

Targets

Replicates: 150323\_QPrESTmid

- IGF2
  - R.GFYFSRPASRV [24, 33]
    - 594.3014++ (dotop 0)
      - precursor - 594.3014++ (rank 1)
      - precursor [M+1] - 594.8029++ (rank 2)
      - precursor [M+2] - 595.3042++ (rank 3)
      - Y [b]8 - 983.5057+
      - F [b]7 - 820.4424+
      - S [b]6 - 673.3740+
      - P [b]4 - 430.2409+
      - S [b]5 - 602.2609+
      - R [b]6 - 758.3620+
      - P [b]7 - 855.4148+
    - 599.3056++ (heavy) (dotop 0.54)
      - precursor - 599.3056++ (rank 1)[1]
      - precursor [M+1] - 599.8071++ (rank 2)
      - precursor [M+2] - 600.3084++ (rank 3)
      - Y [b]8 - 993.5140+
      - F [b]7 - 830.4507+
      - S [b]6 - 683.3823+
      - P [b]4 - 440.2491+
      - S [b]5 - 602.2609+
      - R [b]6 - 758.3620+
      - P [b]7 - 855.4148+
  - R.GIVEECCFRS [40, 48]
    - 585.2575++ (dotop 0.96, dotop 0.61, rdopt 1, total ratio 0.06)
      - precursor - 585.2575++ (rank 1)[1] (ratio 0.06)
      - precursor [M+1] - 585.7589++ (rank 2)[2] (ratio 0.07)
      - precursor [M+2] - 586.2588++ (rank 3)[3] (ratio 0.06)
      - E [b]6 - 900.3338+ (rank 1)[4] (ratio 0.08)
      - E [b]5 - 771.2913+ (rank 3)[5] (ratio 0.08)
      - R [b]1 - 175.1190+ (rank 2)
    - 590.2617++ (heavy) (dotop 0.98, dotop 0.61)
      - precursor - 590.2617++ (rank 1)[1]
      - precursor [M+1] - 590.7631++ (rank 2)[2]
      - precursor [M+2] - 591.2627++ (rank 3)[3]
      - E [b]6 - 910.3421+ (rank 1)[4]
      - E [b]5 - 781.2995+ (rank 3)[5]
      - R [b]1 - 185.1272+ (rank 2)
- IGFBP3
  - R.ETEYGPQR.R [72, 79]
    - 506.2136++ (dotop 0.55, dotop 0, total ratio 0.003)
      - precursor - 506.2136++ (rank 1)[1] (ratio 0.003)
      - precursor [M+1] - 506.7151++ (rank 2)
      - precursor [M+2] - 507.2153++ (rank 3)
      - Y [b]5 - 652.2872+ (rank 2)
      - G [b]4 - 489.2238+ (rank 1)
      - P [b]3 - 432.2024+ (rank 3)
    - 511.2178++ (heavy) (dotop 0.96, dotop 0.97)
      - precursor - 511.2178++ (rank 1)[1]
      - precursor [M+1] - 511.7192++ (rank 2)[2]
      - precursor [M+2] - 512.2192++ (rank 3)[3]
      - Y [b]5 - 662.2954+ (rank 2)[5]
      - G [b]4 - 499.2321+ (rank 1)[4]
      - P [b]3 - 442.2106+ (rank 3)[6]
- IGFBP5

150323\_QPrESTmid

Peptide Settings

Digestion Prediction Filter Library Modifications

Structural modifications:  
 Carbamidomethyl (C) Edit list...  
 Carbamidomethyl Cysteine  
 Phospho (Y)  
 Phospho (S,T)  
 Deamidated (NQ)

Max variable mods: 3 Max neutral losses: 1

Isotope label type: heavy

Isotope modifications:  
 Label:13C(6)15N(2) (C-term K) Edit list...  
 Label:13C(6)15N(4) (C-term R)

Internal standard type: heavy

OK Cancel

