Scan number 7345 Raw file nak_cdc48UbconjugatecontrolSAX4_22may10_13
Method ITMS; CID Pepti... 93.48

Protein Group 145

m/z (full)

m/z (zoom)
Protein Group 178

m/z (full)

m/z (zoom)

Relative Abundance

Intensity
Protein Group 354

Scan number 27946 Raw file nak_UBX2LSAX3_8feb11_03
Method ITMS; CID Pepti... 81.62

m/z (full)

m/z (zoom)

Relative Abundance

Intensity
Scan number 22389
Raw file nak_UBX2HSAX4_8feb11_08
Method ITMS; CID Pepti... 82.67

Protein Group 498

m/z (full)

Relative Abundance

m/z (zoom)

Intensity
Protein Group 501

Scan number 13055 Raw file nak_cdc48UbconjugatesSAX4_22may10_5
Method ITMS; CID Pepti... 161.02

m/z (full)

m/z (zoom)

Relative Abundance

Intensity [10^4]
Scan number 22247 Raw file nak_ubx4LightSAX5_11nov05
Method ITMS; CID Pepti... 117.7
Scan number 31993 Raw file nak_cdc48UbconjugatesSAX5_21may16
Method ITMS; CID Pepti... 143.39

m/z (full)

m/z (zoom)

Relative Abundance

Intensity

Protein Group 602
Scan number: 25948  
Raw file: nak_ubx4LightSAX4_11nov04  
Method: ITMS; CID  
Pepti...: 90.69
Protein Group 672

Scan number 33747  Raw file  nak_ubx4LightSAX2_11nov02
Method ITMS; CID Pepti... 91.62

m/z (full)

m/z (zoom)
Method: ITMS; CID

Protein Group 722
Scan number 14421 Raw file nak_ubx4LightSAX5_11nov05
Method ITMS; CID Pepti... 91.47

Protein Group 806

m/z (full)

m/z (zoom)

Relative Abundance

Intensity
Scan number 35722
Raw file nak_cdc48UbconjugatesSAX2_21may13
Method ITMS; CID Pepti... 75.48

Protein Group 849

A I I G A P L L G G P V L R _

m/z (full)

Relative Abundance

0 100 200 300

Intensity

0 100 200 300

m/z (zoom)
Protein Group 868

Scan number 15021
Raw file nak_UBX2HSAX6_8feb11_10
Method ITMS; CID Pepti... 194.48
Scan number 13027 Raw file nak_cdc48UbconjugatecontrolSAX3_21may22 Method ITMS; CID Pepti... 112.75

Protein Group 874
Scan number 25643 Raw file nak_UBX6SAX2L_14jan11_12
Method ITMS; CID Peptide 158.04

Protein Group 879

m/z (full)

m/z (zoom)
Scan number 14525 Raw file nak_ubx5L_SAX6_08Sep6
Method ITMS; CID Pepti... 55.01

Protein Group 929

m/z (full)

m/z (zoom)

Relative Abundance

Intensity

m/z (full)

m/z (zoom)
Scan number 14968
Raw file nak_UBX6SAX1L_14Jan11_11
Method ITMS; CID Pepti... 127.83

Protein Group 1111

m/z (full)

m/z (zoom)
Scan number 13614 Raw file nak_UBX3LSAX6_28jan11_24
Method ITMS; CID Pepti... 199.37

Protein Group 1168

m/z (full)

Relative Abundance

m/z (zoom)

Intensity
Scan number 11193 Raw file nak_ubx4LightSAX3_11nov03

Method ITMS; CID Pepti... 196.88

Protein Group 1286
Protein Group 1361

Scan number 14808
Raw file nak_UBX3HSAX5_28jan11_31
Method ITMS; CID Pepti... 107.21
Scan number 15317 Raw file nak_WTonetooneSAX2_4feb11_02
Method ITMS; CID Pepti... 72.7

Protein Group 1634

m/z (full)

m/z (zoom)

Relative Abundance

Intensity

0 200 400 600 800 1000 1200 1400 1600

0 200 400 600 800 1000 1200 1400 1600
Scan number 8552
Raw file nak_WTonetooneSAX6_4feb11_06
Method ITMS; CID Peptid... 85.54

Protein Group 1733

m/z (full)

m/z (zoom)
Scan number 33399
Raw file nak_cdc48UbconjugatesSAX4_21may15
Method ITMS; CID Pepti... 41.11

Protein Group 1757

m/z (full)

Relative Abundance

Intensitry

m/z (zoom)
Protein Group 1806

Scan number 31253
Raw file nak_UBX6SAX3L_14jan11_13
Method ITMS; CID Pepti... 34.22

m/z (full)

m/z (zoom)

Relative Abundance

Intensity
Scan number 28809
Raw file nak_ubx4LightSAX5_11nov05
Method ITMS; CID Pept... 200.55

Protein Group 1823

m/z (full)

m/z (zoom)

Relative Abundance

Intensity
Scan number 16977
Raw file nak_cdc48UbconjugatesSAX6_14may10_10
Method ITMS; CID Pepti... 193.96

Protein Group 1882

m/z (full)

m/z (zoom)
Scan number 12443 Raw file nak_cdc48UbconjugatesSAX6_21may17
Method ITMS; CID Pepti... 102.64
_ E ... 20 40 60 80 100 120
Relative Abundance
0 50 100 150 200 250
Intensity
200 400 600 800 1000 1200 1400 1600
m/z (zoom)
Method: ITMS; CID

Peptide Mass: 165.7

Protein Group 2080

Scan number 36182 Raw file nak_cdc48UbconjugatecontrolSAX1_22may10_10

m/z (full)

Relative Abundance

m/z (zoom)

Intensity
Protein Group 2114
Scan number: 39190
Raw file: nak_cdc48UbconjugatesSAX2_14may10_06
Method: ITMS; CID
Peptide: 138.94

Protein Group 2249

A₁₋₃₋₄₋₅₋₆₋₇₋₈₋₉₋₁₀₋₁₁₋₁₂-
L₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋₋

m/z (full)

m/z (zoom)

Relative Abundance

Intensity
Scan number 25603 Raw file nak_ubx4LightSAX6_11nov06
Method ITMS; CID Pepti... 85.96

Protein Group 2347

m/z (full)

m/z (zoom)

Relative Abundance

Intensity
Protein Group 2359

Scan number 26107 Raw file nak_ubx4HeavySAX6_11nov11
Method ITMS; CID Pepti... 182.82

m/z (full)

m/z (zoom)

Relative Abundance

Intensity
Scan number 18206
Raw file nak_cdc48UbconjugatesSAX4_14may10_08
Method ITMS; CID Pepti... 147.09

Protein Group 2383

m/z (full)

Relative Abundance

m/z (zoom)

Intensity

m/z (zoom)
Scan number 29108 Raw file nak_cdc48UbconjugatesSAX4_21may15
Method ITMS; CID Pepti...
119.27

Protein Group 2445

m/z (full)

m/z (zoom)

Relative Abundance

Intensity

b₂₃ b₂₆ b₈₂⁺ b₅ b₅-H₂O b₈-H₂O b₉-H₂O b₉
y₂ y₇ y₈ y₉ y₁₀

200 300 400 500 600 700 800 900 1000 1100 1200 1300

0 20 40 60 80 100 120

200 300 400 500 600 700 800 900 1000 1100 1200 1300

0 100 200 300 400 500
Scan number 24193 Raw file nak_UBX2HSAX3_8feb11_07
Method ITMS; CID Pepti... 177.78

Relative Abundance
0 500 1000 1500
Intensity
200 300 400 500 600 700 800 900 1000 1100
m/z (zoom)

m/z (full)
Scan number 29177
Raw file nak_cdc48UbconjugatecontrolSAX4_22may10_13
Method ITMS; CID Pepti... 140.49

Protein Group 2553

m/z (full)

m/z (zoom)
Scan number 3779 Raw file nak_cdc48UbconjugatesSAX5_22may10_6
Method ITMS; CID Pepti... 77.64

Protein Group 2580

m/z (full)

m/z (zoom)

Relative Abundance

Intensity
Scan number 10035 Raw file nak_UBX2HSAX3_8feb11_07
Method ITMS; CID Pepti... 96.64

Protein Group 2596

m/z (full)

200 300 400 500 600 700 800 900 1000 1100 1200 1300

m/z (zoom)

0 500 1000 1500 2000 2500 3000 3500 4000 4500 5000 5500 6000

Relative Abundance

0 20 40 60 80 100 120

Intensity

0 500 1000 1500

y₁₁²⁺
A
-NH₃
y₁₀
Wb₂
y₉
Gb₃
y₈
Gb₄
y₇
D
-H₂O
y₆
Vb₆ G
-H₂O
N
-NH₃
-H₂O
D
y₂
E Tb₁₁ K _

a₂ 230.13
y₂ 256.17
b₂ 258.12
b₃ 315.15
y₃-H₂O 367.21
b₄ 372.17
b₅-H₂O 469.18
y₄-H₂O 482.23
y₁₀²⁺ 500.23
b₆ 586.26
y₁₁²⁺ 593.27
y₆ 671.31
b₈-NH₃ 740.3
y₇ 770.38
y₈ 885.4
y₉ 942.43
y₁₀-NH₃ 982.42
y₁₀ 999.45
b₁₁ 1102.44
Scan number 11116

Raw file nak_cdc48UbconjugatesSAX5_14may10_09

Method ITMS; CID

Peptidome 71.56
Scan number: 19097
Raw file: nak_UBX3HSAX4_28jan11_30
Method: ITMS; CID
Peptid... 110.53