>	sp P56524 HDAC4_HUMAN	PEHAGRIQS IWSRLQETGLRGKCECIRGRKA	738

Q8WUI4:Sequence conflict

<input type="checkbox"/>	sp Q9BY41 HDAC8_HUMAN	- - - - SIEYGLGYDCP - - - - ATEGIFDYAA - -	140
<input type="checkbox"/>	sp O15379 HDAC3_HUMAN	- - - - LNAFNVGDDCP - - - - VFPGLFEFCSS - -	132
<input type="checkbox"/>	sp Q92769 HDAC2_HUMAN	- - - - MQR FNVGEDCP - - - - VFDGLFEFCQ - -	139
<input type="checkbox"/>	sp Q13547 HDAC1_HUMAN	- - - - MQR FNVGEDCP - - - - VFDGLFEFCQ - -	138
<input type="checkbox"/>	sp Q9UBN7 HDAC6_HUMAN	HRE - - - - - - - - - - SSNFDSIYICPSTFA	608
<input checked="" type="checkbox"/>	sp Q8WUI4 HDAC7_HUMAN	AGLLAQRMFVMLPCGGVGVDTDTIWNELHSS	667
<input checked="" type="checkbox"/>	sp Q9UKV0 HDAC9_HUMAN	LGDDSQKFFSSLP CGGLGVDSDTIWNELH	780
<input checked="" type="checkbox"/>	sp Q9UQL6 HDAC5_HUMAN	LGPISQKMYAVLP CGGIGVDSDTVWNEMHSS	830
<input checked="" type="checkbox"/>	sp P56524 HDAC4_HUMAN	LGLAS - V FVRLPCGGVGVDSDTIWNEVHSA	800

Q8WUI4:Sequence conflict

<input type="checkbox"/>	sp Q9BY41 HDAC8_HUMAN	WHHAKKDEASGFCYLNDAVLGILRLRR - - -	201
<input type="checkbox"/>	sp O15379 HDAC3_HUMAN	LHHAKKFEASGFCYVNDIVIGILELLK - - -	193
<input type="checkbox"/>	sp Q92769 HDAC2_HUMAN	LHHAKKSEASGFCYVNDIVLAILELLK - - -	200
<input type="checkbox"/>	sp Q13547 HDAC1_HUMAN	LHHAKKSEASGFCYVNDIVLAILELLK - - -	199
<input type="checkbox"/>	sp Q9UBN7 HDAC6_HUMAN	GHHAEQDAACGFCFFNSVAVAAARHAQTISGH	672
<input checked="" type="checkbox"/>	sp Q8WUI4 HDAC7_HUMAN	GHHADHSTAMGFCFFNSVAIACRQLQQQS -	730
<input checked="" type="checkbox"/>	sp Q9UKV0 HDAC9_HUMAN	GHHAEESTAMGFCFFNSVAITAKYL RDQL -	843
<input checked="" type="checkbox"/>	sp Q9UQL6 HDAC5_HUMAN	GHHAEESTAMGFCFFNSVAITAKLLQQKL -	893
<input checked="" type="checkbox"/>	sp P56524 HDAC4_HUMAN	GHHAEESTPMGFCYFNSVAVAAKLLQQRL -	863

Q8WUI4:Sequence conflict

<input type="checkbox"/>	sp Q9BY41 HDAC8_HUMAN	KFSPG - FFPG - - TGDVSDVGLGKG	258
<input type="checkbox"/>	sp O15379 HDAC3_HUMAN	KYGN - FFPG - - TGDMEVGAESEGRYYCLNV	250
<input type="checkbox"/>	sp Q92769 HDAC2_HUMAN	KYGEY - - FPG - - TGDLRD I GAGKGKYAVNFF	256
<input type="checkbox"/>	sp Q13547 HDAC1_HUMAN	KYGEY - - FPG - - TGDLRD I GAGKGKYAVNYF	255
<input type="checkbox"/>	sp Q9UBN7 HDAC6_HUMAN	RYDHGTF FPMGDEGASSQIGRAAGTGFTVNV	733
<input checked="" type="checkbox"/>	sp Q8WUI4 HDAC7_HUMAN	RHDDGNFFPG - - SGAVDEVGA	792
<input checked="" type="checkbox"/>	sp Q9UKV0 HDAC9_HUMAN	RYDEGNFFPG - - SGAPNEVG TGLGEGYNINI	905
<input checked="" type="checkbox"/>	sp Q9UQL6 HDAC5_HUMAN	RYDNGNFFPG - - SGAPEEVGGGPVGYNVNV	955
<input checked="" type="checkbox"/>	sp P56524 HDAC4_HUMAN	RYDDGNFFPG - - SGAPDEVGTGPVG FNVNM	925

Q8WUI4:Sequence conflict



- ☐ sp|Q9BY41|HDAC8_HUMAN
- ☐ sp|O15379|HDAC3_HUMAN
- ☐ sp|Q92769|HDAC2_HUMAN
- ☐ sp|Q13547|HDAC1_HUMAN
- ☐ sp|Q9UBN7|HDAC6_HUMAN
- ☒ sp|Q8WUI4|HDAC7_HUMAN
- ☒ sp|Q9UKV0|HDAC9_HUMAN
- ☒ sp|Q9UQL6|HDAC5_HUMAN
- ☒ sp|P56524|HDAC4_HUMAN

AVVLQLGADTIAGD - - PMCSFNMTPTVGIGKCL319
CIVLQCGADSLGCD - - RLGC FNLSIRGHGECV311
AVVLQCGADSLSGD - - RLGC FNLT VKGHA KCV317
AVVLQCGSDSLSGD - - RLGC FNLT IKGHAKCV316
LVLVSAGFDAARGD - - PLGGCQVSPEGY AHLT795
LVLVSAGFDA AEGHPAPLGGYHVSAKCFGYMT856
MVLVSAGFDAL EGH TPPLGGYKVTAKCFGHL T969
VVLVSAGFDAVEGHL SPLGGYSVTARCFGHL T1019
VVLVSSGFD AVEGHPTPLGGYNLSARCFGYL T989

Q8WUI4:Sequence conflict

- ☐ sp|Q9BY41|HDAC8_HUMAN
- ☐ sp|O15379|HDAC3_HUMAN
- ☐ sp|Q92769|HDAC2_HUMAN
- ☐ sp|Q13547|HDAC1_HUMAN
- ☐ sp|Q9UBN7|HDAC6_HUMAN
- ☒ sp|Q8WUI4|HDAC7_HUMAN
- ☒ sp|Q9UKV0|HDAC9_HUMAN
- ☒ sp|Q9UQL6|HDAC5_HUMAN
- ☒ sp|P56524|HDAC4_HUMAN

GVILGKTLSS EIPDHEFFFTAYGPDYVLEIT F377
SLLVEEAISEELPYSEYFEYFAPDFTLHPD\372
AVALDCEIPNELPYNDYFEYFGPDFKLHIS F377
AVALDTEIPNELPYNDYFEYFGPDFKLHIS F376
RSL LGDPP - - - - - PL LTLPRP F840
AALLGNRVDP - L - - - - - SEEGWKQK F904
NALLGNELEP - L - - - - - AEDILHQS F1011
SALLSVELQP - L - - - - - DEAVLQQK F1067
SALLGNELD P - L - - - - - PEKVLQQR F1037

Q8WUI4:Sequence conflict

- ☐ sp|Q9BY41|HDAC8_HUMAN
- ☐ sp|O15379|HDAC3_HUMAN
- ☐ sp|Q92769|HDAC2_HUMAN
- ☐ sp|Q13547|HDAC1_HUMAN
- ☐ sp|Q9UBN7|HDAC6_HUMAN
- ☒ sp|Q8WUI4|HDAC7_HUMAN
- ☒ sp|Q9UKV0|HDAC9_HUMAN
- ☒ sp|Q9UQL6|HDAC5_HUMAN
- ☒ sp|P56524|HDAC4_HUMAN

- - - - - .377
PSVQIH - - - - - DVPADLLTYDRTDEADA F412
PGVQMQ - - - - - AIPEDAVHEDSGDEDEGE[420
PGVQMQ - - - - - AIPEDA IPEESGDEDEDED[419
DREGPSSSKLVTKKAPQPAKPR - - - - LAERM899
PDSWV - - PR - - - - VP - GADKE - - - - EVEA\942
- - - - - .1011
LGRSL - - RE - - - - AQ - AGETE - - - - EAET\1114
AGRSL - - IE - - - - AQ - TCENE - - - - EAET\1076

Q8WUI4:Sequence conflict

- ☐ sp|Q9BY41|HDAC8_HUMAN
- ☐ sp|O15379|HDAC3_HUMAN
- ☐ sp|Q92769|HDAC2_HUMAN
- ☐ sp|Q13547|HDAC1_HUMAN
- ☐ sp|Q9UBN7|HDAC6_HUMAN
- ☒ sp|Q8WUI4|HDAC7_HUMAN
- ☒ sp|Q9UKV0|HDAC9_HUMAN
- ☒ sp|Q9UQL6|HDAC5_HUMAN
- ☒ sp|P56524|HDAC4_HUMAN

- - - - - .377
- - - - - FYDGDHDNDK - - - - .428
- - - - - FSDSEDEGE GGRNVADHK - - - K(463
- - - - - FSDSEEEGE GGRKNSSNFK - - - K(461
TAVVALTQDQPSEAATGGATLAQTISEAAI(963
- - - - LVEEEEP MNL - - - - .952
- - - - .1011
- - - - P MEQEPAL - - - - .1122
- - - - P MEEEPPL - - - - .1084

Q8WUI4:Sequence conflict

- ☐ sp|Q9BY41|HDAC8_HUMAN
- ☐ sp|O15379|HDAC3_HUMAN

- - - - - 377
- - - - - 428
EDKSKD NSGEKTDTKG TKSE - - - - -

<input type="checkbox"/>	sp Q92769 HDAC2_HUMAN	EEKTKEE - - - KPEAKGVKEE - - - - -	488
<input type="checkbox"/>	sp Q13547 HDAC1_HUMAN	QTTSEDAVGGATLGQTTSEEAVGGATLAQTT	482
<input type="checkbox"/>	sp Q9UBN7 HDAC6_HUMAN	- - - - -	1027
<input checked="" type="checkbox"/>	sp Q8WUI4 HDAC7_HUMAN	- - - - -	952
<input checked="" type="checkbox"/>	sp Q9UKV0 HDAC9_HUMAN	- - - - -	1011
<input checked="" type="checkbox"/>	sp Q9UQL6 HDAC5_HUMAN	- - - - -	1122
<input checked="" type="checkbox"/>	sp P56524 HDAC4_HUMAN	- - - - -	1084

Q8WUI4:Sequence conflict

<input type="checkbox"/>	sp Q9BY41 HDAC8_HUMAN	- - - - -	377
<input type="checkbox"/>	sp O15379 HDAC3_HUMAN	- - - - -	428
<input type="checkbox"/>	sp Q92769 HDAC2_HUMAN	- - - - -	488
<input type="checkbox"/>	sp Q13547 HDAC1_HUMAN	- - - - -	482
<input type="checkbox"/>	sp Q9UBN7 HDAC6_HUMAN	DHQTPPTSPVQGTTPQISPSTLIGSLRTLLEL	1091
<input checked="" type="checkbox"/>	sp Q8WUI4 HDAC7_HUMAN	- - - - -	952
<input checked="" type="checkbox"/>	sp Q9UKV0 HDAC9_HUMAN	- - - - -	1011
<input checked="" type="checkbox"/>	sp Q9UQL6 HDAC5_HUMAN	- - - - -	1122
<input checked="" type="checkbox"/>	sp P56524 HDAC4_HUMAN	- - - - -	1084

Q8WUI4:Sequence conflict

<input type="checkbox"/>	sp Q9BY41 HDAC8_HUMAN	- - - - -	377
<input type="checkbox"/>	sp O15379 HDAC3_HUMAN	- - - - -	428
<input type="checkbox"/>	sp Q92769 HDAC2_HUMAN	- - - - -	488
<input type="checkbox"/>	sp Q13547 HDAC1_HUMAN	- - - - -	482
<input type="checkbox"/>	sp Q9UBN7 HDAC6_HUMAN	MQGSRGLTDQAIFYAVTPLPWCPHLVAVCPI	1155
<input checked="" type="checkbox"/>	sp Q8WUI4 HDAC7_HUMAN	- - - - -	952
<input checked="" type="checkbox"/>	sp Q9UKV0 HDAC9_HUMAN	- - - - -	1011
<input checked="" type="checkbox"/>	sp Q9UQL6 HDAC5_HUMAN	- - - - -	1122
<input checked="" type="checkbox"/>	sp P56524 HDAC4_HUMAN	- - - - -	1084

Q8WUI4:Sequence conflict

<input type="checkbox"/>	sp Q9BY41 HDAC8_HUMAN	- - - - -	377
<input type="checkbox"/>	sp O15379 HDAC3_HUMAN	- - - - -	428
<input type="checkbox"/>	sp Q92769 HDAC2_HUMAN	- - - - -	488
<input type="checkbox"/>	sp Q13547 HDAC1_HUMAN	- - - - -	482
<input type="checkbox"/>	sp Q9UBN7 HDAC6_HUMAN	YINGHMLQHHGNSGHPLVLSYIDLSAWCYYC	1215
<input checked="" type="checkbox"/>	sp Q8WUI4 HDAC7_HUMAN	- - - - -	952
<input checked="" type="checkbox"/>	sp Q9UKV0 HDAC9_HUMAN	- - - - -	1011
<input checked="" type="checkbox"/>	sp Q9UQL6 HDAC5_HUMAN	- - - - -	1122
<input checked="" type="checkbox"/>	sp P56524 HDAC4_HUMAN	- - - - -	1084

Q8WUI4:Sequence conflict



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