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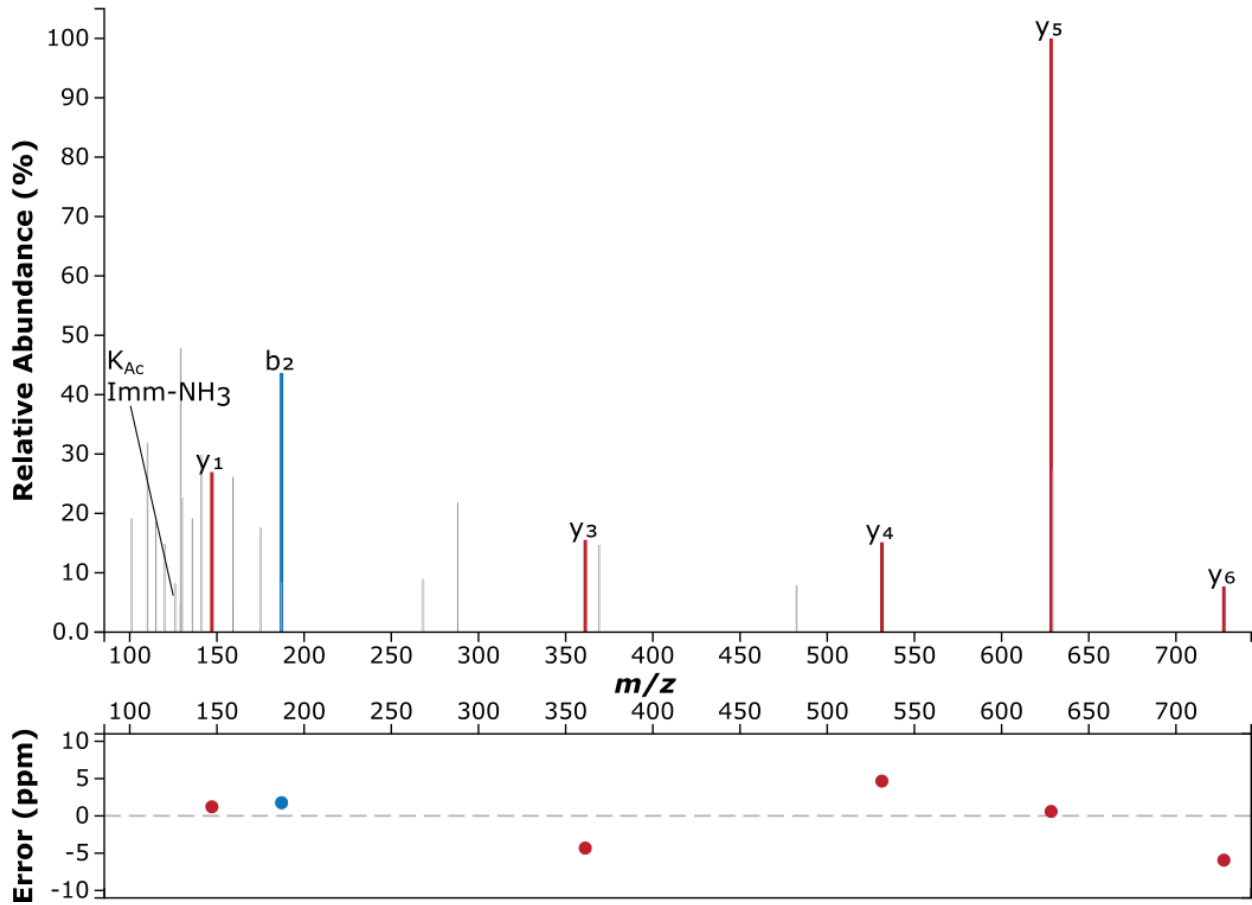
Acetyl

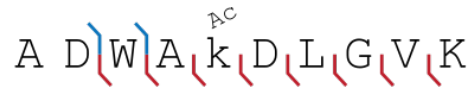


Precursor m/z: 457.2713

Charge: +2

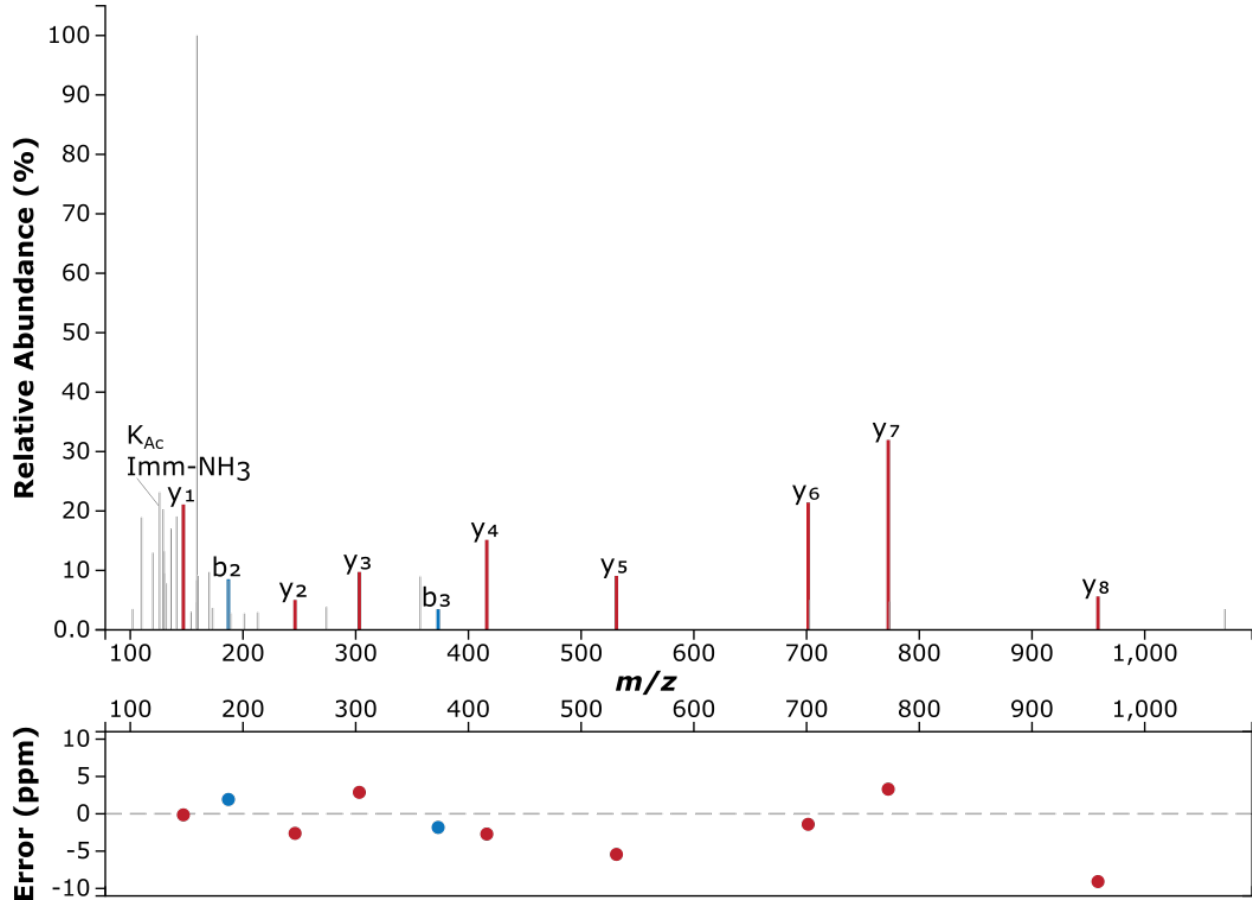
6-hydroxycyclohex-1-ene-1-carboxyl-CoA dehydrogenase (Q2LXU3)





Precursor m/z: 572.8035 Charge: +2

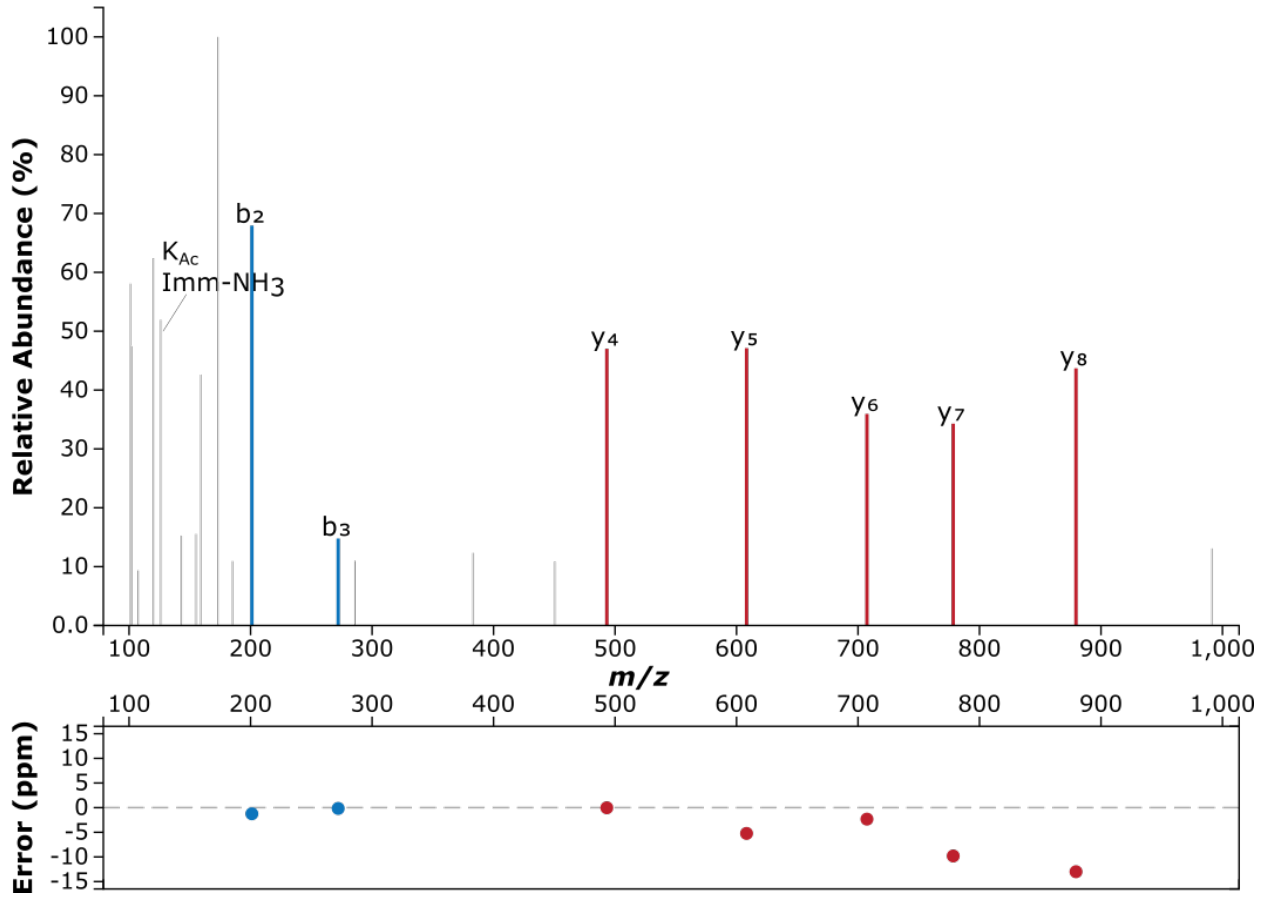
Iron-sulphur-binding reductase (Q2LWR1)

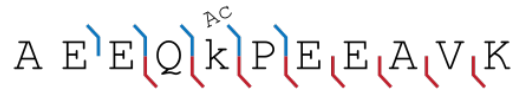


A E A E k T I Q T A V D N F G R

Precursor m/z: 597.9672
3-oxoacyl-reductase (Q2LXS6)

Charge: +3

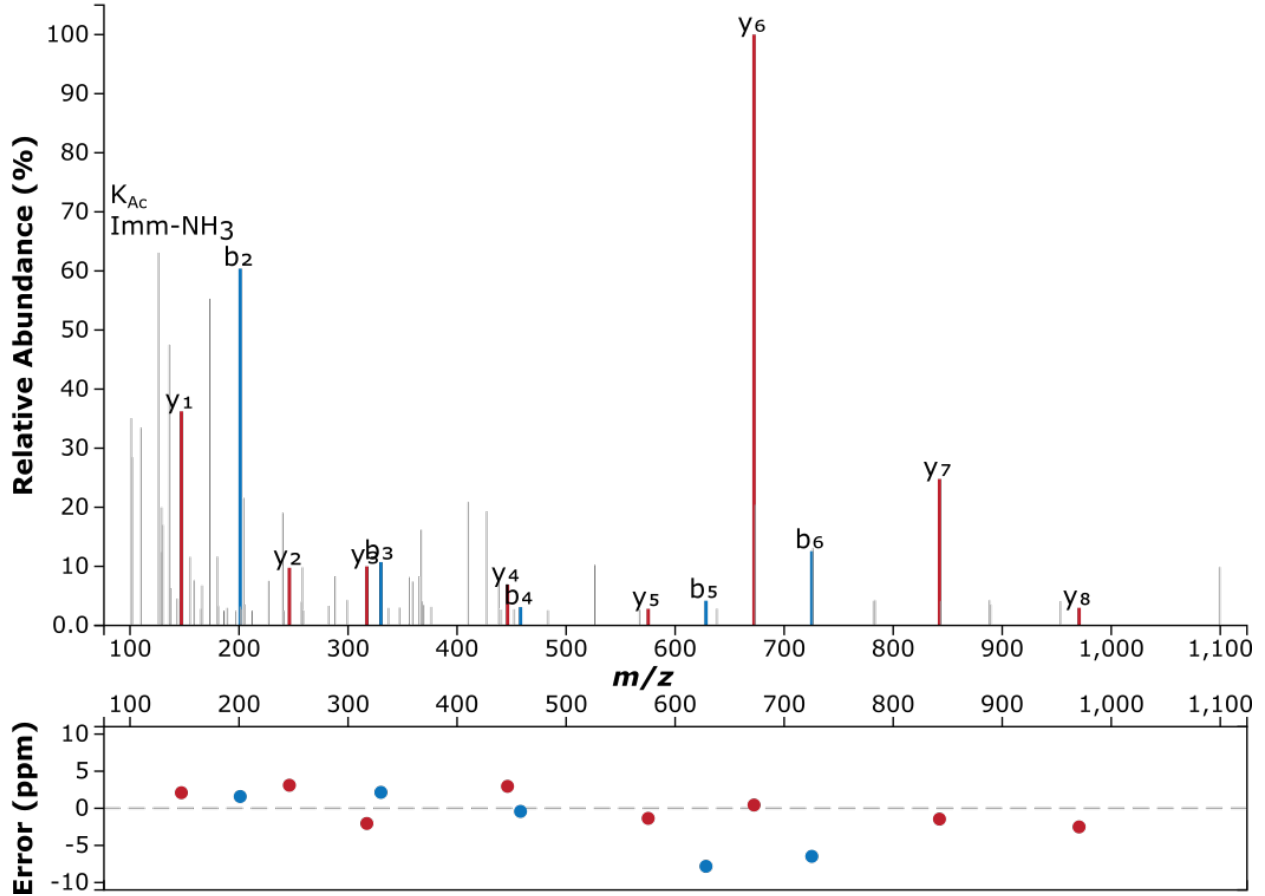




Precursor m/z: 650.3250

Charge: +2

Hypothetical exported protein (Q2LUP5)

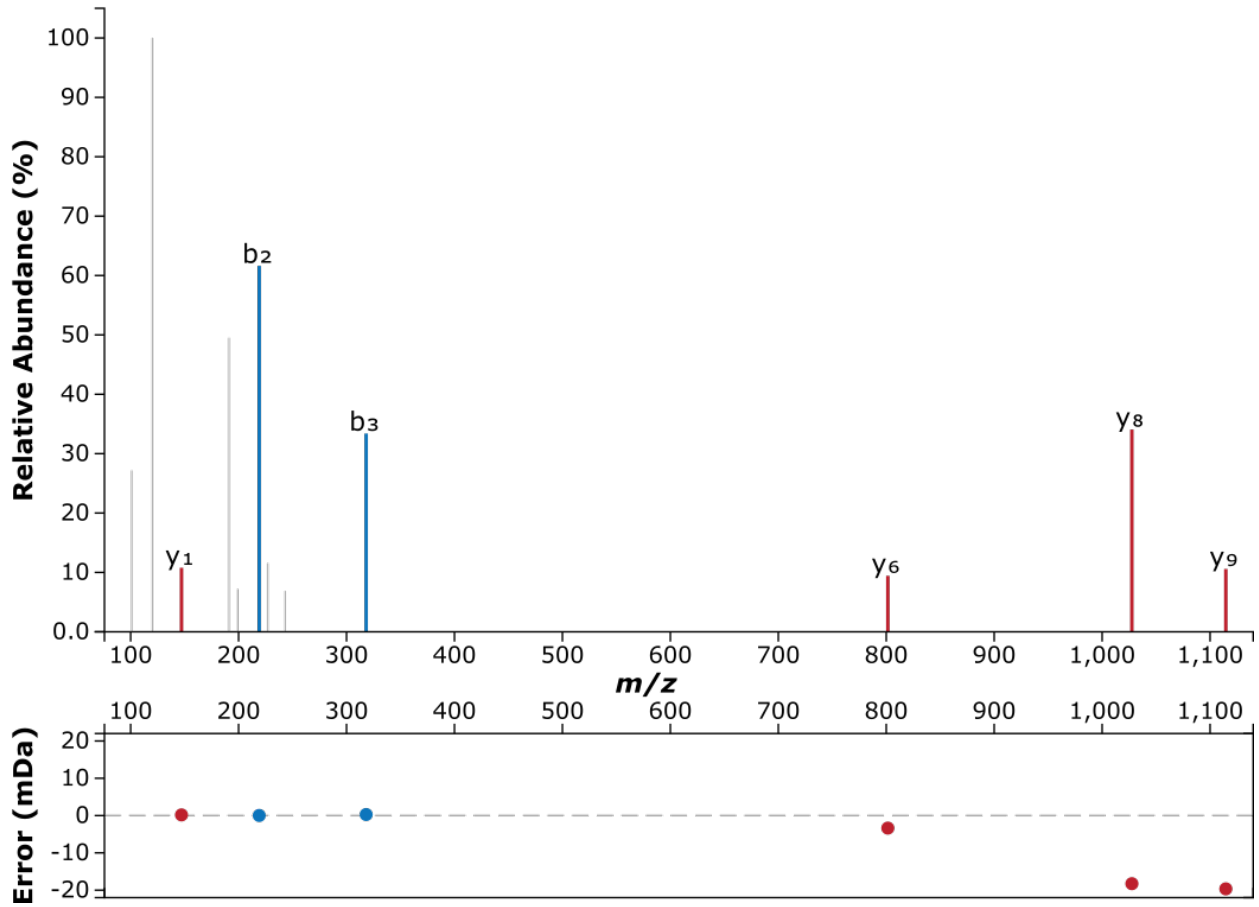


A F¹V¹V L N Q G F A Q₁S₁P E₁L E^{AC} k E L₁K

Precursor m/z: 763.7423

Charge: +3

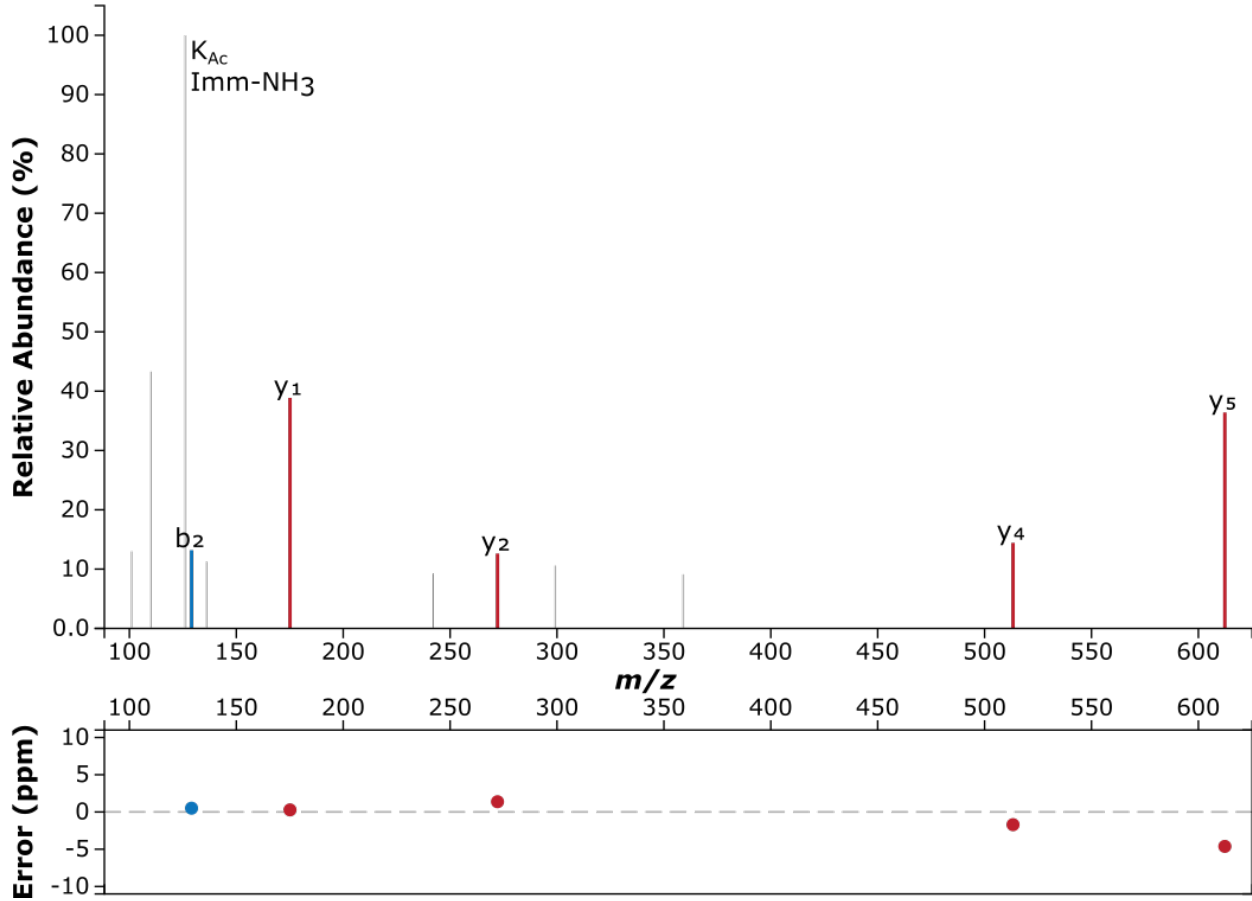
4-hydroxybenzoate-CoA ligase / benzoate-CoA ligase (Q2LRH0)





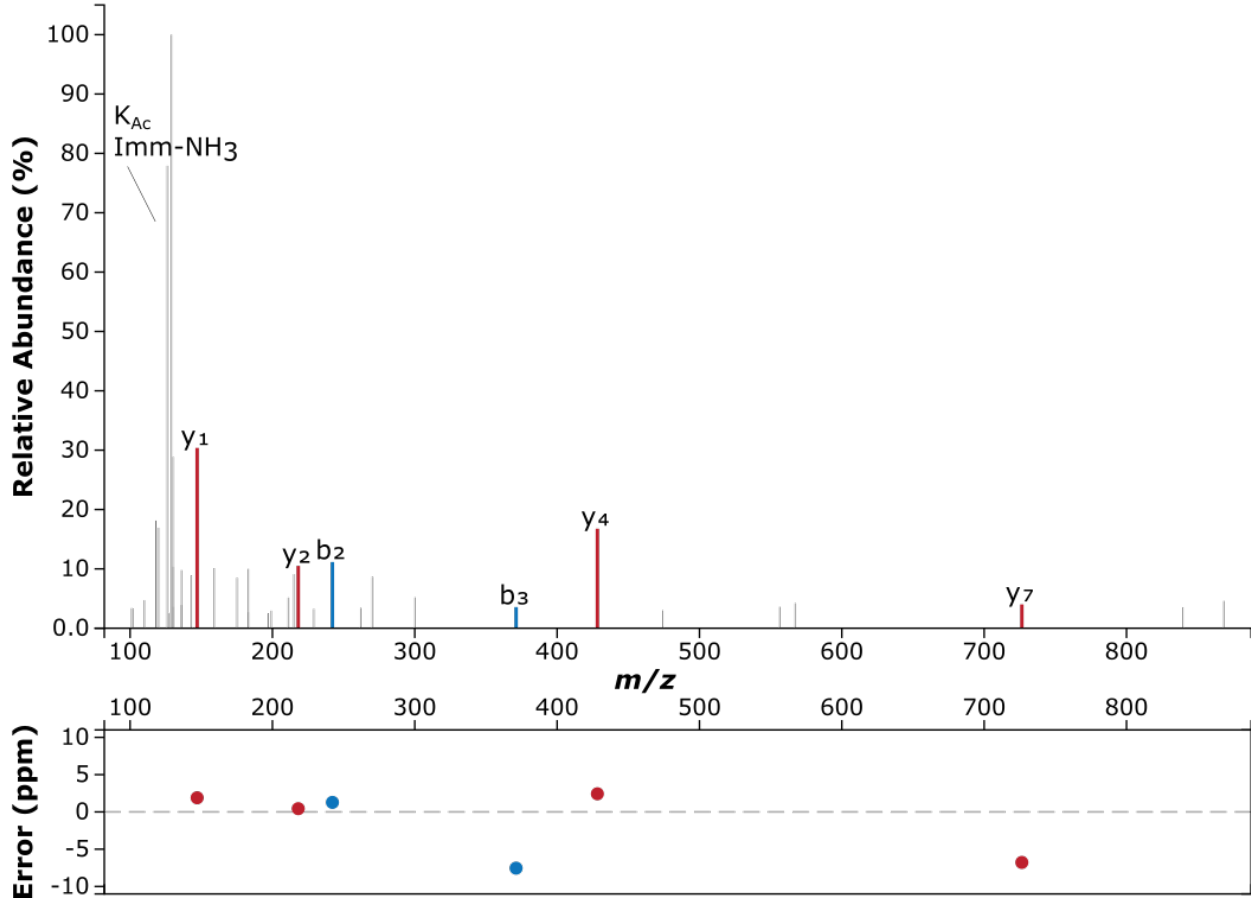
Precursor m/z: 491.2956
Aspartokinase (Q2LTK0)

Charge: +2





Precursor m/z: 403.9237 Charge: +3
 Acetyl-CoA acetyltransferase (Q2LWQ5)

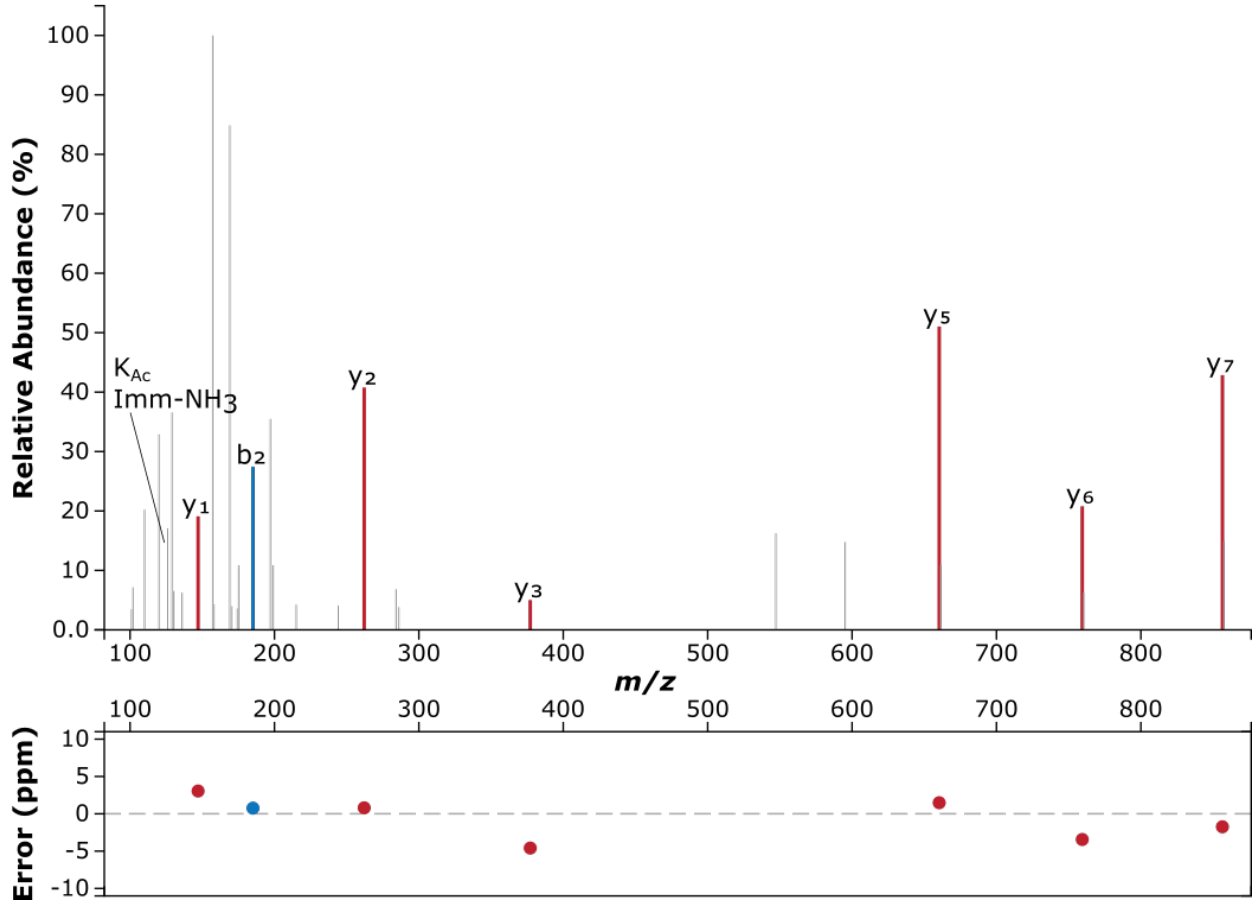




Precursor m/z: 520.8029

Charge: +2

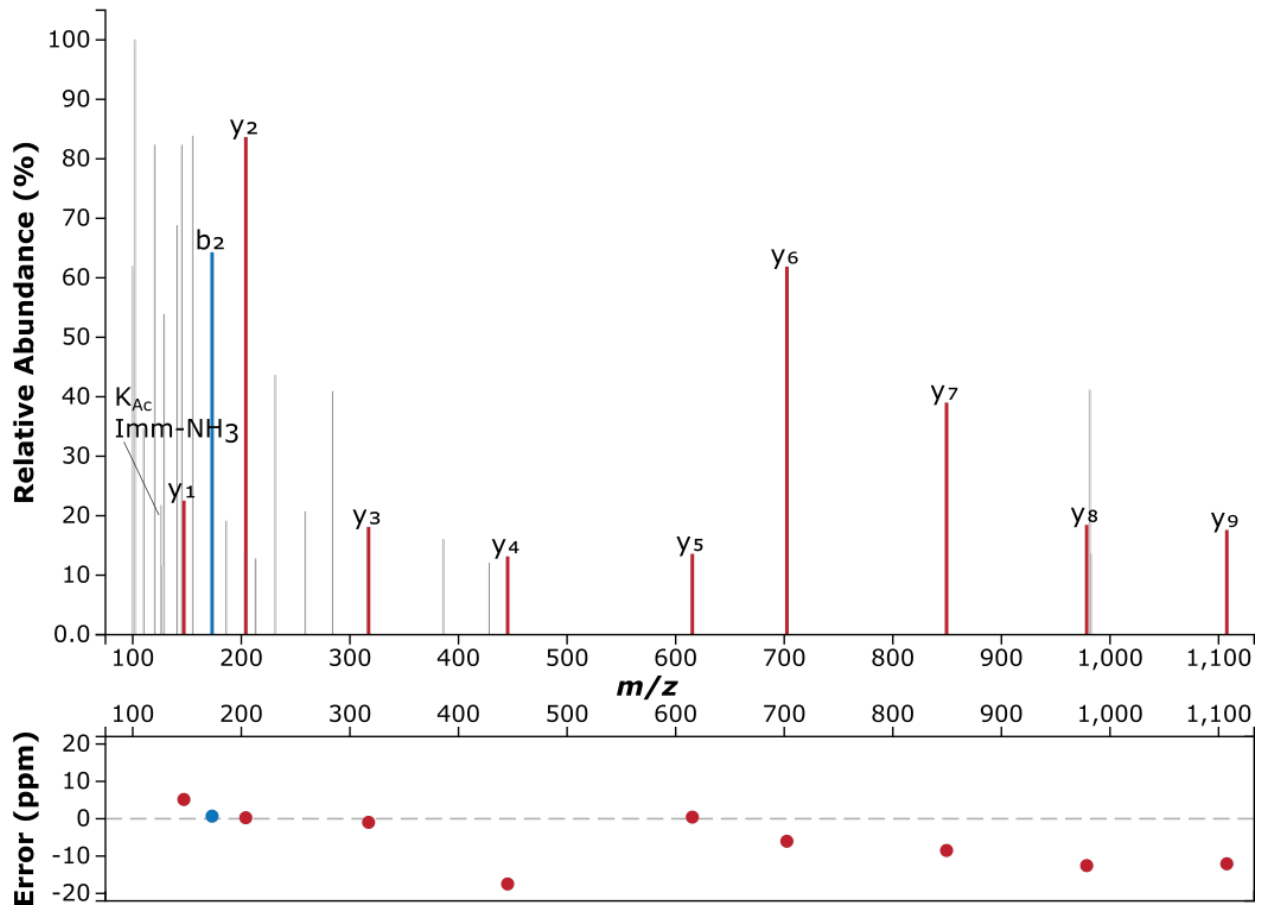
6-oxocyclohex-1-ene-1-carboxyl-CoA hydrolase (Q2LXU2)





Precursor m/z: 640.3301 Charge: +2

3-hydroxyacyl-CoA dehydrogenase (Q2LUN4)

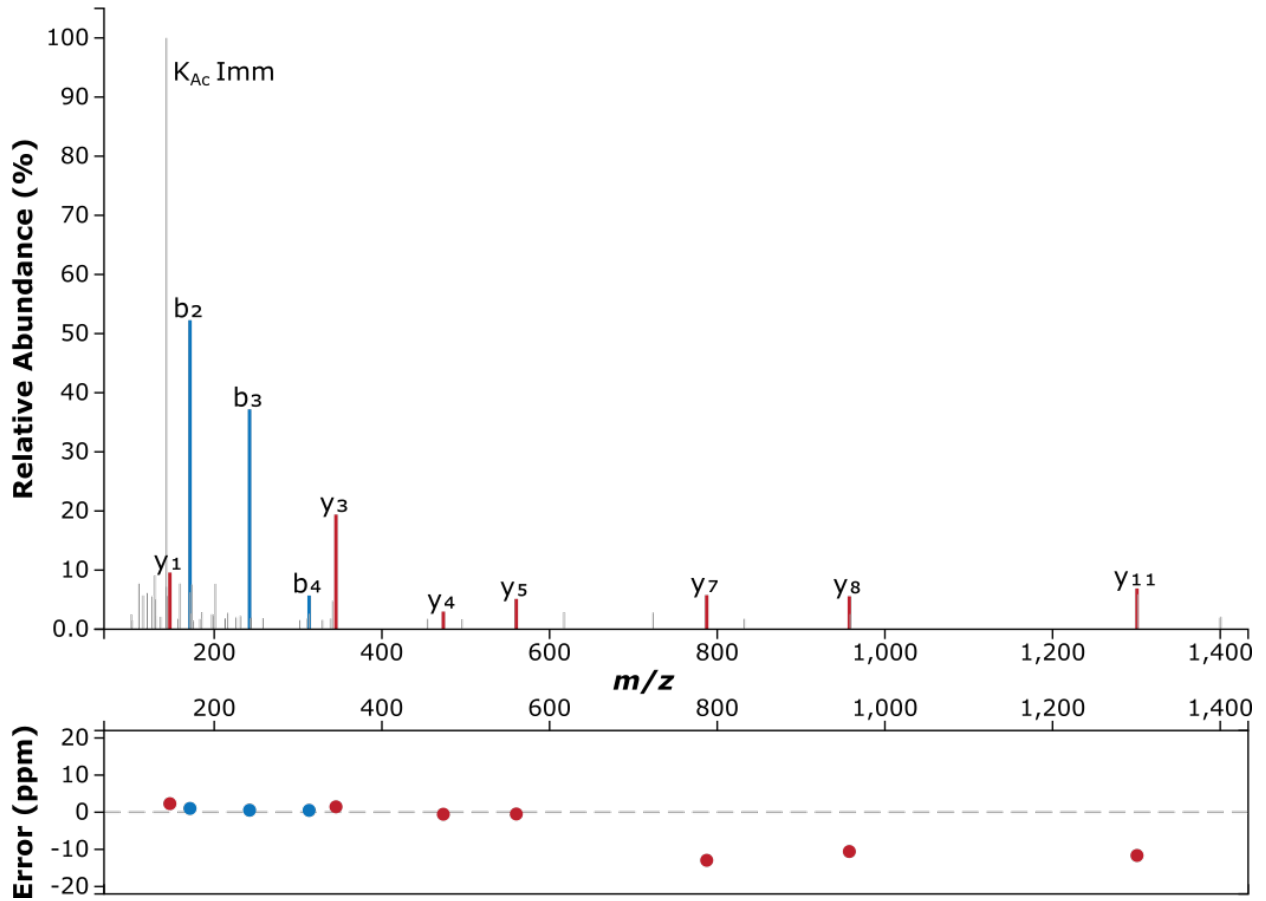




Precursor m/z: 604.3594

Charge: +3

60 kDa chaperonin 2 (Q2LTG7)

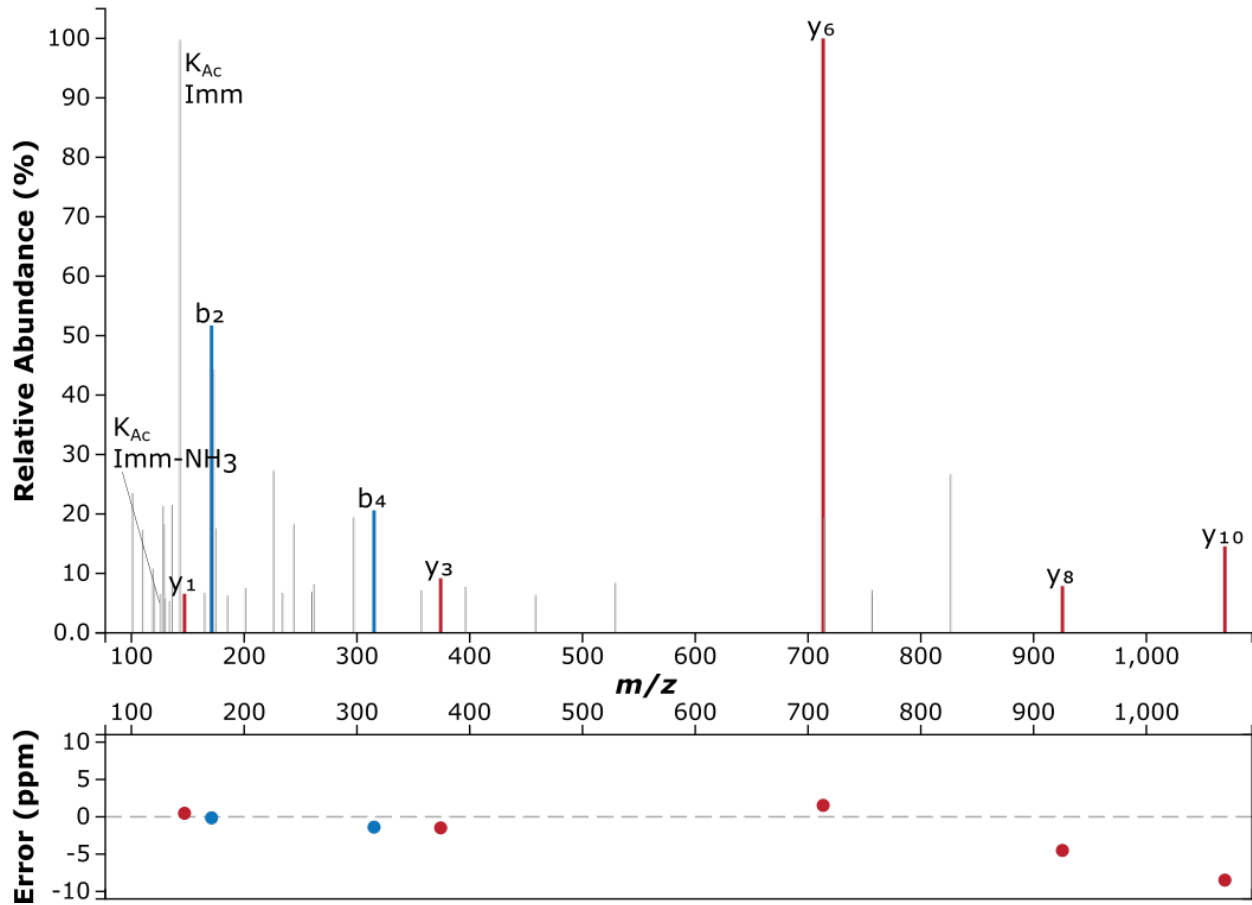


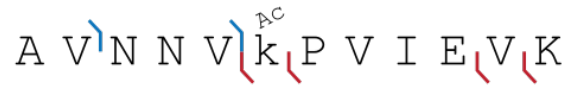


Precursor m/z: 620.3690

Charge: +2

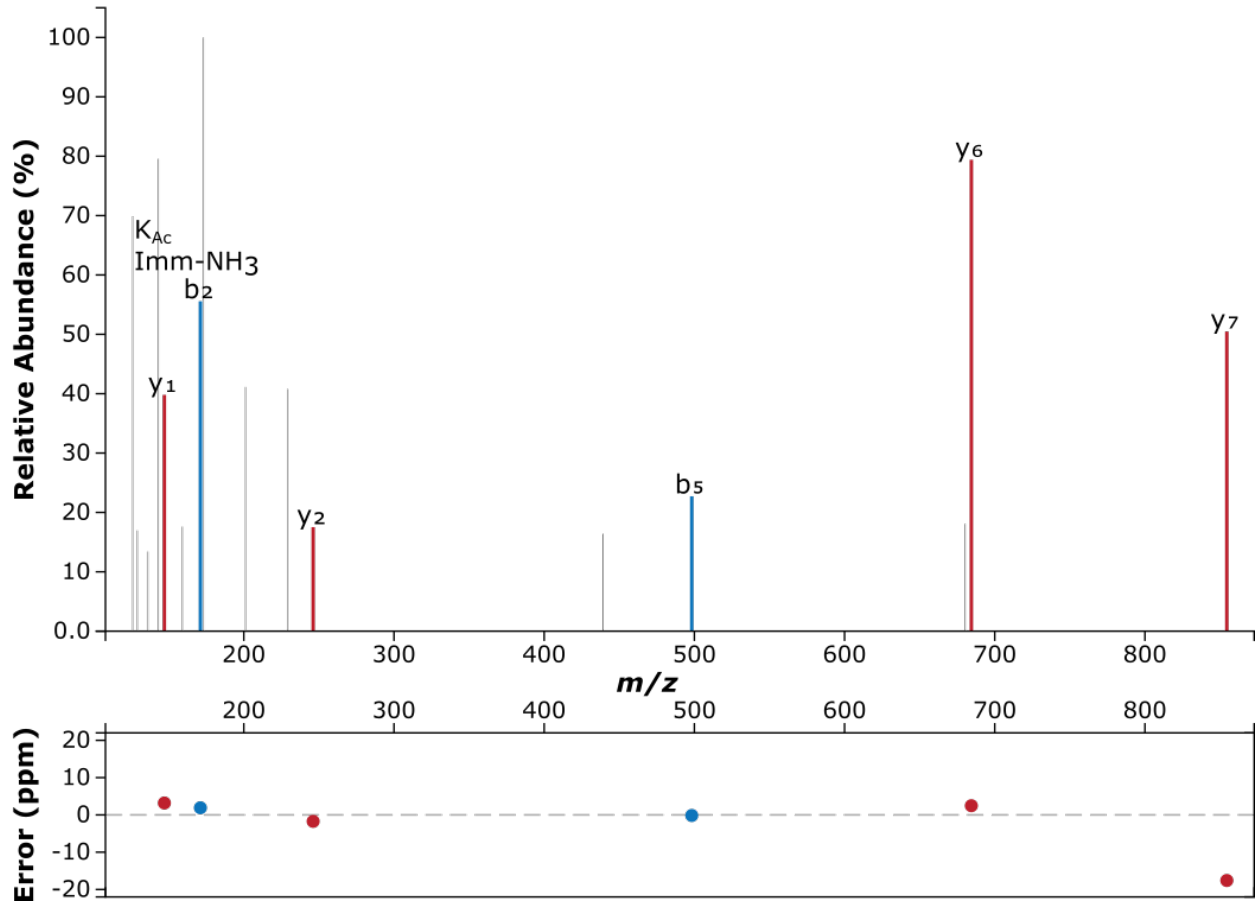
Glyceraldehyde-3-phosphate dehydrogenase (Q2LWW2)





Precursor m/z: 676.4008 Charge: +2

30S ribosomal protein S7 (Q2LQA4)

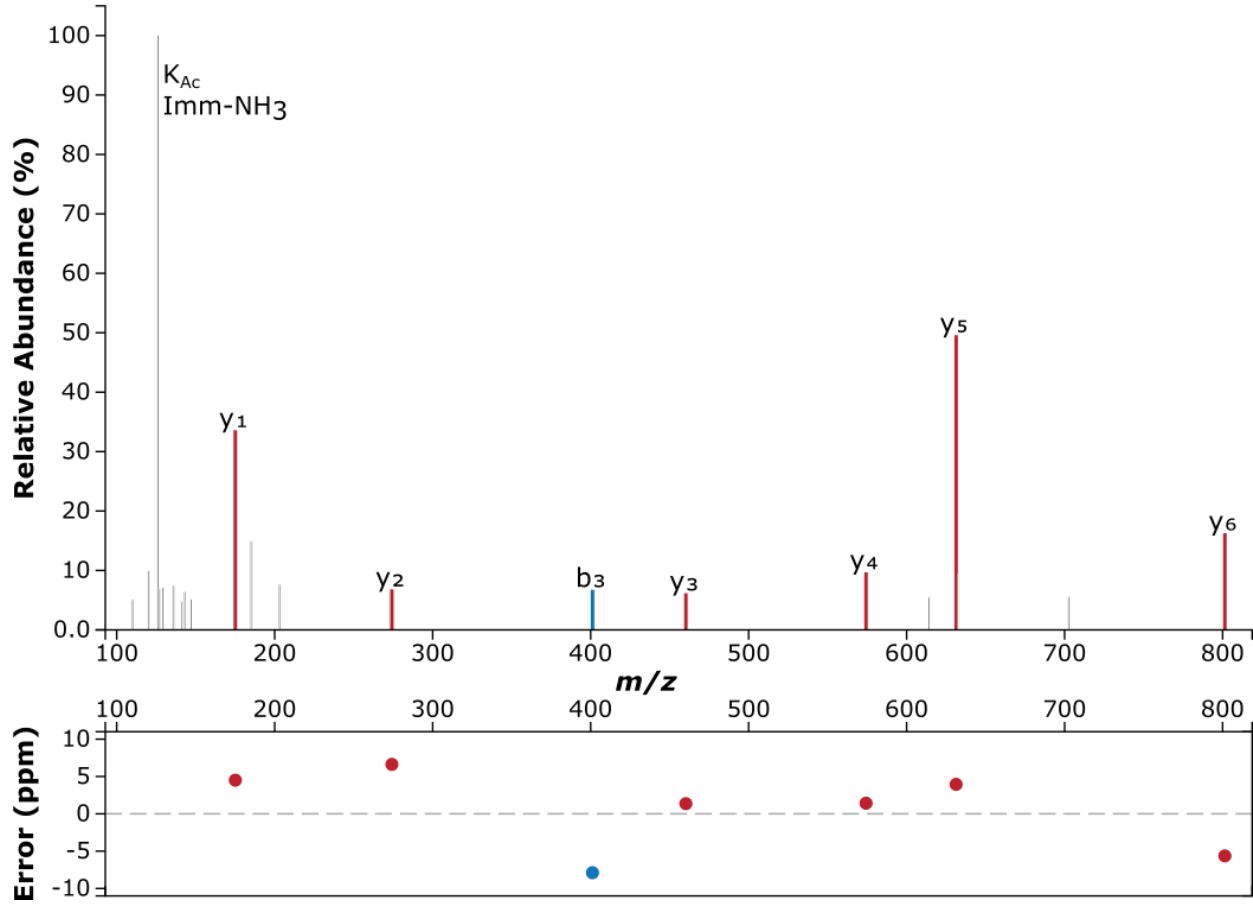


D D ^{AC}k | G | N | W | V | R

Precursor m/z: 516.2489

Charge: +2

6-oxocyclohex-1-ene-1-carboxyl-CoA hydrolase (Q2LXU2)

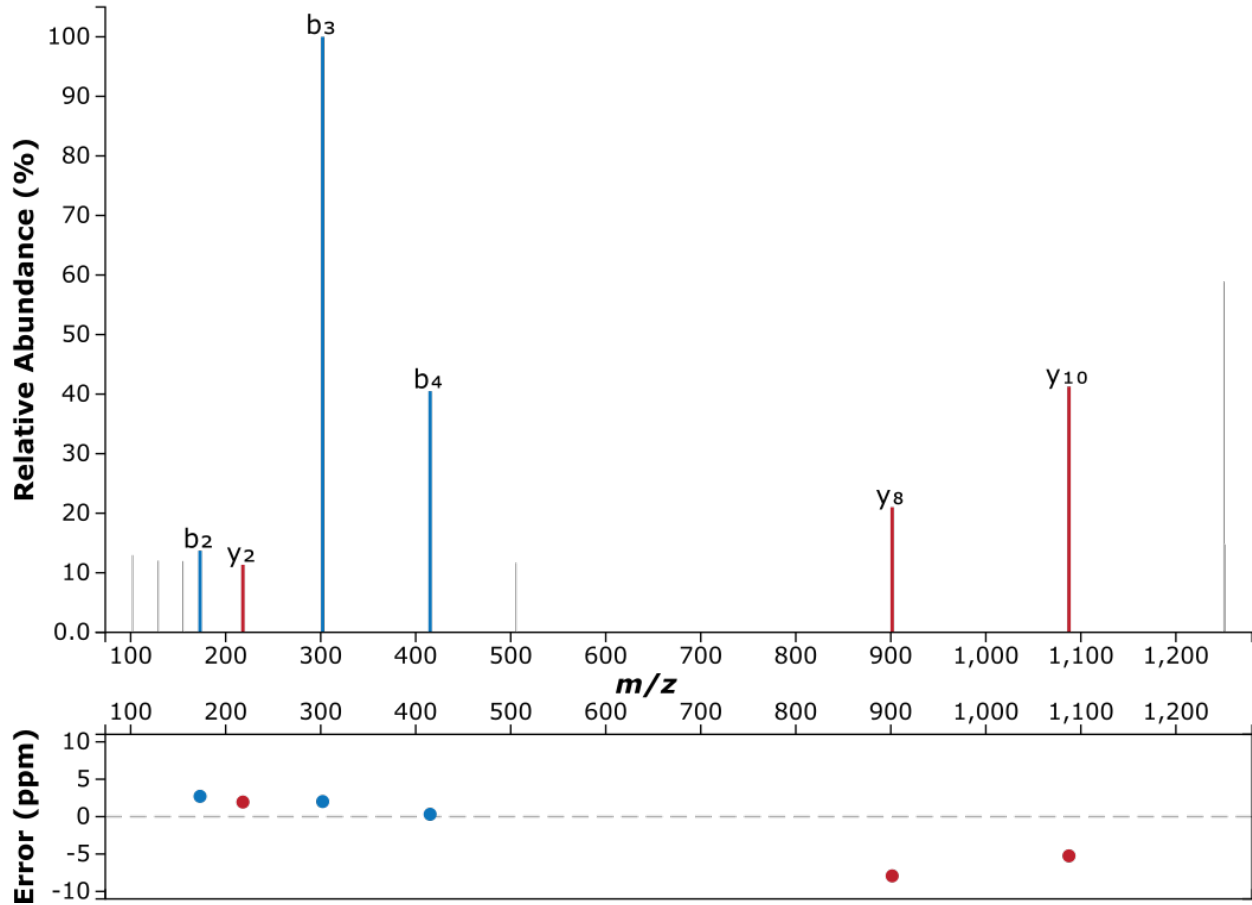


D G E I V Y G E P ^{AC} k T G E E A K

Precursor m/z: 882.4203

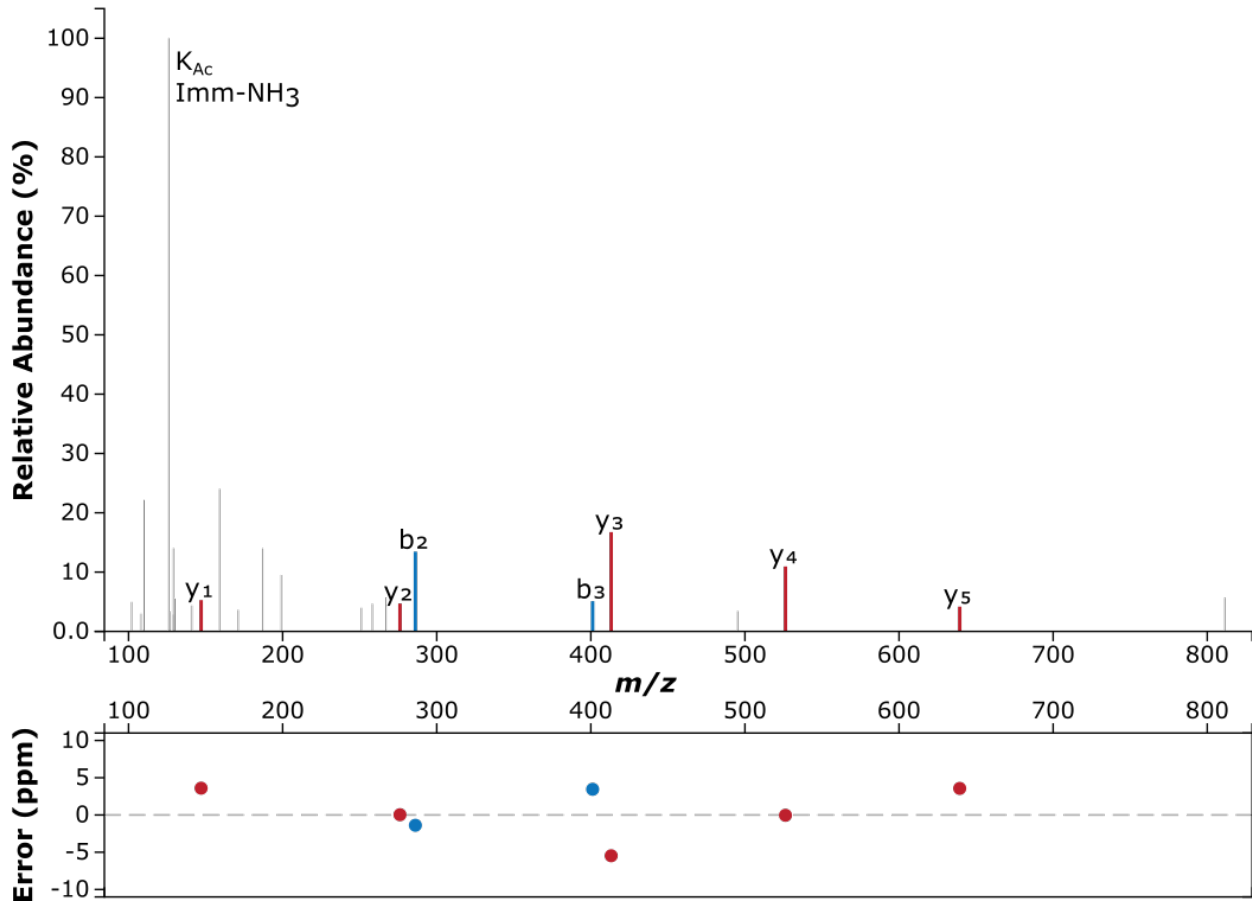
Charge: +2

6-oxocyclohex-1-ene-1-carbonyl-CoA hydrolase (Q2LXU2)





Precursor m/z: 428.2140 Charge: +3
 Xaa-pro aminopeptidase (Q2LRF2)

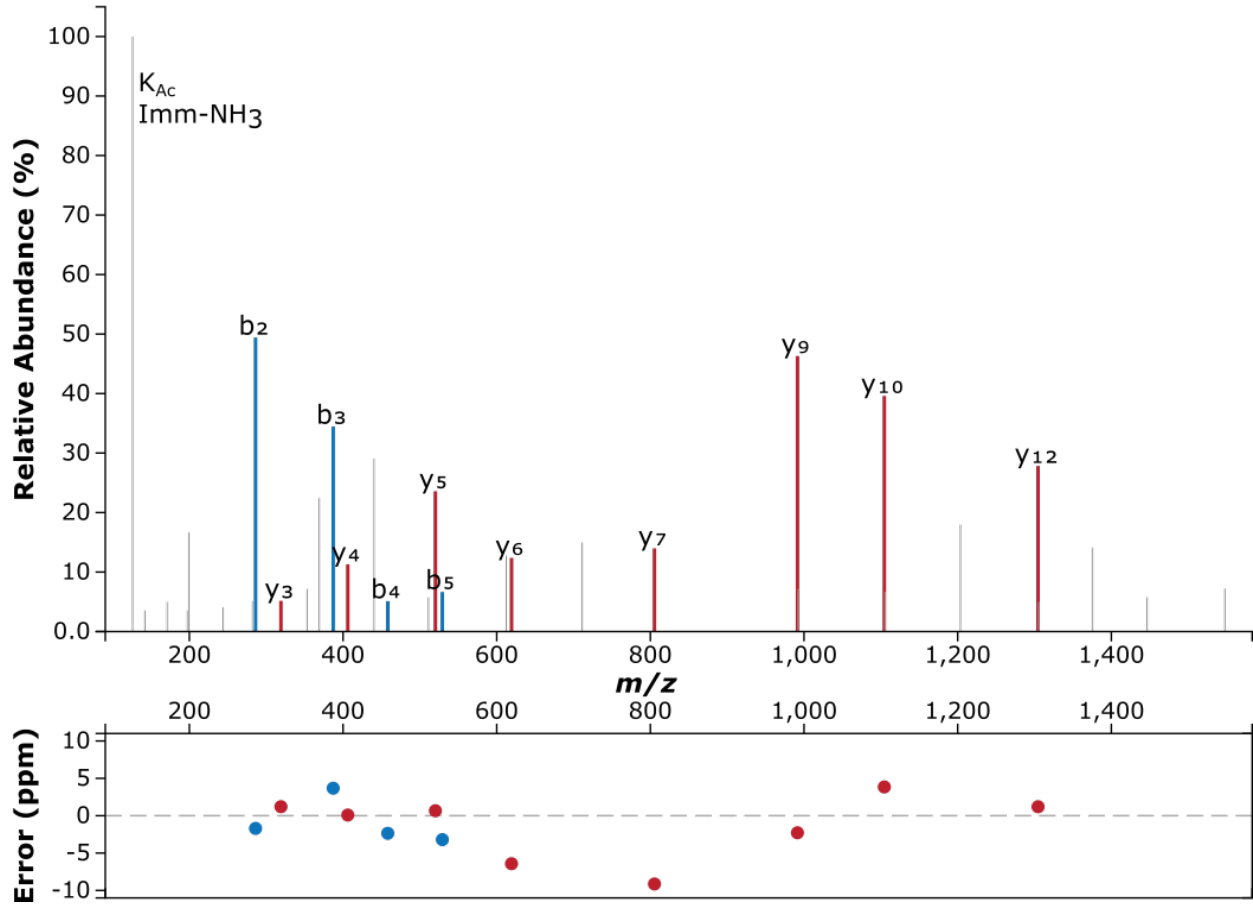


D ¹³C T A A T V I G E W V N S G D K

Precursor m/z: 916.9549

Charge: +2

4-hydroxybenzoate-CoA ligase / benzoate-CoA ligase (Q2LRH0)

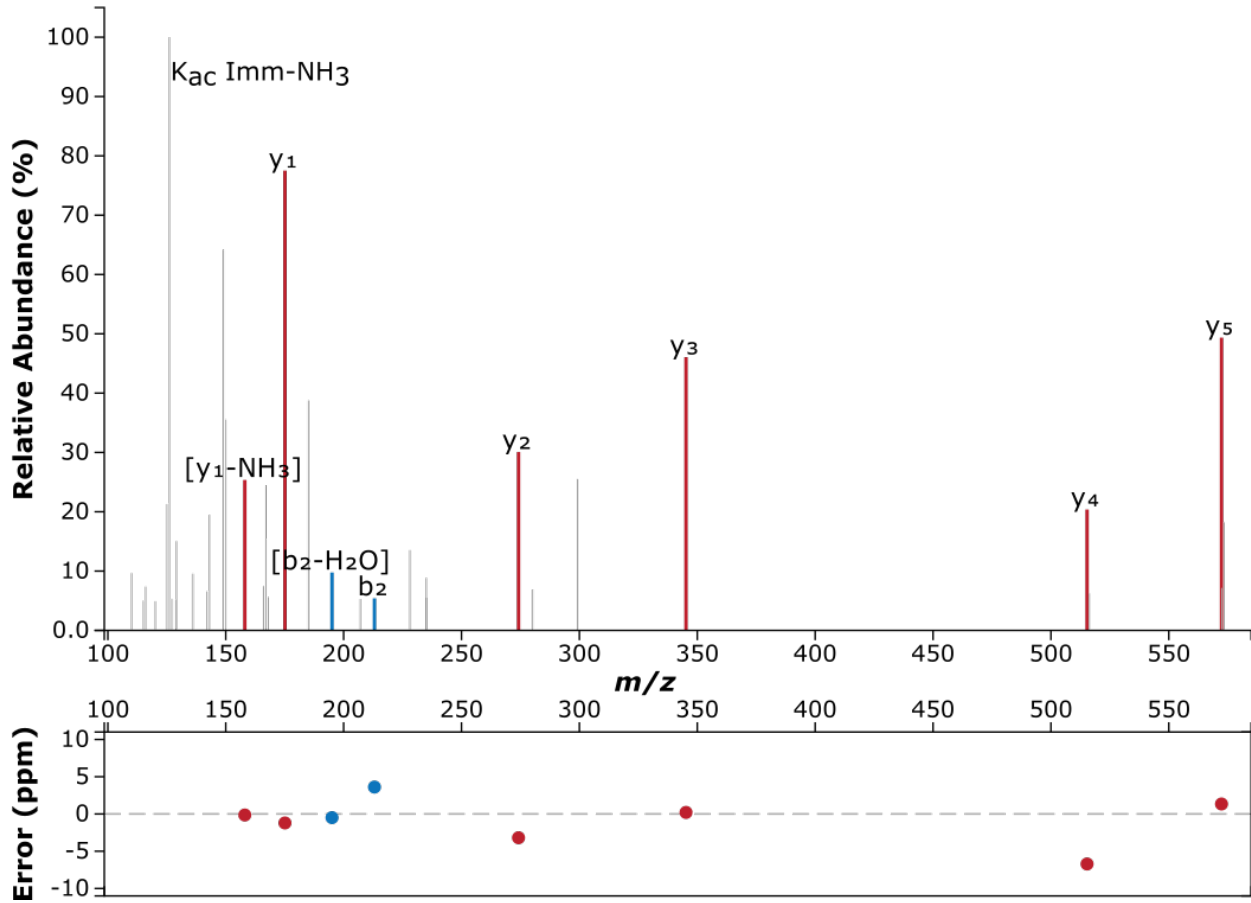




Precursor m/z: 392.7192

Charge: +2

Ion-translocating oxidoreductase complex subunit B (Q2LPK2)

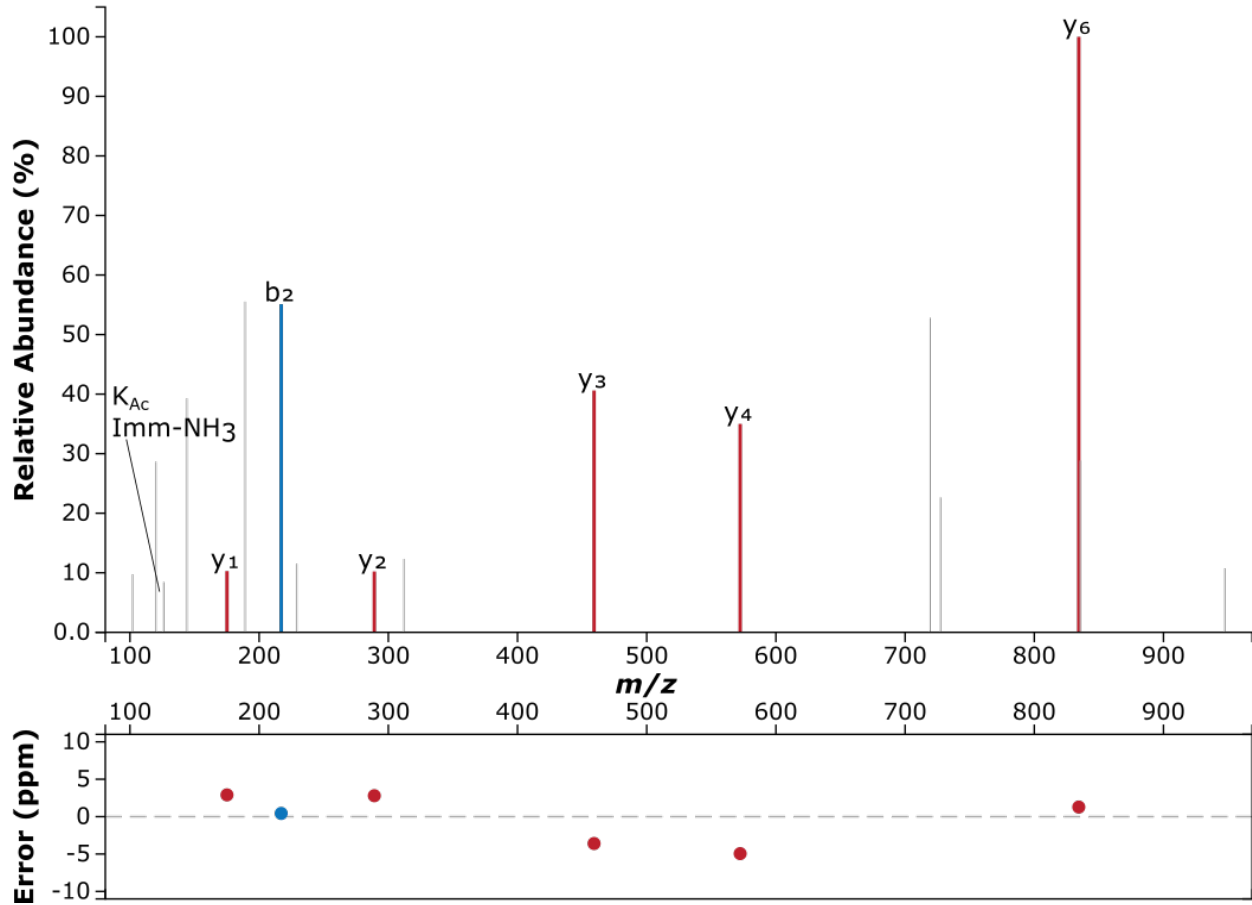


D T I D F I k N R

Precursor m/z: 582.3064

Charge: +2

6-oxocyclohex-1-ene-1-carboxyl-CoA hydrolase (Q2LXU2)

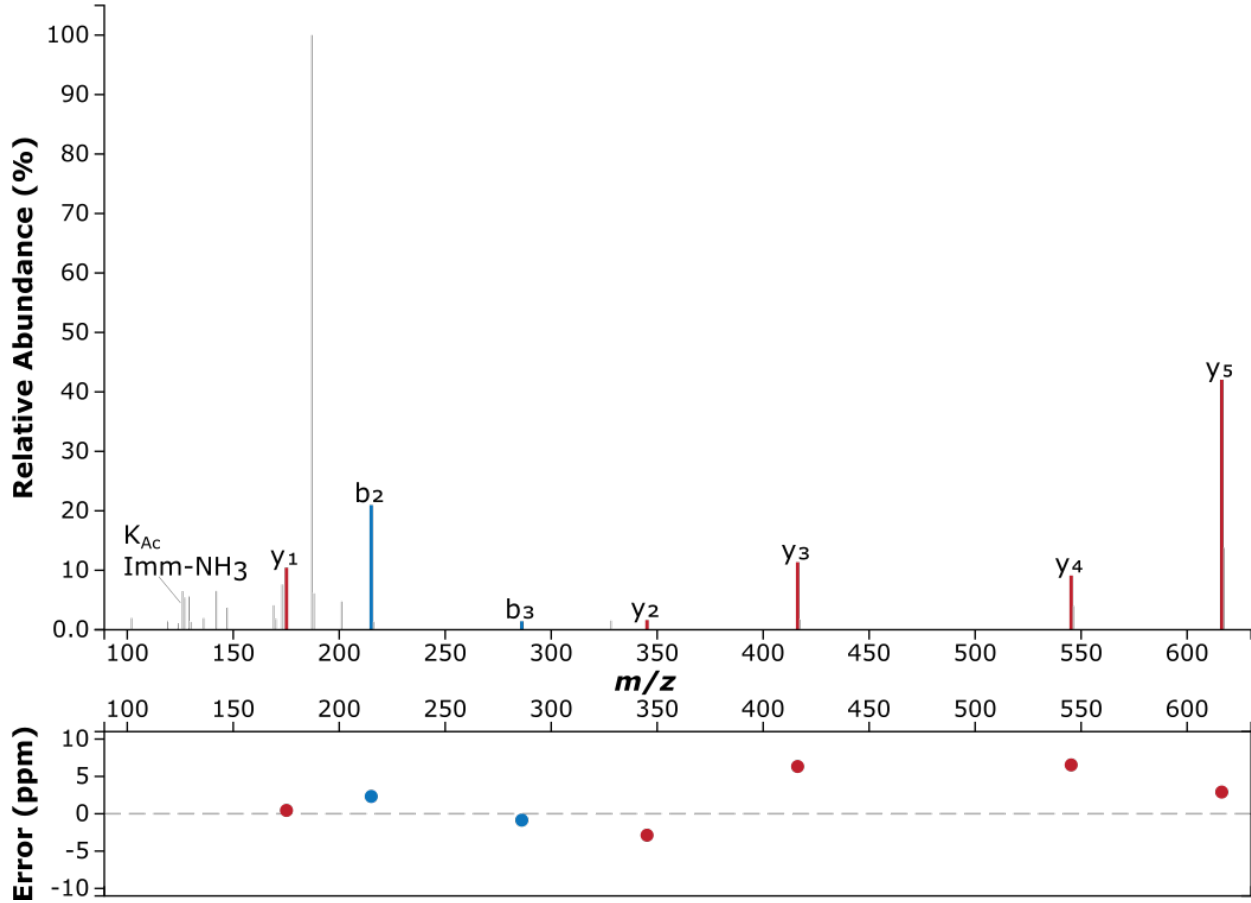




Precursor m/z: 415.7220

Charge: +2

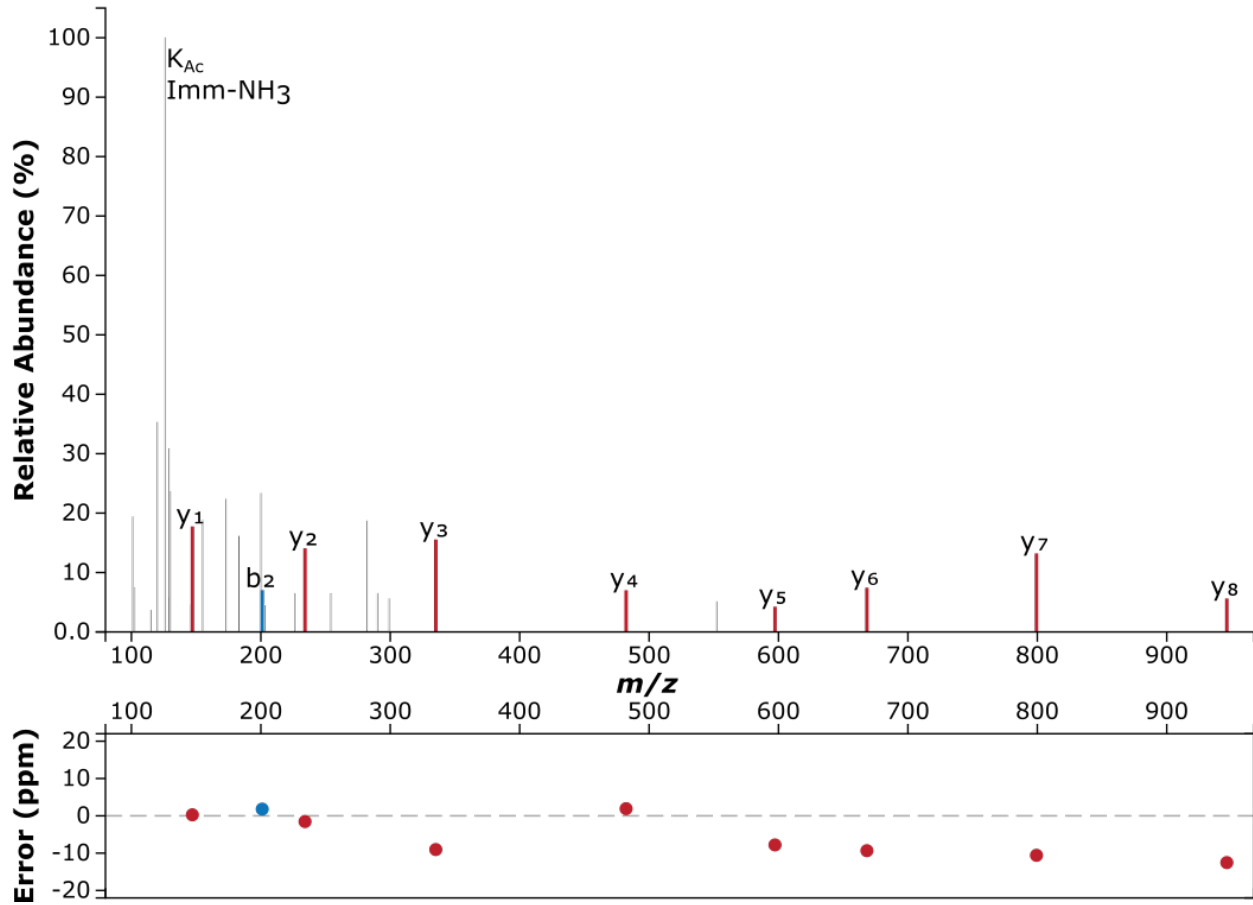
ATP synthase subunit b (Q2LR00)



E A A Q k^{AC} F M A D F T S K

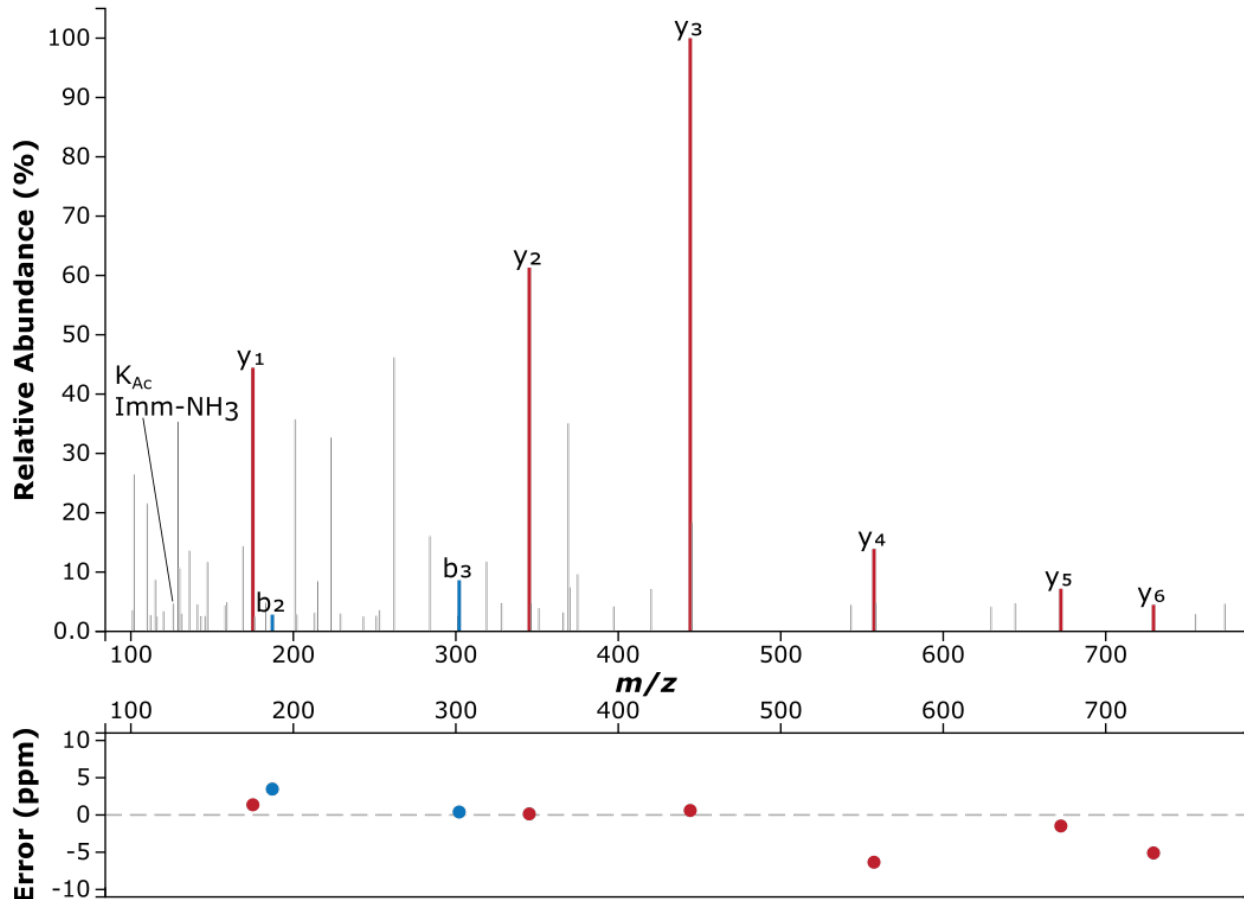
Precursor m/z: 758.3610 Charge: +2

Putative enoyl-CoA hydratase (Q2LXU6)





Precursor m/z: 429.7376 Charge: +2
 ATP synthase subunit alpha 1 (Q2LQZ7)

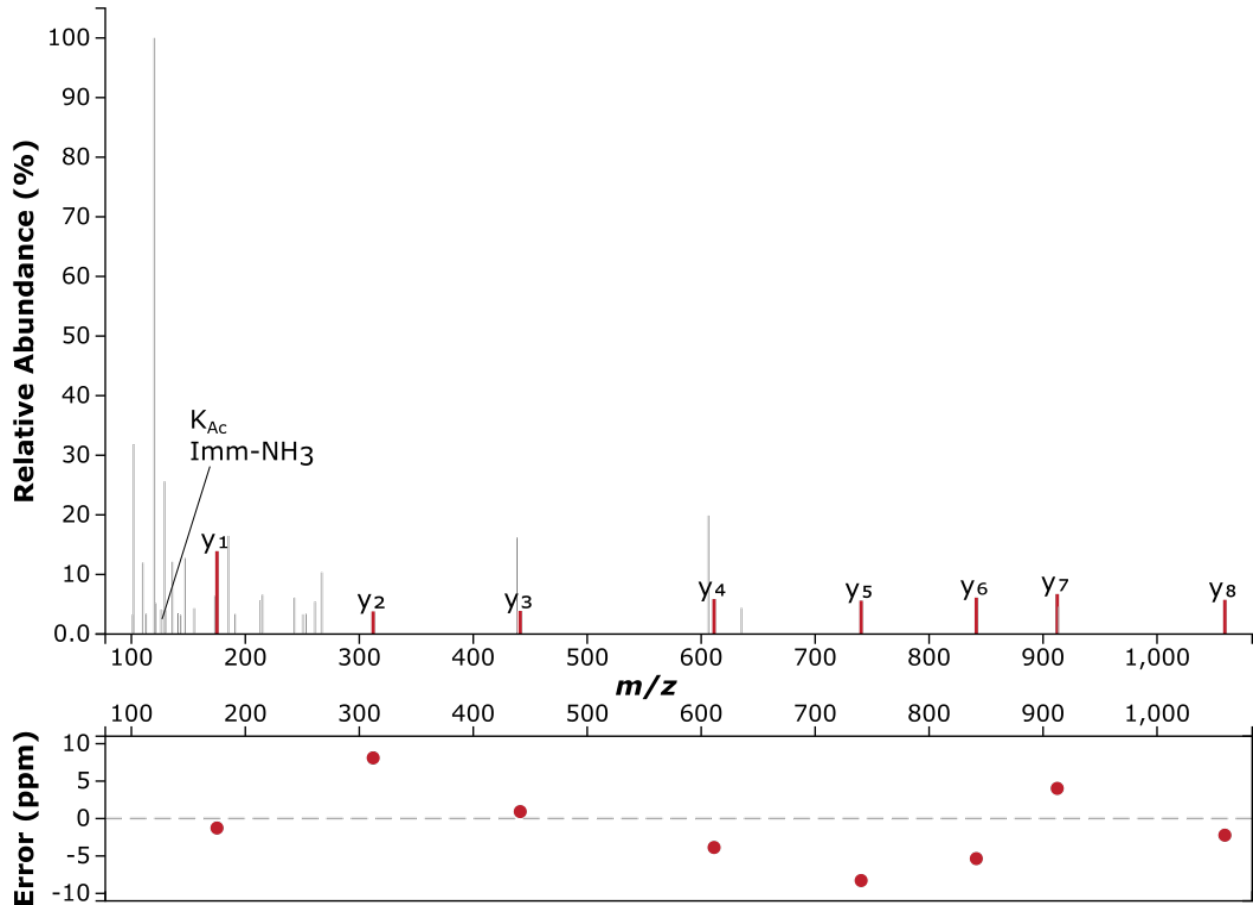


E G I R P A P G T E L F F F A T E ^{Ac} k E H R

Precursor m/z: 582.5499

Charge: +4

2,3-bisphosphoglycerate-independent phosphoglycerate mutase 1 (Q2LWQ4)

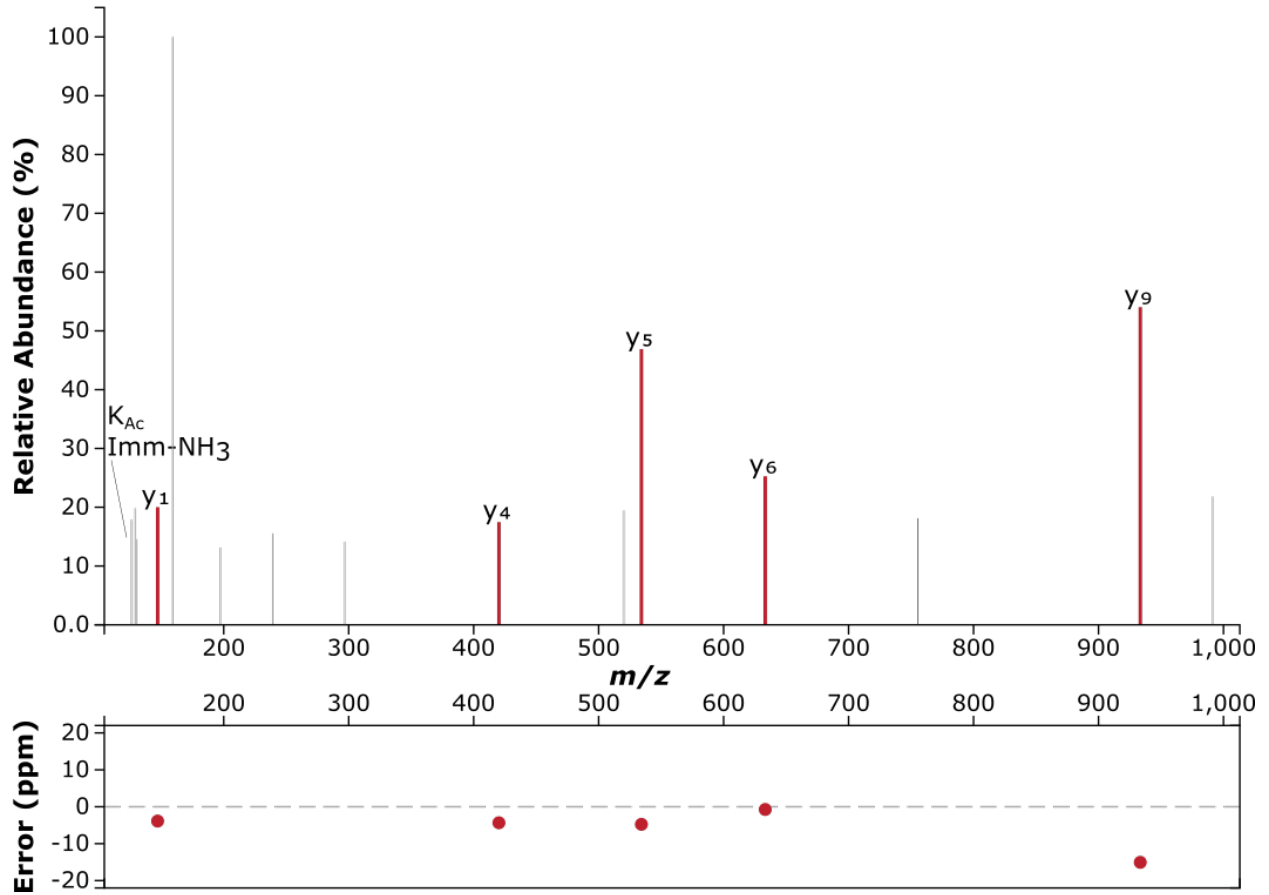


E k T A S T I I ^{Ac} G G W V N T G D K

Precursor m/z: 606.9793

Charge: +3

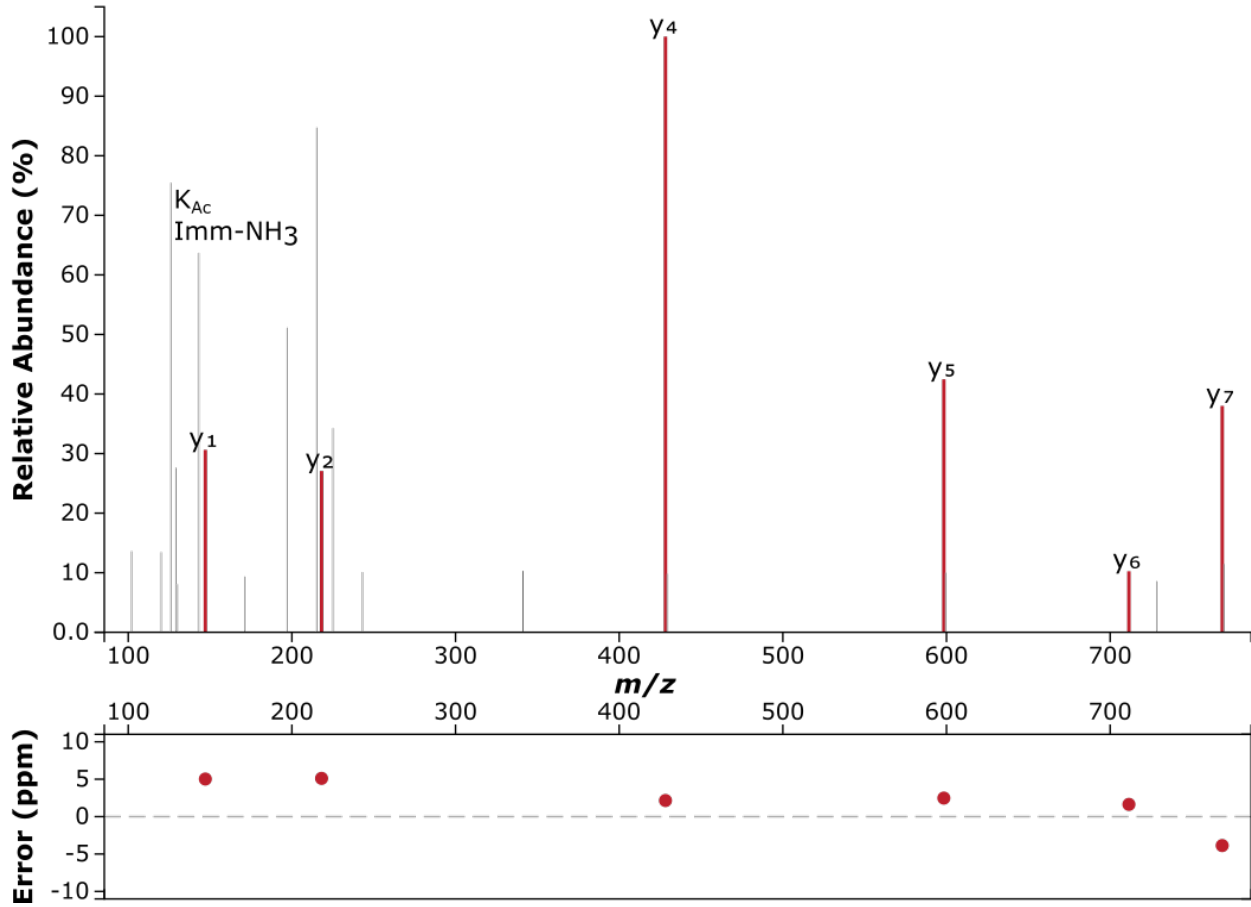
4-hydroxybenzoate-CoA ligase / benzoate-CoA ligase (Q2LRH7)



E L G L ^{PC} k P L A K

Precursor m/z: 505.8159 Charge: +2

Acetyl-CoA acetyltransferase (Q2LWQ5)

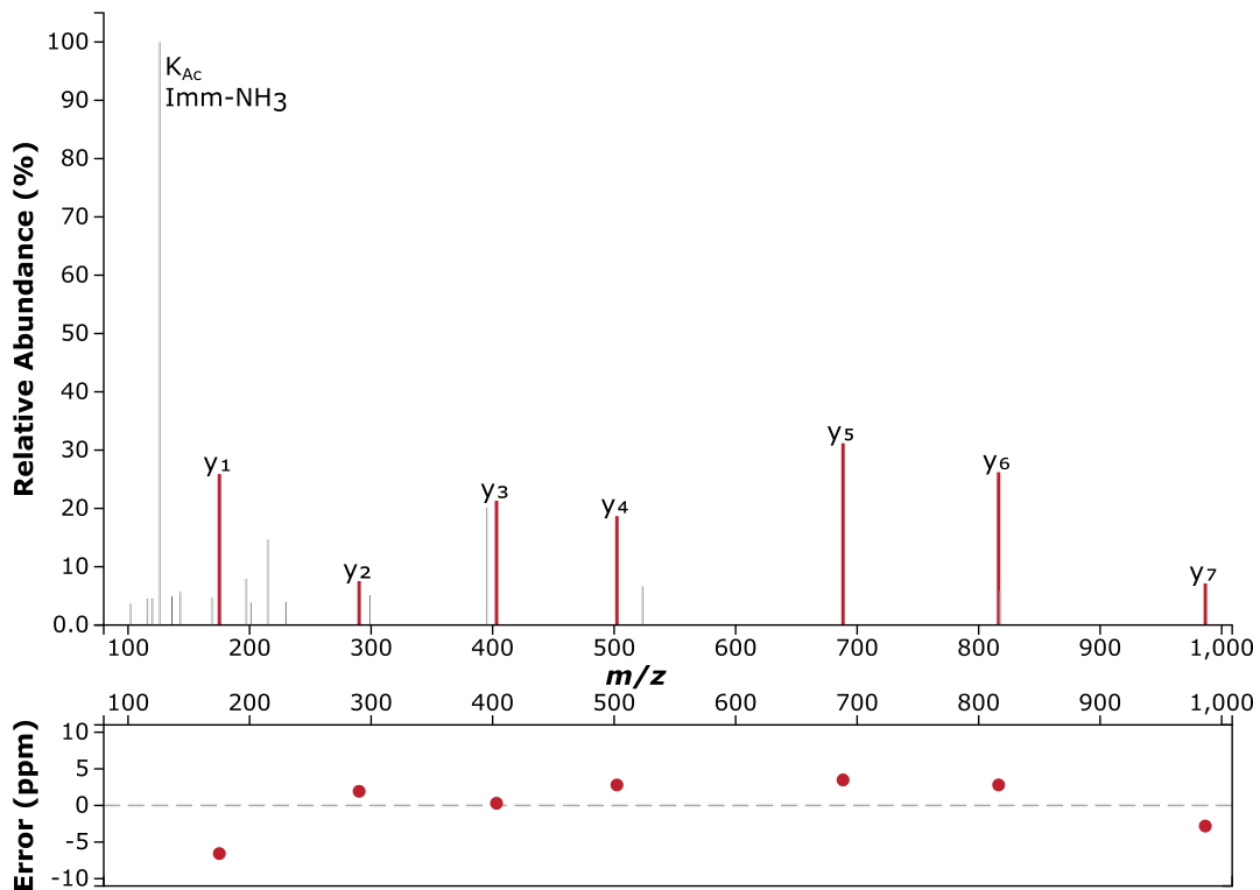


E L ^{Ac} k Q W V L D R

Precursor m/z: 614.8379

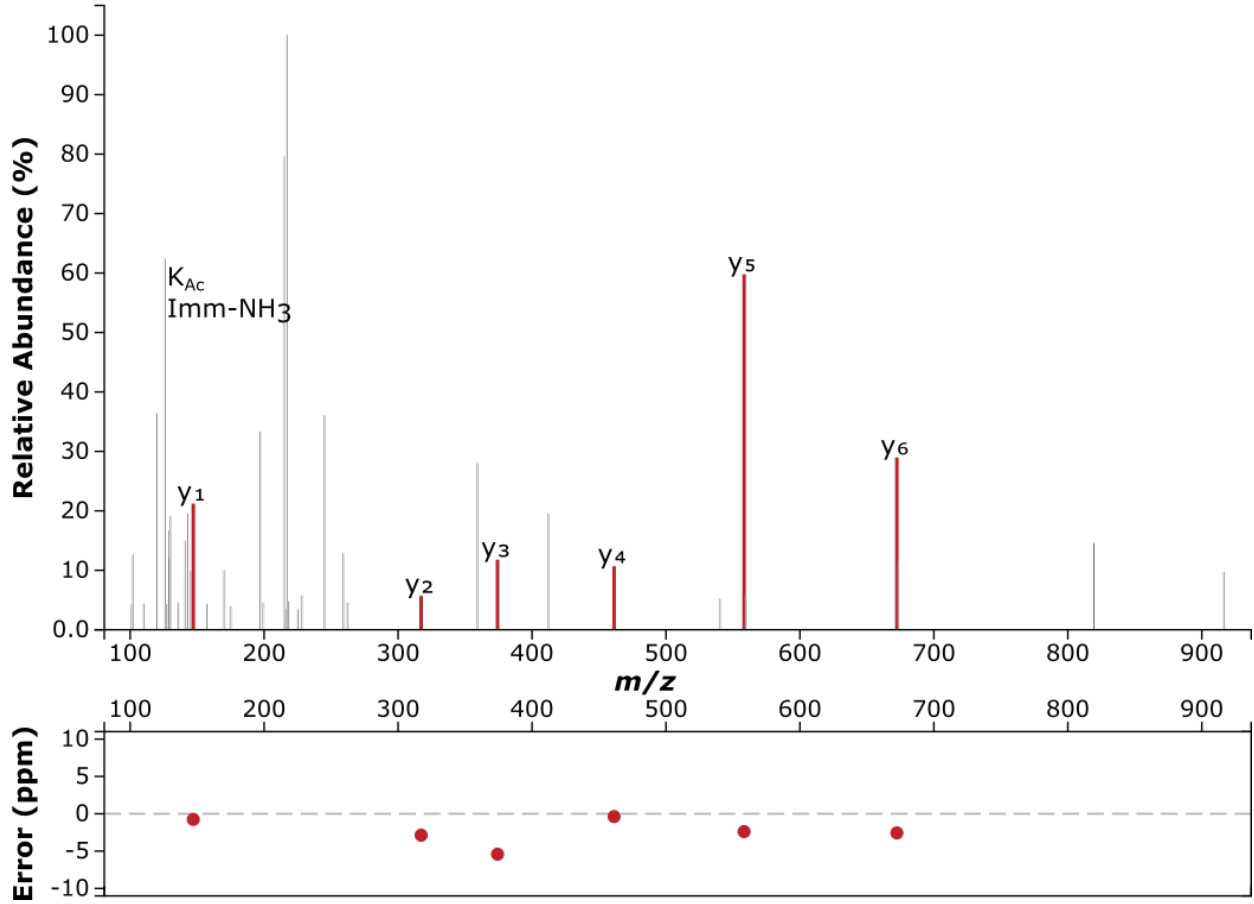
Charge: +2

4-hydroxybenzoate-CoA ligase / benzoate-CoA ligase (Q2LRH7)



E L P F N P S G k K^{AC}

Precursor m/z: 579.8113 Charge: +2
Long-chain-fatty-acid-CoA ligase (Q2LR69)

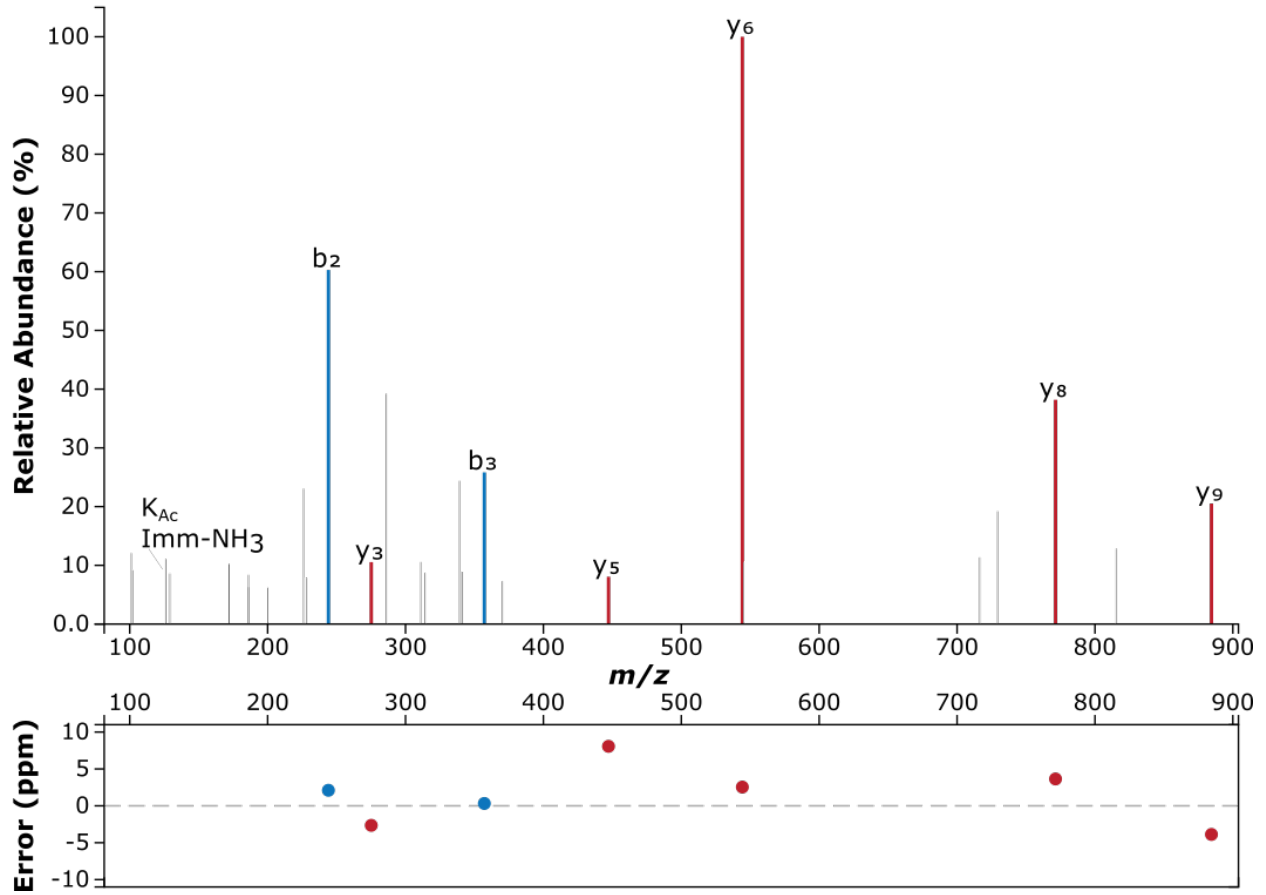




Precursor m/z: 620.8302

Charge: +2

Acyl-CoA dehydrogenase, short-chain specific (Q2LUY8)

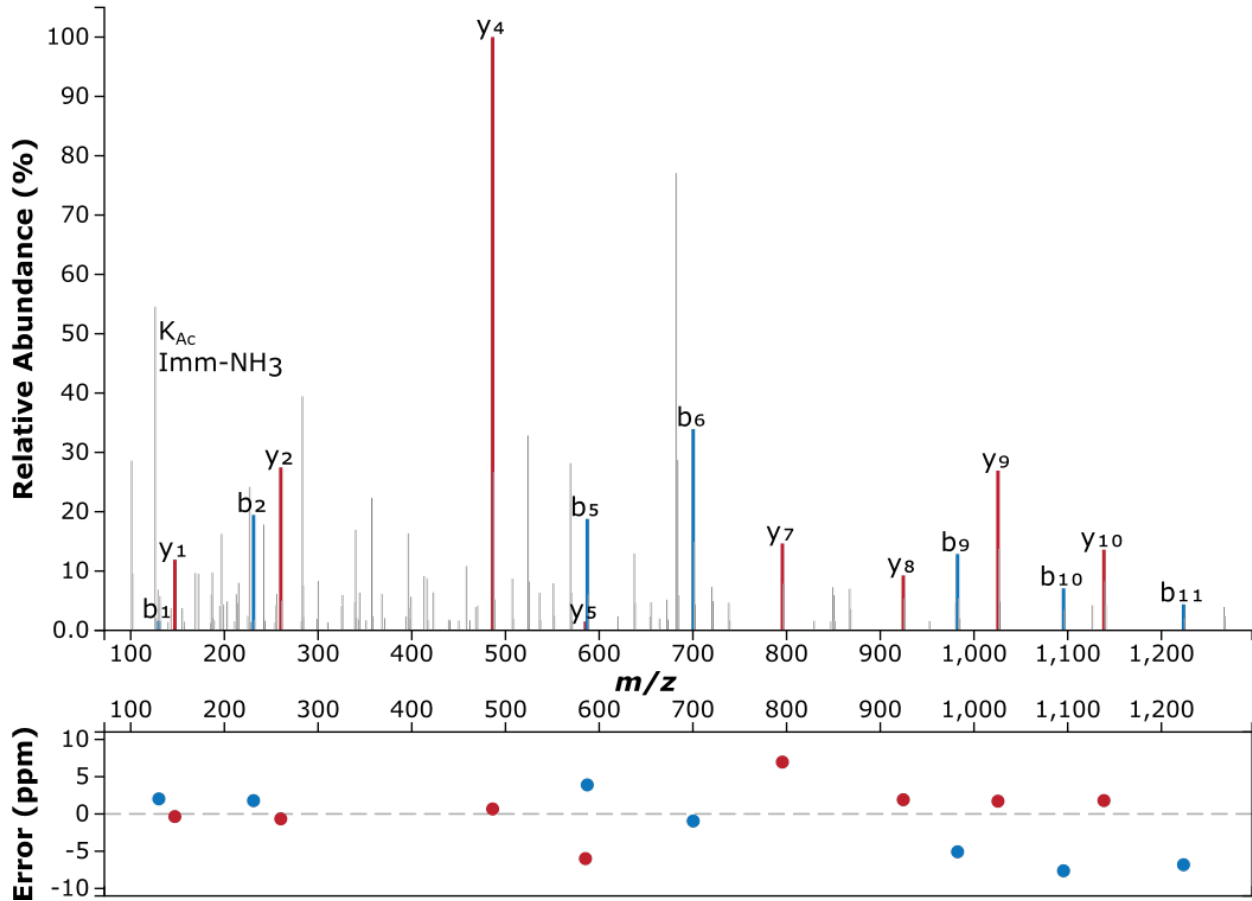


E T G E k I P G Q L Q L T E I P V P E L K

Precursor m/z: 787.7686

Charge: +3

6-hydroxycyclohex-1-ene-1-carboxyl-CoA dehydrogenase (Q2LXU3)

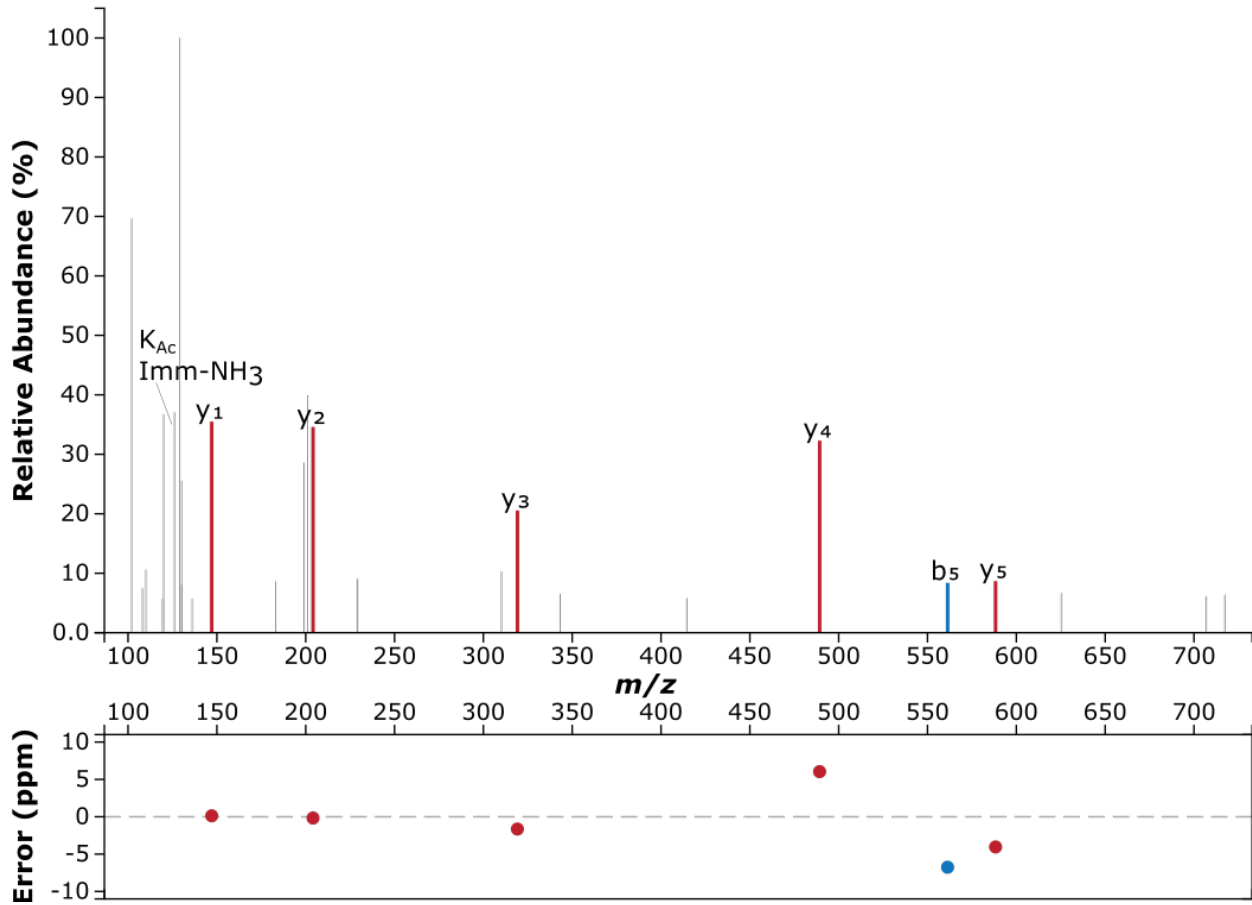


F K G E V E V ^{AC} k ^{AC} D G K

Precursor m/z: 426.5627

Charge: +3

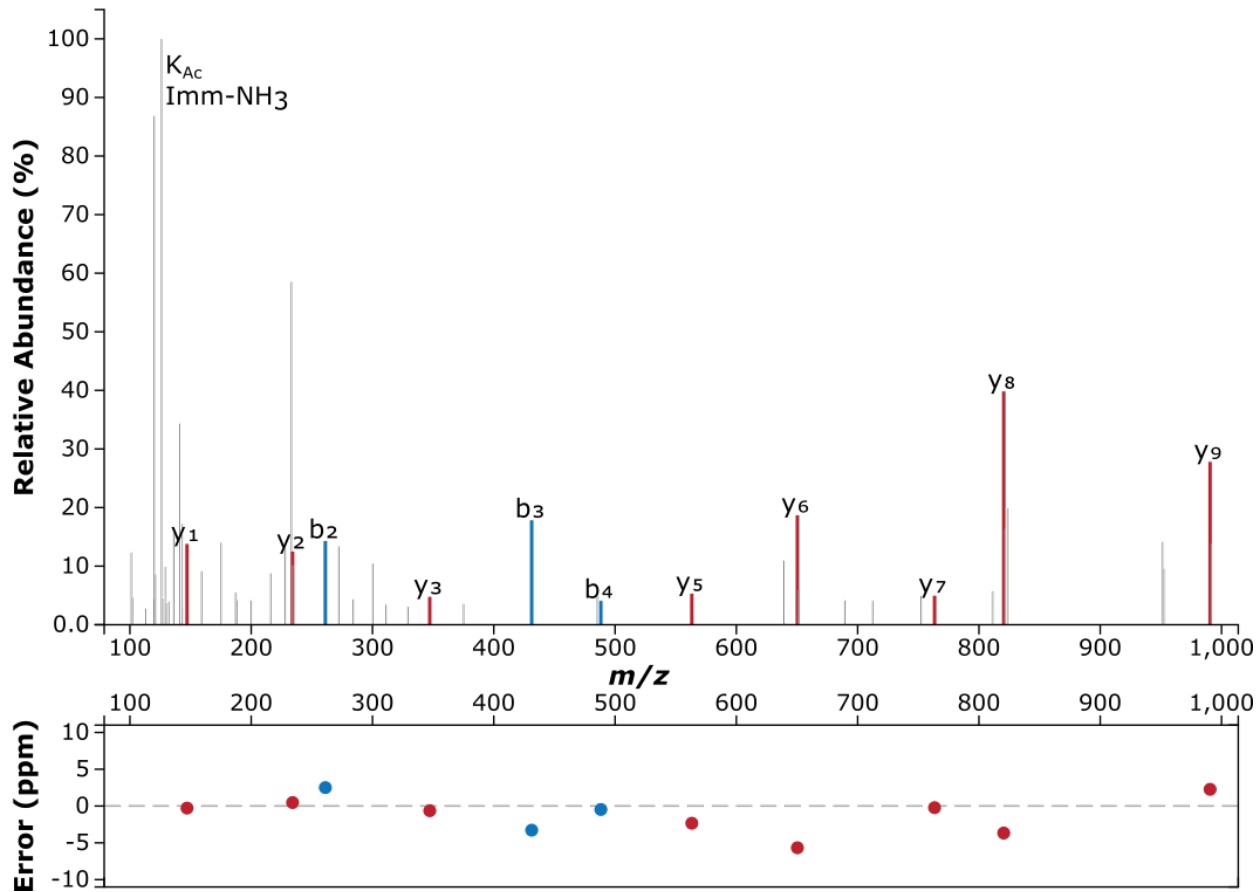
Glyceraldehyde-3-phosphate dehydrogenase (Q2LWW2)





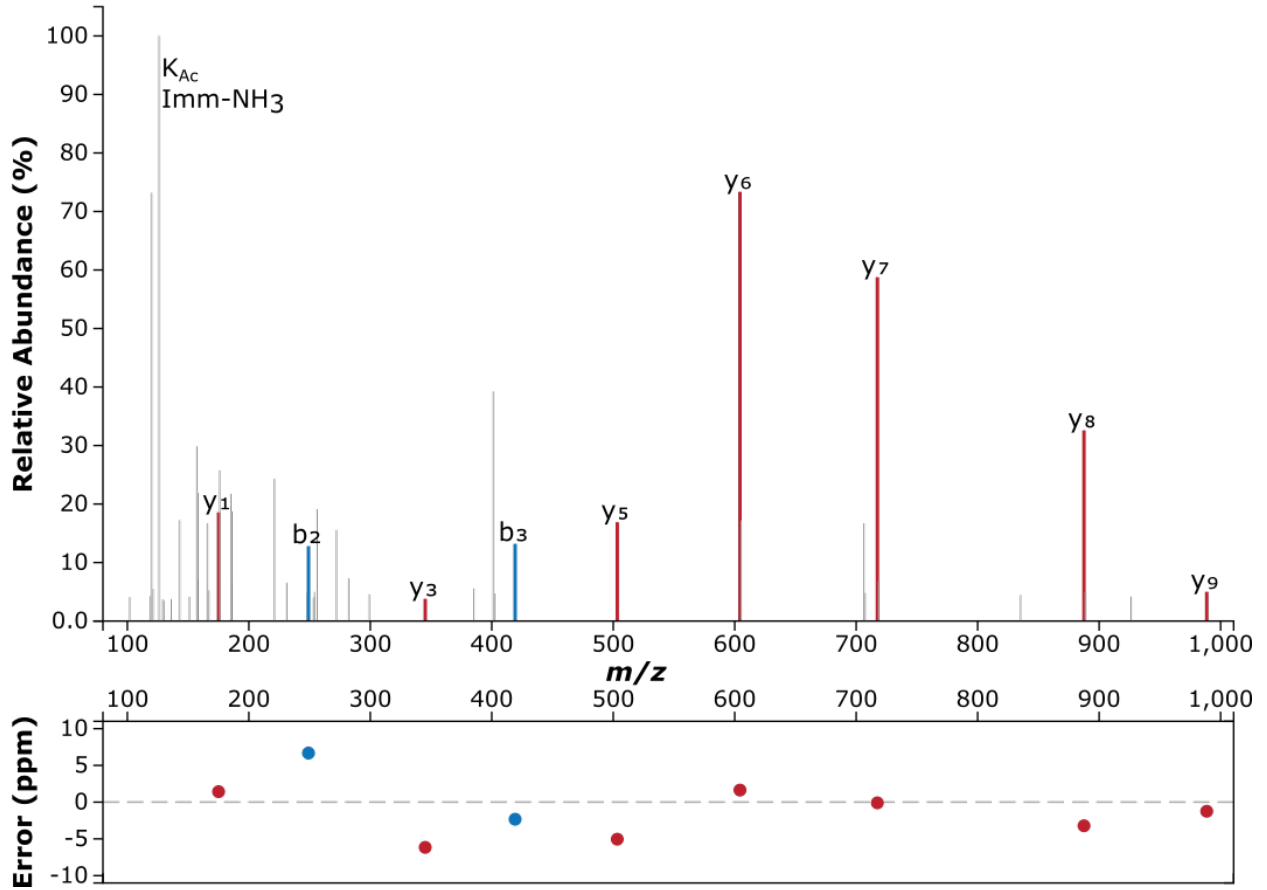
Precursor m/z: 625.8532 Charge: +2

Phosphoserine aminotransferase (Q2LSJ6)





Precursor m/z: 568.3271 Charge: +2
 Inosine-5'-monophosphate dehydrogenase (Q2LTT4)

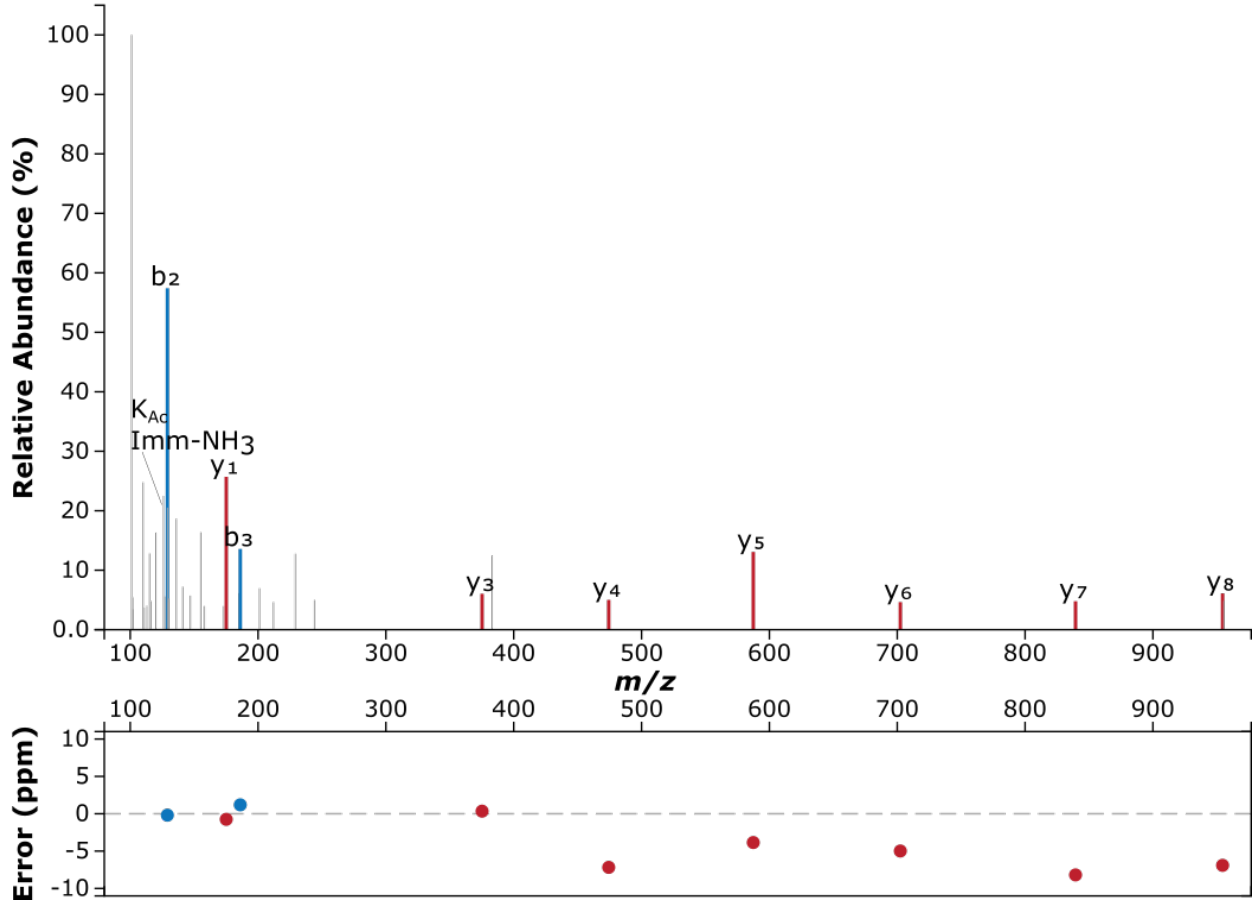


G A G A D P k^{PC} D H D L V S L R

Precursor m/z: 531.6057

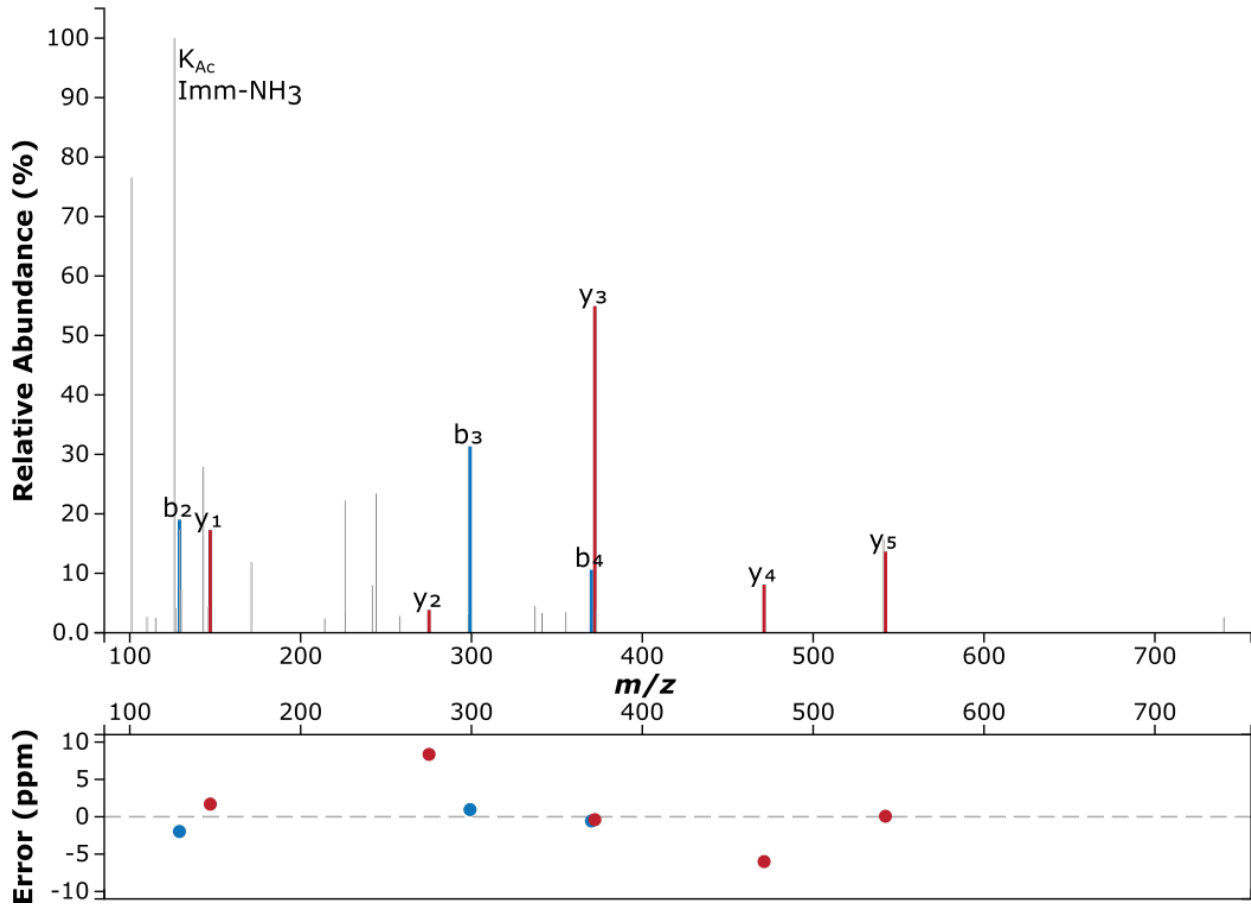
Charge: +3

4-hydroxybenzoate-CoA ligase / benzoate-CoA ligase (Q2LRH0)





Precursor m/z: 420.7505 Charge: +2
 Acetyl-CoA synthetase (Q2LWR8)

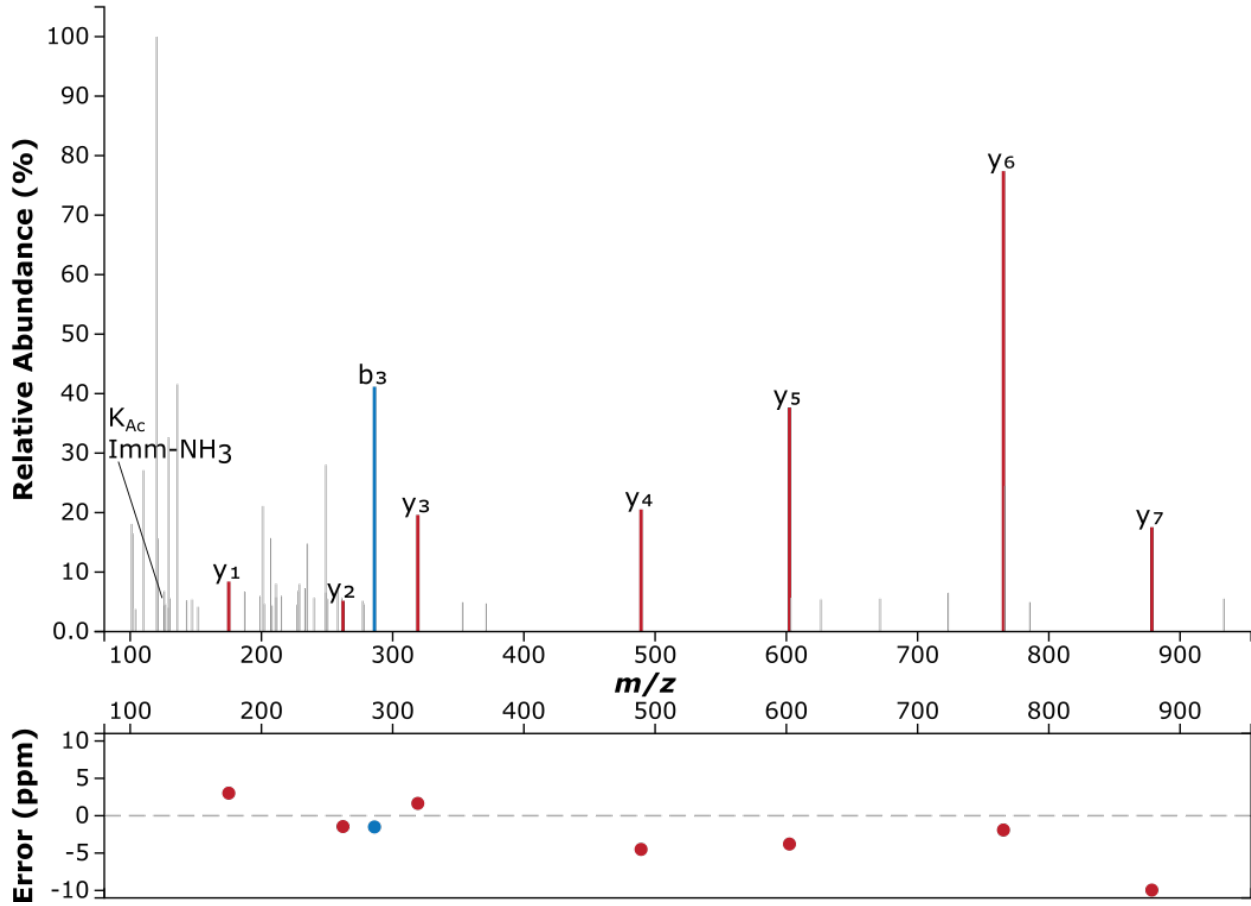




Precursor m/z: 582.3246

Charge: +2

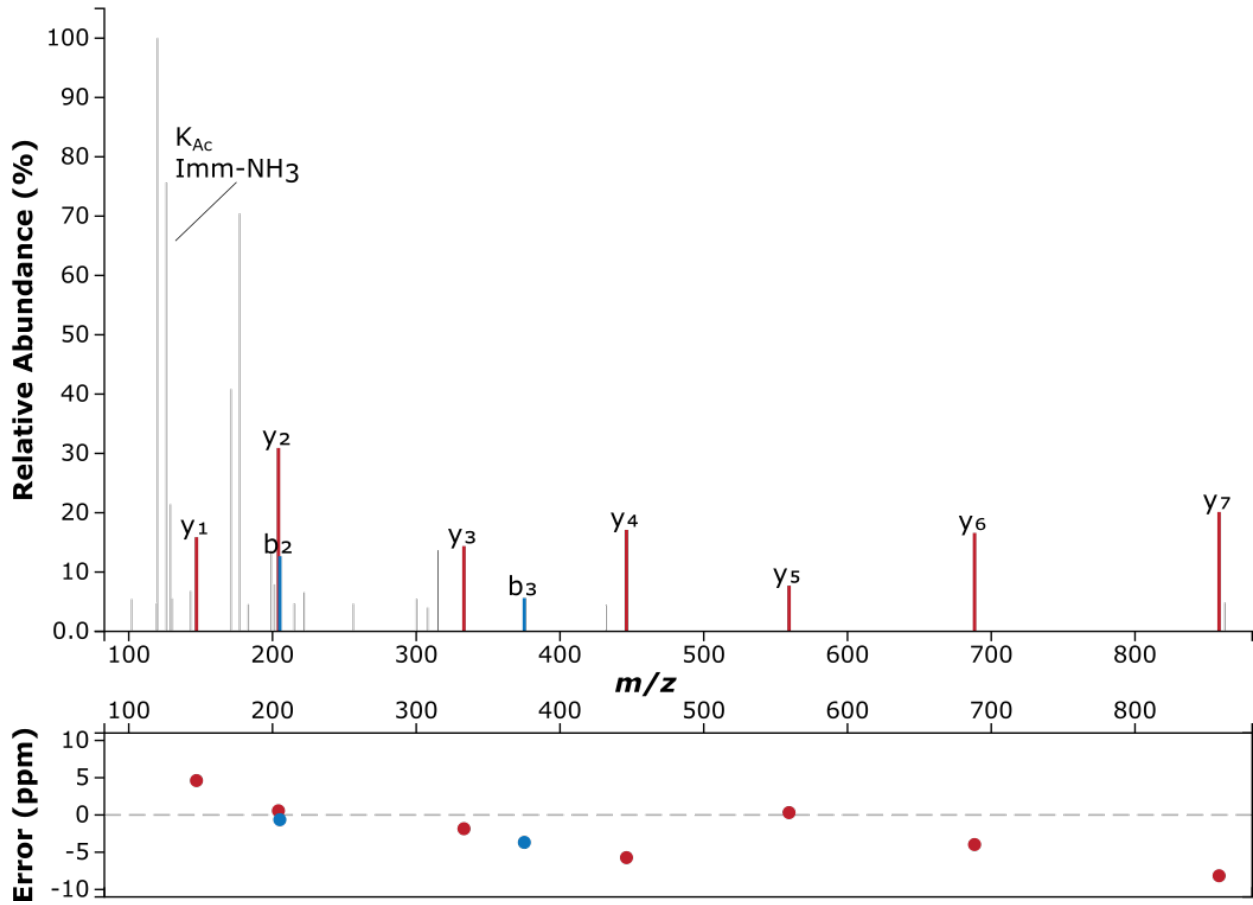
Molecular chaperone (Q2LYH4)





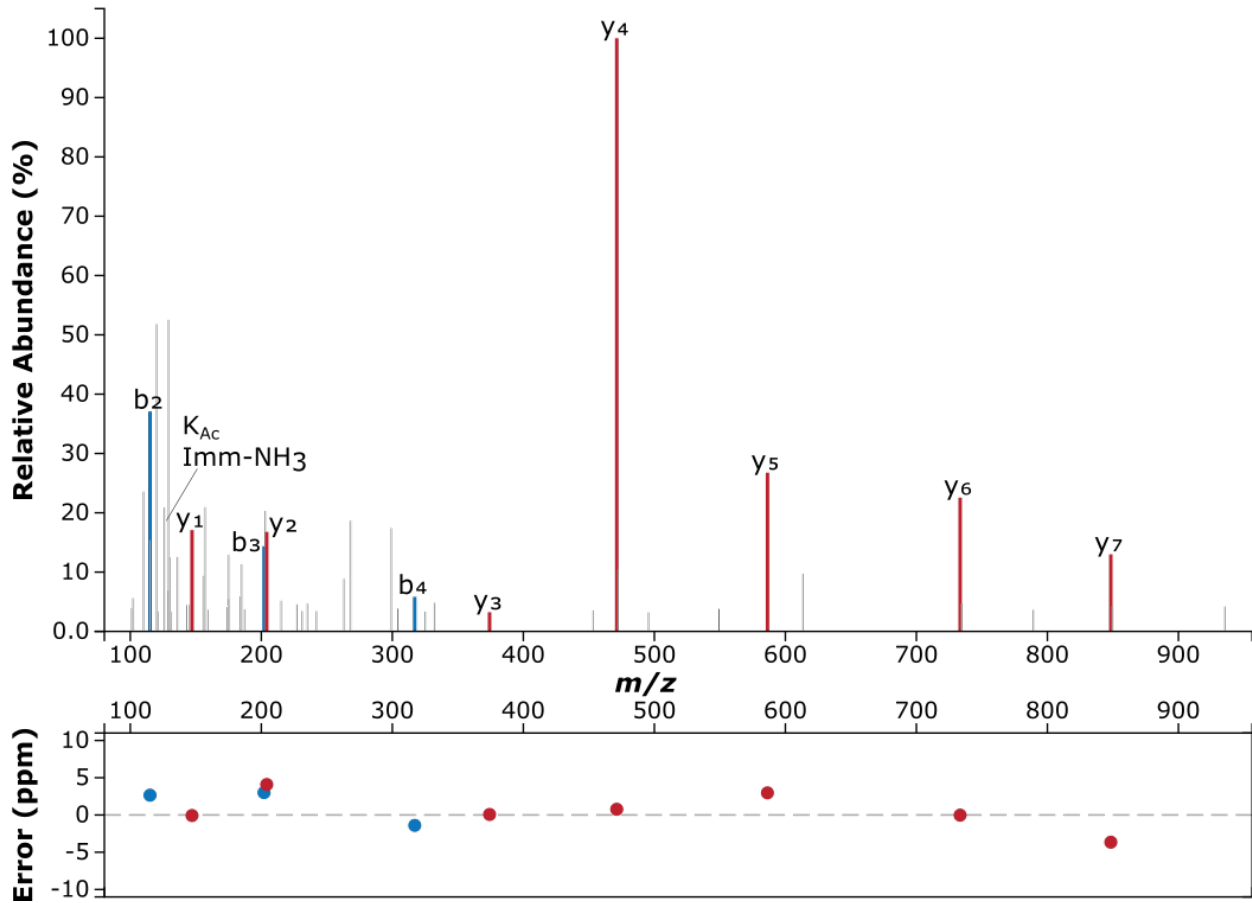
Precursor m/z: 531.7951 Charge: +2

ATP synthase subunit beta (Q2LR05)





Precursor m/z: 525.2485 Charge: +2
 Glutamate dehydrogenase (Q2LQG2)

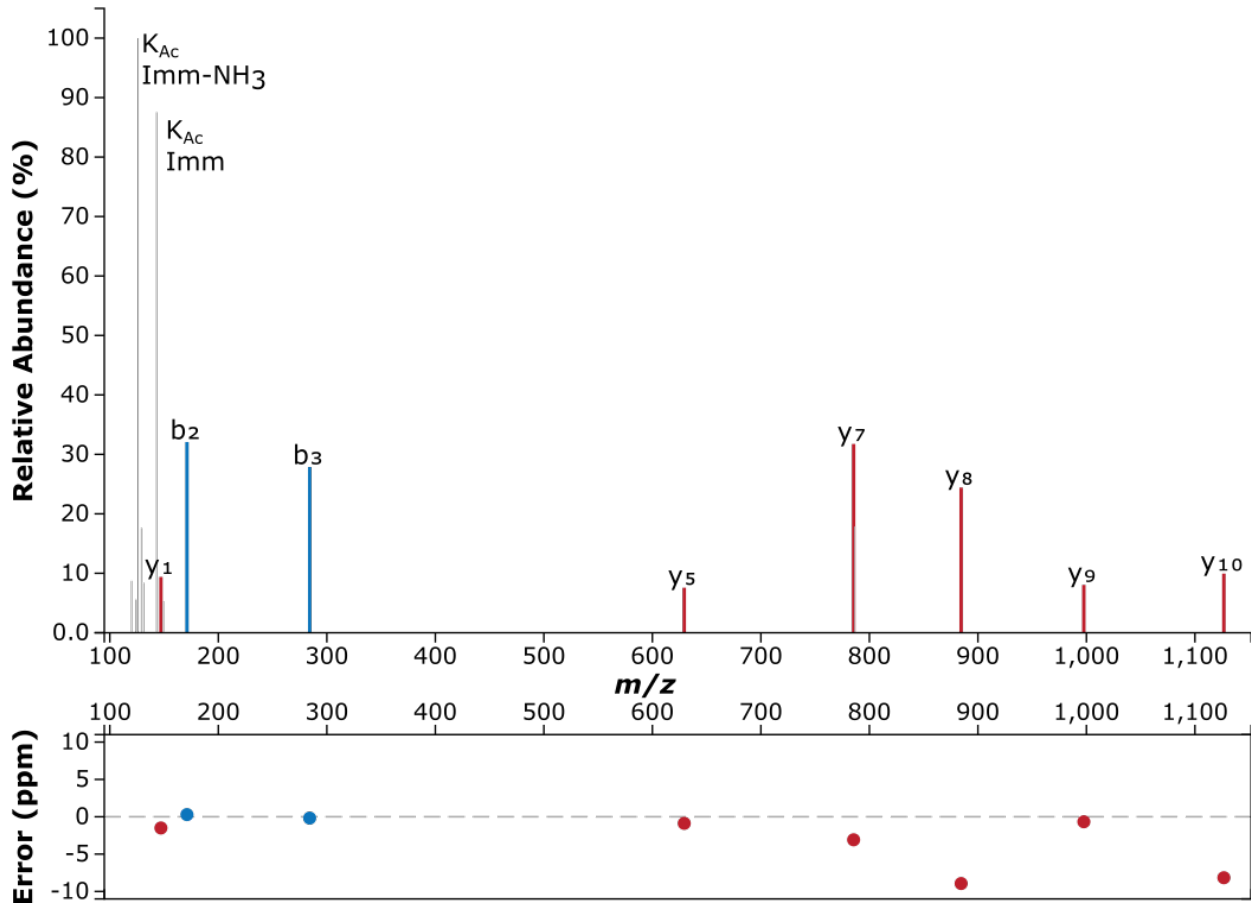


G I I ^{AC} k T G D E V E I V G V R P T Q K

Precursor m/z: 694.3915

Charge: +3

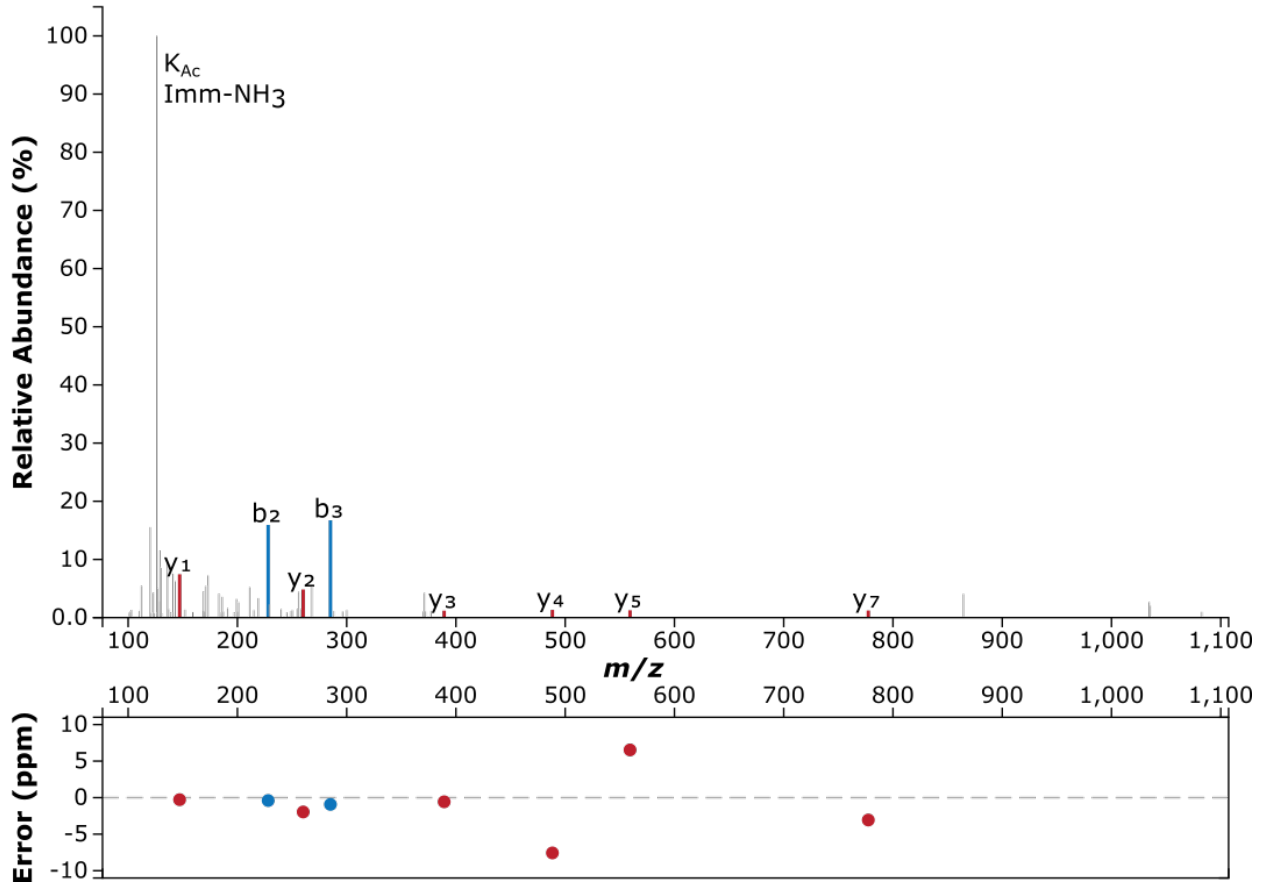
Elongation factor Tu (Q2LQA3)





Precursor m/z: 631.3612 Charge: +2

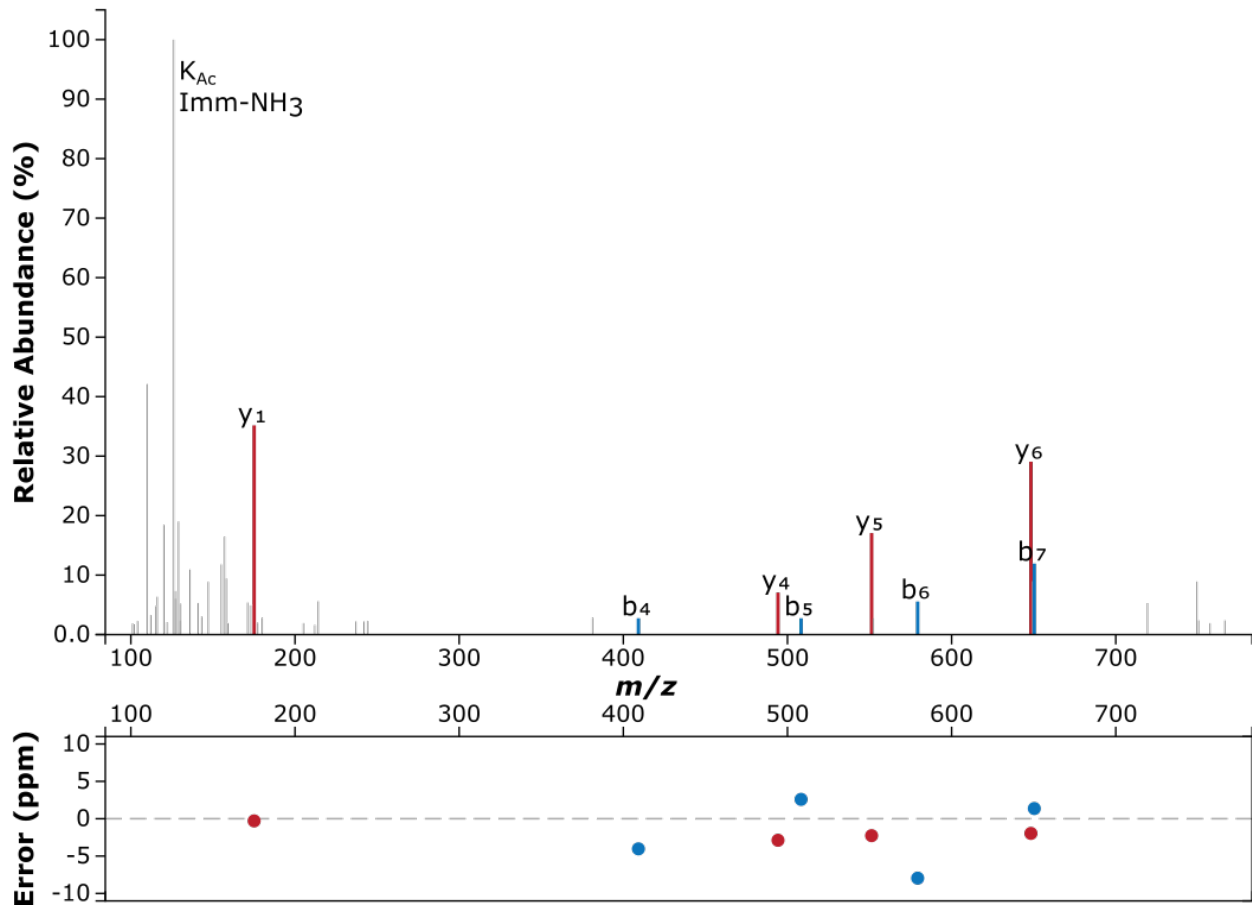
Acyl-CoA dehydrogenase, short-chain specific (Q2LUY8)





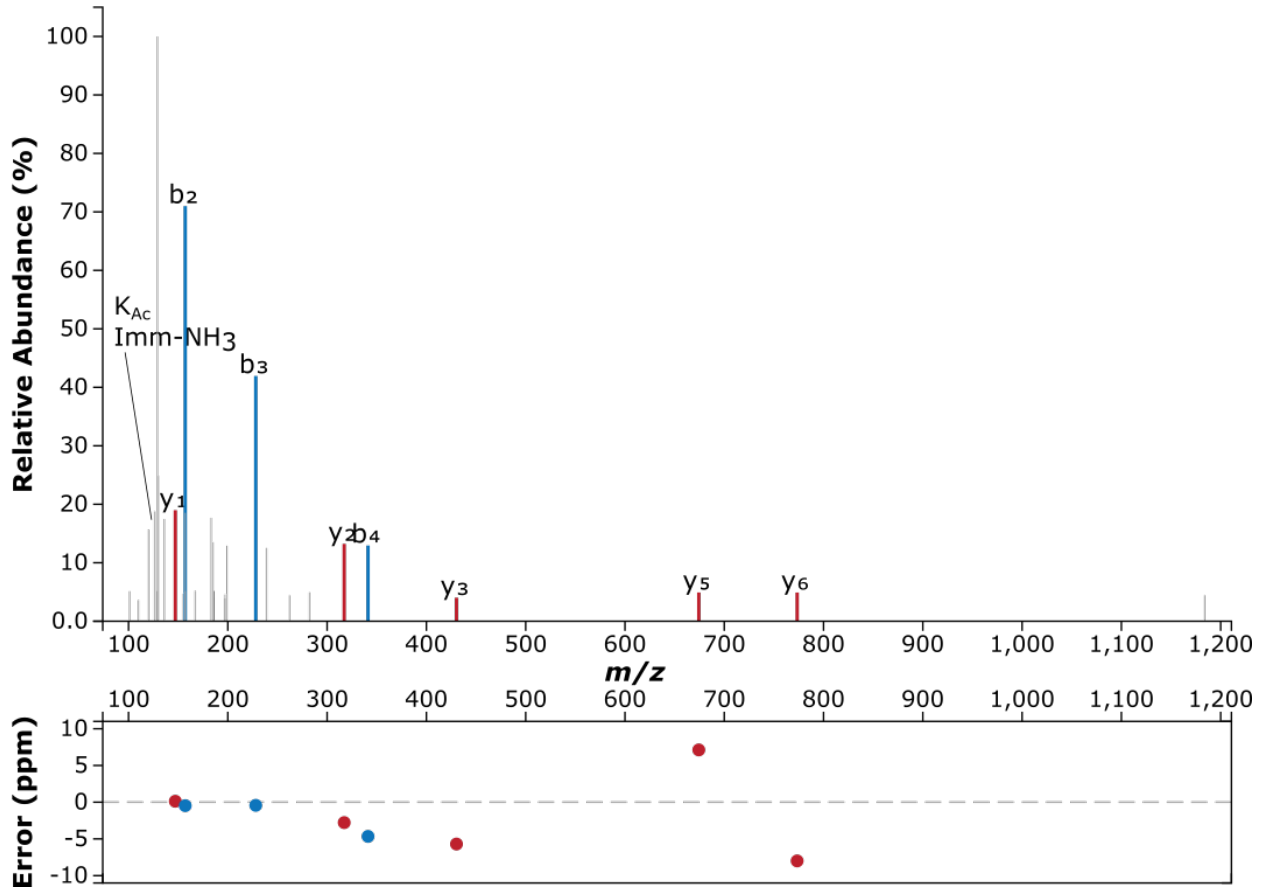
Precursor m/z: 546.6301 Charge: +3

60 kDa chaperonin 2 (Q2LTG7)





Precursor m/z: 606.8635 Charge: +2
 60 kDa chaperonin 3 (Q2LU42)

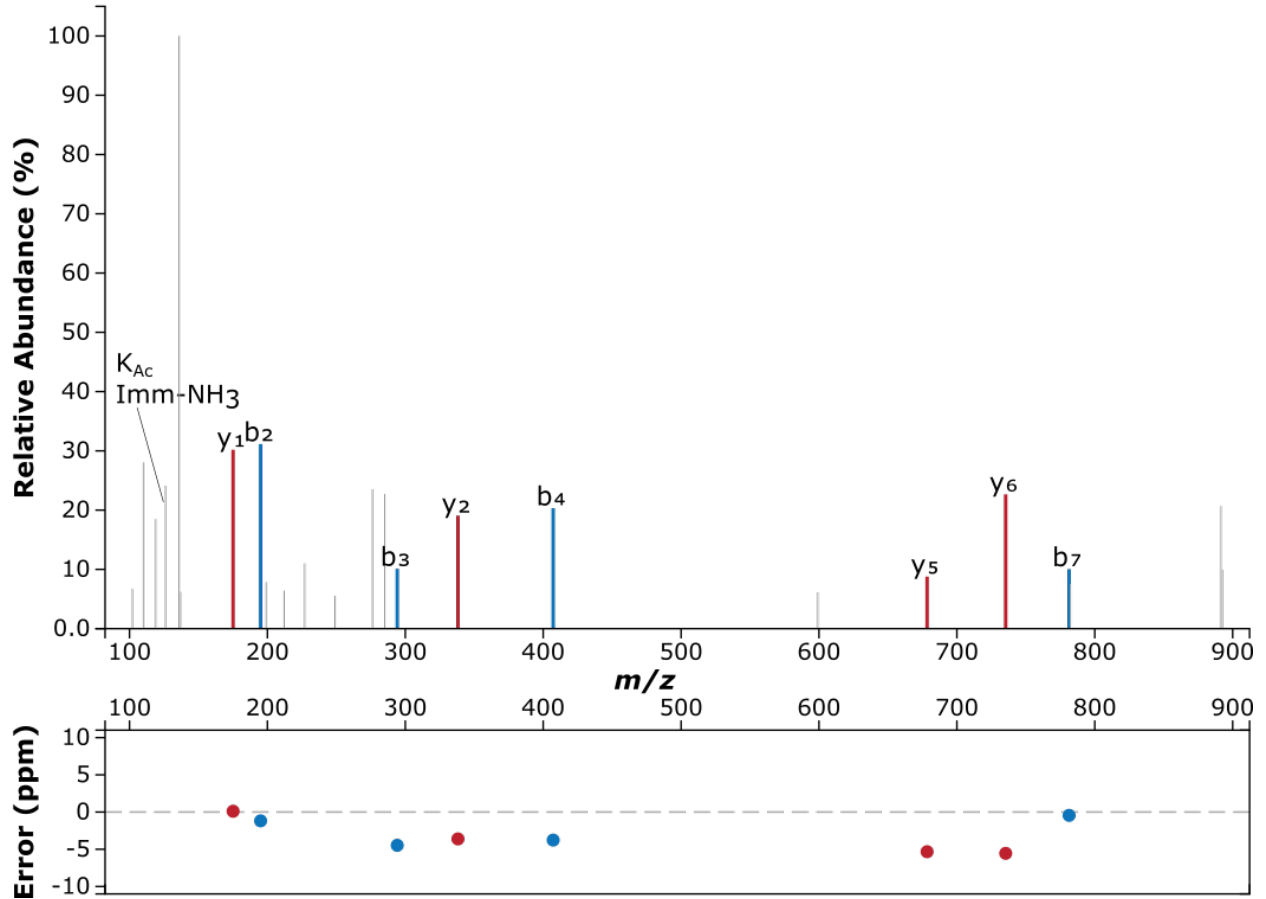


H G V L P N Y P E Y T N E Y I G V G G K L Y R

Precursor m/z: 894.4501

Charge: +3

Glutaconyl-CoA decarboxylase A subunit (Q2LUY7)

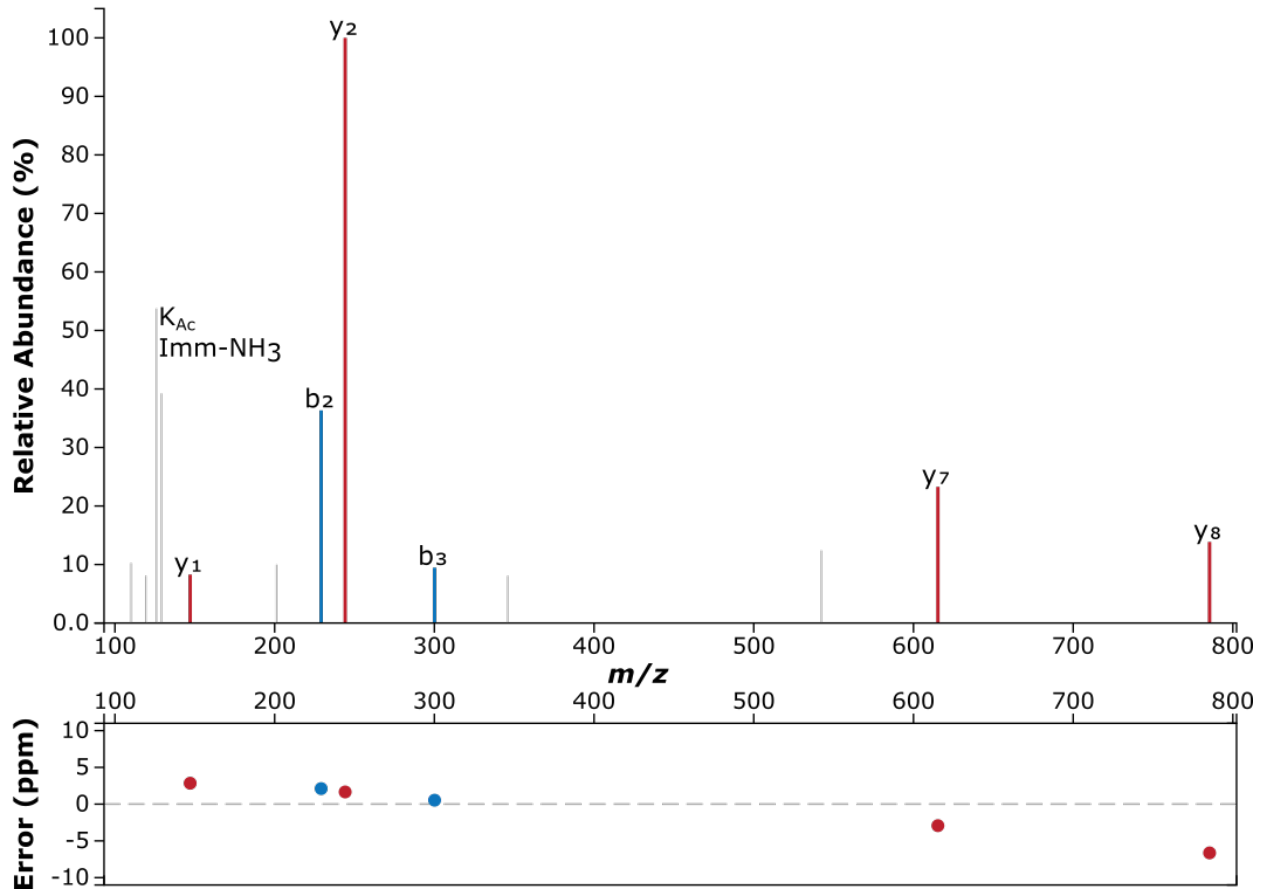


I D A E ^{Ac} k G A G A D P K

Precursor m/z: 607.3066

Charge: +2

4-hydroxybenzoate-CoA ligase / benzoate-CoA ligase (Q2LRH0)

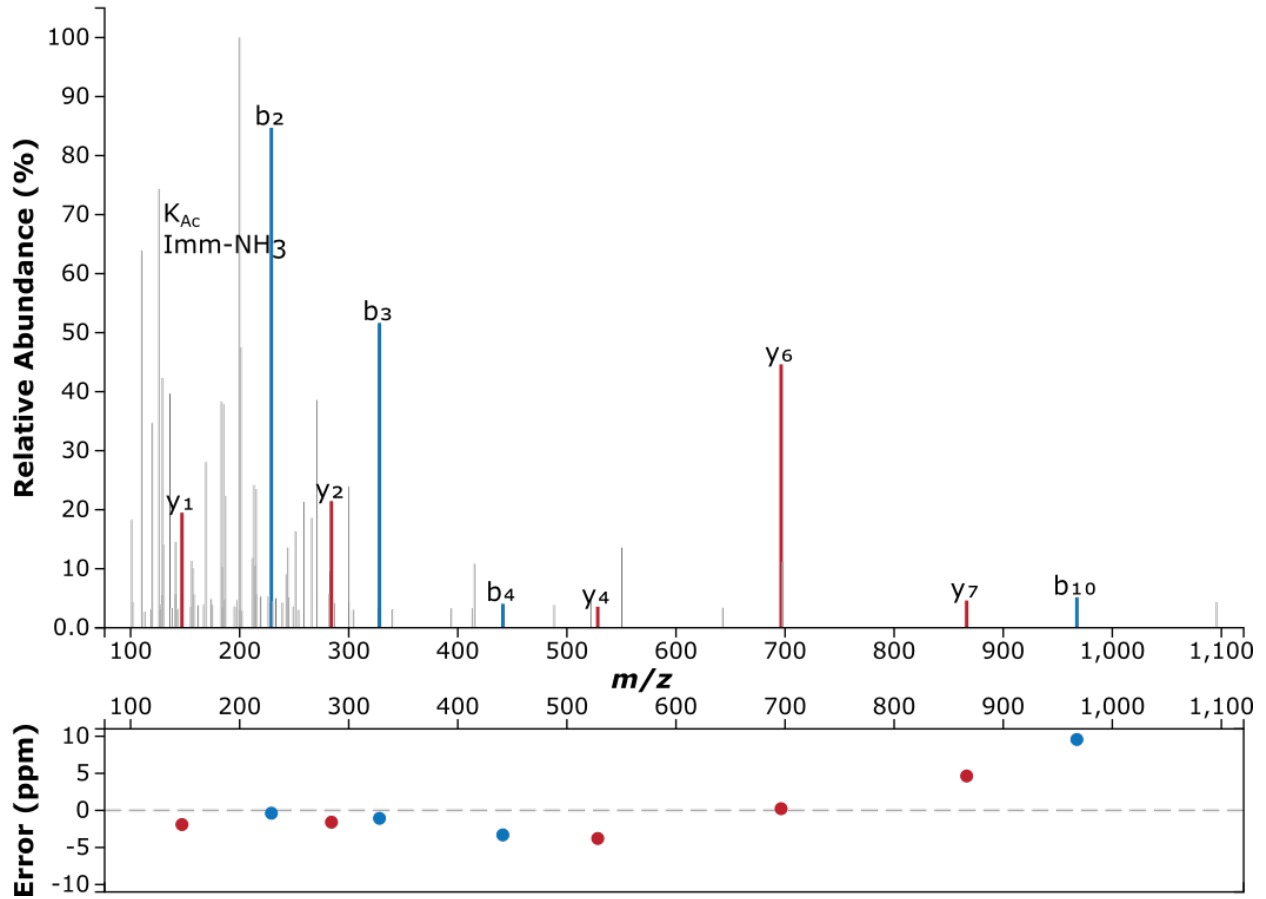


I D V L V N N A G A T ^{Ac} k P A M L H K

Precursor m/z: 645.3558

Charge: +3

3-oxoacyl-reductase (Q2LXS6)

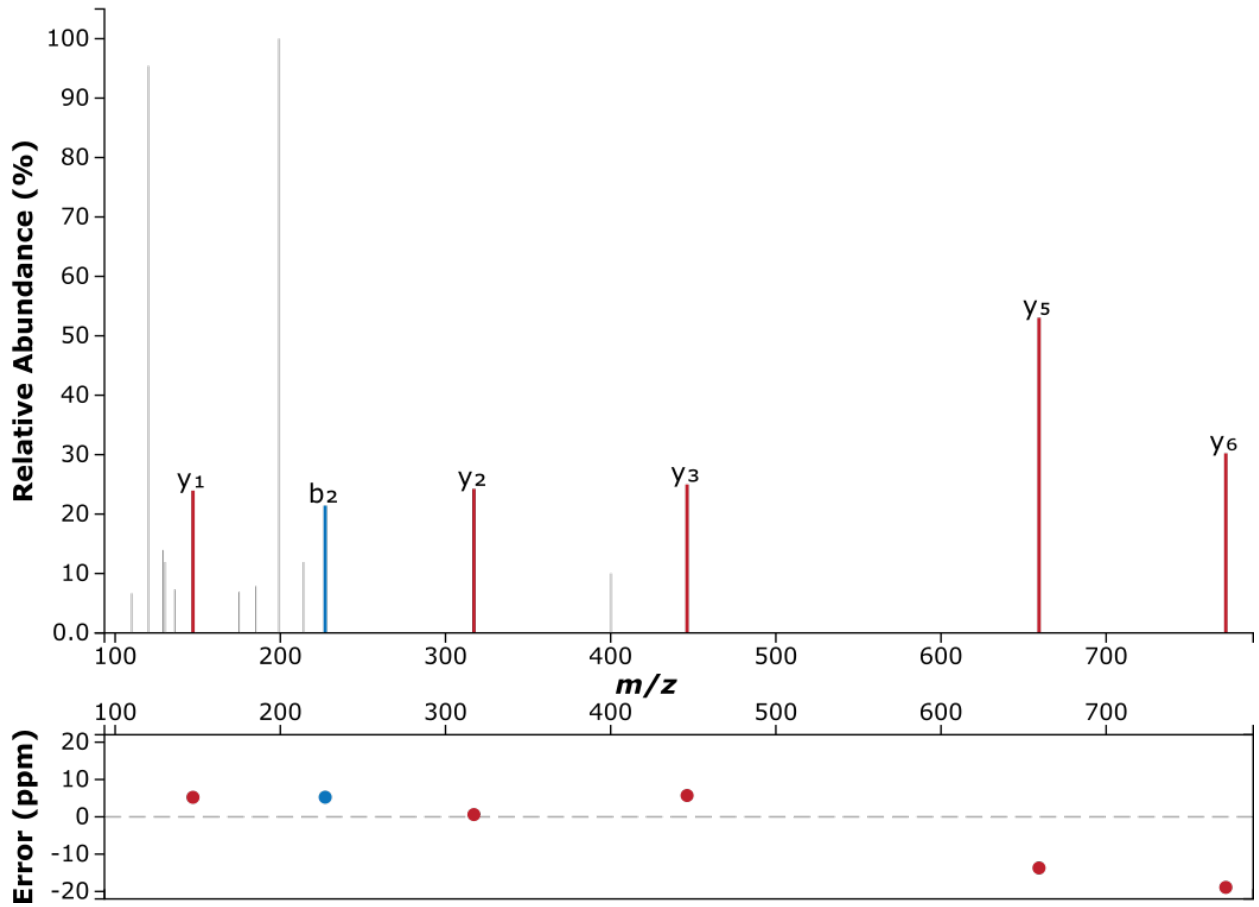




Precursor m/z: 443.2794

Charge: +2

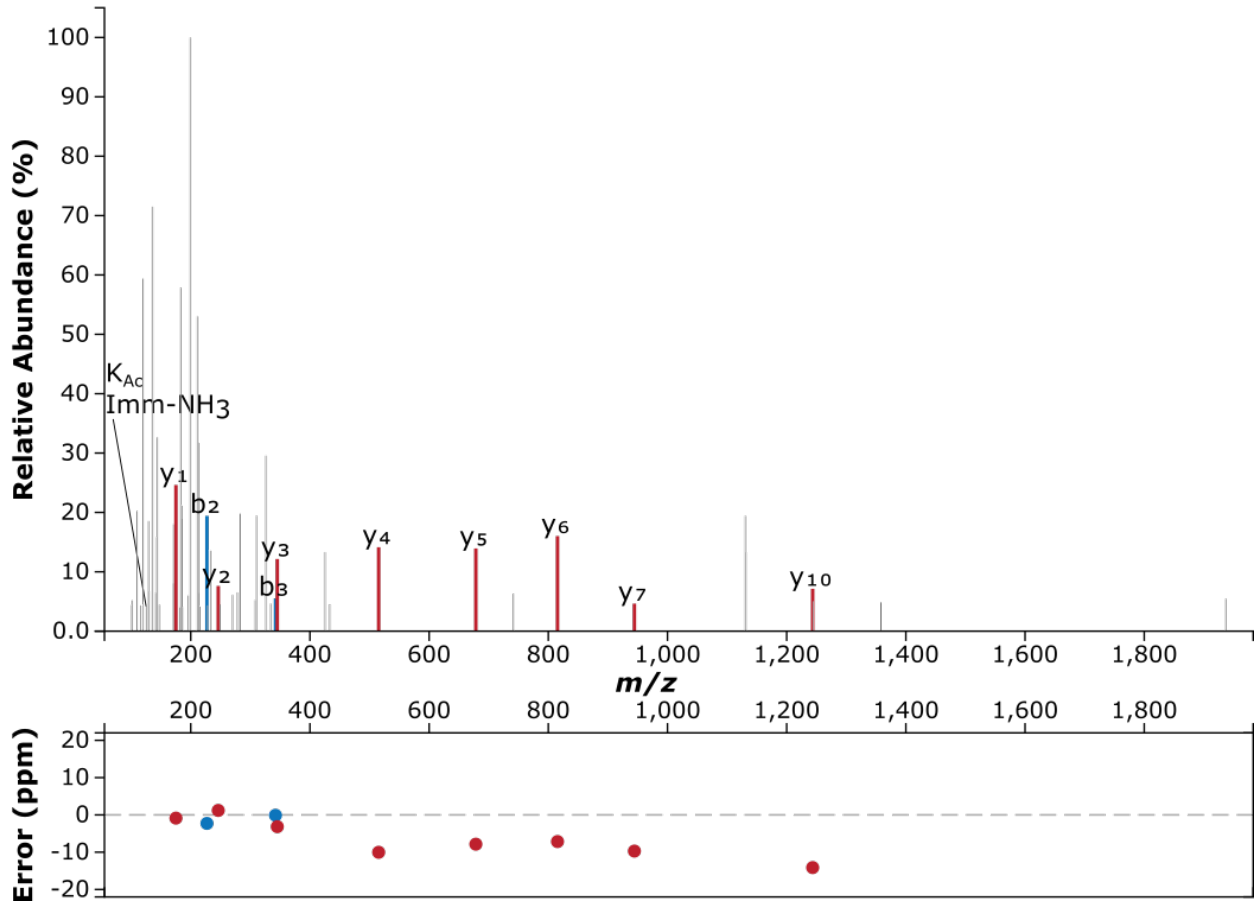
Alpha-1,4 glucan phosphorylase (Q2LXZ0)



I L D P I V L G E E H Y ^{Ac} k V A R

Precursor m/z: 632.0193 Charge: +3

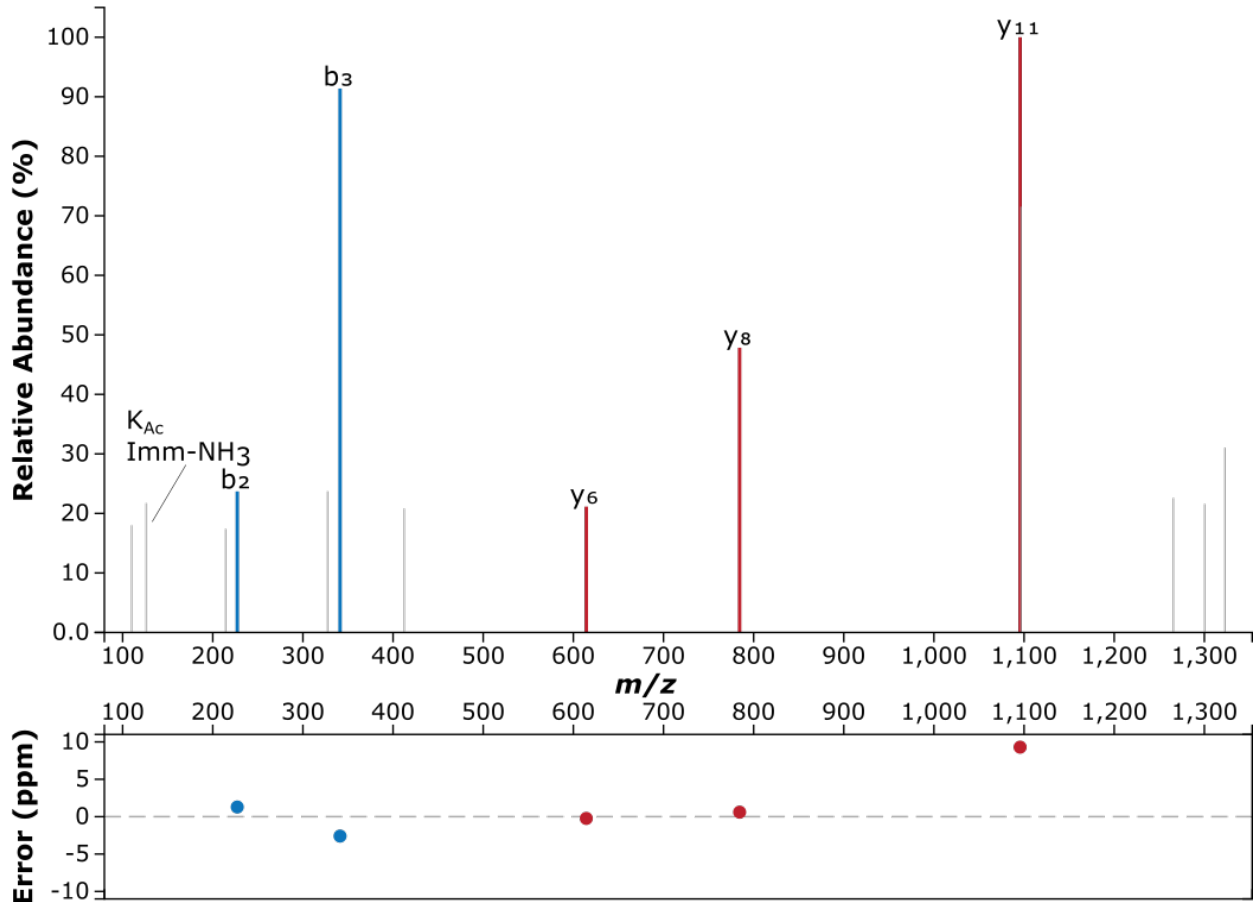
ATP synthase subunit beta (Q2LR05)



I L N V V G k P V D G L G P I E A K

Precursor m/z: 931.0433 Charge: +2

ATP synthase subunit beta (Q2LR05)

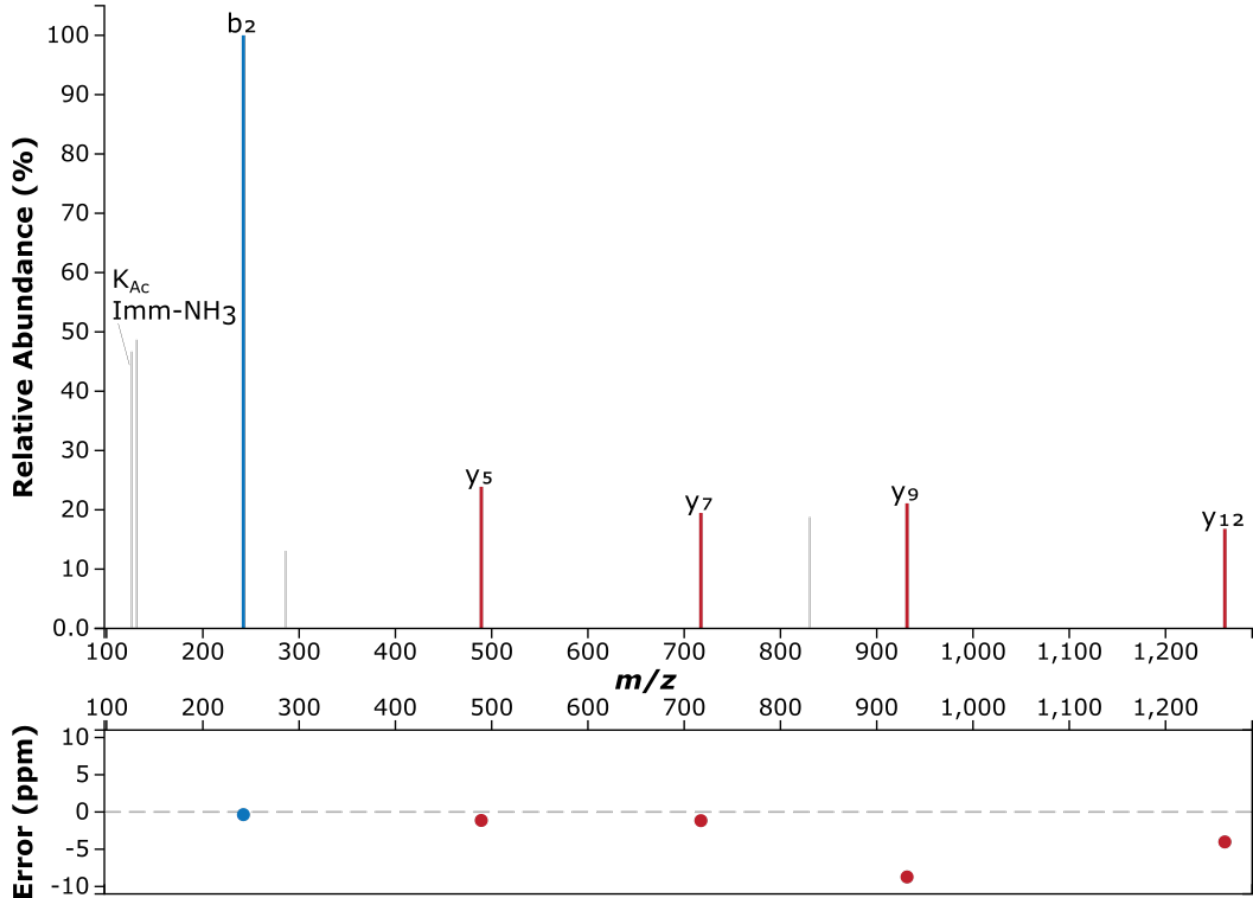


I Q I D k^{Ac} D N T T I I D G A G E R

Precursor m/z: 950.9842

Charge: +2

60 kDa chaperonin 2 (Q2LTG7)

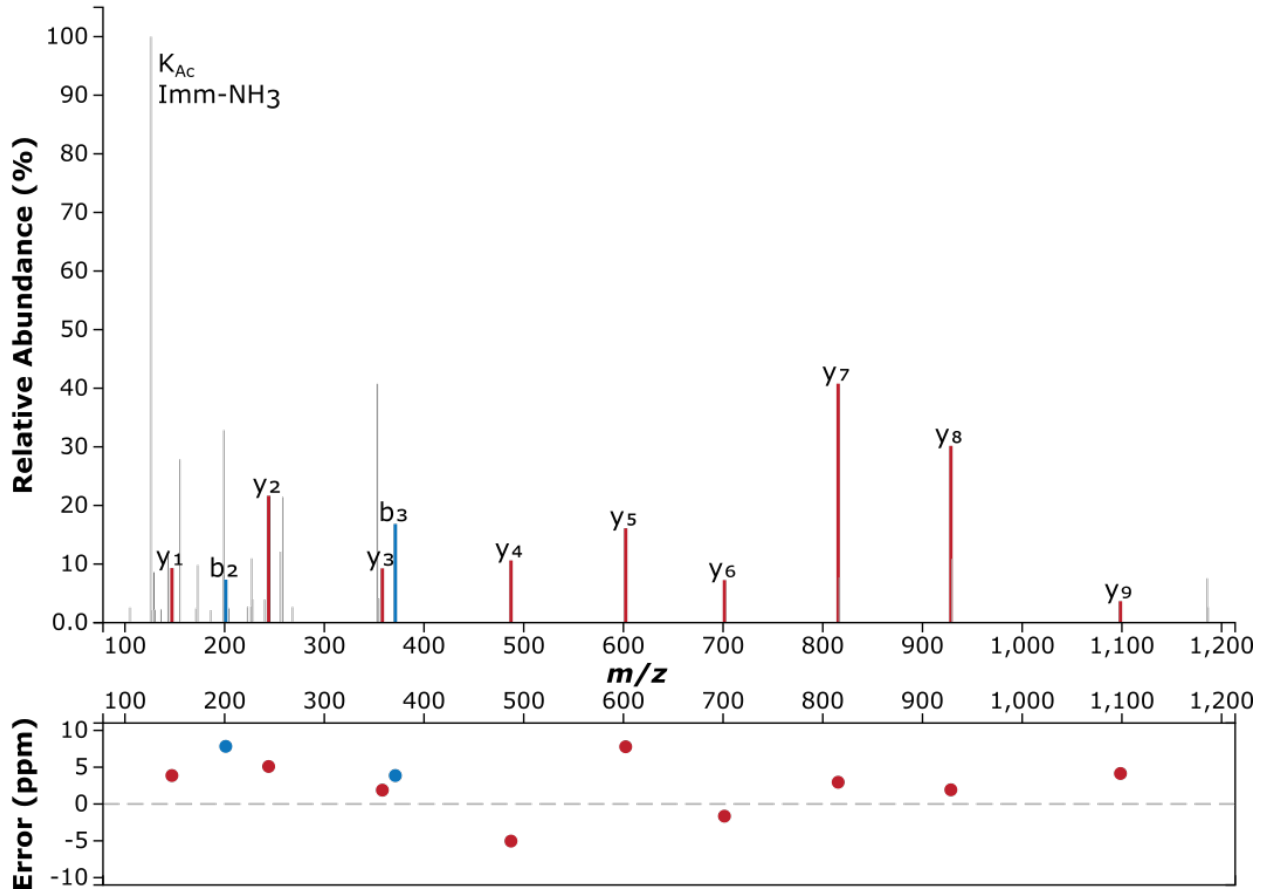




Precursor m/z: 649.8512

Charge: +2

Thioredoxin (Q2LVG3)

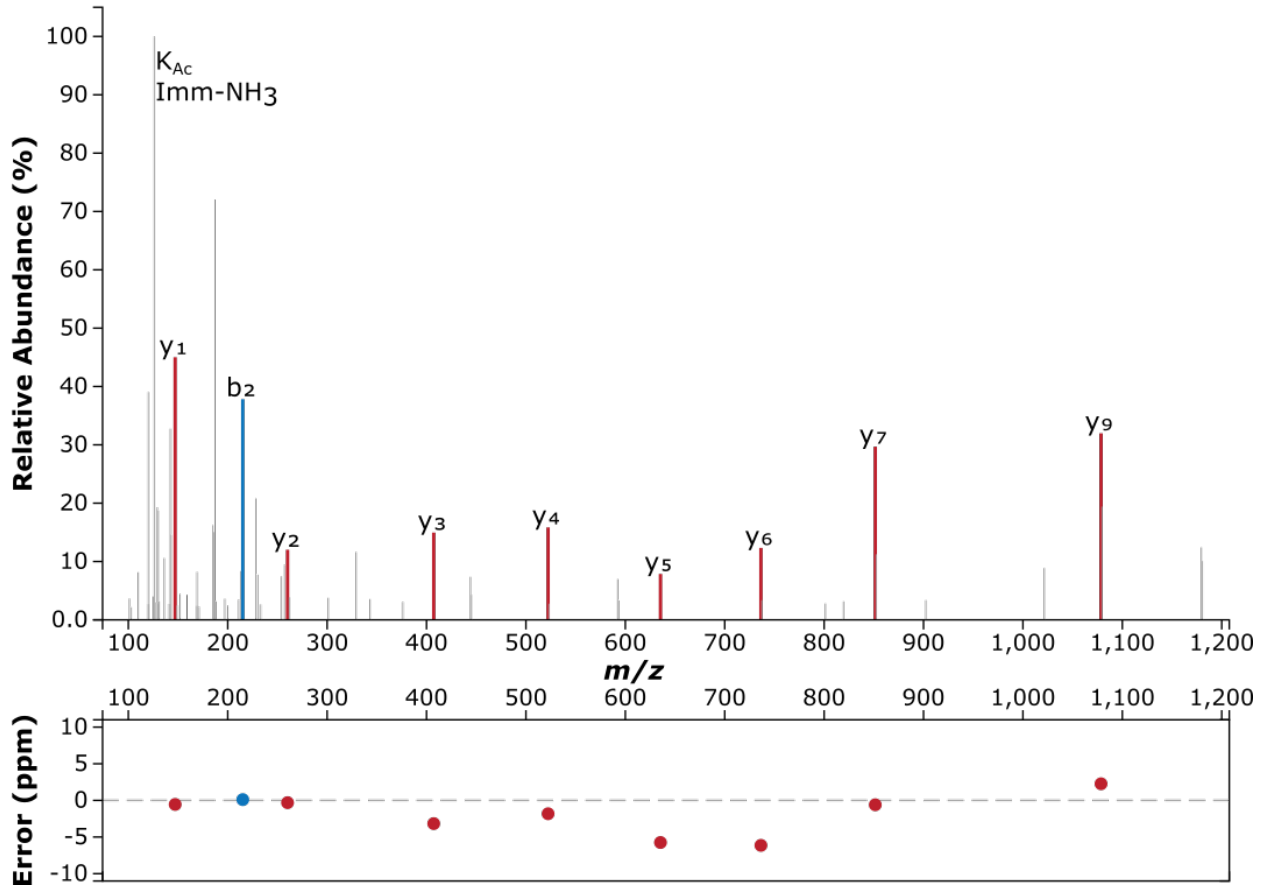




Precursor m/z: 646.8584

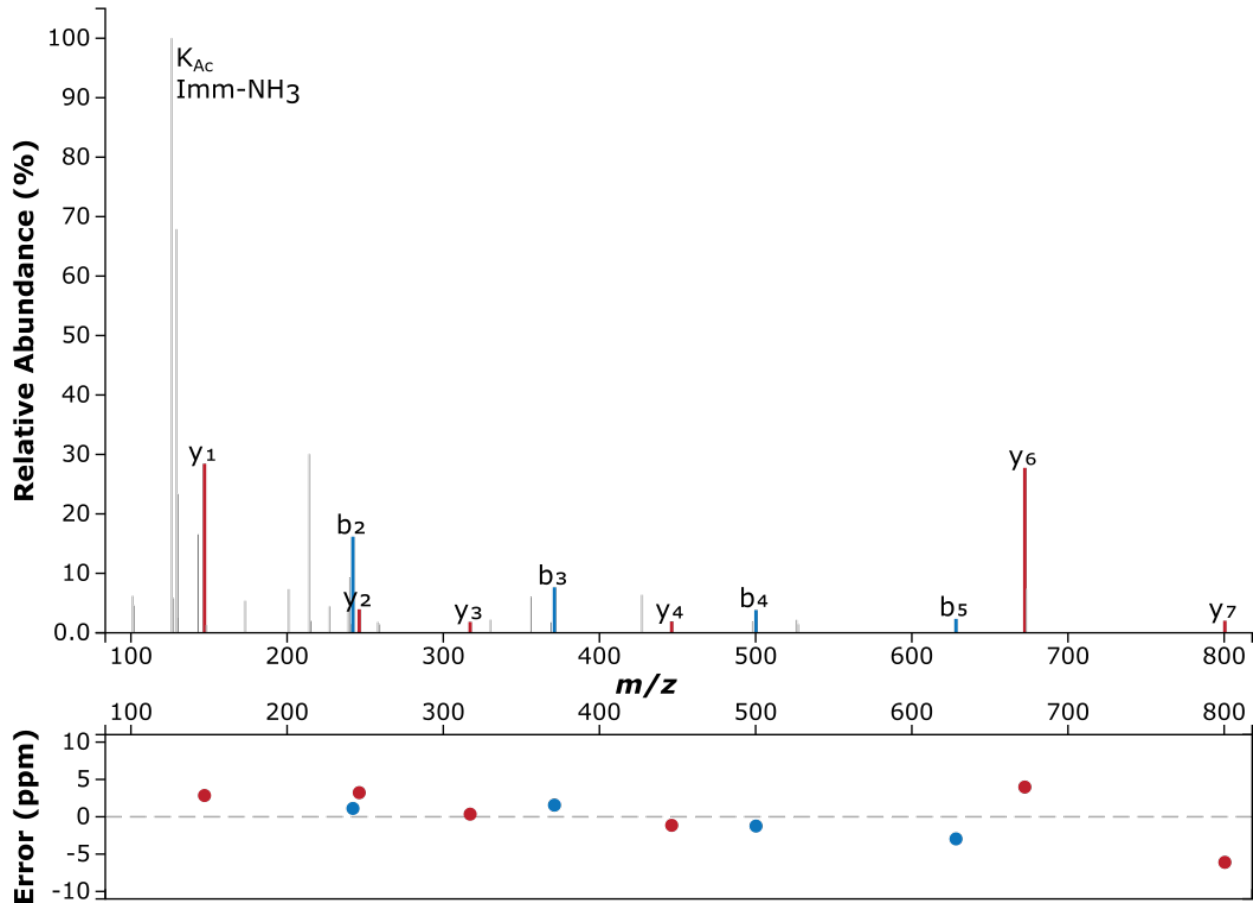
Charge: +2

6-oxocyclohex-1-ene-1-carboxyl-CoA hydrolase (Q2LXU2)





Precursor m/z: 476.5841 Charge: +3
 Hypothetical exported protein (Q2LUP5)

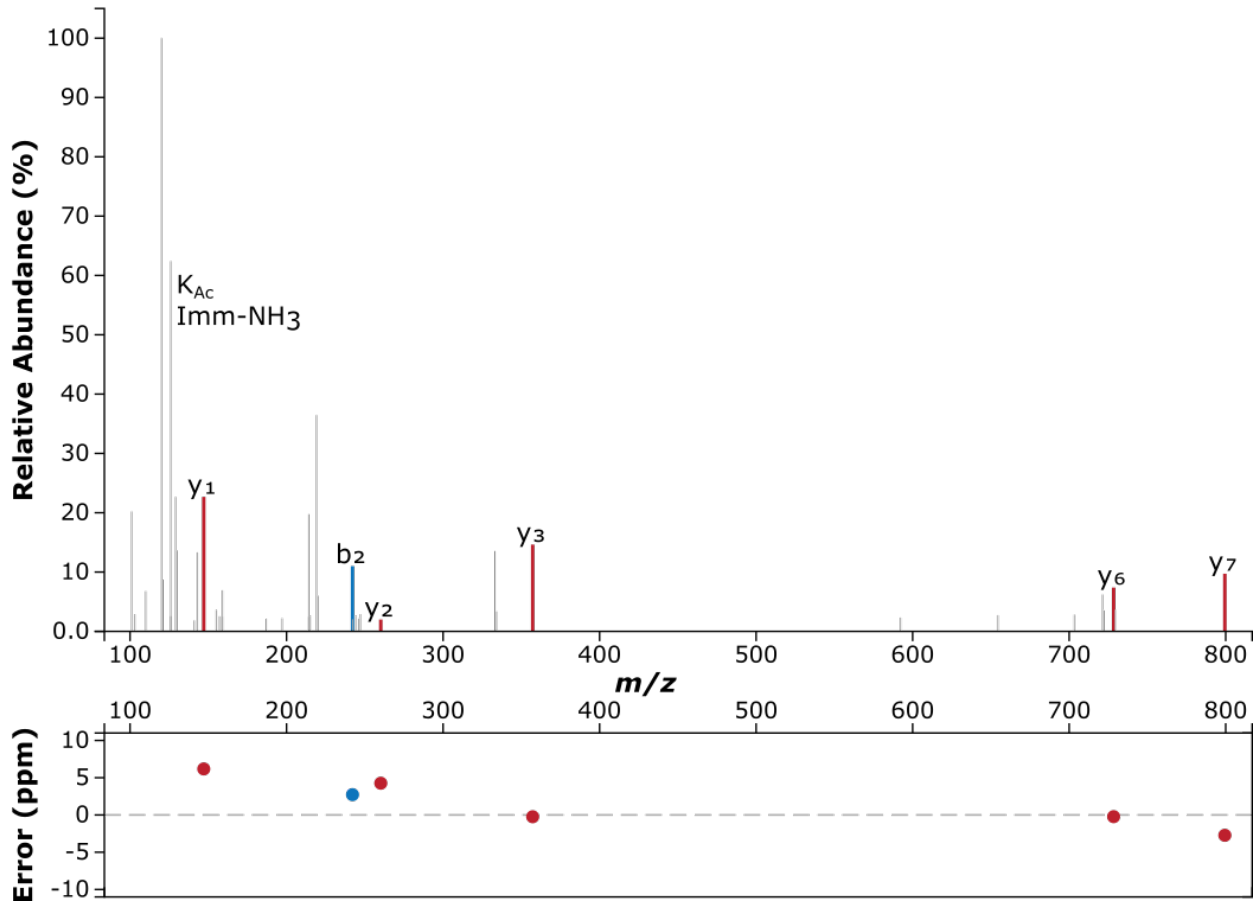




Precursor m/z: 485.2844

Charge: +2

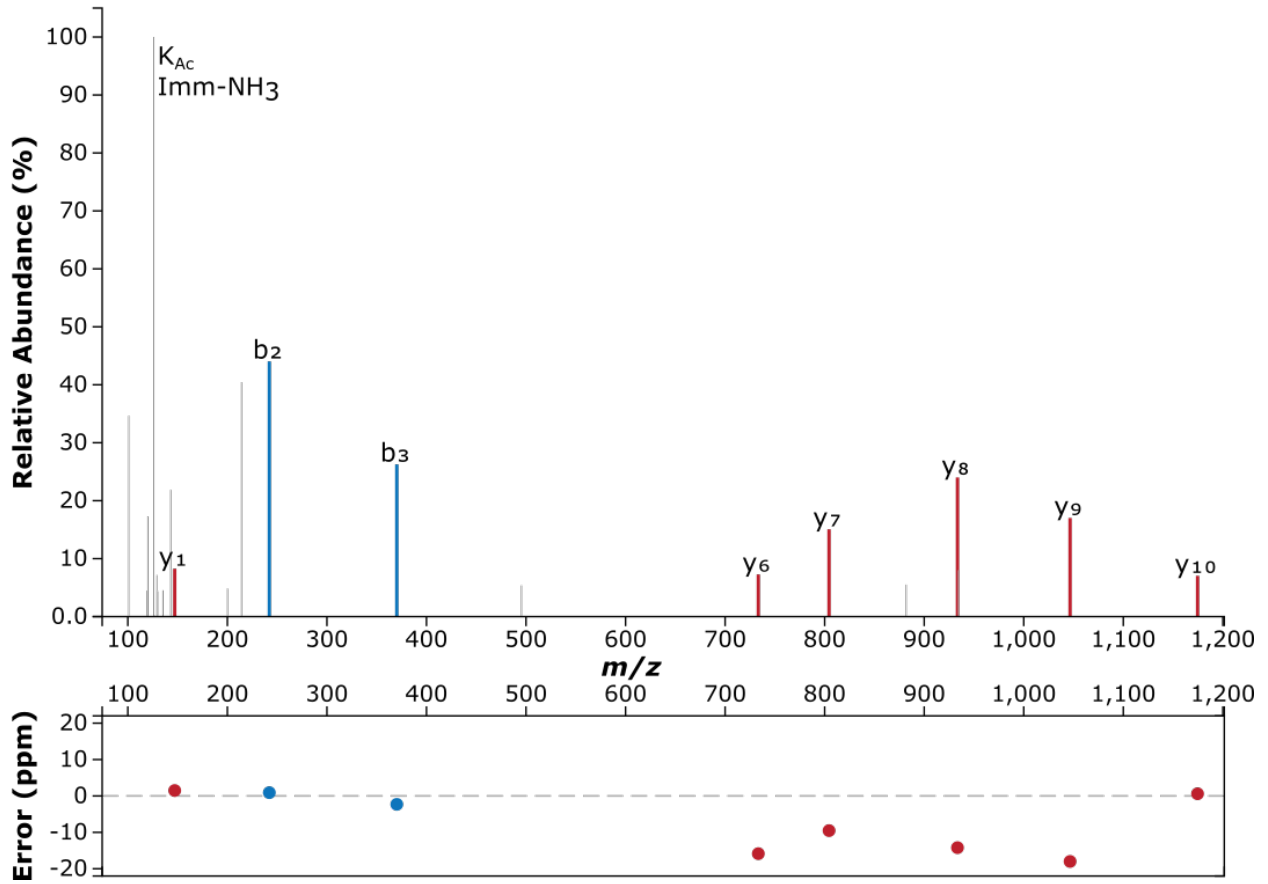
60 kDa chaperonin 2 (Q2LTG7)





Precursor m/z: 708.3725 Charge: +2

ATP synthase subunit b (Q2LR00)

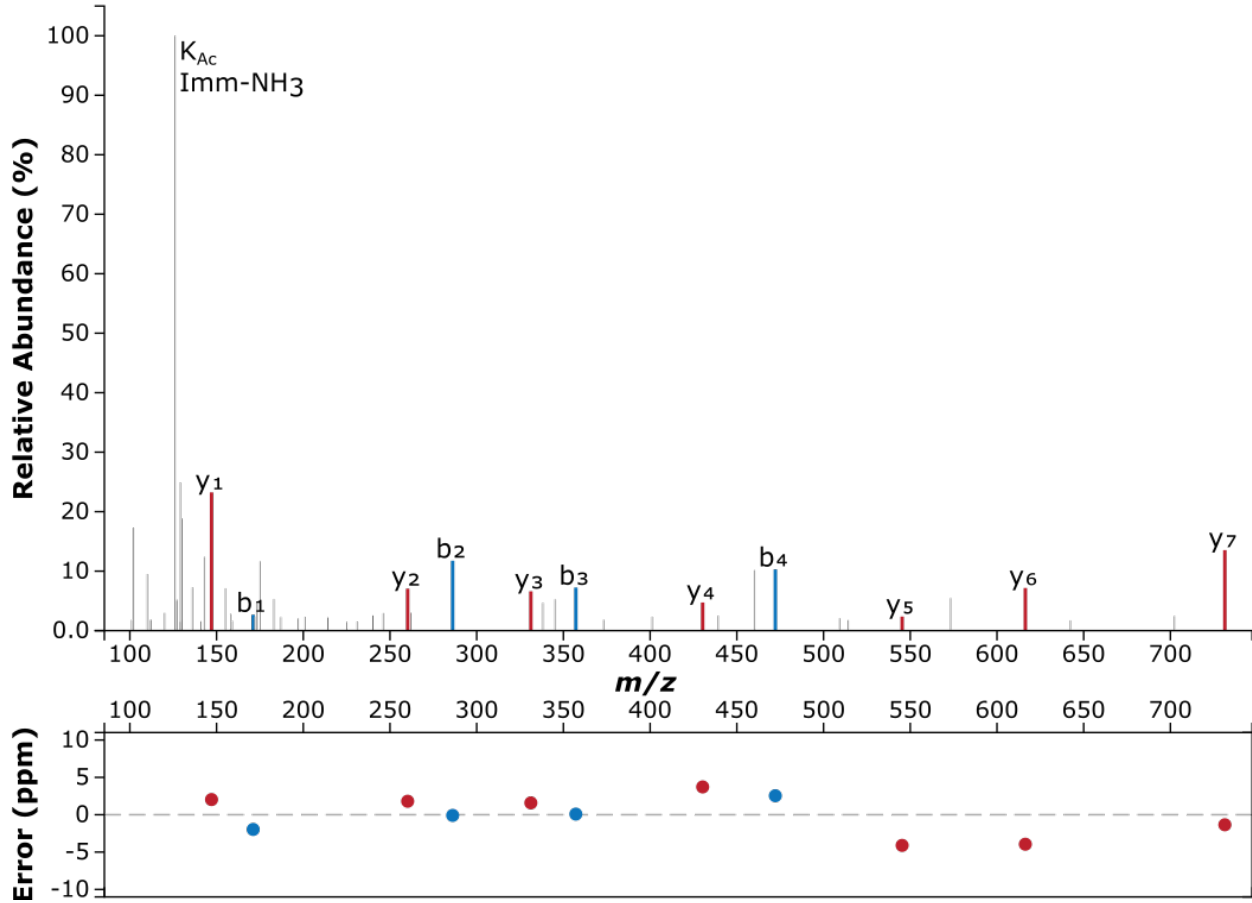




Precursor m/z: 451.2531

Charge: +2

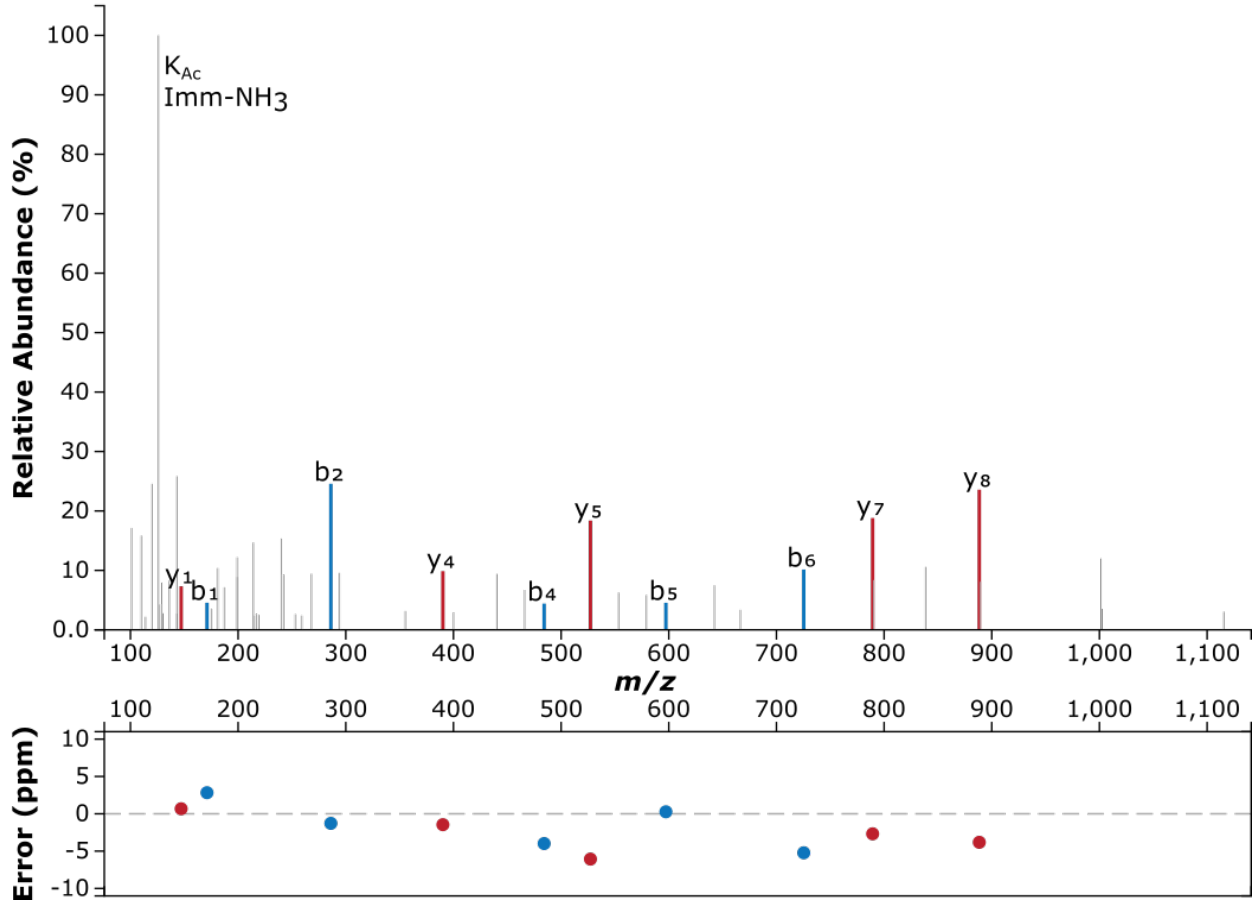
Acetyl-CoA synthetase (Q2LWR8)





Precursor m/z: 613.6597 Charge: +3

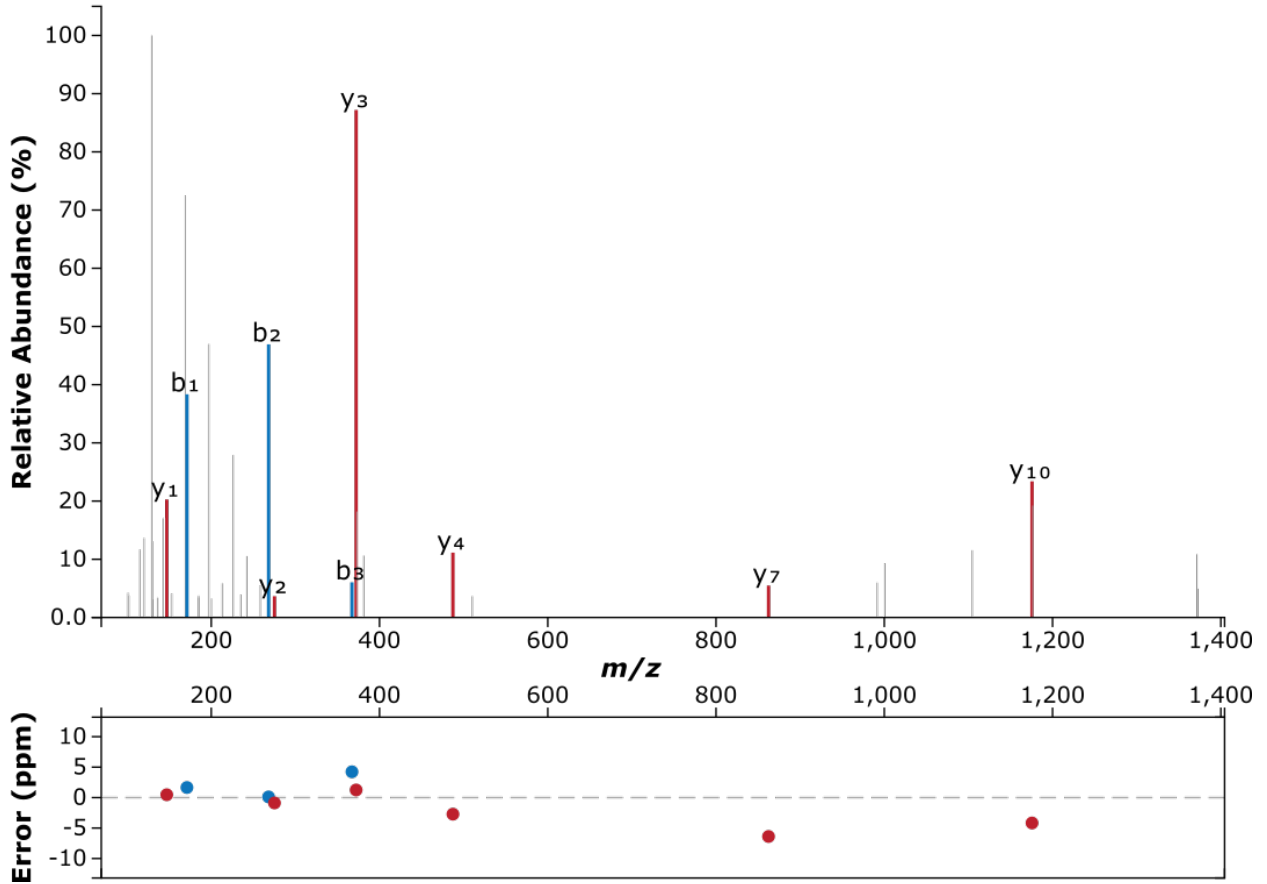
Putative enoyl-CoA hydratase (Q2LXU6)





Precursor m/z: 771.4141 Charge: +2

Acetyl-CoA acetyltransferase (Q2LXR7)

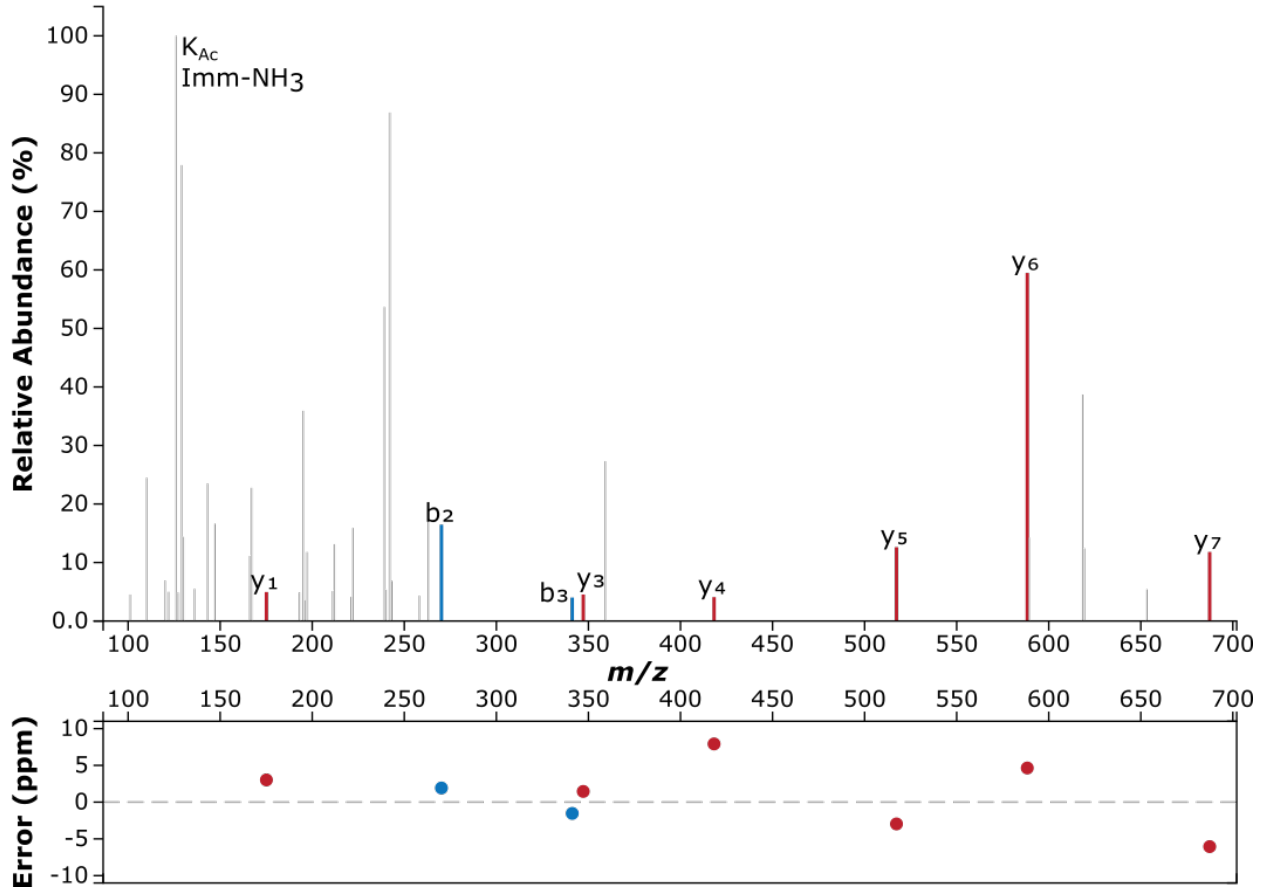




Precursor m/z: 429.2638

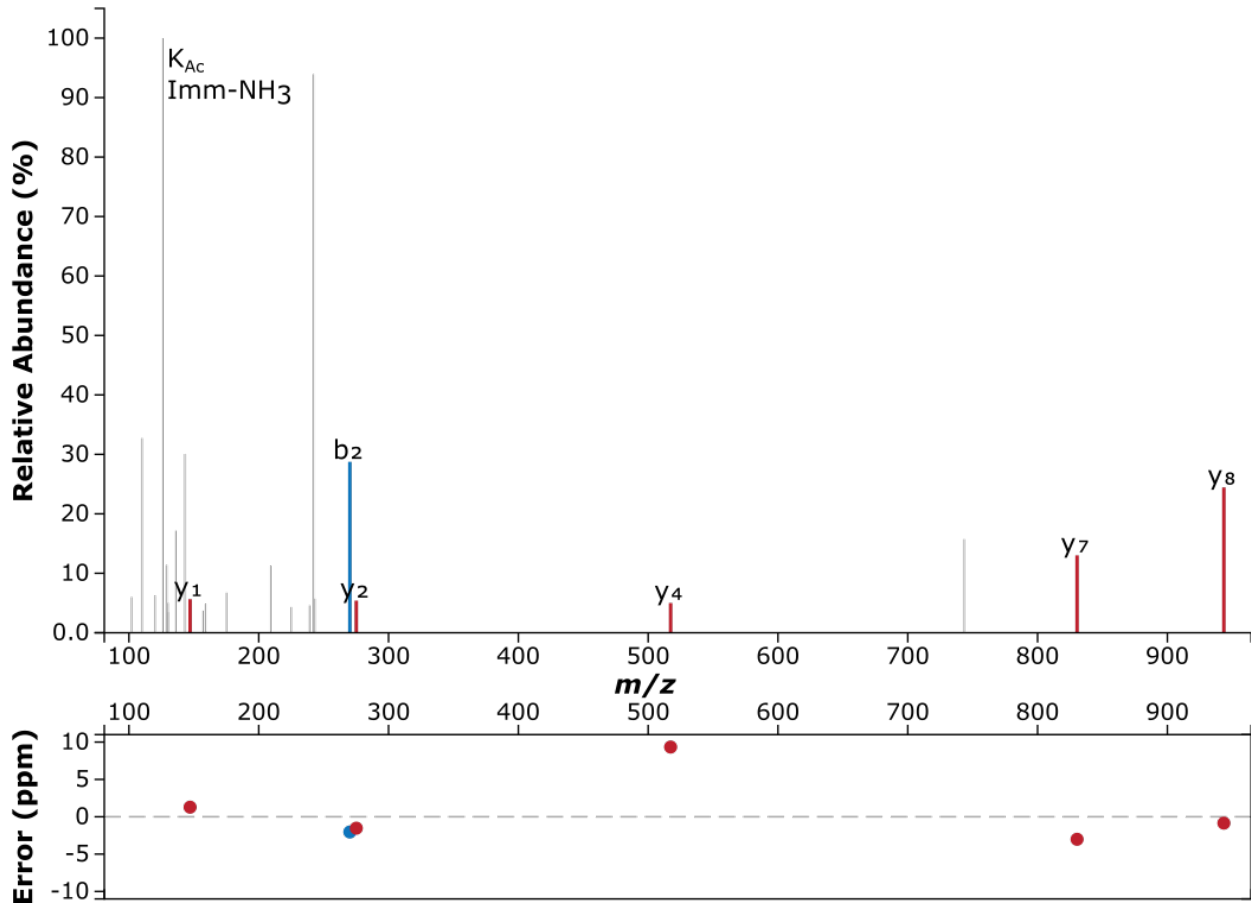
Charge: +2

Fe-S oxidoreductase (Q2LT65)





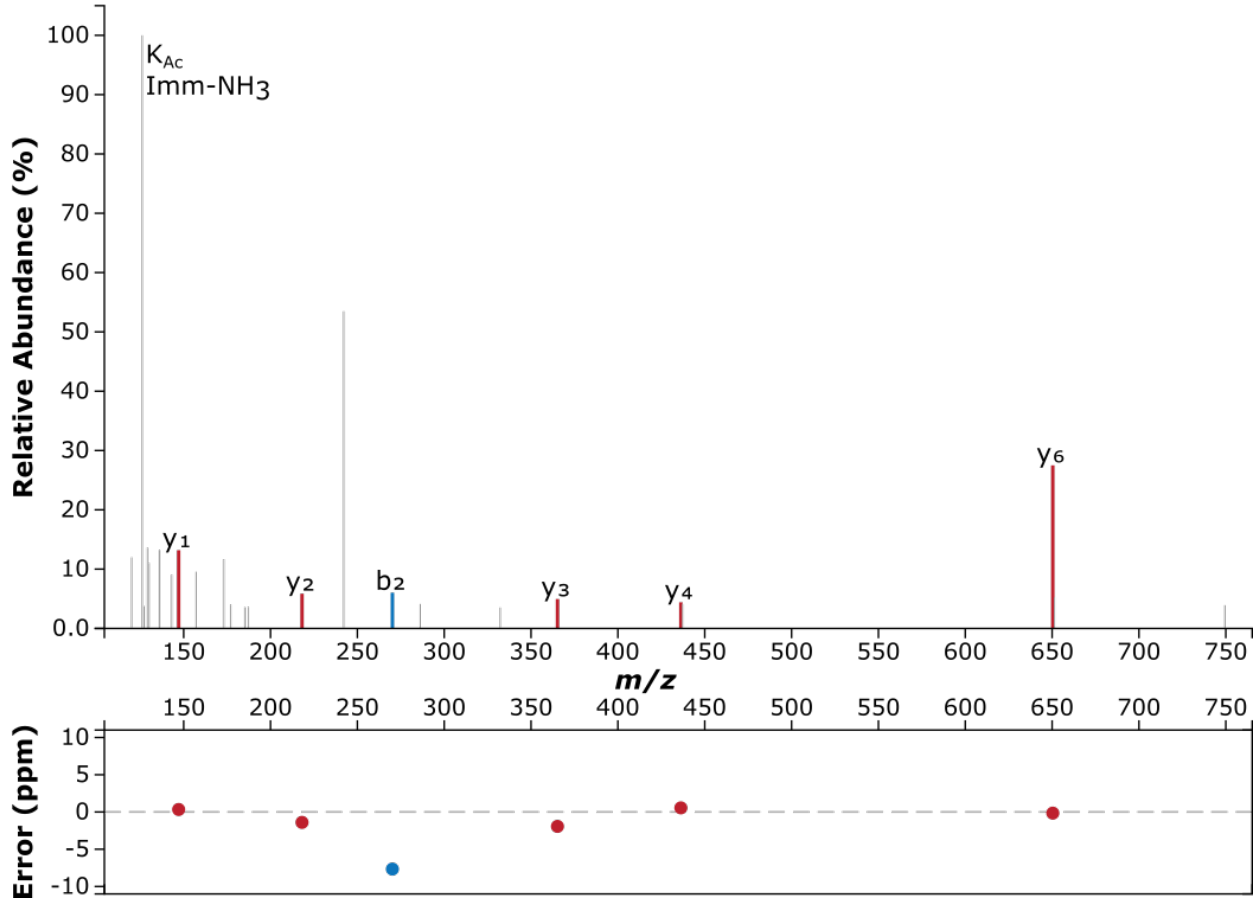
Precursor m/z: 606.8453 Charge: +2
 ATP synthase subunit alpha 1 (Q2LQZ7)





Precursor m/z: 460.2842 Charge: +2

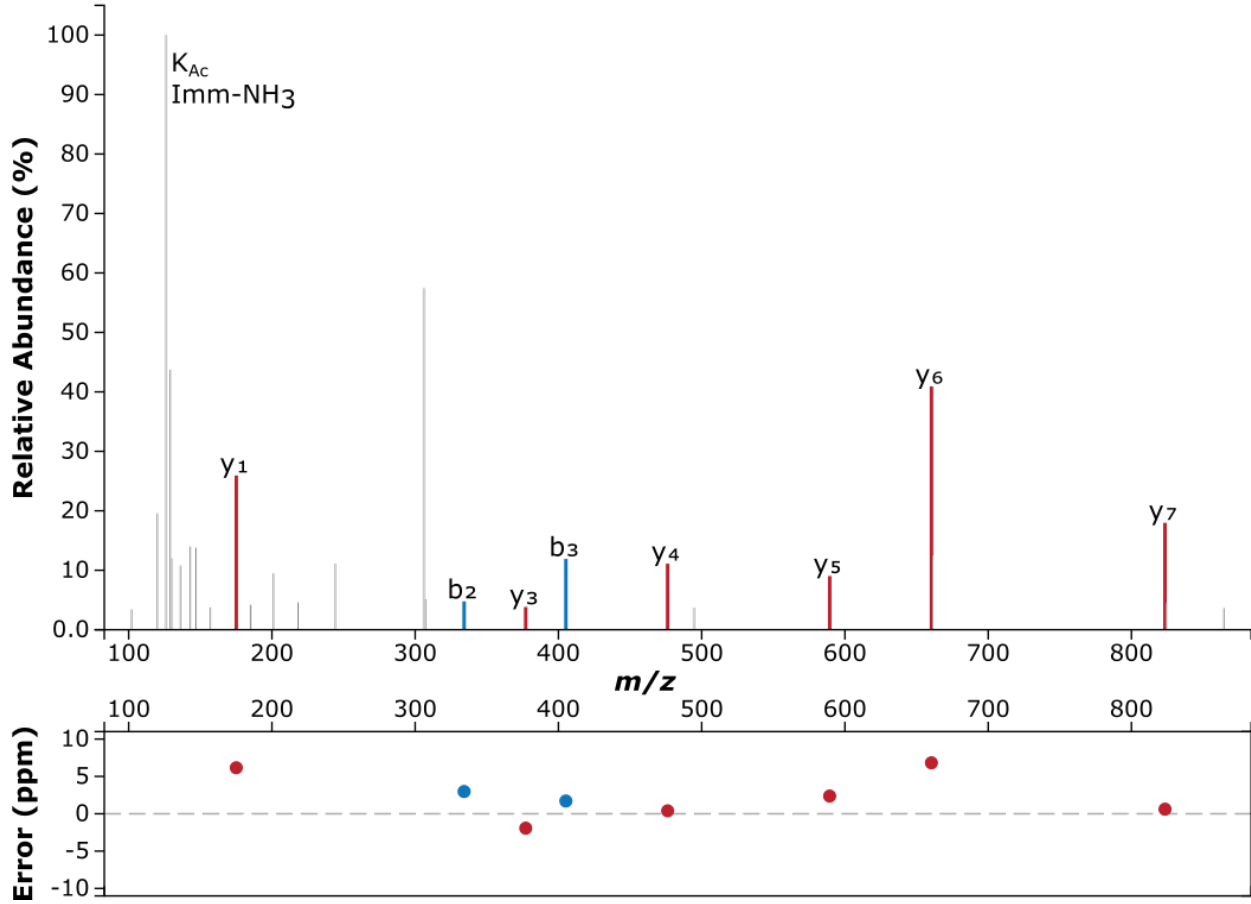
Iron-sulphur-binding reductase (Q2LWR1)





Precursor m/z: 497.2718 Charge: +2

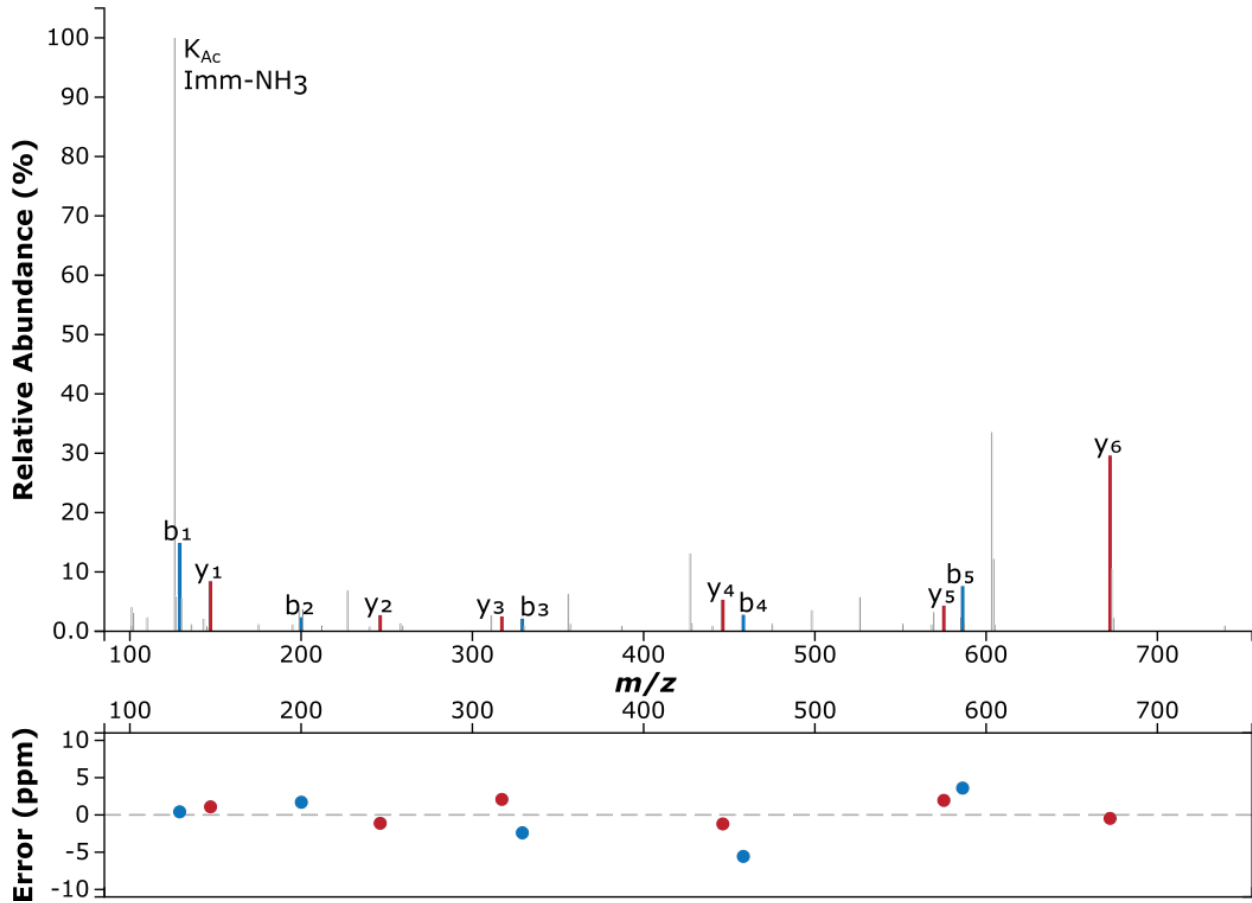
ATP synthase gamma chain (Q2LQZ6)





Precursor m/z: 476.5841 Charge: +3

Hypothetical exported protein (Q2LUP5)

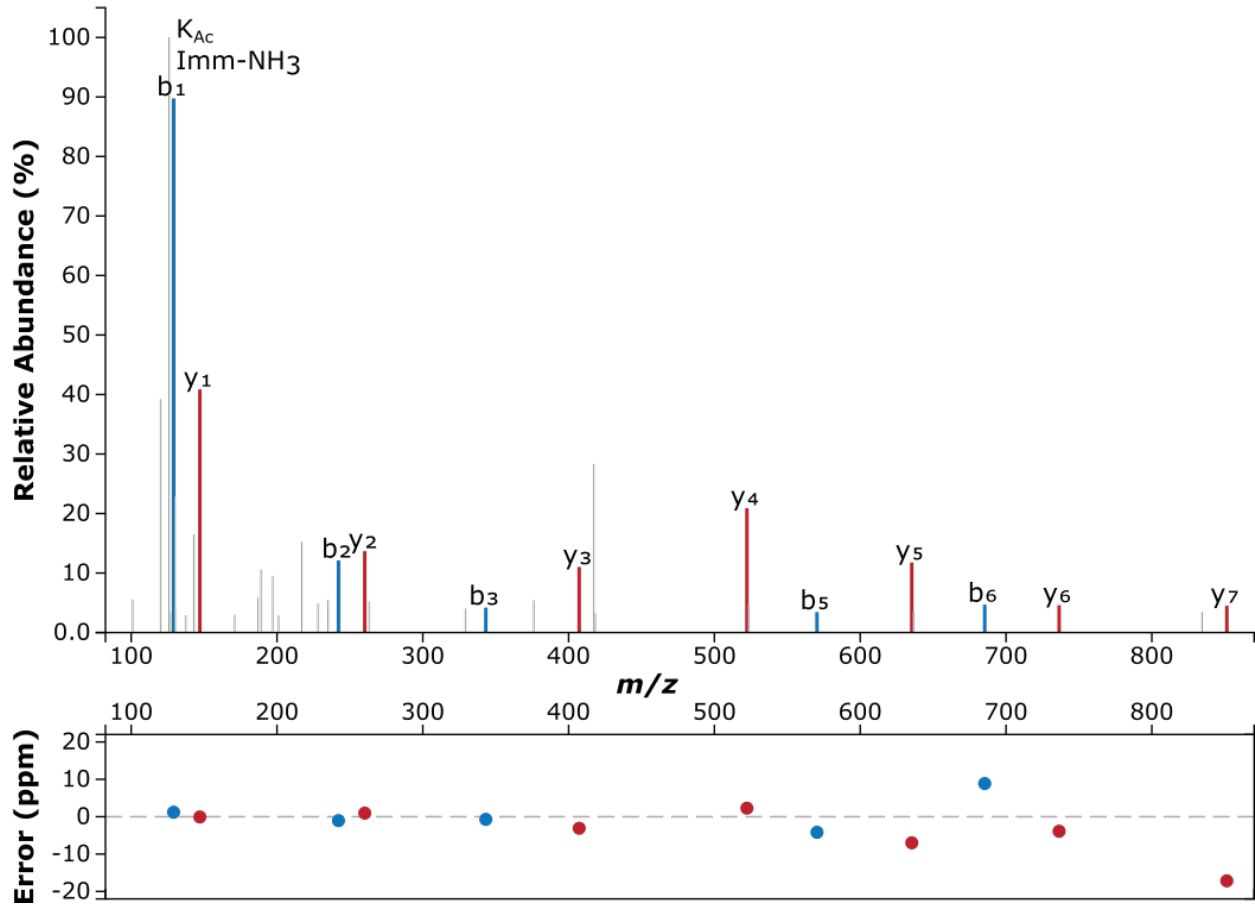




Precursor m/z: 474.2730

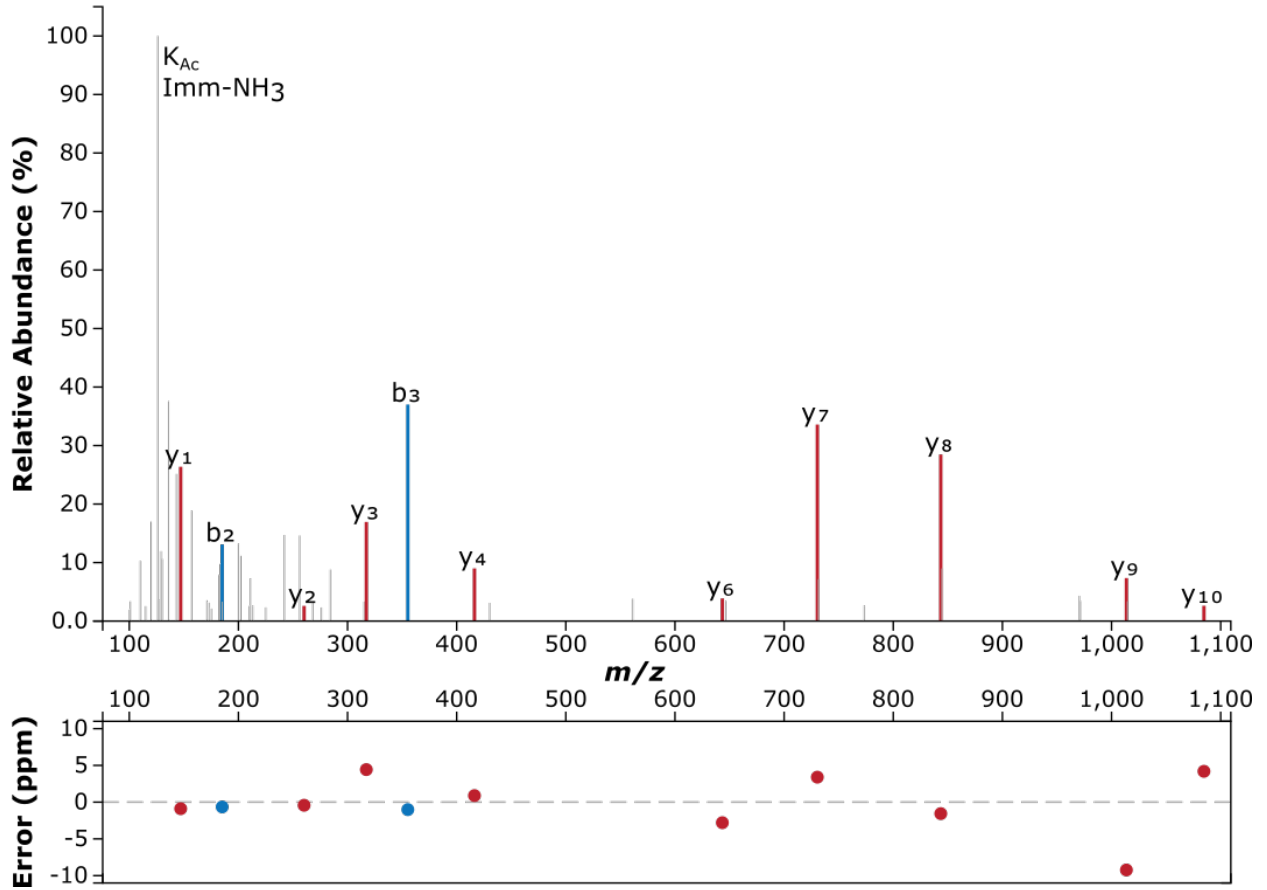
Charge: +3

6-oxocyclohex-1-ene-1-carboxyl-CoA hydrolase (Q2LXU2)





Precursor m/z: 599.3819 Charge: +2
 4-hydroxy-tetrahydrodipicolinate synthase (Q2LTA1)

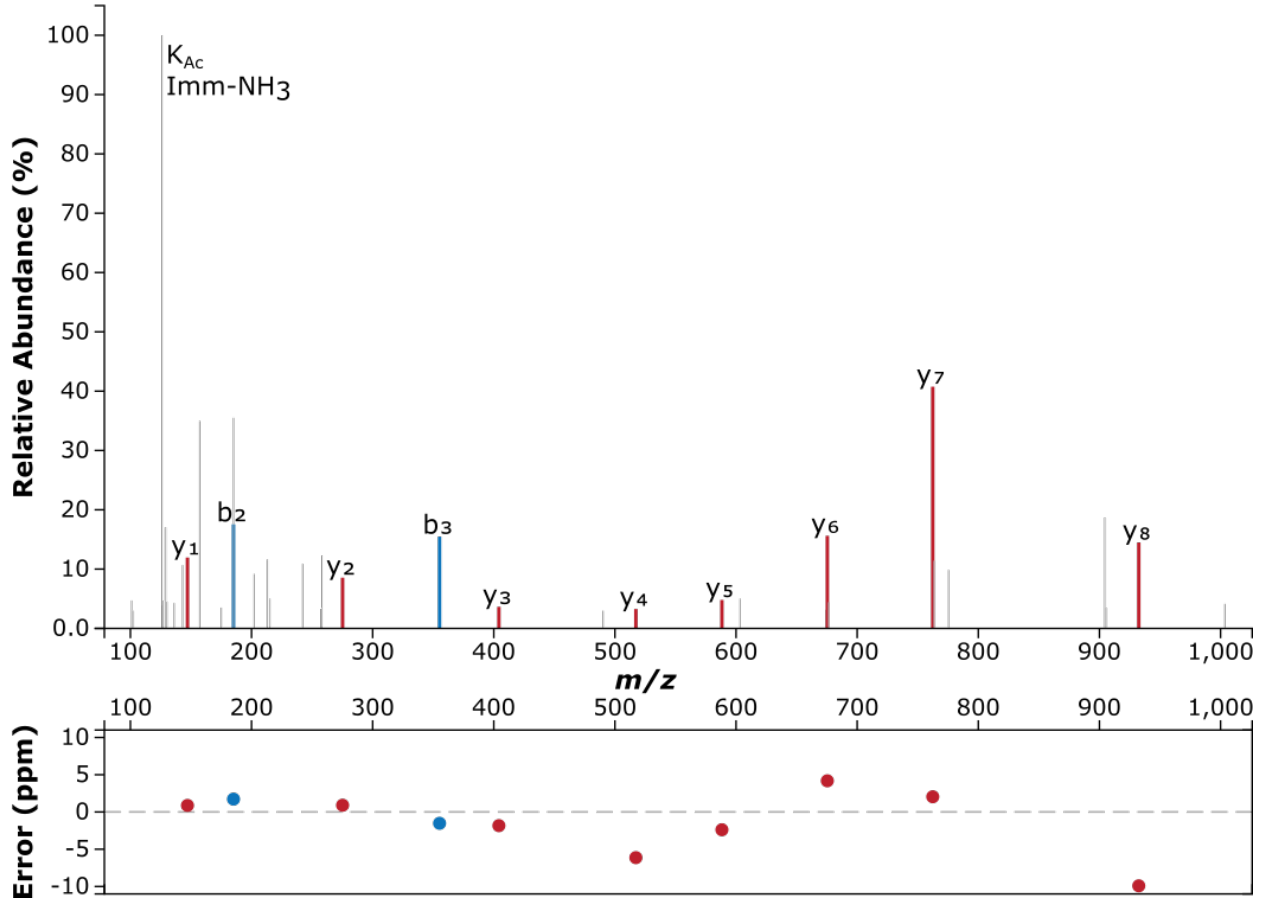




Precursor m/z: 558.8166

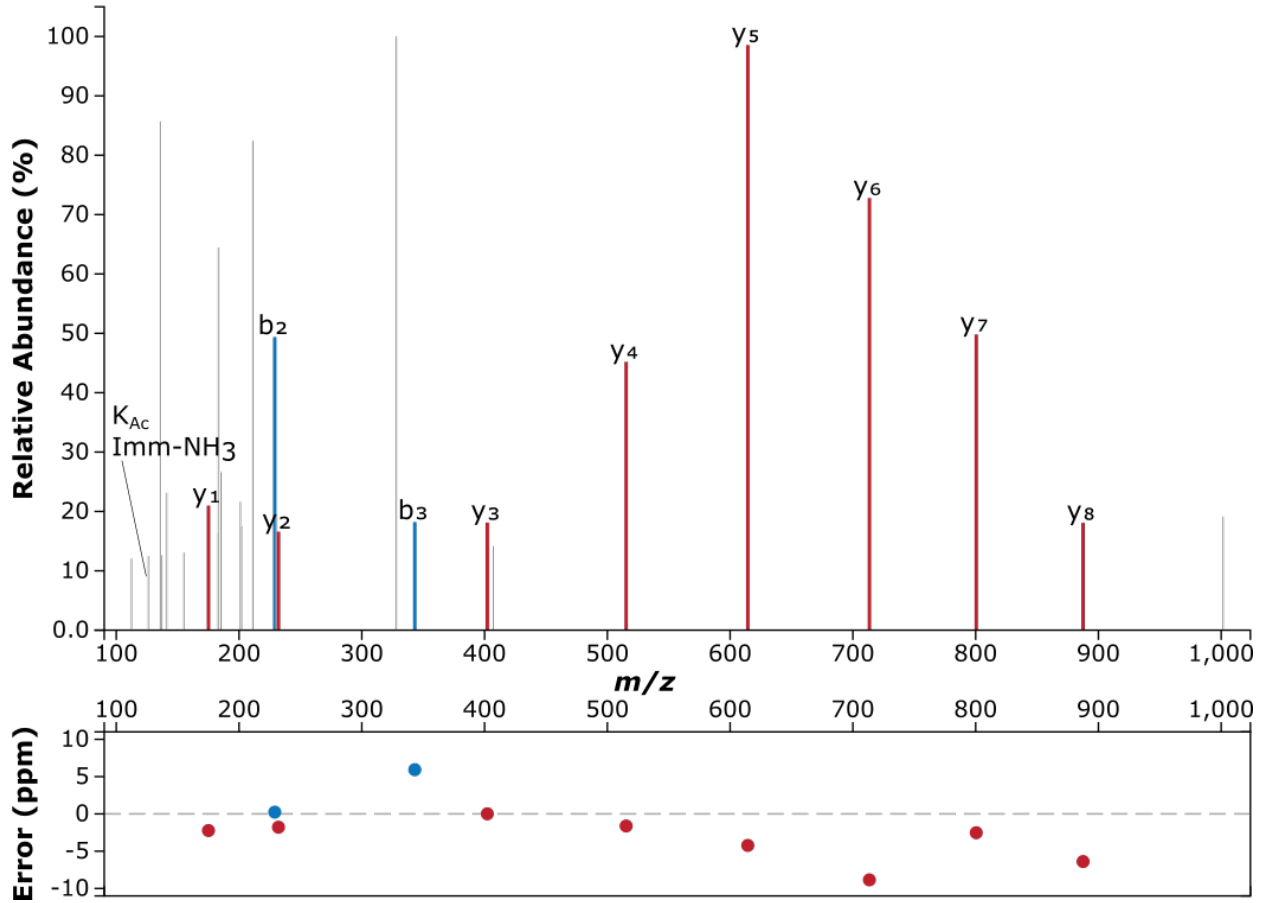
Charge: +2

ATP synthase subunit b (Q2LR00)





Precursor m/z: 615.3460 Charge: +2
 Mechanosensitive ion channel (Q2LX89)

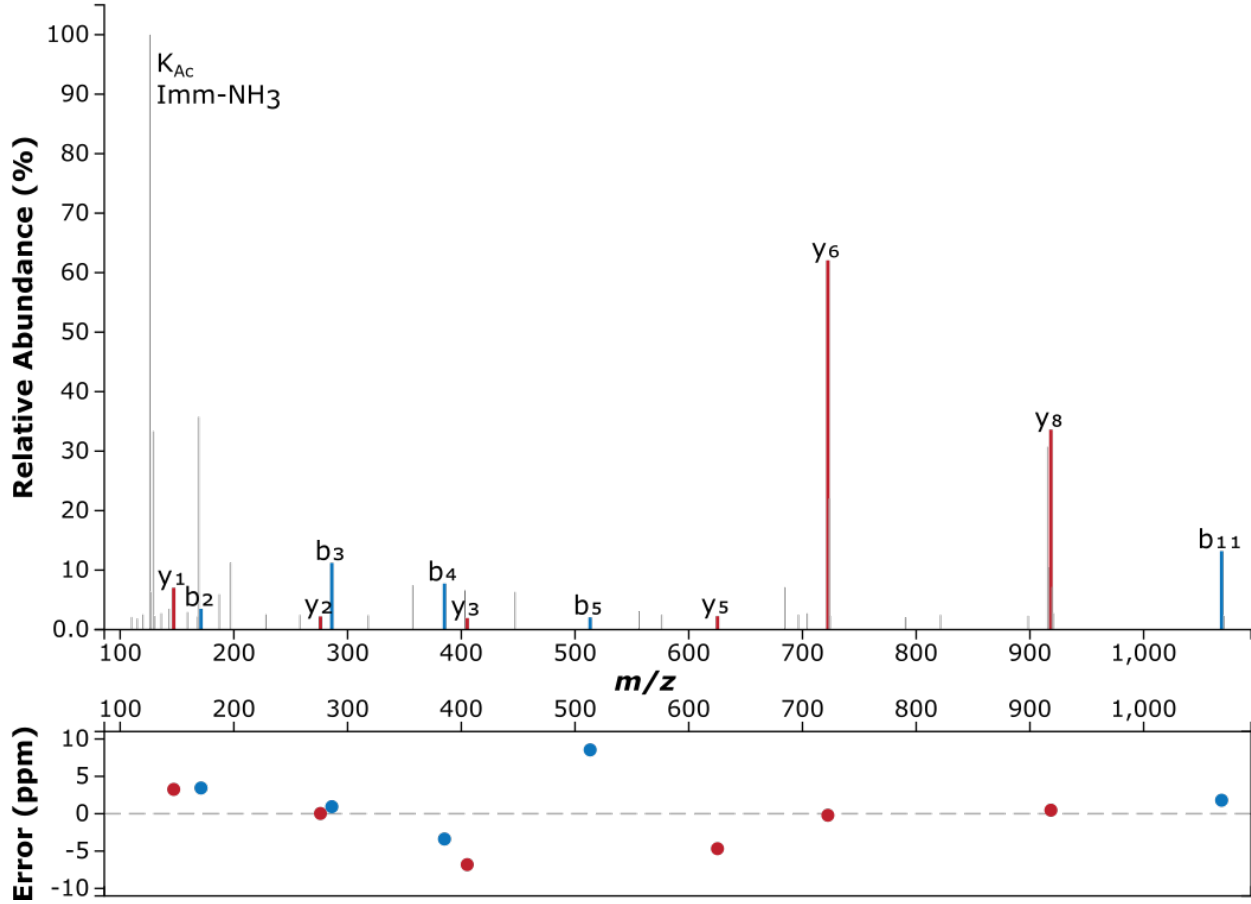


L G D V K P G S S G k^{Ac} P V P G Y E E K

Precursor m/z: 662.6775

Charge: +3

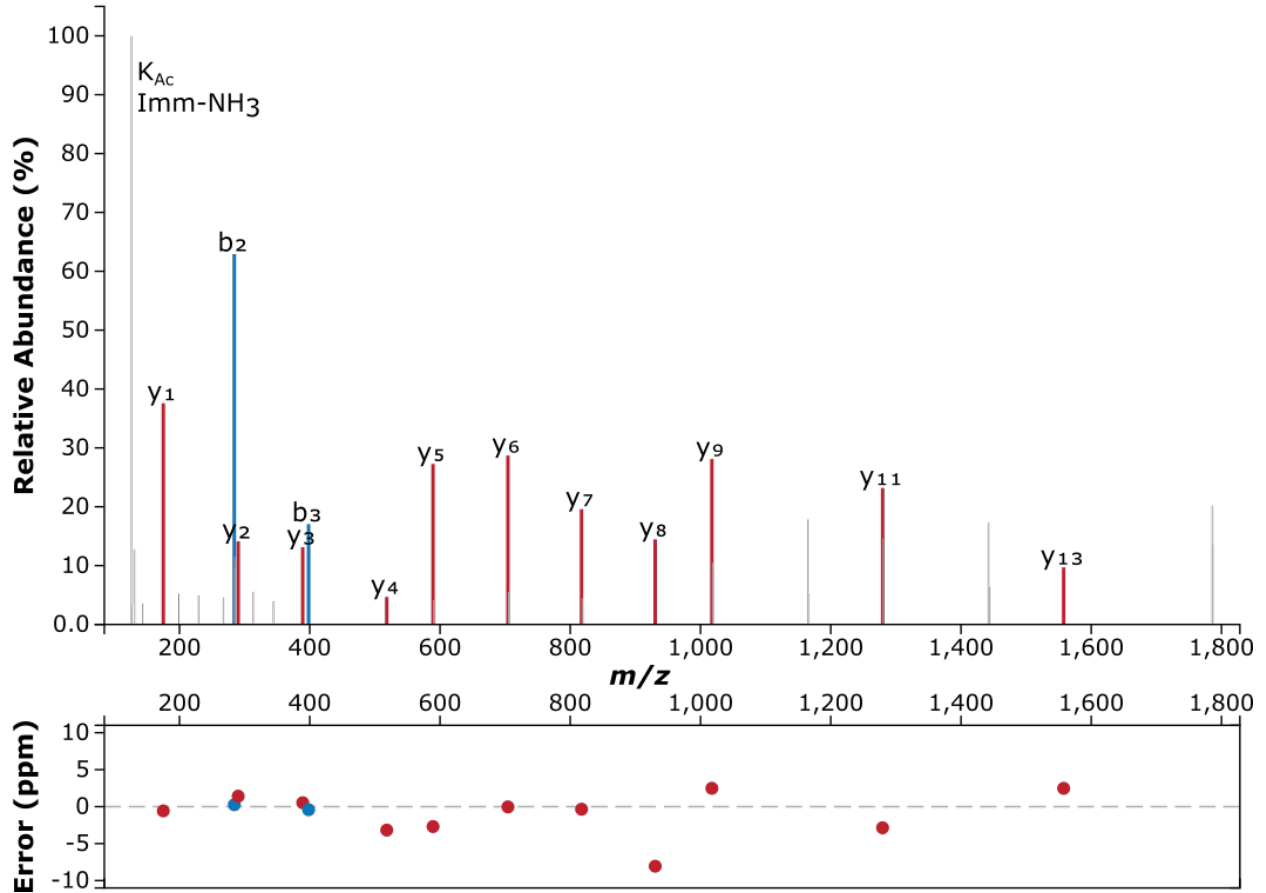
4-hydroxybenzoate-CoA ligase / benzoate-CoA ligase (Q2LRH0)





Precursor m/z: 1,034.9947 Charge: +2

6-oxocyclohex-1-ene-1-carboxyl-CoA hydrolase (Q2LXU2)

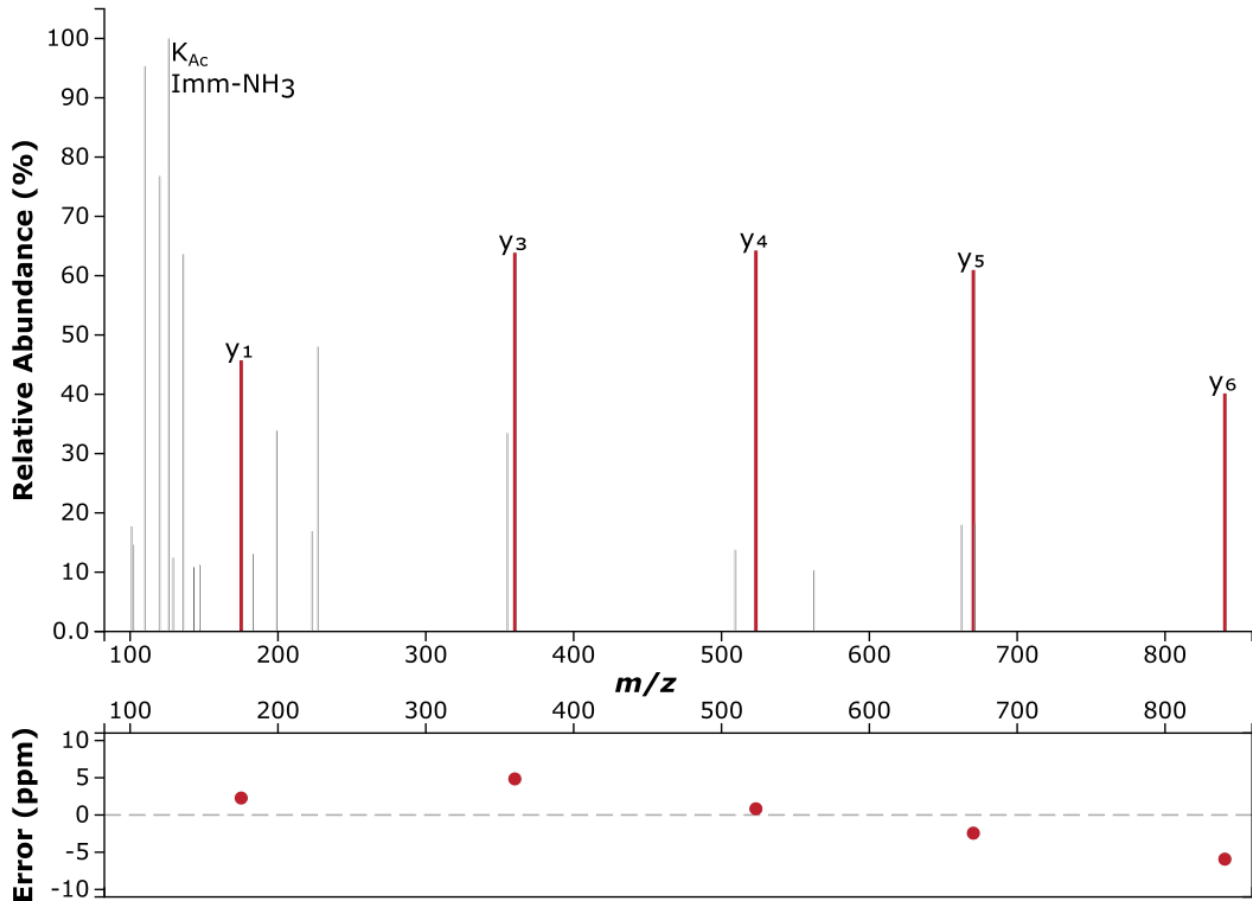


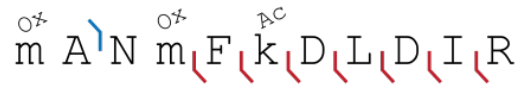
L P E Q H^{AC} k F Y A N R

Precursor m/z: 482.2492

Charge: +3

Glutaconyl-CoA decarboxylase A subunit (Q2LUY7)

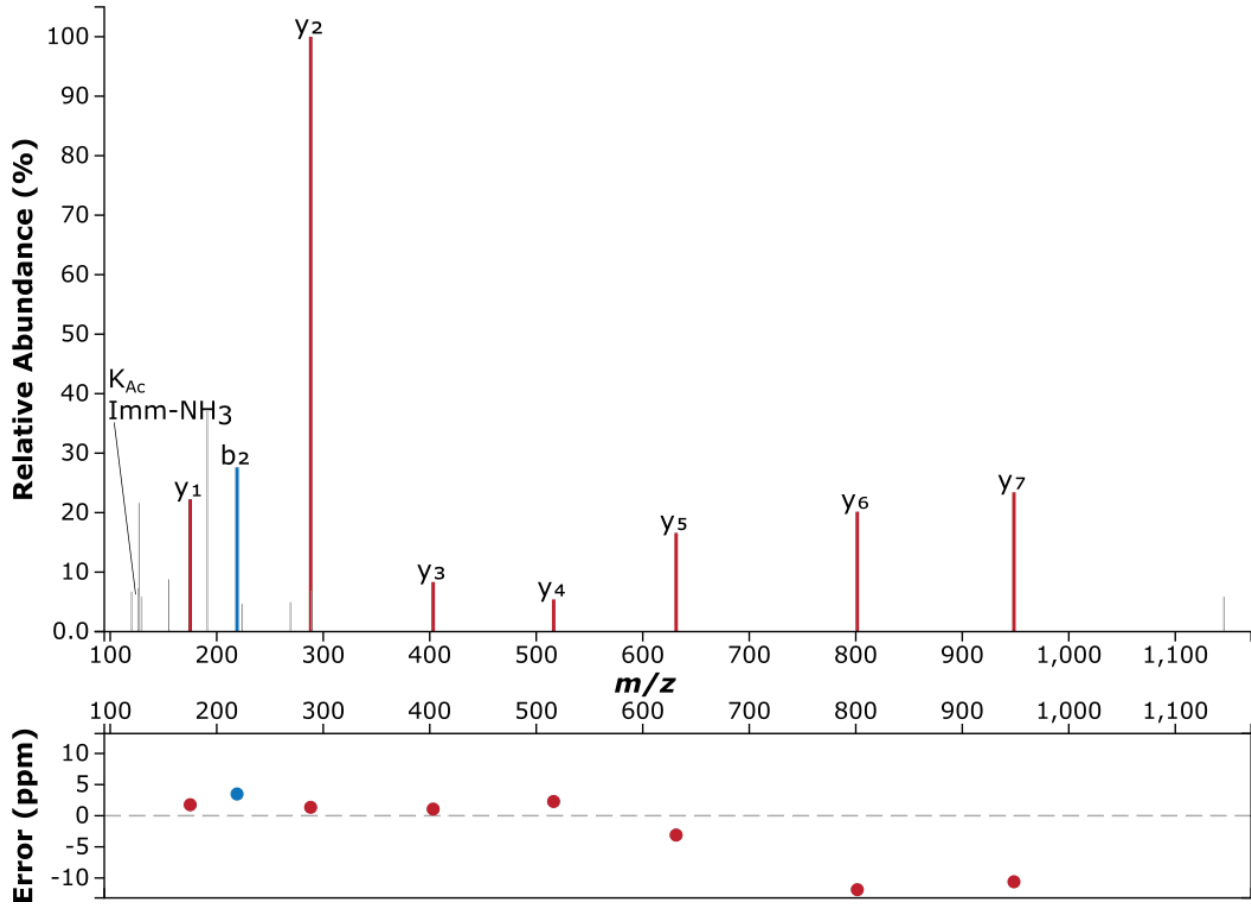




Precursor m/z: 476.5601

Charge: +3

4-hydroxybenzoate-CoA ligase / benzoate-CoA ligase (Q2LRH0)

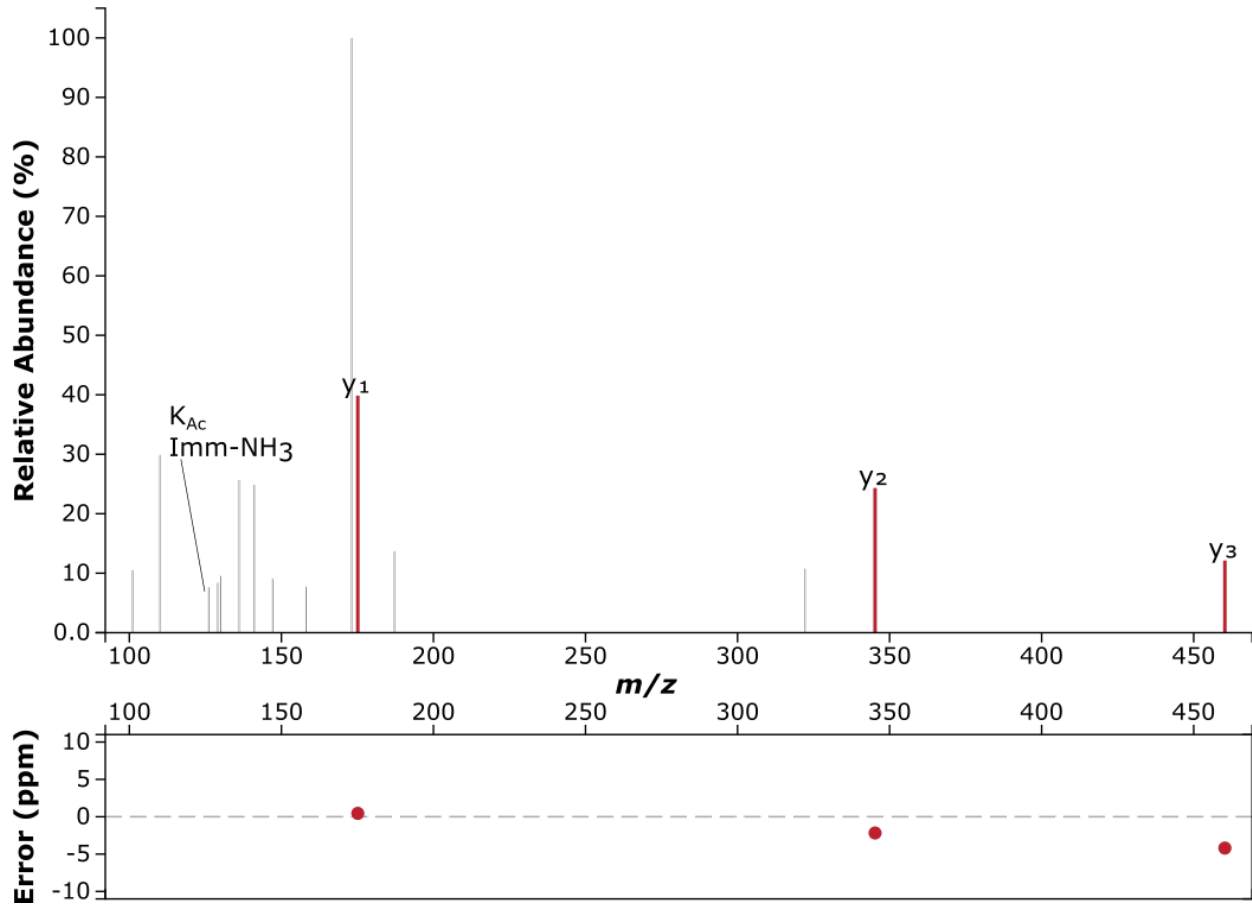




Precursor m/z: 446.7133

Charge: +2

Transcriptional regulator AsnC family (Q2LQV4)

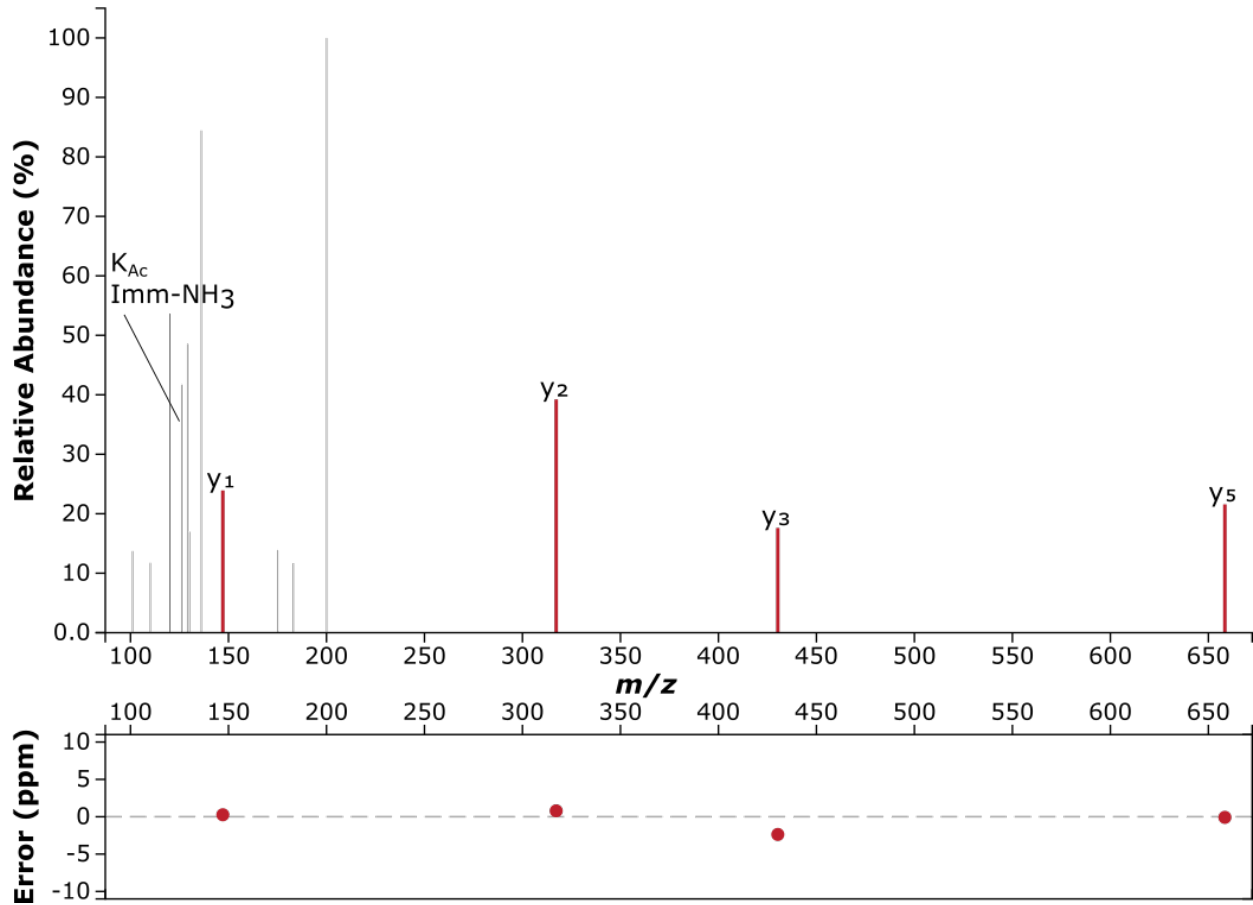


N L E V L k K

Precursor m/z: 443.2738

Charge: +2

3-dehydroquinase synthase (Q2LUD2)

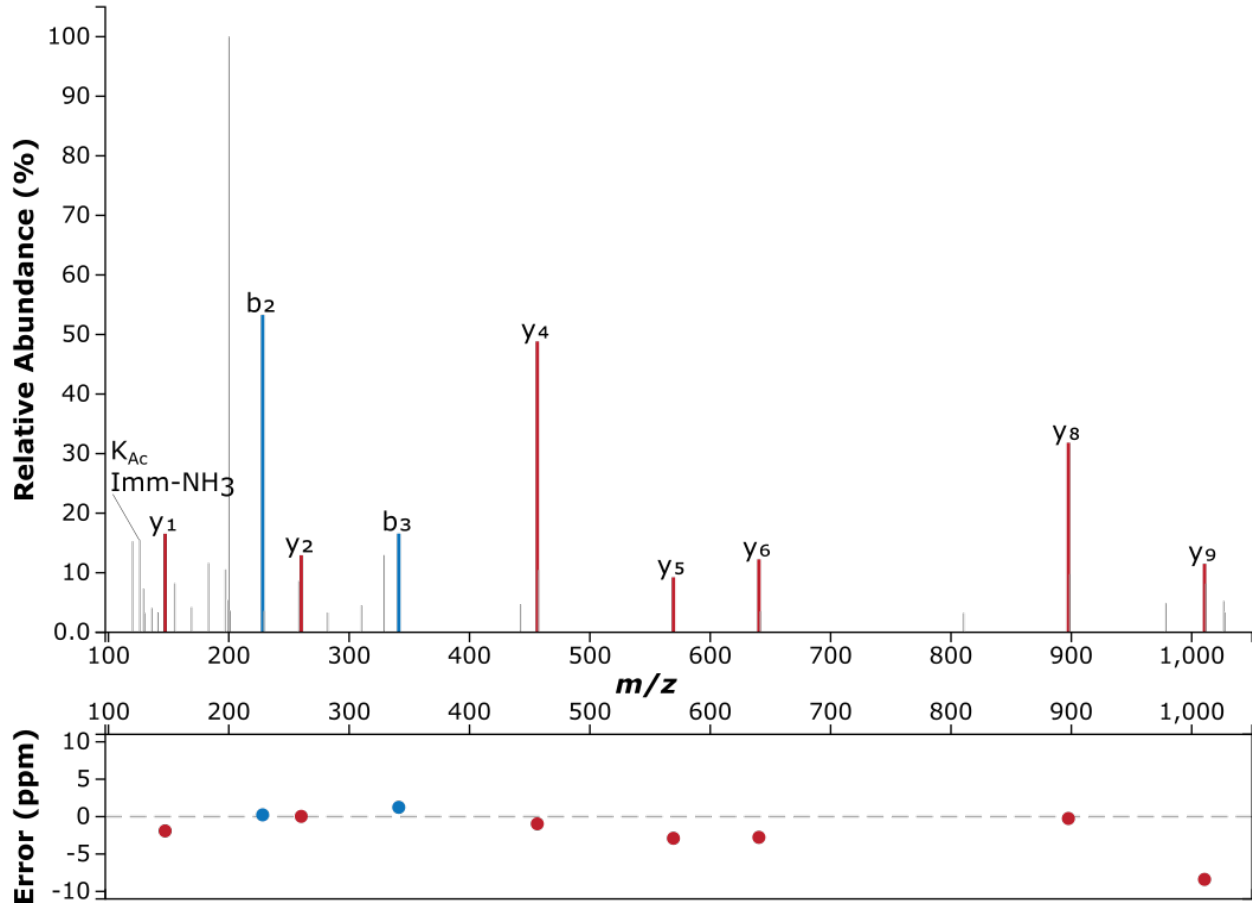




Precursor m/z: 619.3975

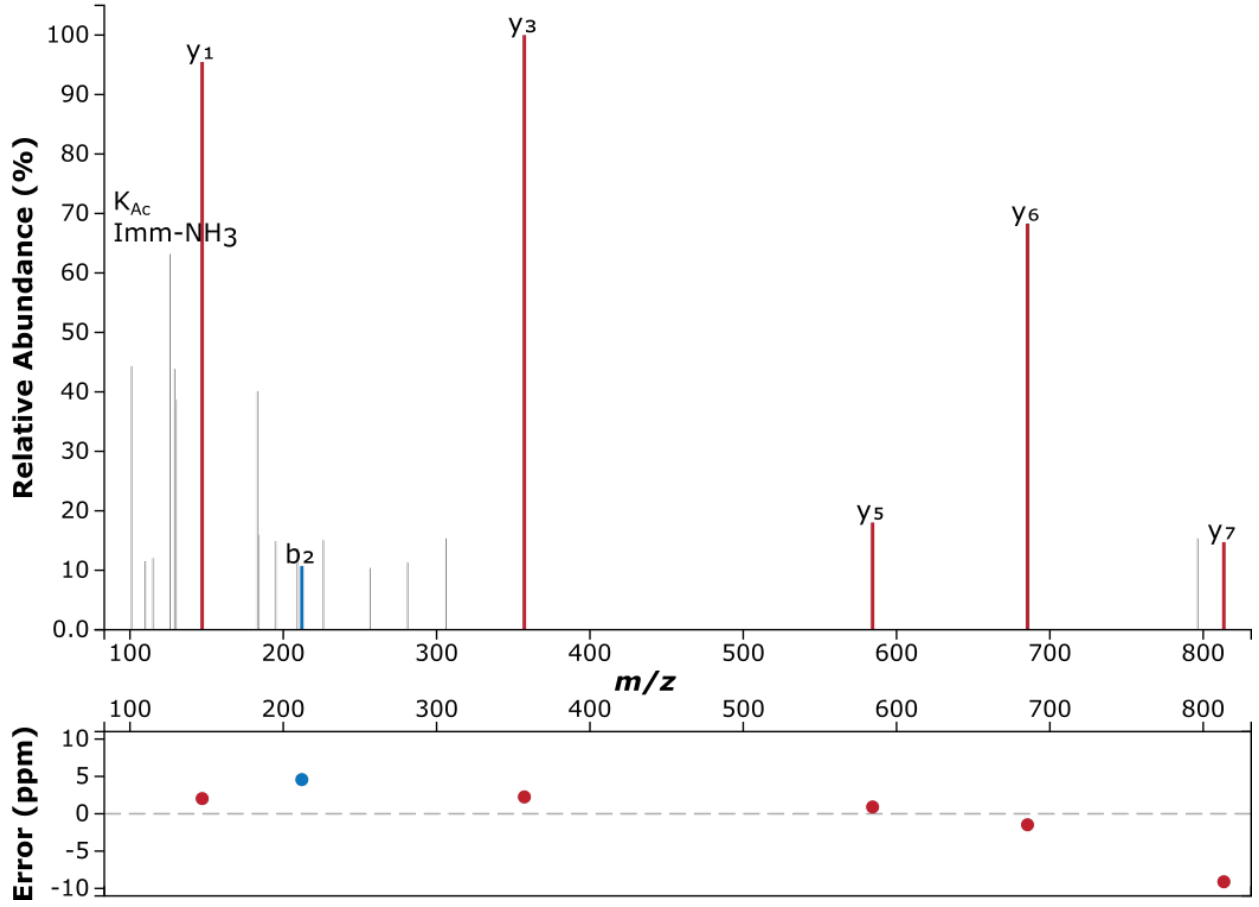
Charge: +2

6-oxocyclohex-1-ene-1-carboxyl-CoA hydrolase (Q2LXU2)





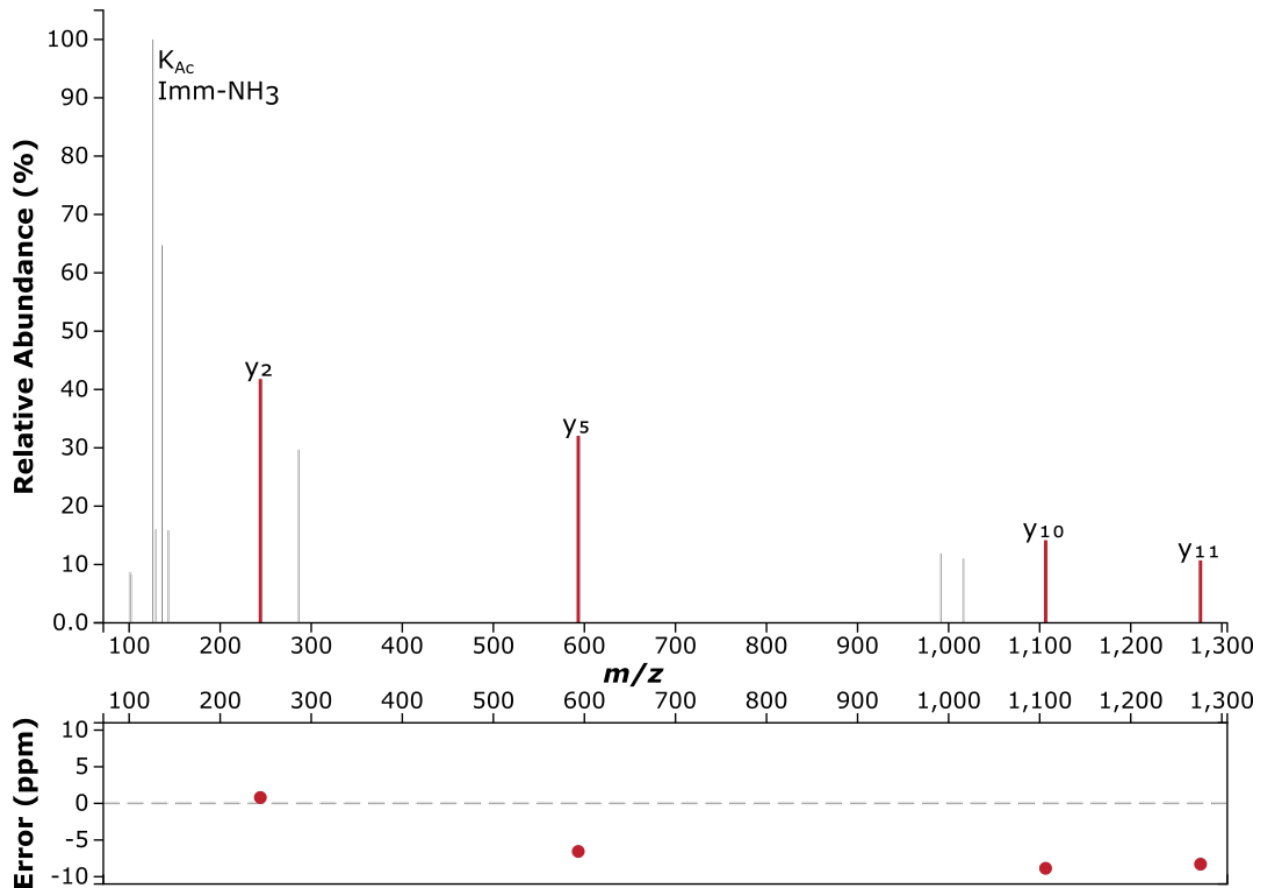
Precursor m/z: 512.7929 Charge: +2
 DNA-binding protein HU (Q2LW81)



N P Q V Y T D T Y V ^{Ac} k D G E I V Y G E P K

Precursor m/z: 819.7322 Charge: +3

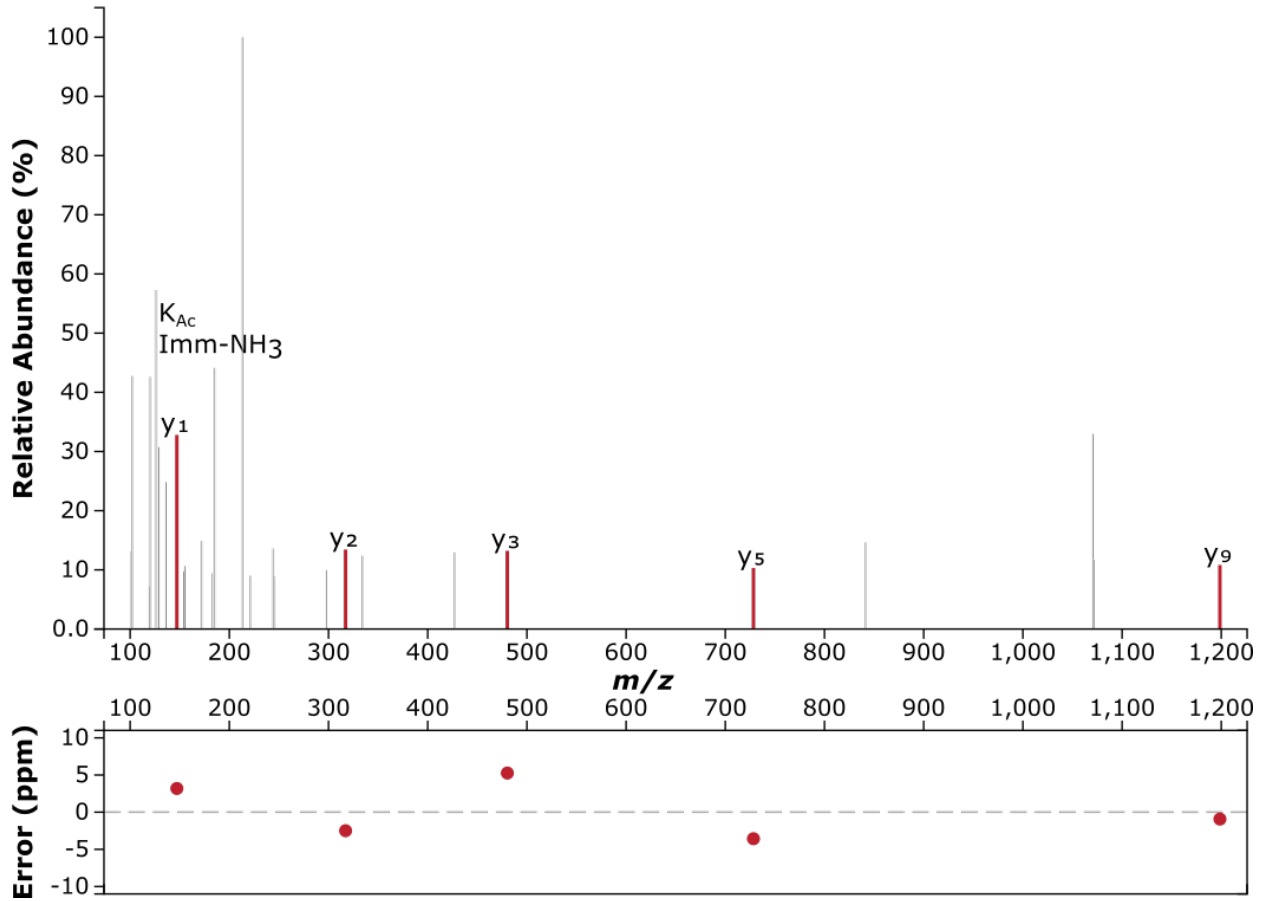
6-oxocyclohex-1-ene-1-carboxyl-CoA hydrolase (Q2LXU2)



Q A D E I L T F Y ^{Ac} k K

Precursor m/z: 699.3692 Charge: +2

Acetyl-CoA synthetase (Q2LWR8)

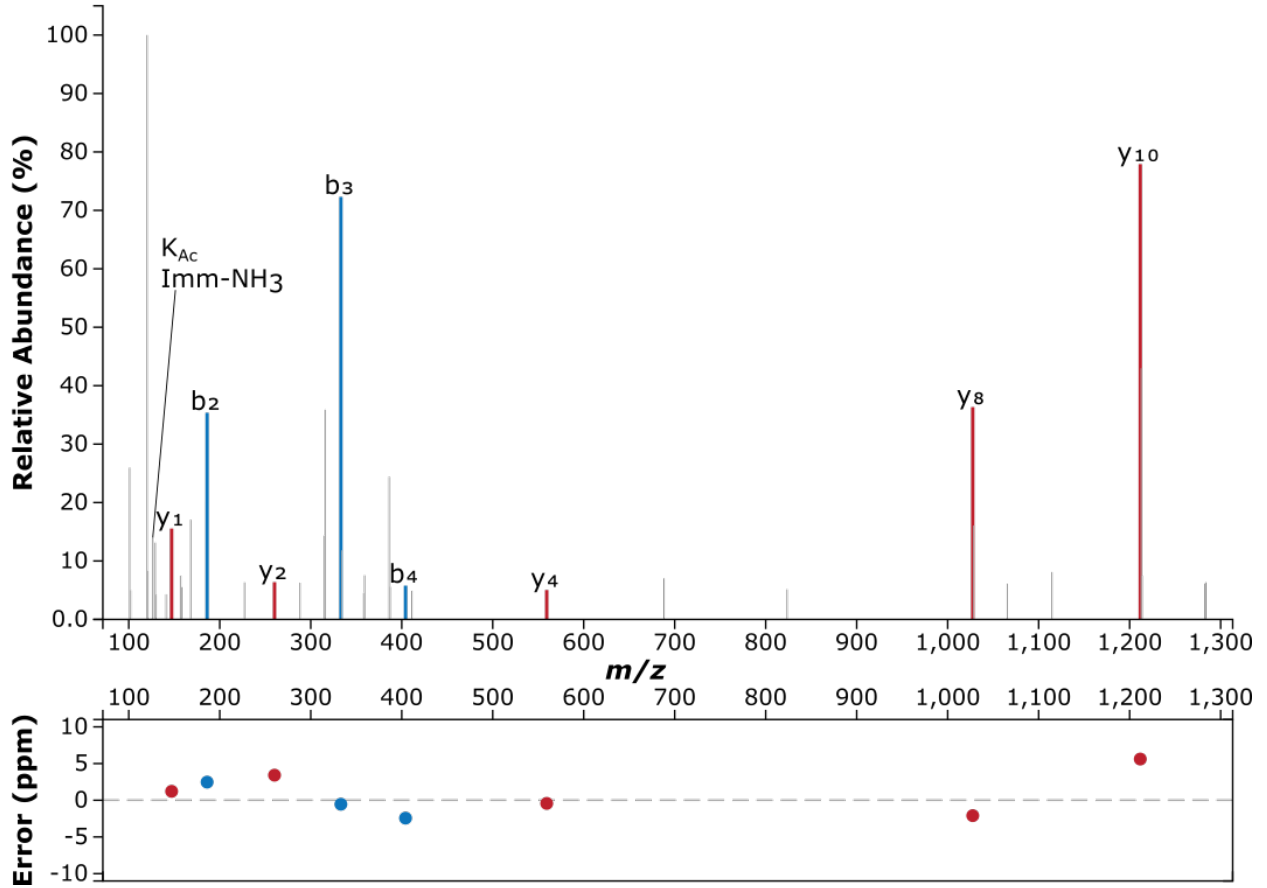


Q G F A P S P E L E ^{Ac} k E L K

Precursor m/z: 807.9223

Charge: +2

4-hydroxybenzoate-CoA ligase / benzoate-CoA ligase (Q2LRH7)

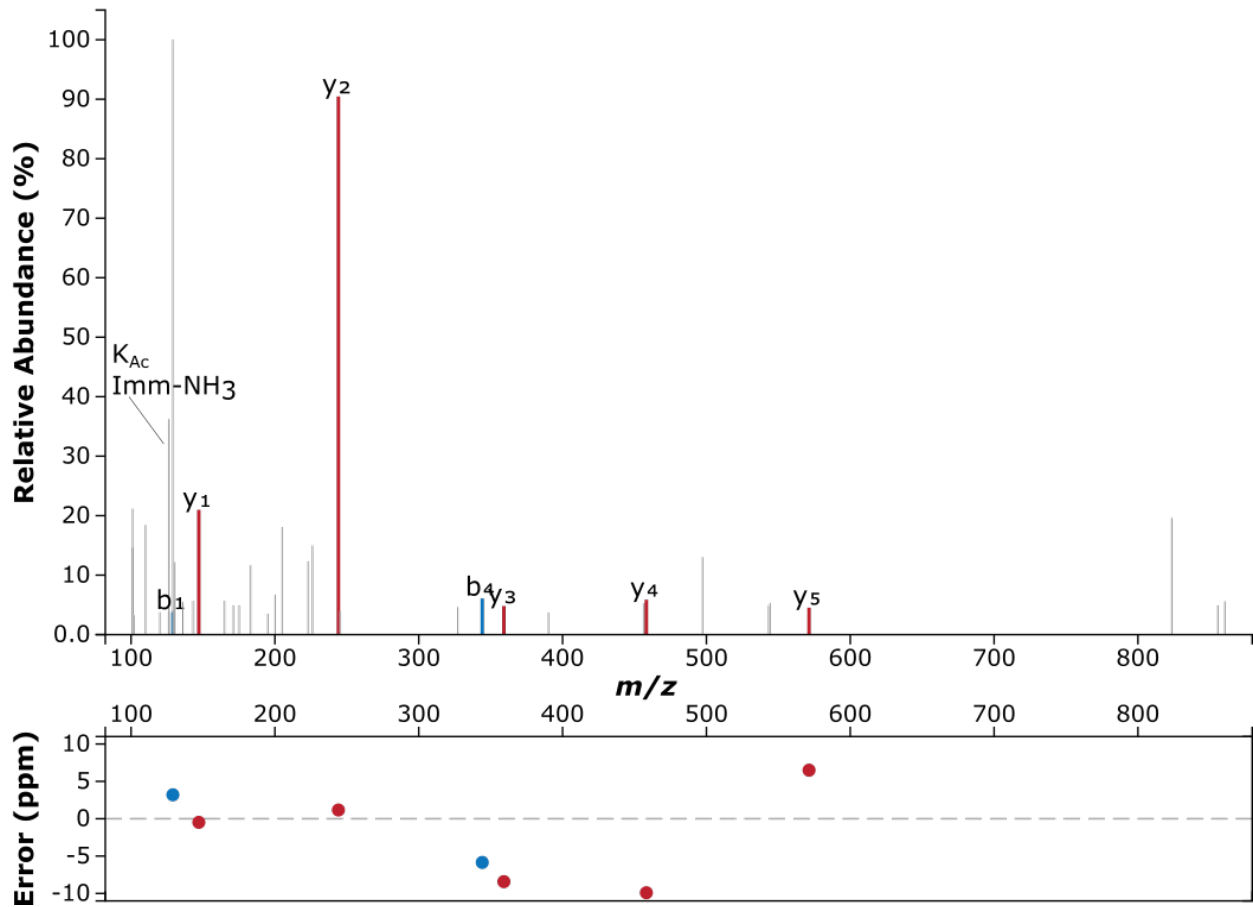




Precursor m/z: 542.8035

Charge: +2

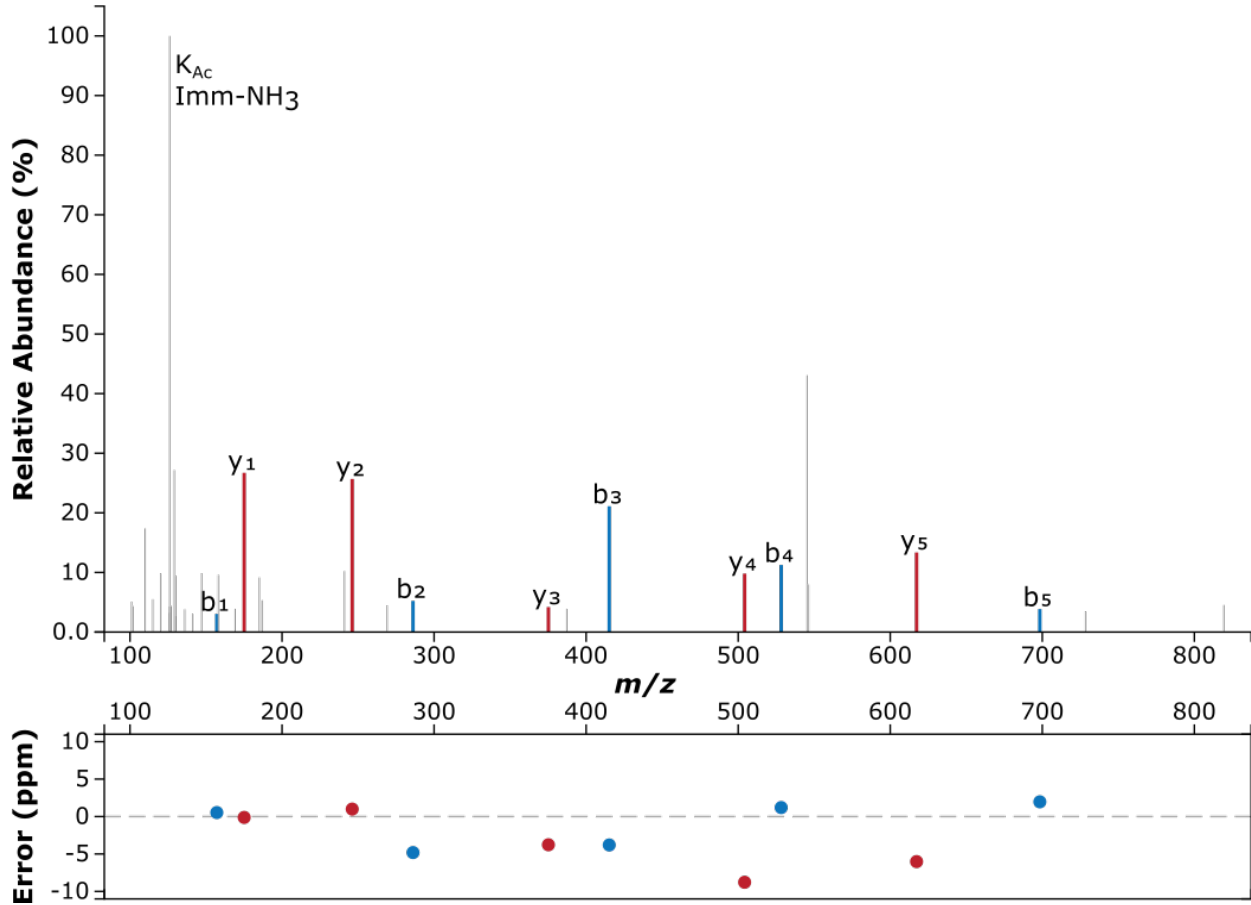
DNA-directed RNA polymerase subunit beta (Q2LQ87)





Precursor m/z: 506.2704 Charge: +3

ATP synthase subunit b 1 (Q2LQZ9)

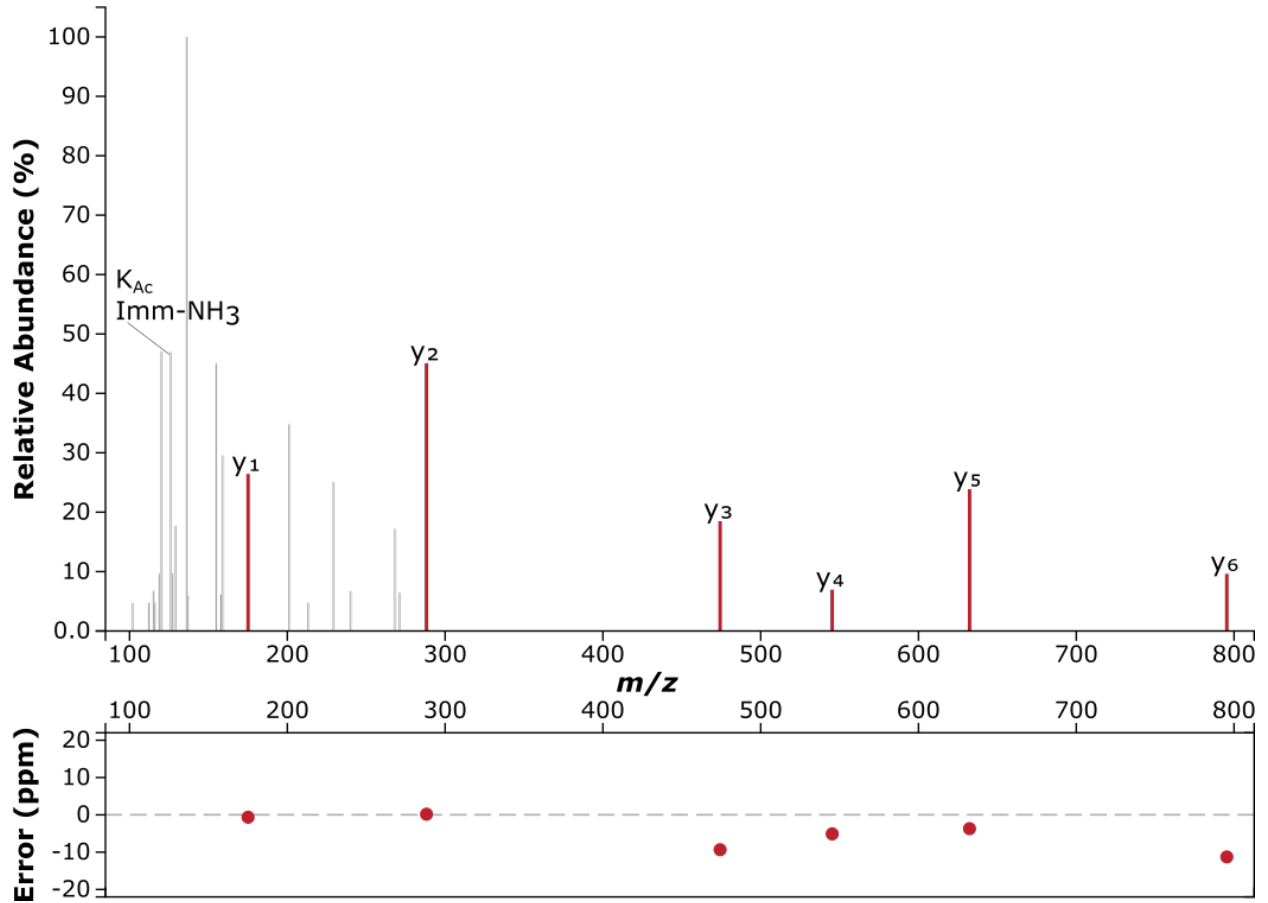


R P L K D P ^{Ac}k G N V V P G L Y S A W I R

Precursor m/z: 577.8286

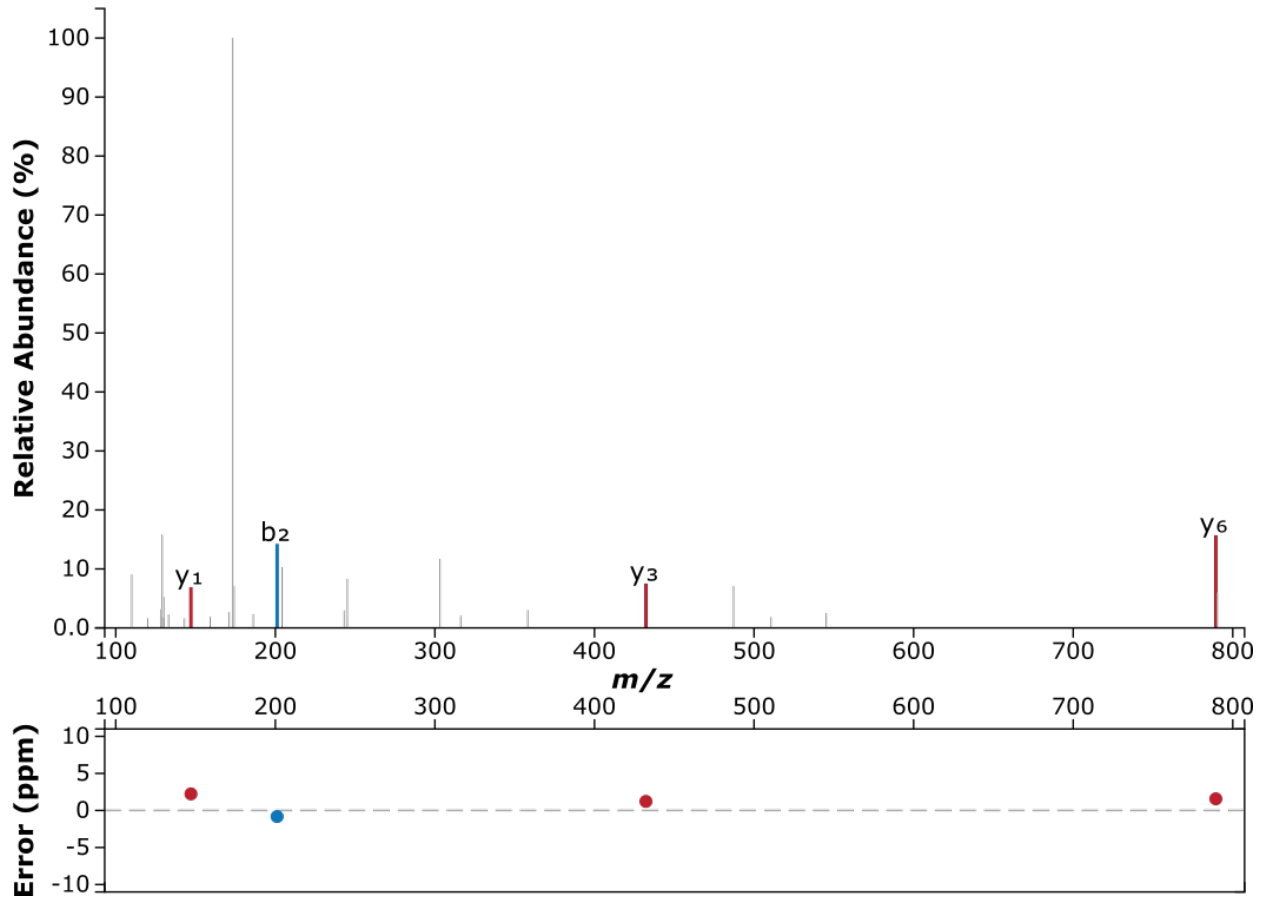
Charge: +4

6-oxocyclohex-1-ene-1-carboxyl-CoA hydrolase (Q2LXU2)



S I E D L D k K

Precursor m/z: 495.2611 Charge: +2
Phosphoglycerate mutase (Q2LTT8)

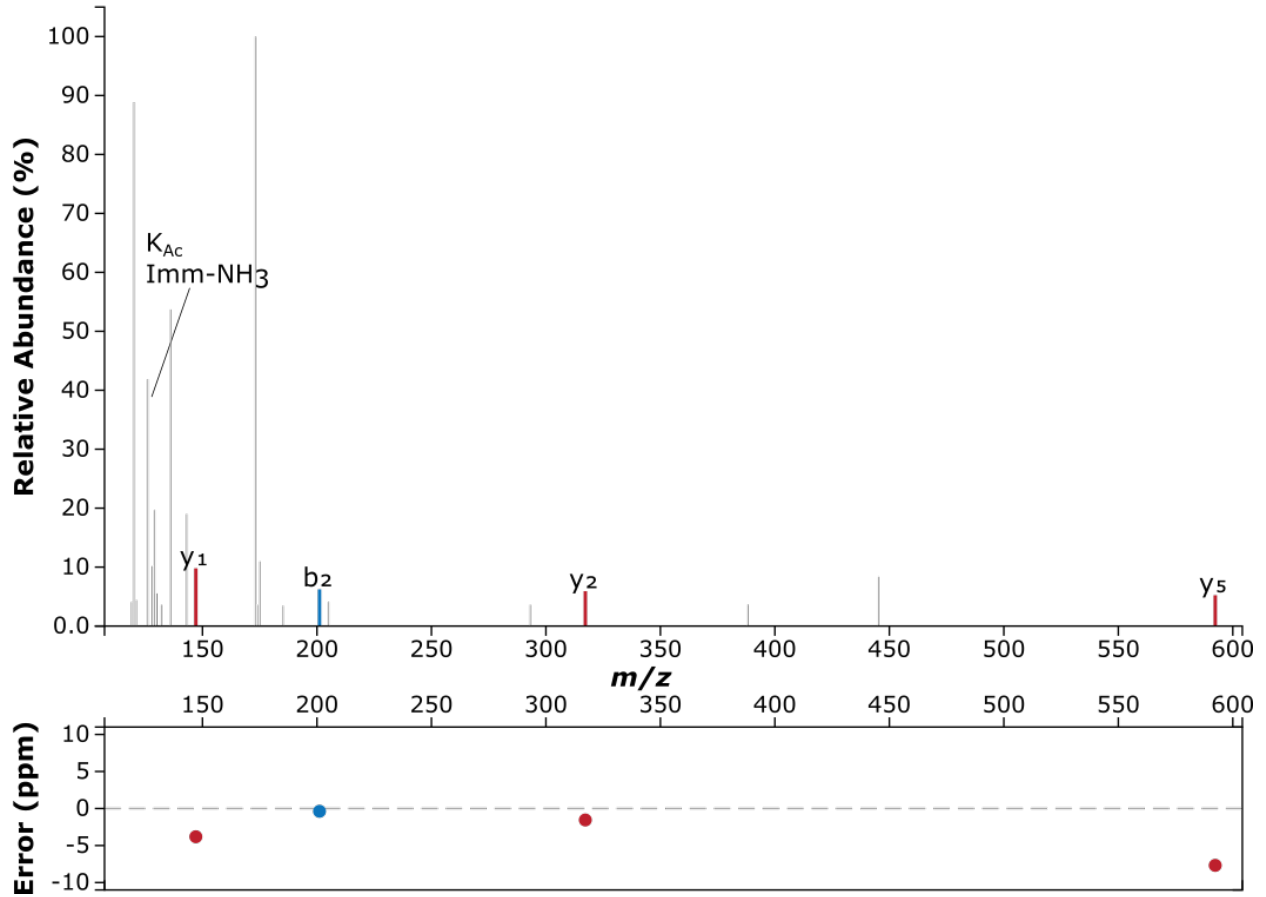




Precursor m/z: 396.7343

Charge: +2

K(+)-insensitive pyrophosphate-energized proton pump (Q2LUK4)

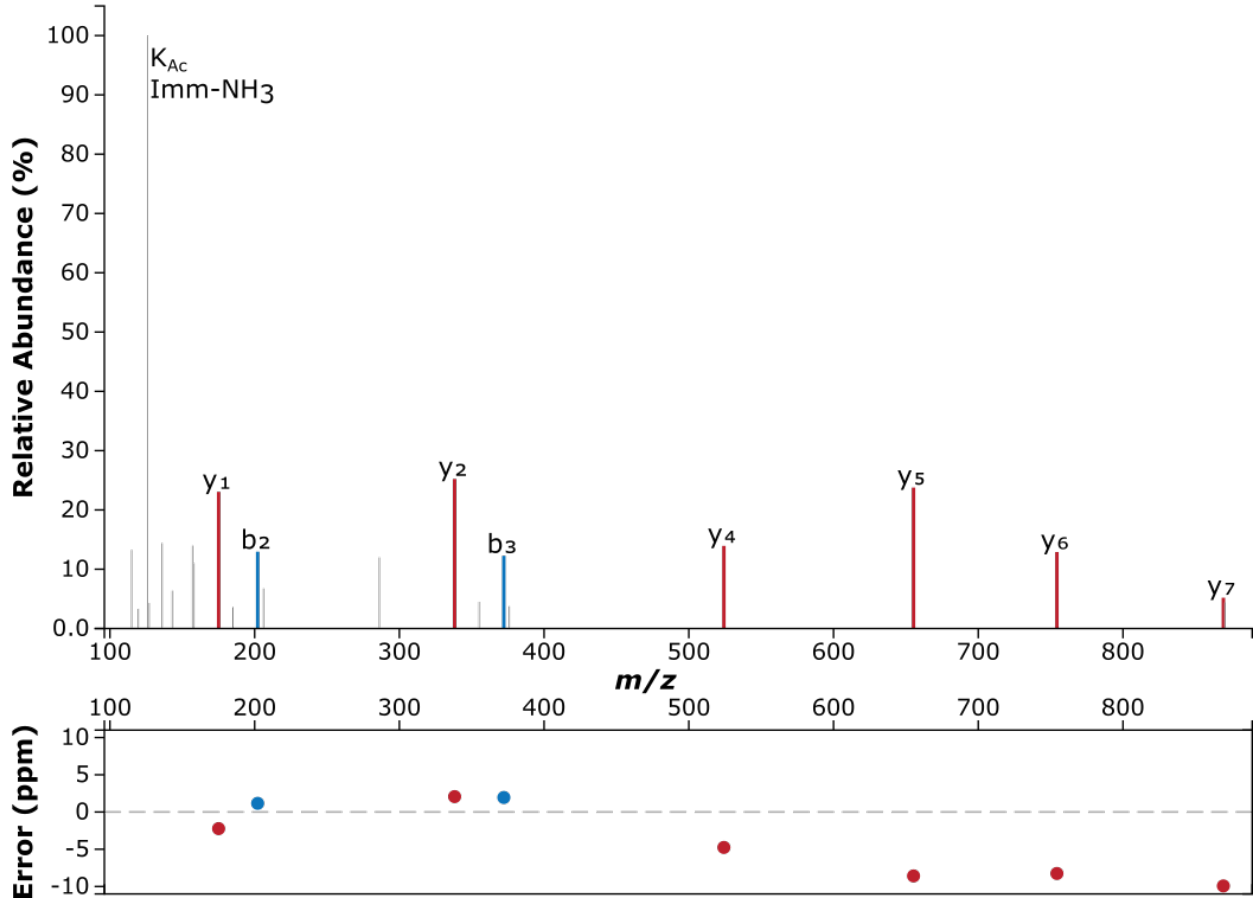


S N ^{AC} k D V M G E Y R

Precursor m/z: 620.7850

Charge: +2

6-hydroxycyclohex-1-ene-1-carboxyl-CoA dehydrogenase (Q2LXU3)

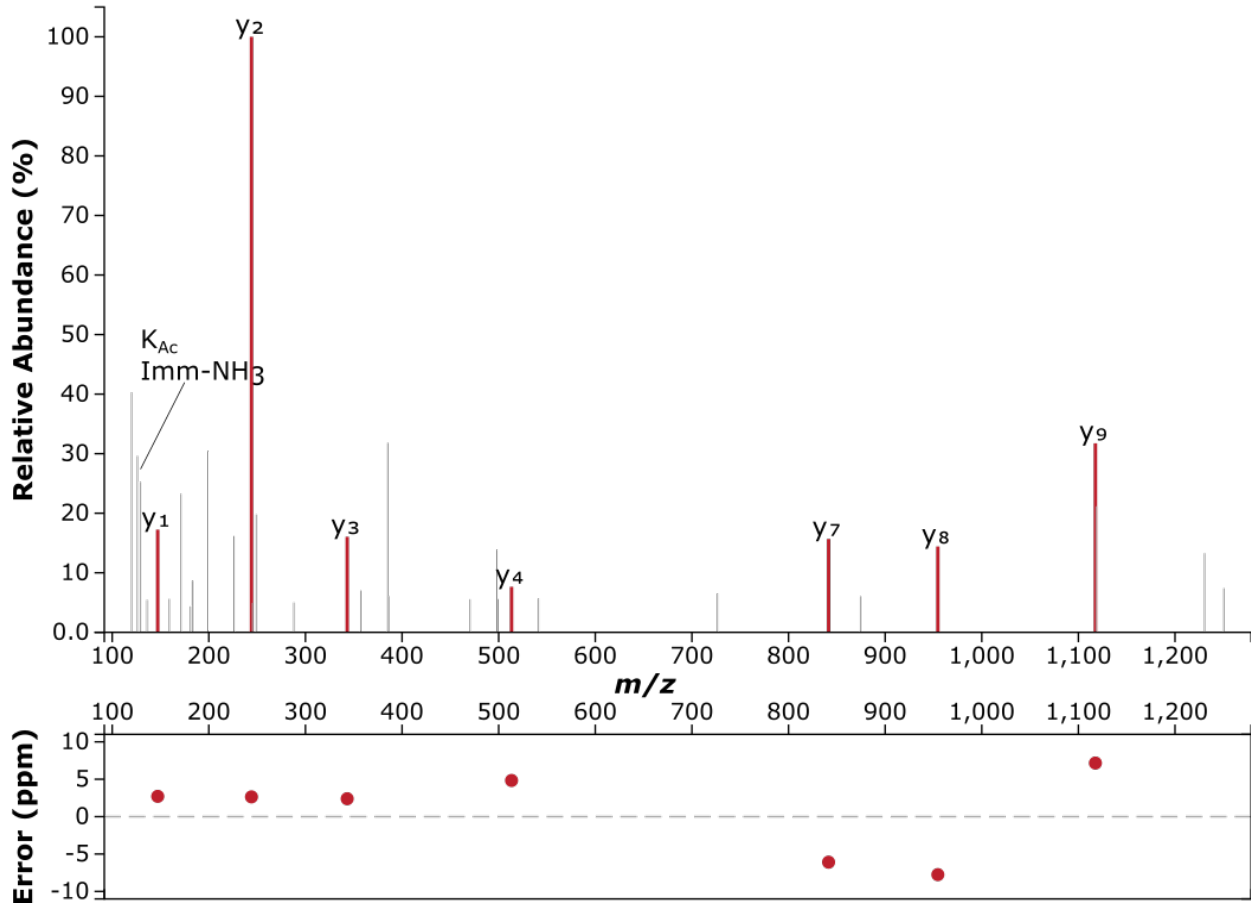


S S P T G E I Y L D N V ^{Ac}k V P K

Precursor m/z: 894.9725

Charge: +2

Acyl-CoA dehydrogenase, short-chain specific (Q2LUY8)

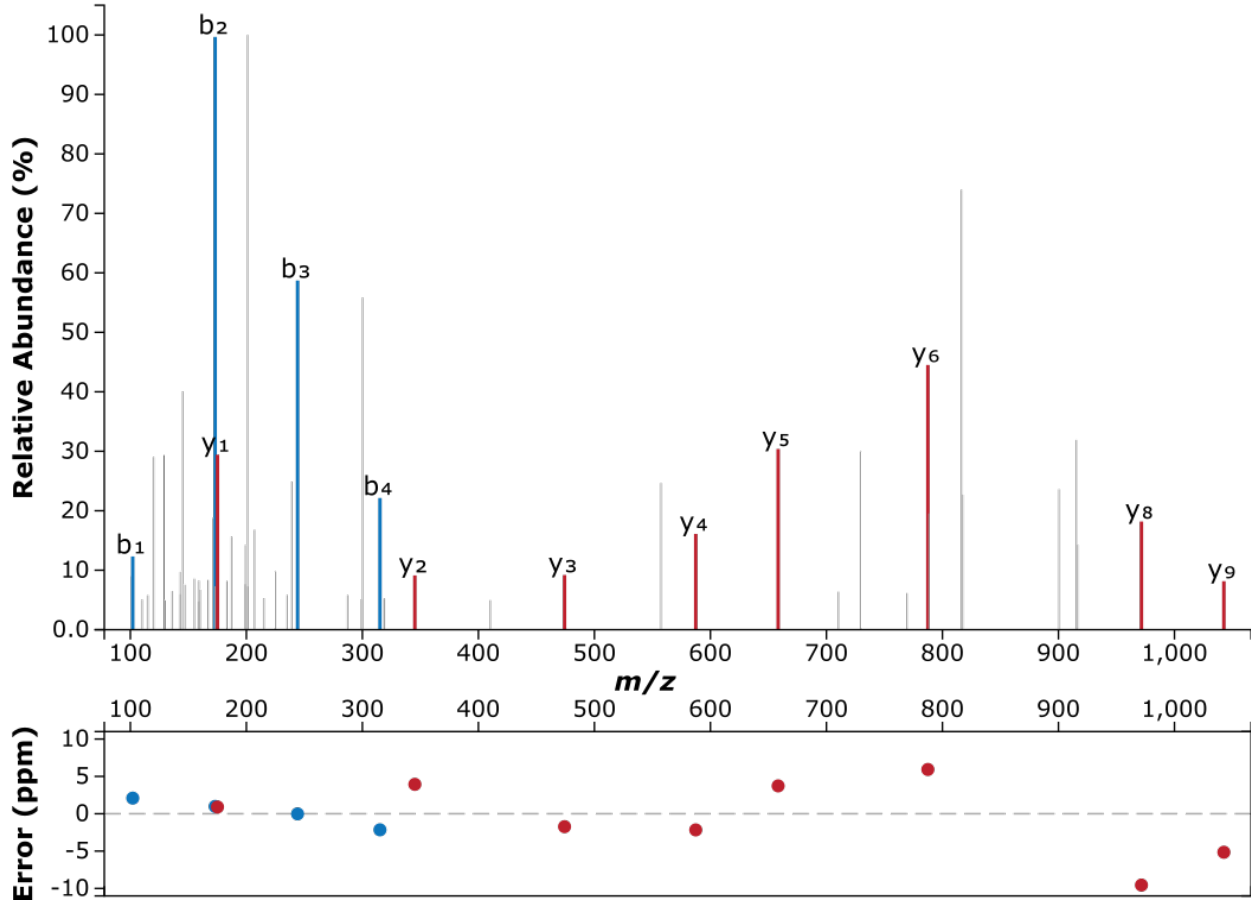




Precursor m/z: 607.8406

Charge: +2

Polyribonucleotide nucleotidyltransferase (Q2LWT4)

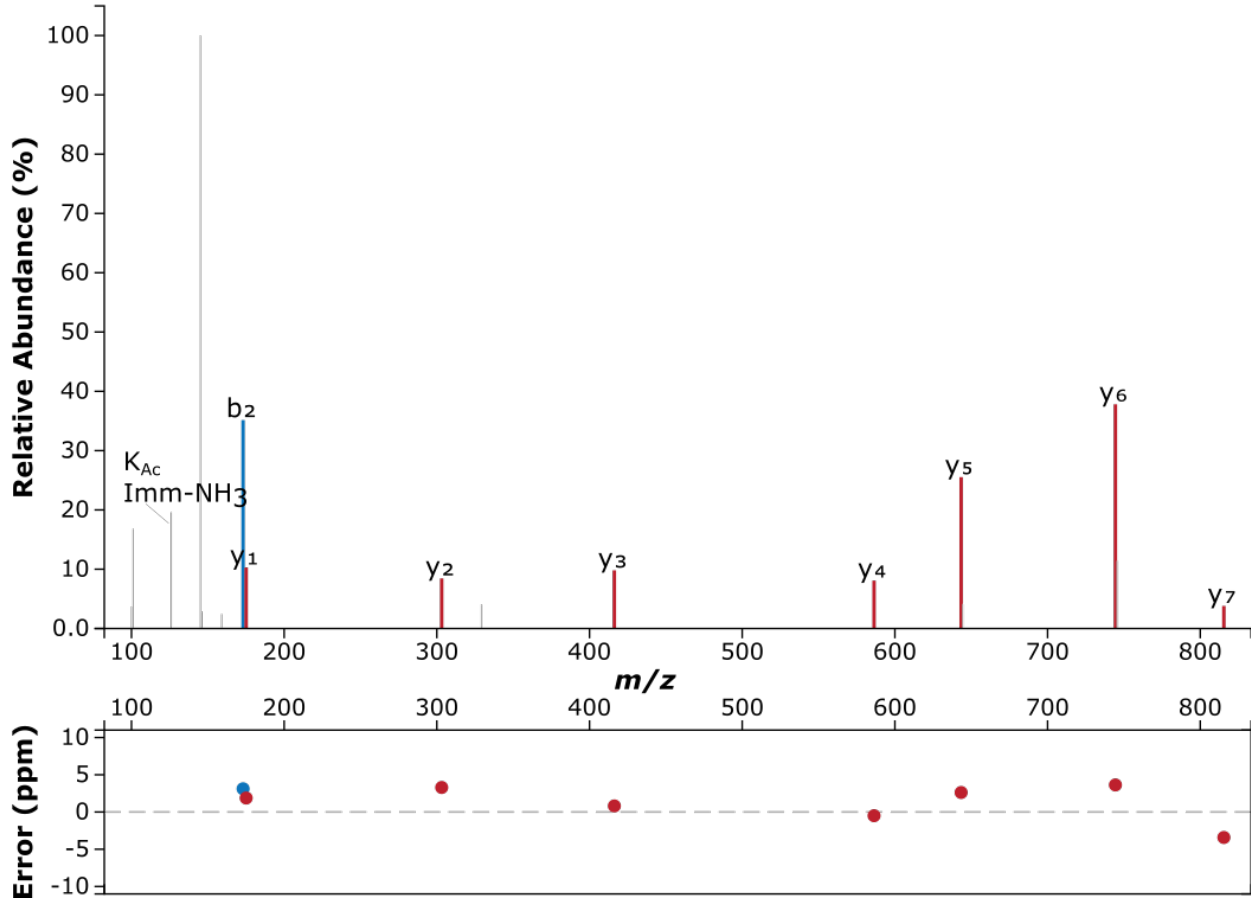




Precursor m/z: 458.7642

Charge: +2

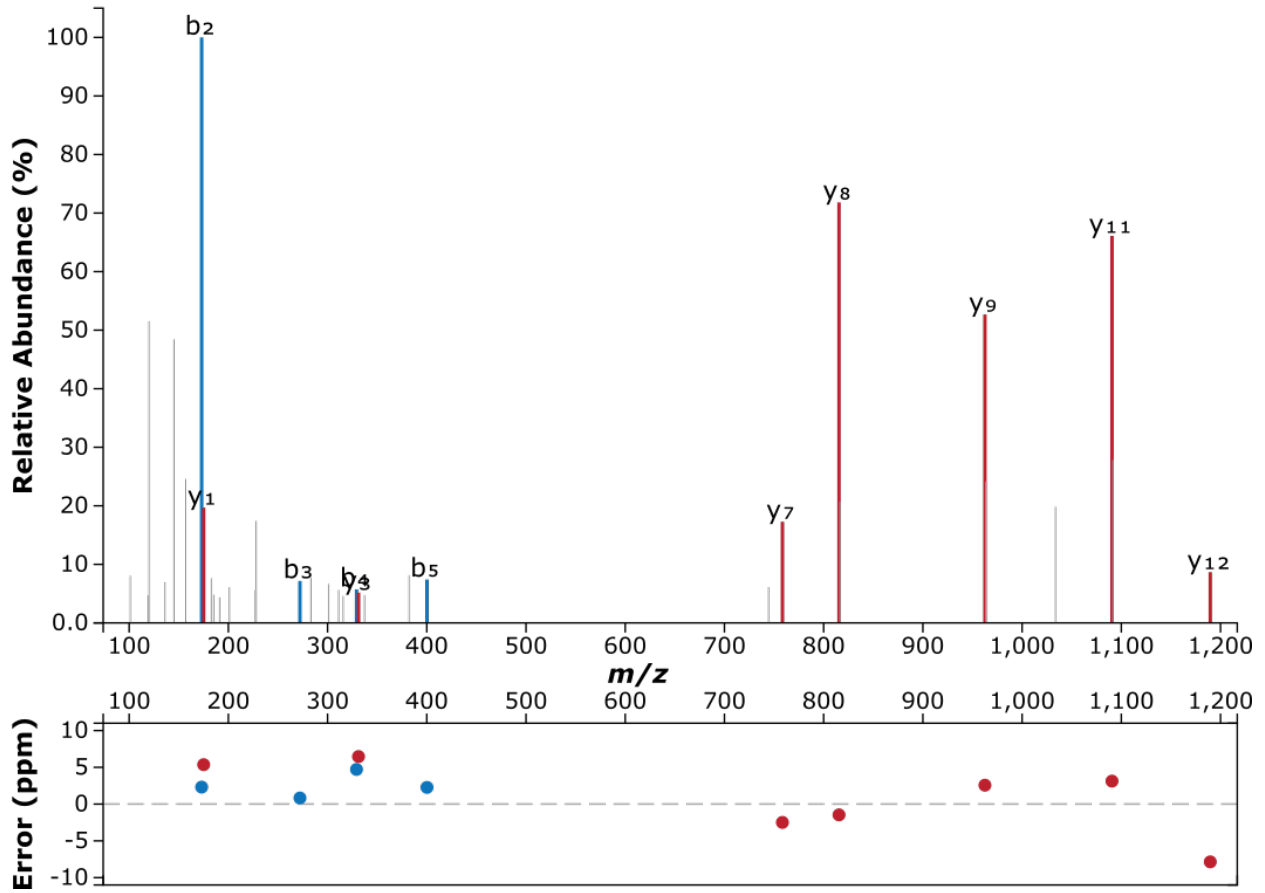
4-hydroxybenzoate-CoA ligase / benzoate-CoA ligase (Q2LRH0, Q2LRH7)





Precursor m/z: 681.3804 Charge: +2

Acetyl-CoA acetyltransferase (Q2LWQ5)

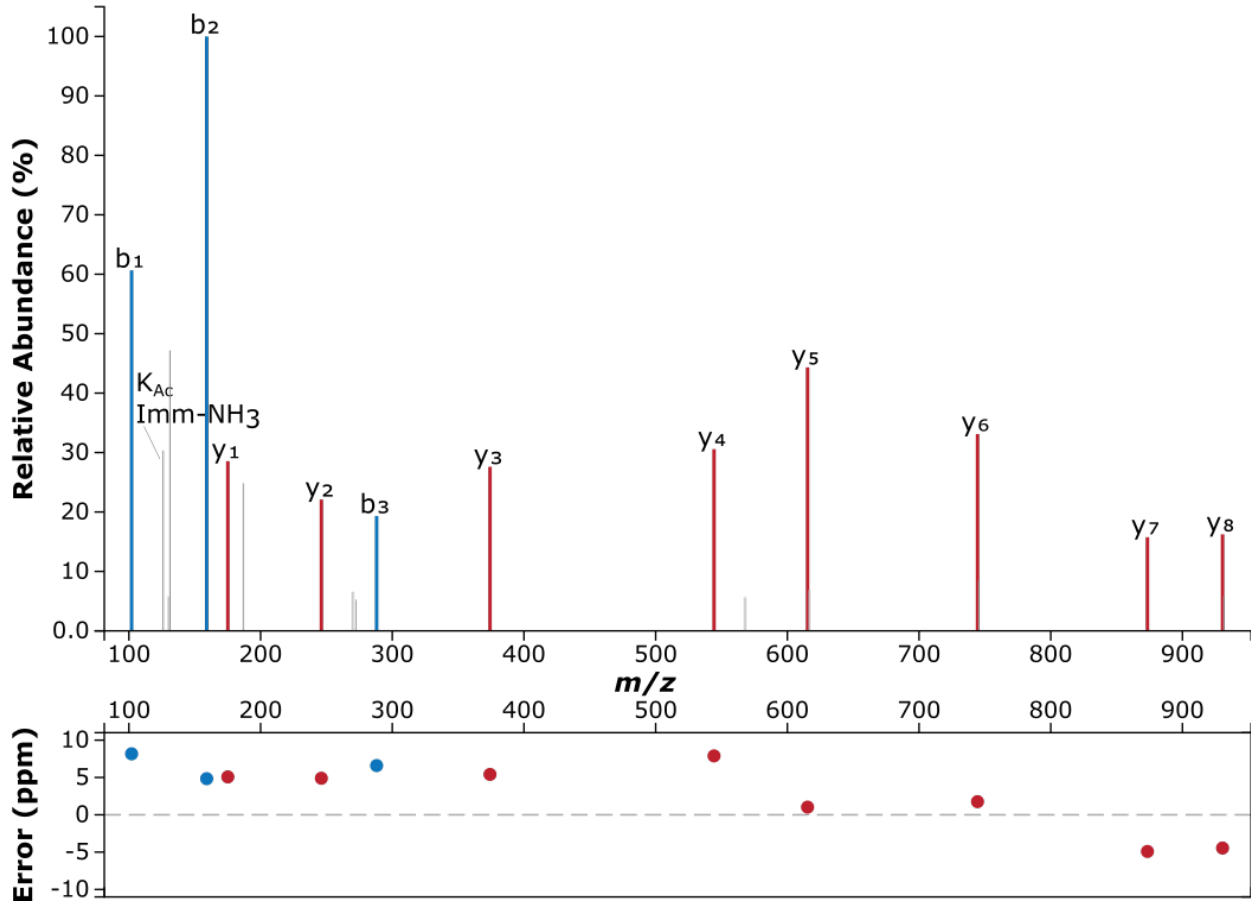




Precursor m/z: 516.2594

Charge: +2

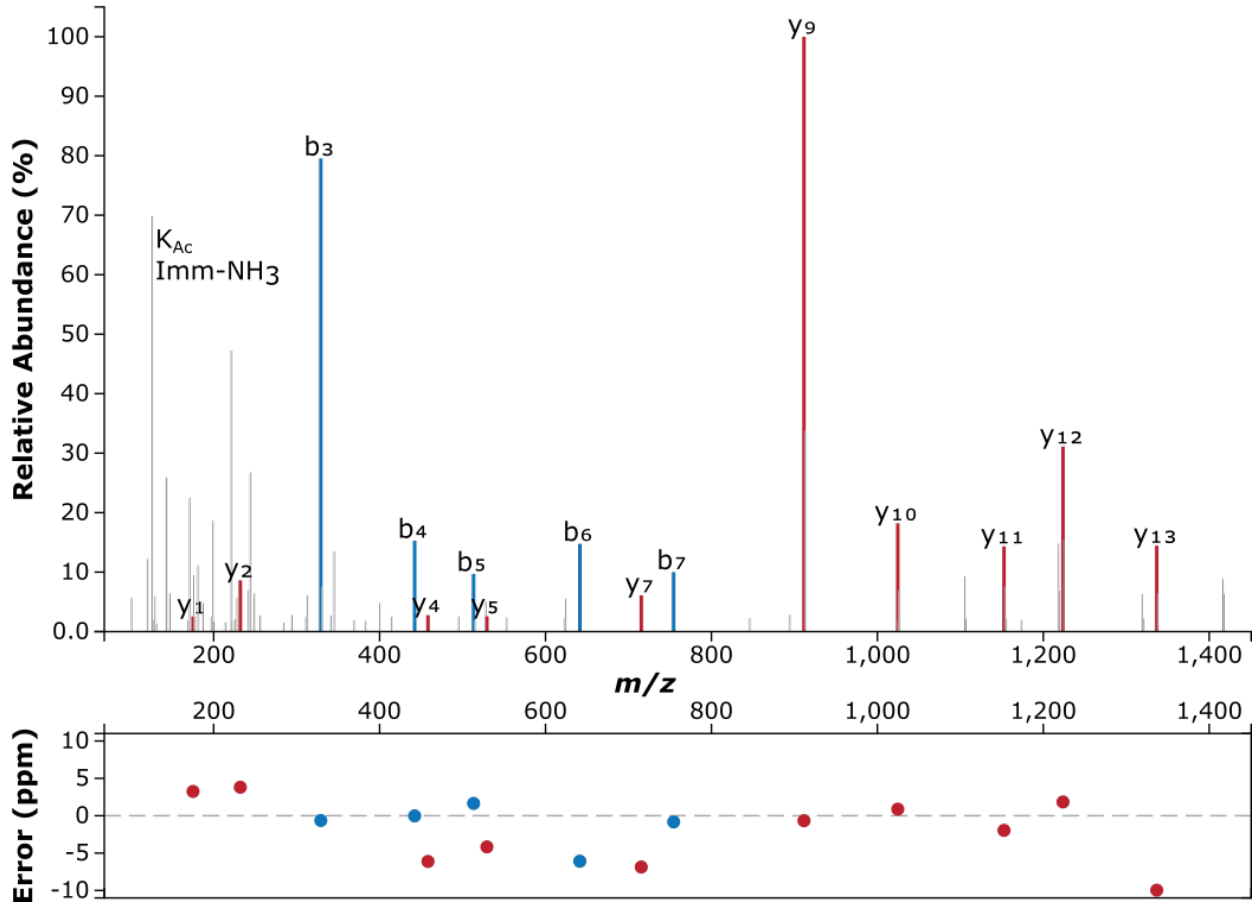
6-oxocyclohex-1-ene-1-carboxyl-CoA hydrolase (Q2LXU2)



T G k^{AC} I A Q I P V G E A L L G R

Precursor m/z: 832.9883 Charge: +2

ATP synthase subunit alpha 1 (Q2LQZ7)

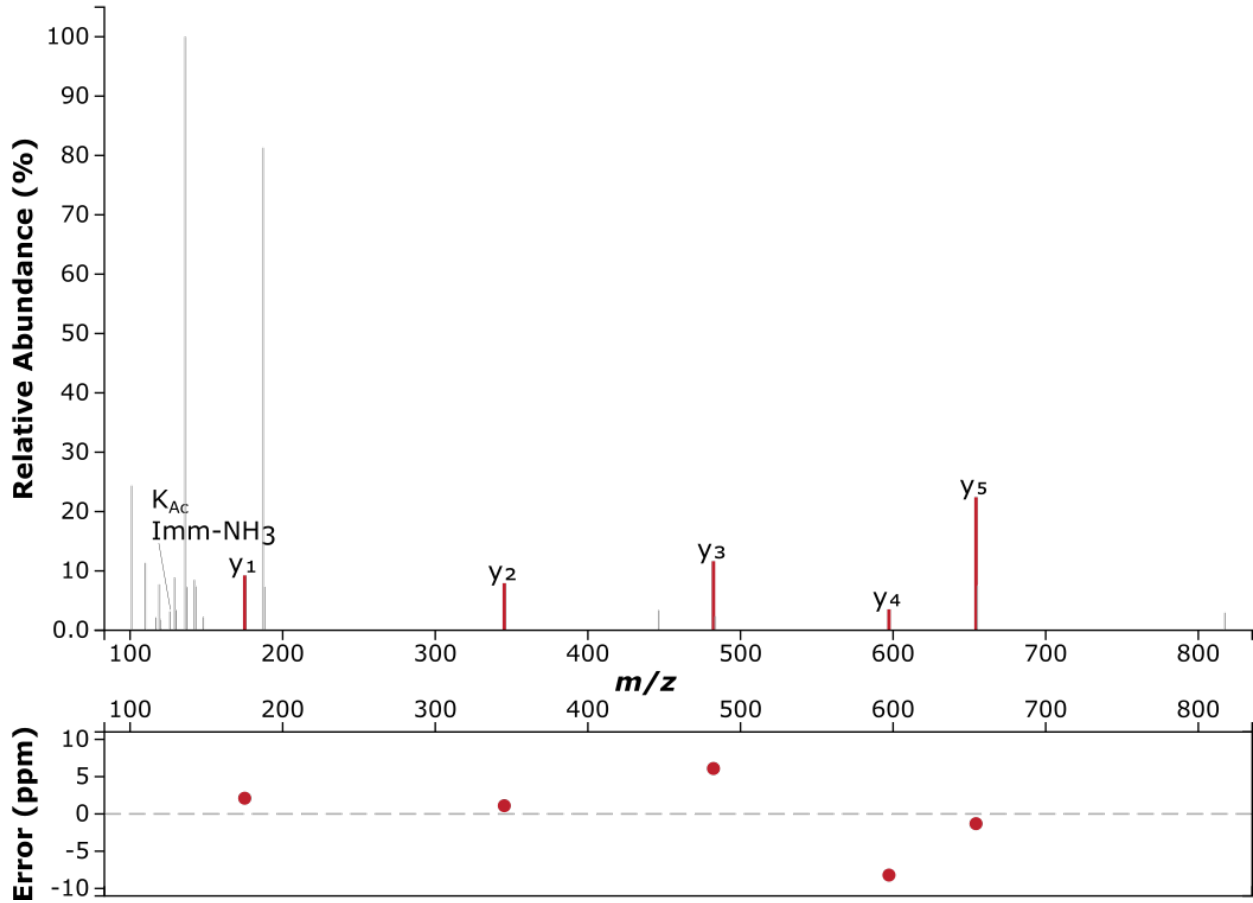


T I Y G D H k R

Precursor m/z: 344.5138

Charge: +3

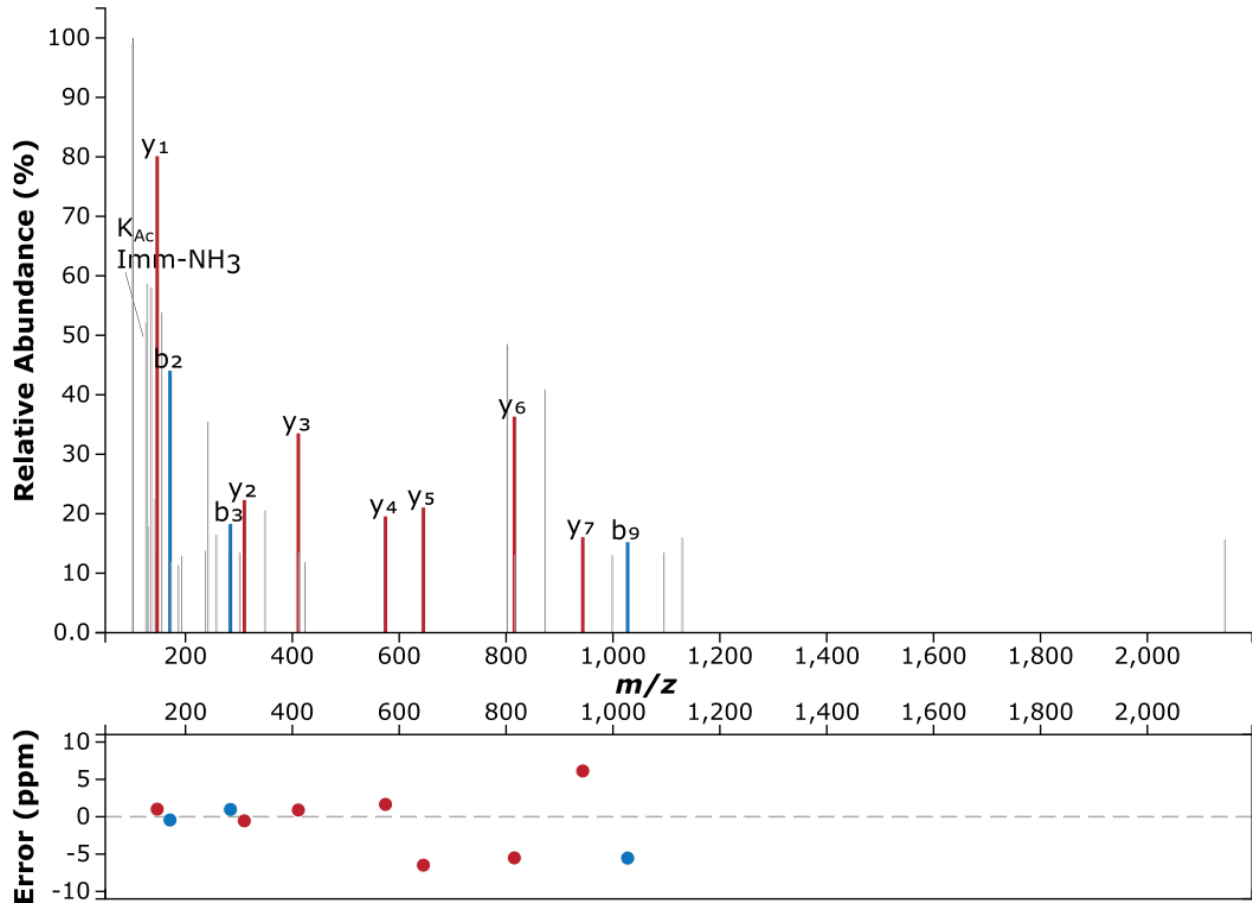
Acetyl-CoA synthetase (Q2LWR8)



V A I Q W E G N E P G E Q ^{Ac} k A Y T Y K

Precursor m/z: 751.6989 Charge: +3

Acetyl-CoA synthetase (Q2LWR8)

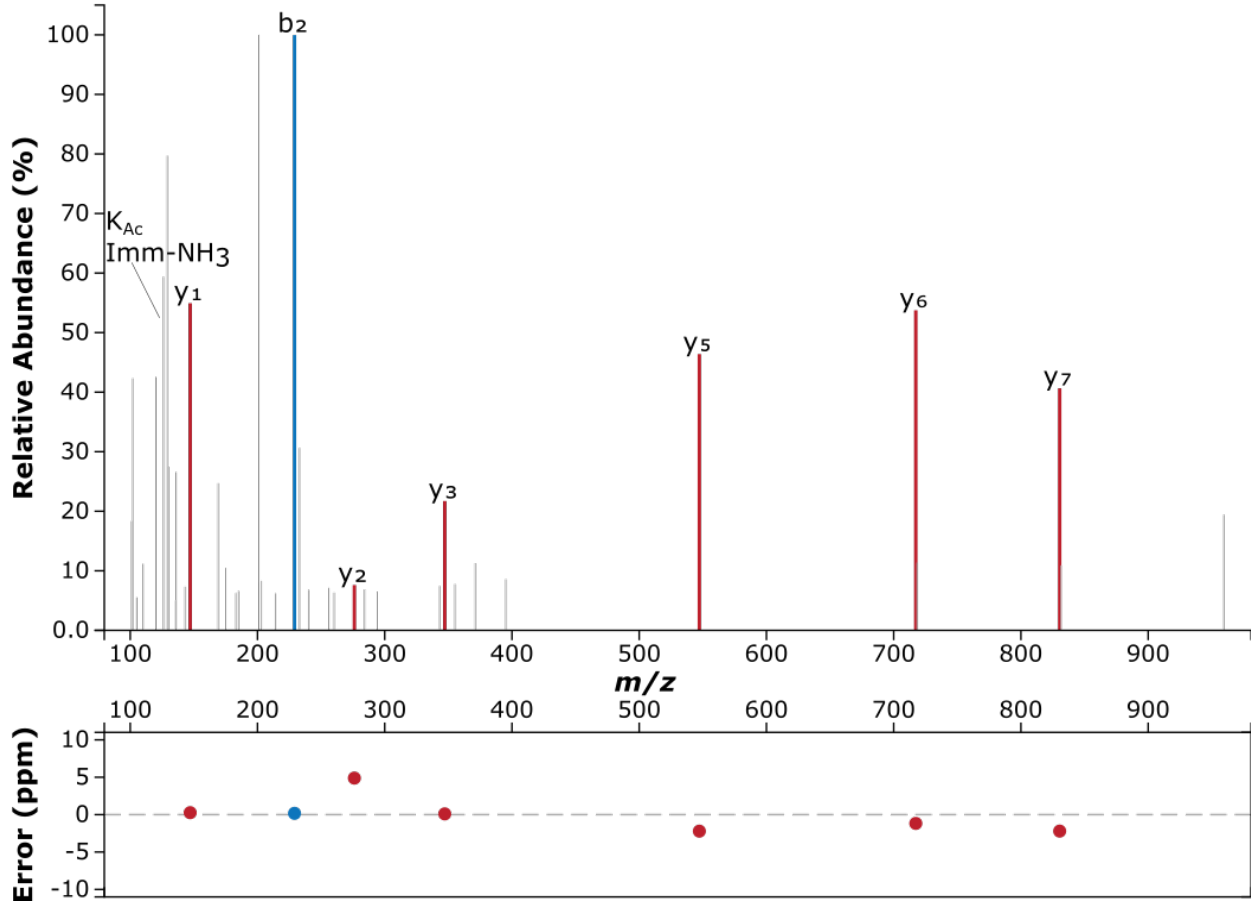




Precursor m/z: 529.7900

Charge: +2

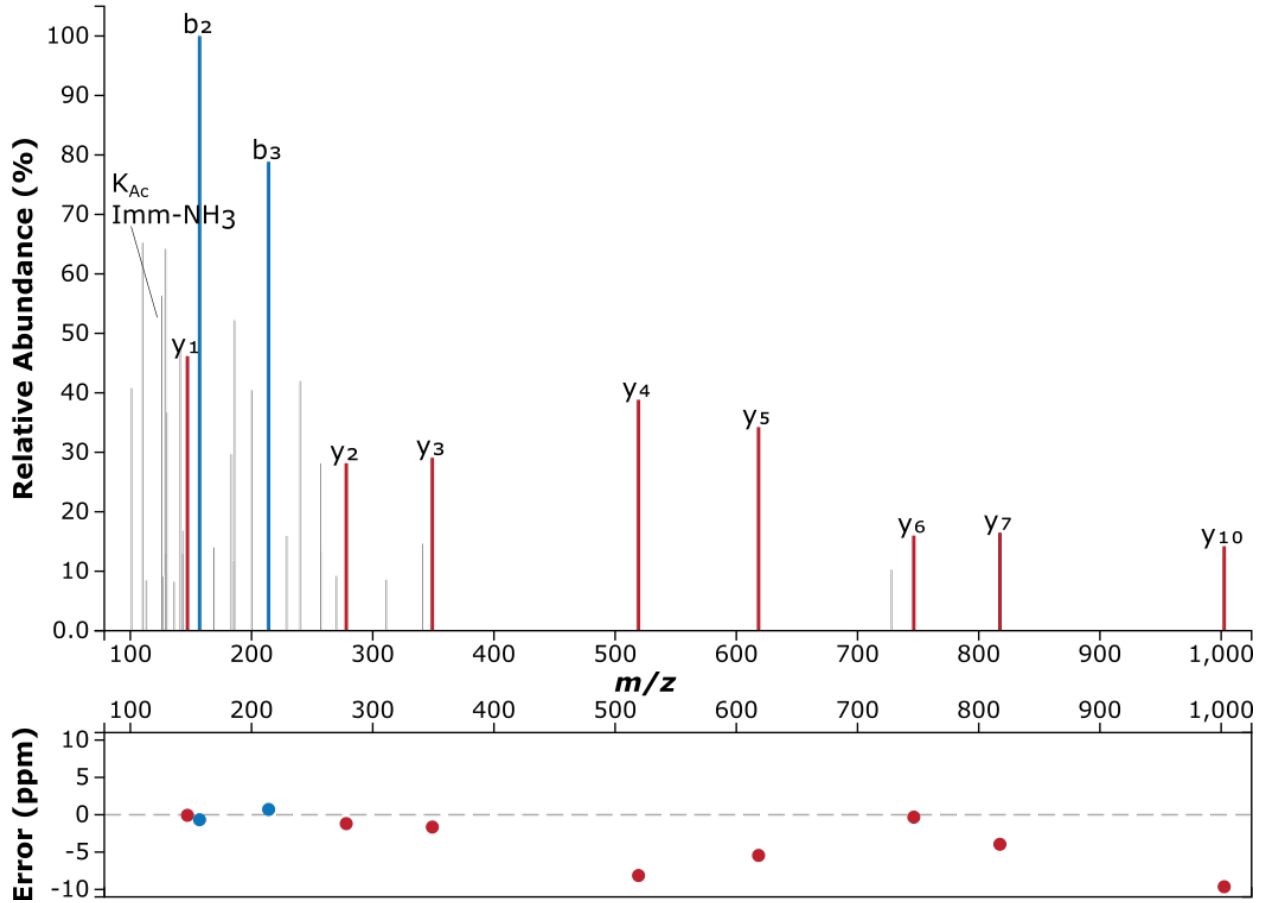
3-oxoacyl-reductase (Q2LXS6)





Precursor m/z: 551.3079 Charge: +2

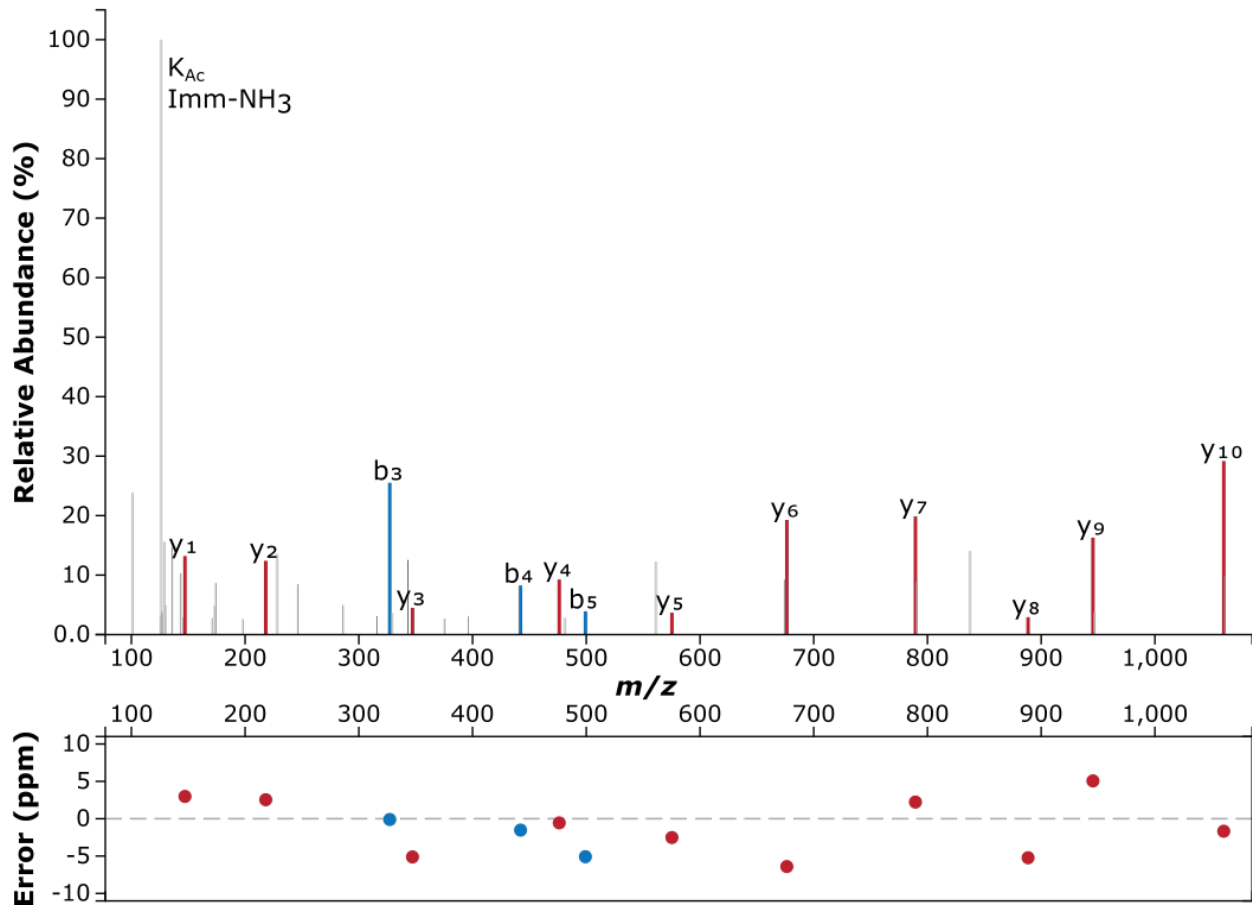
ATP synthase subunit alpha 1 (Q2LQZ7)





Precursor m/z: 693.8774 Charge: +2

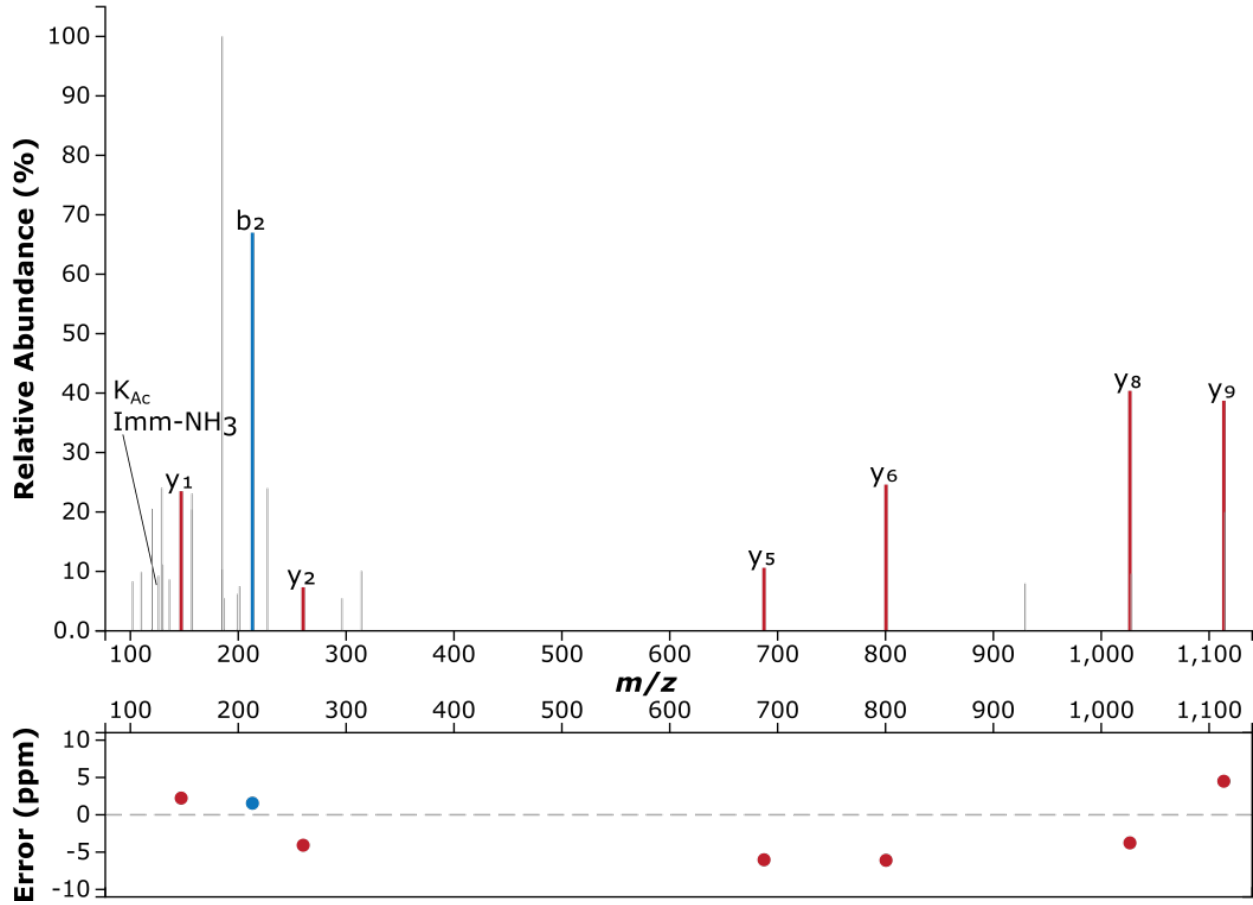
60 kDa chaperonin 3 (Q2LU42)



V I S P E L E Q k I K

Precursor m/z: 663.3874 Charge: +2

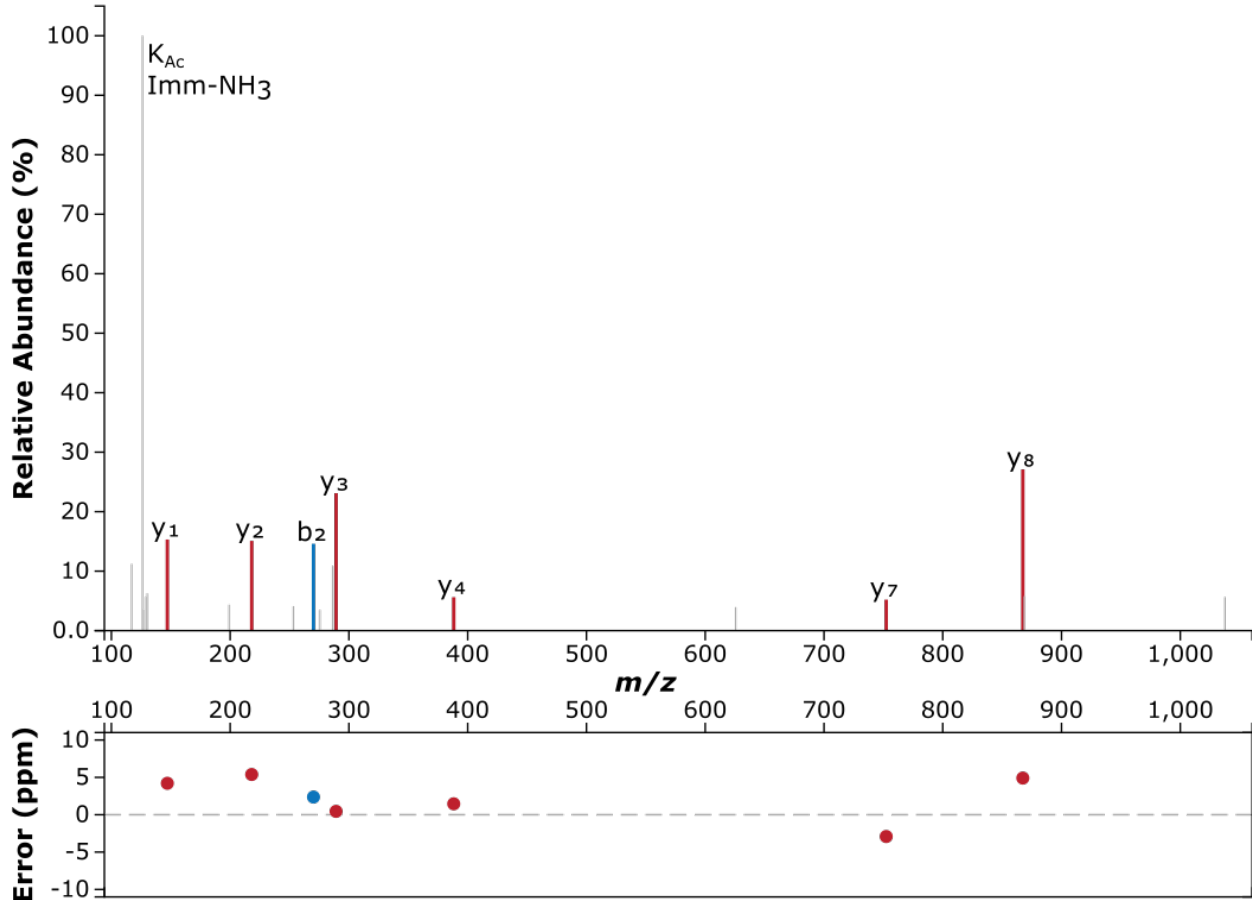
ATP synthase subunit alpha 1 (Q2LQZ7)





Precursor m/z: 568.8009 Charge: +2

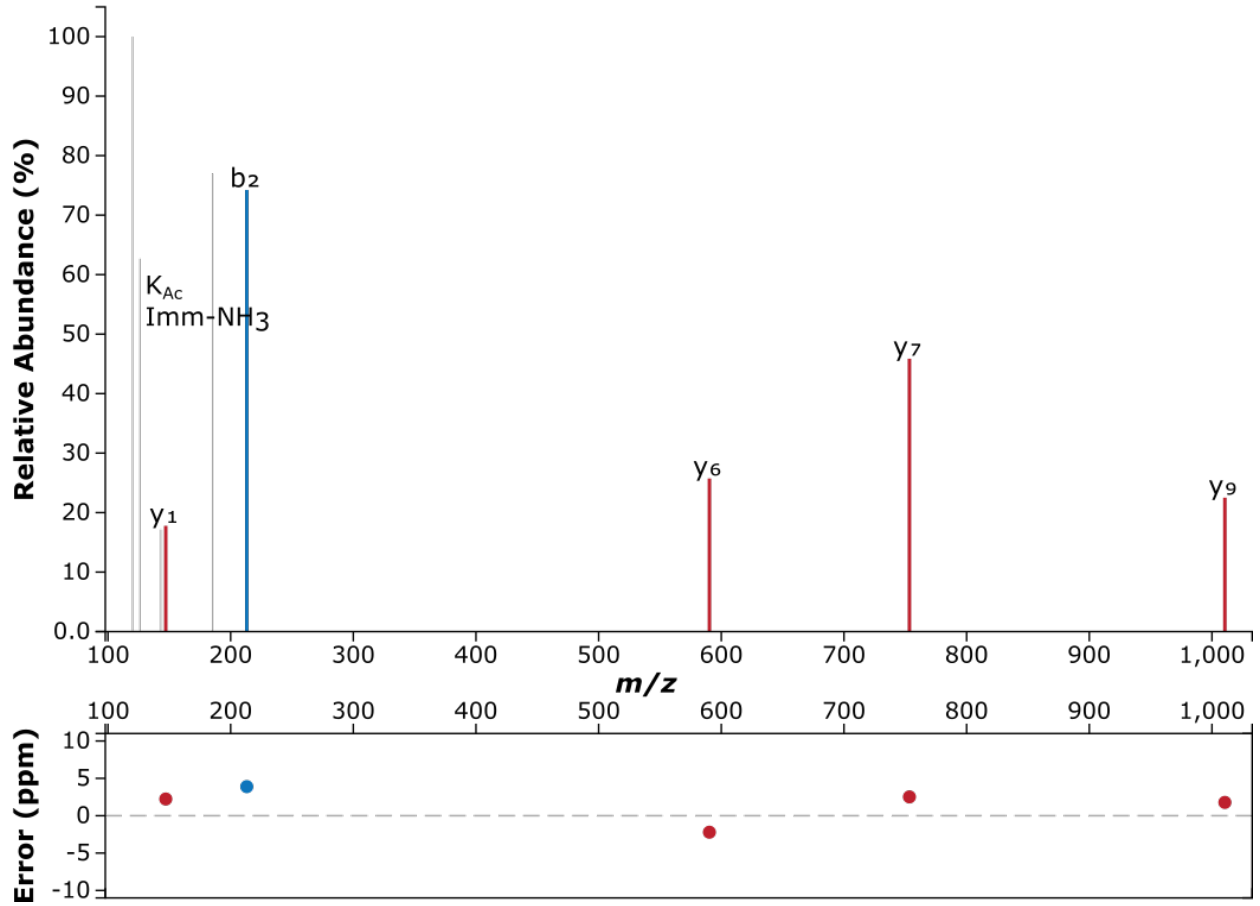
Hypothetical exported protein (Q2LWR5)





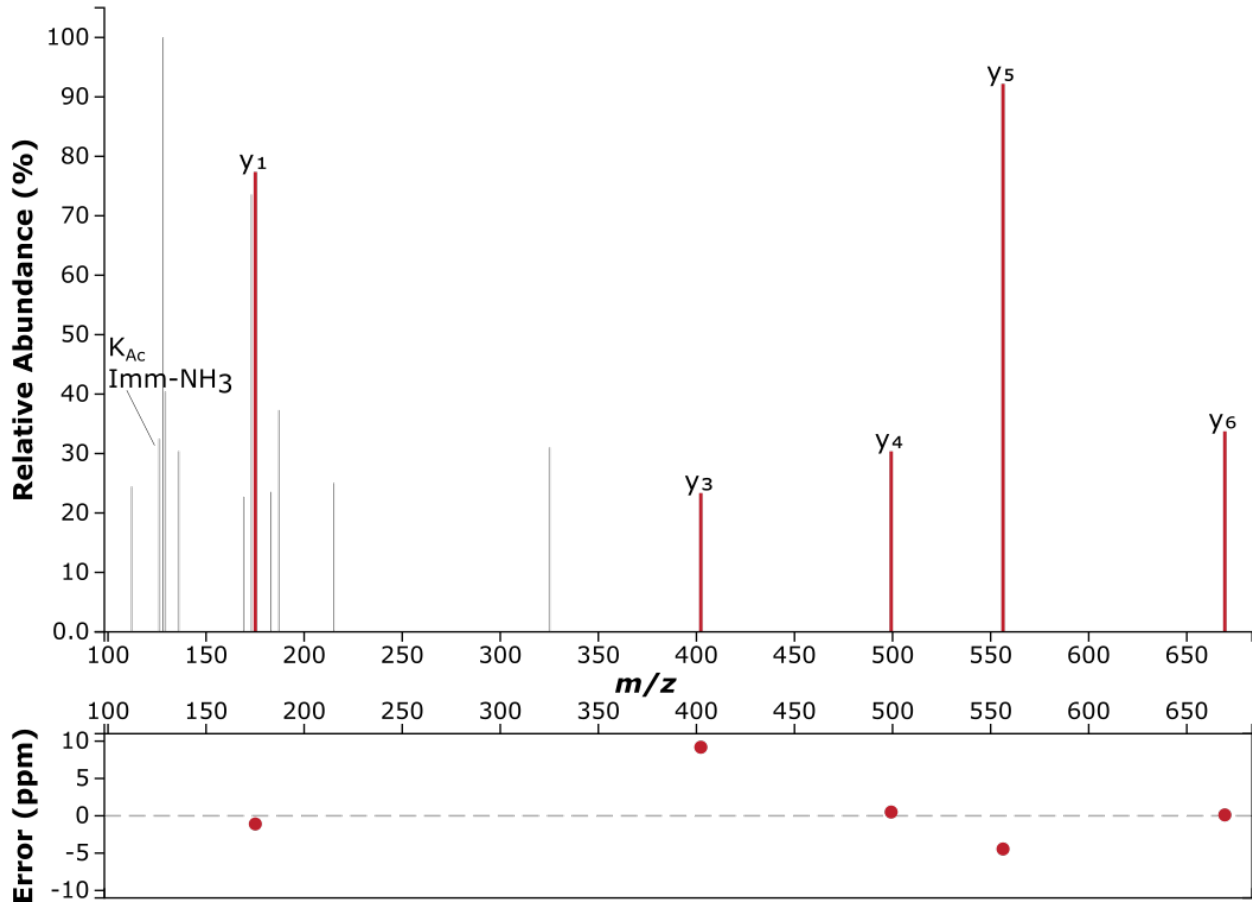
Precursor m/z: 685.3717
 10 kDa chaperonin (Q2LTG3)

Charge: +2





Precursor m/z: 435.2638 Charge: +2
60 kDa chaperonin 3/2 (Q2LU42,Q2LTG7)

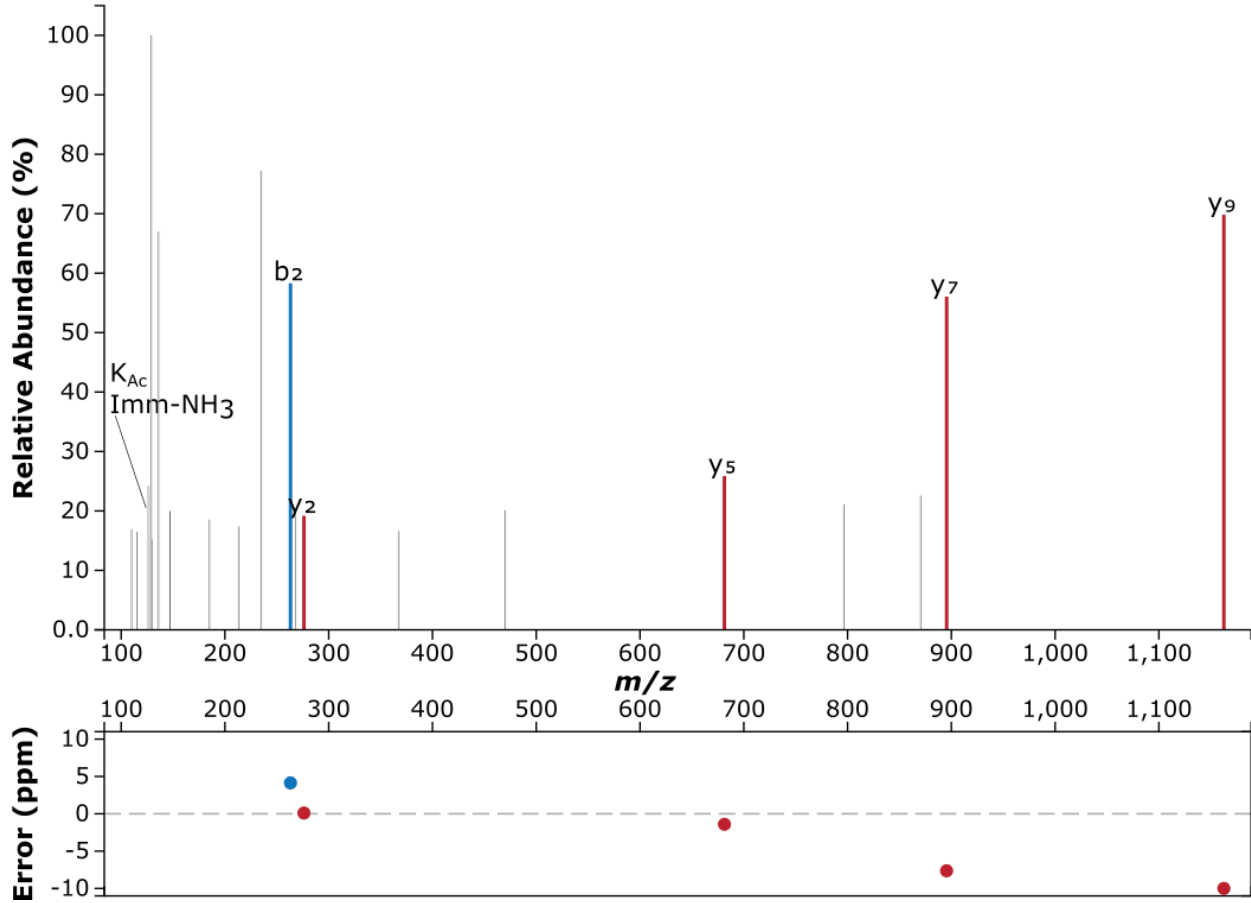




Precursor m/z: 763.3985

Charge: +2

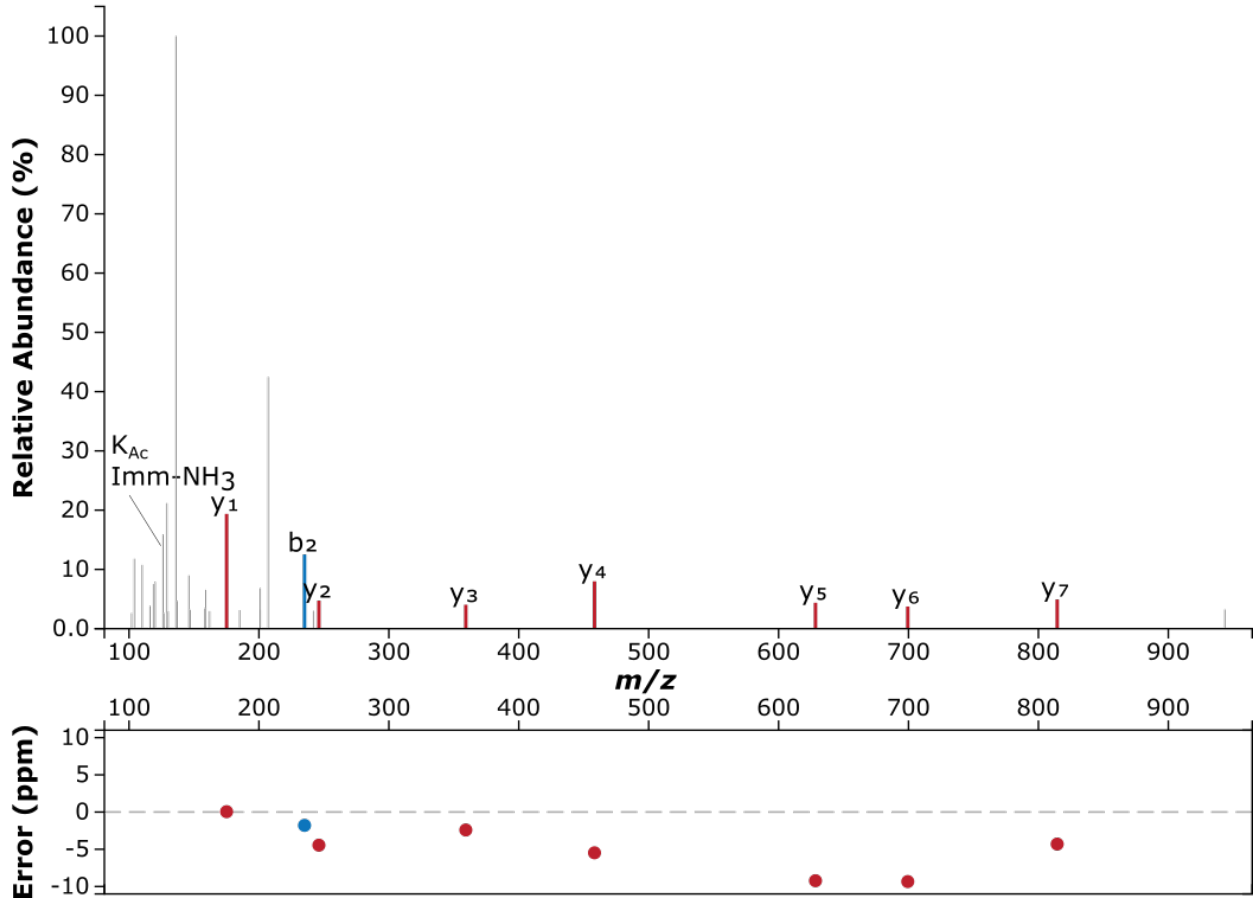
Molecular chaperone (Q2LPJ5)



Y A E D A ^{Ac} k V I A R

Precursor m/z: 589.3142 Charge: +2

Adenylosuccinate synthetase (Q2LSI8)

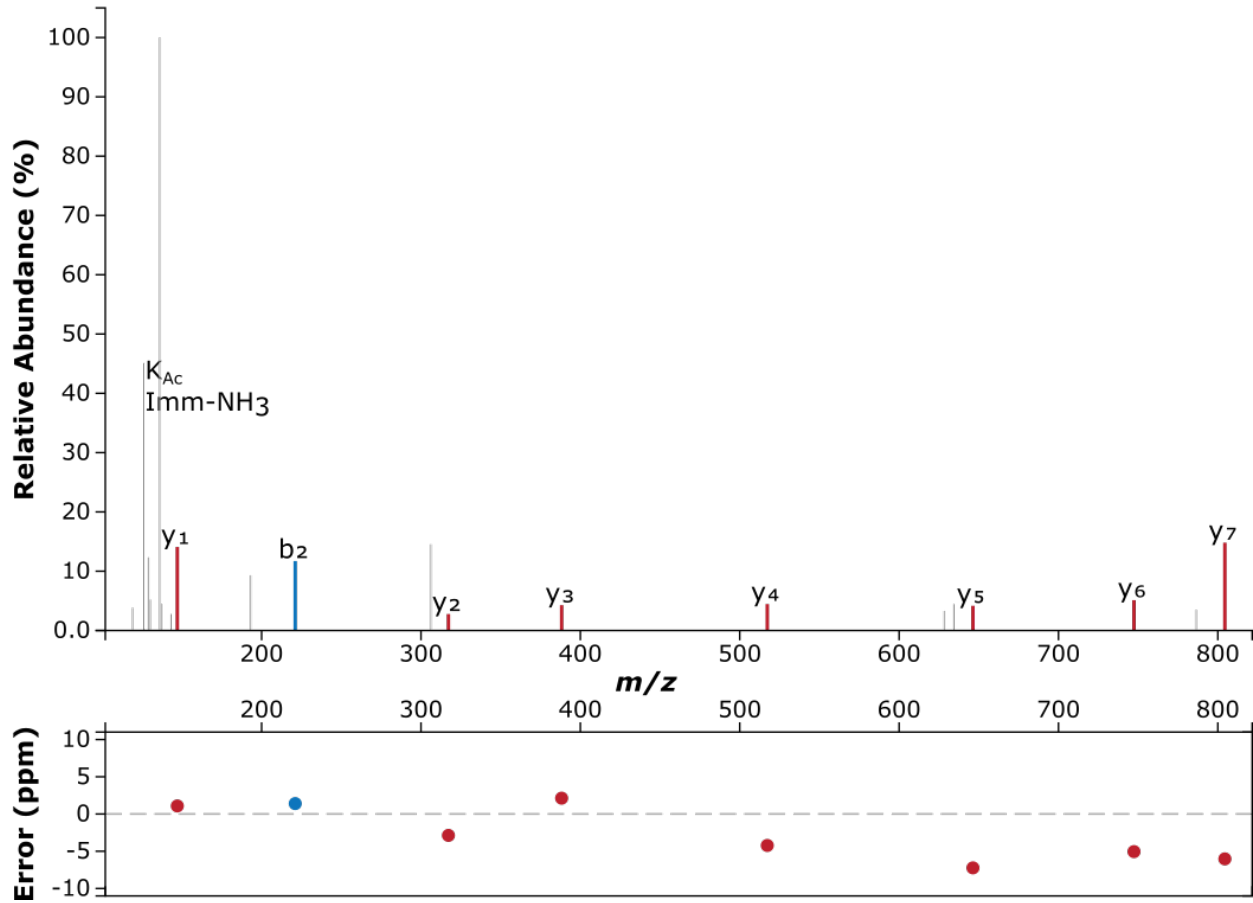




Precursor m/z: 484.2402

Charge: +2

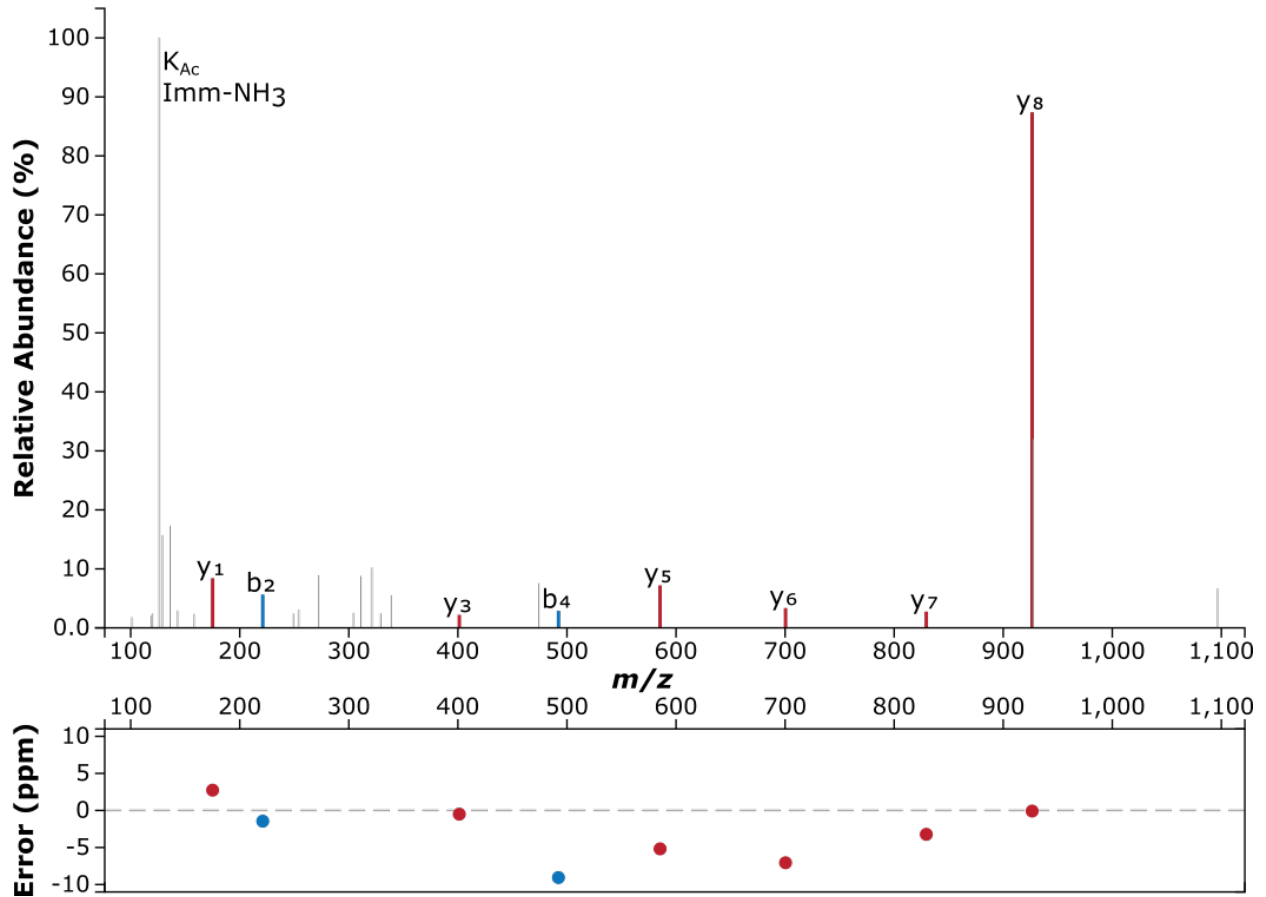
Acyl-CoA dehydrogenase, short-chain specific (Q2LUY8)



Y G T k P E D L A L I R

Precursor m/z: 709.3879 Charge: +2

Acetyl-CoA acetyltransferase (Q2LXR7)

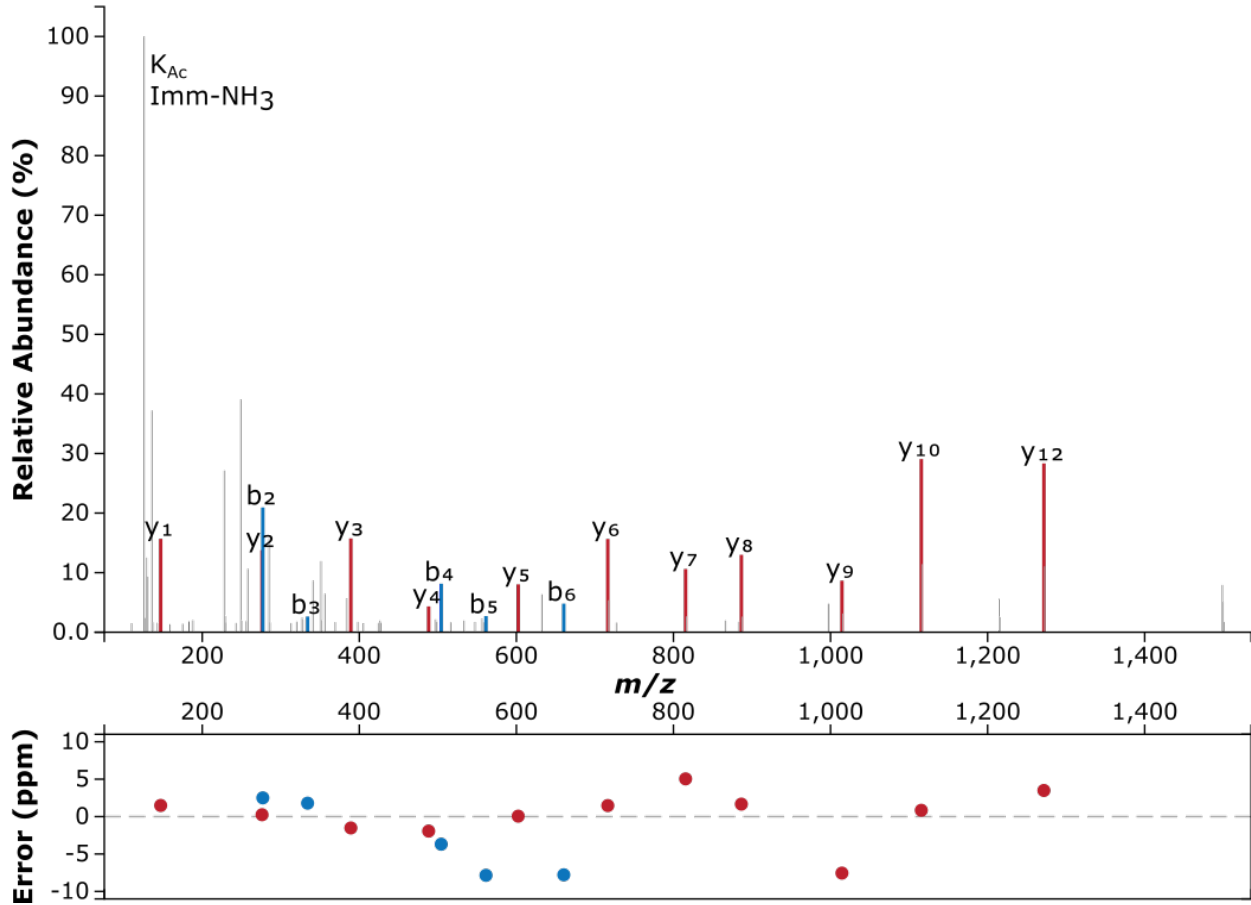




Precursor m/z: 887.9885

Charge: +2

Enolase (Q2LR33)



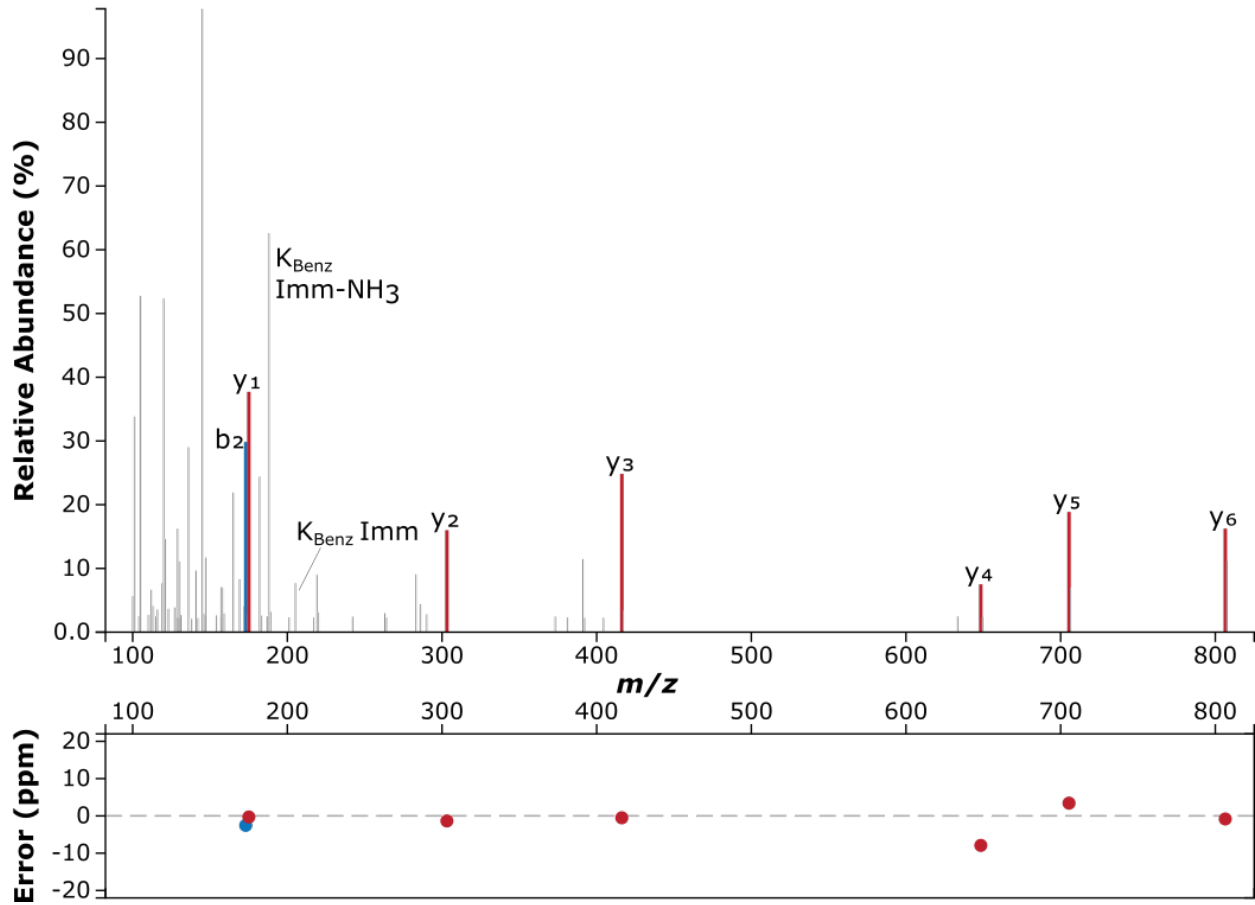
Benzoyl



Precursor m/z: 489.7720

Charge: +2

4-hydroxybenzoate-CoA ligase / benzoate-CoA ligase (Q2LRH0, Q2LRH7)

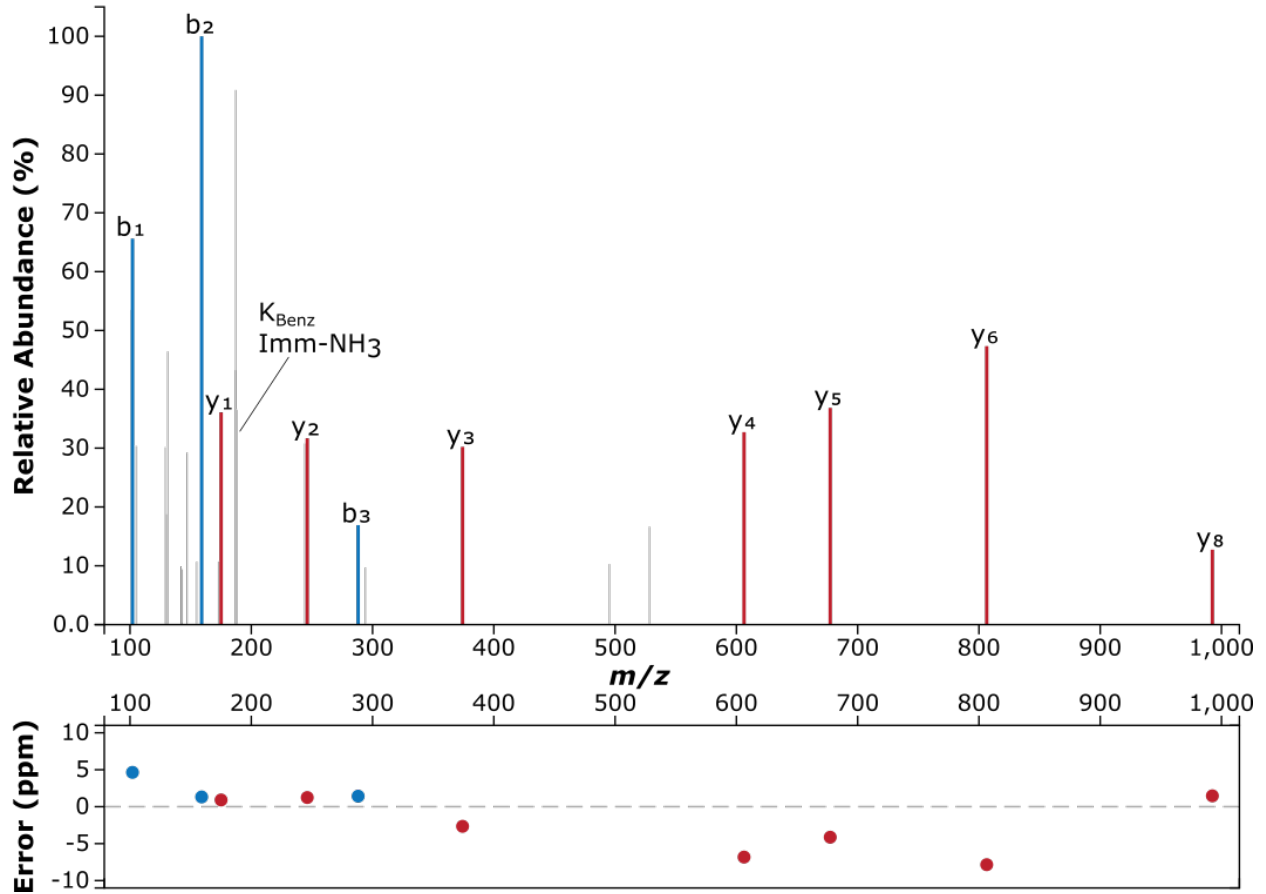


T\G\E\E\A\k\Q\A\R

Precursor m/z: 547.2673

Charge: +2

6-oxocyclohex-1-ene-1-carboxyl-CoA hydrolase (Q2LXU2)

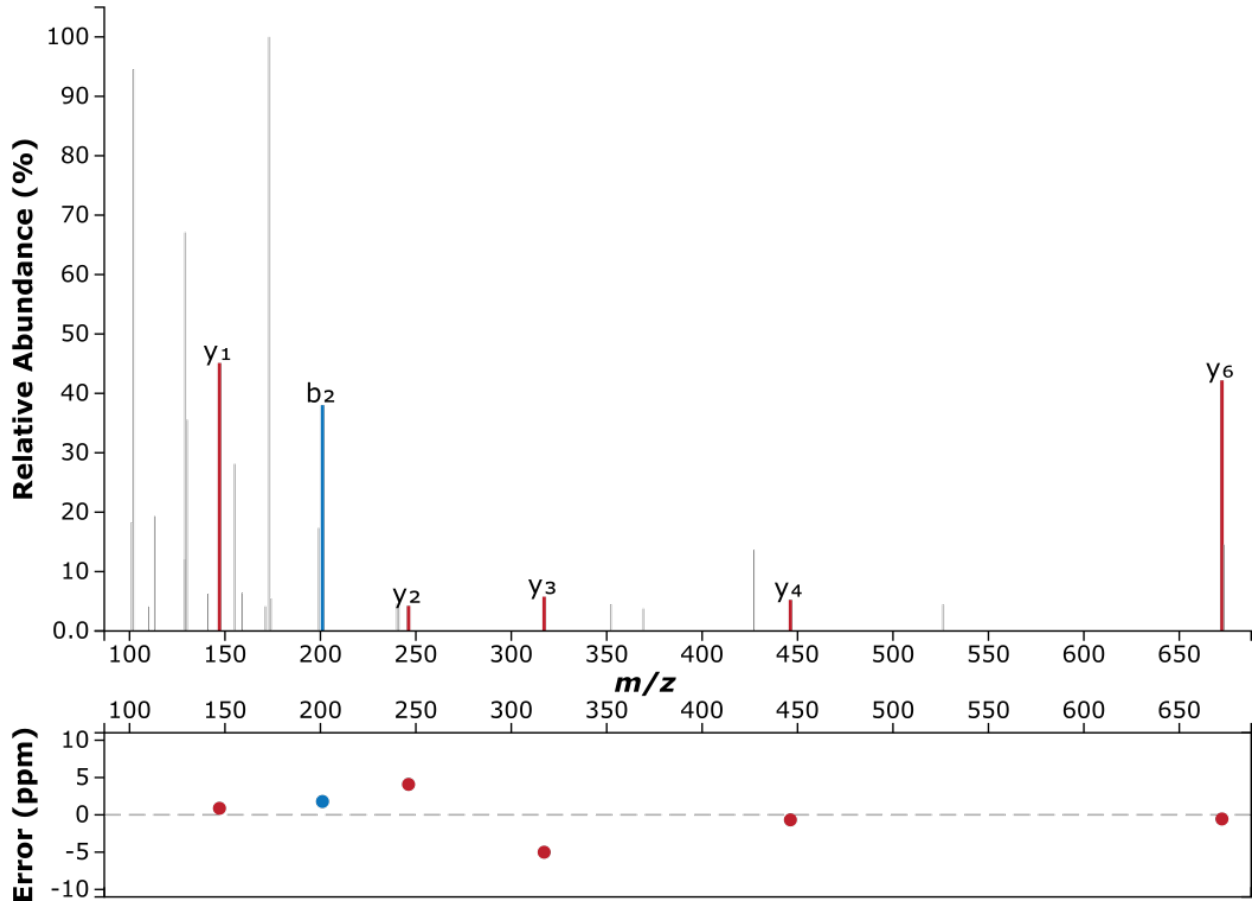


Butyryl



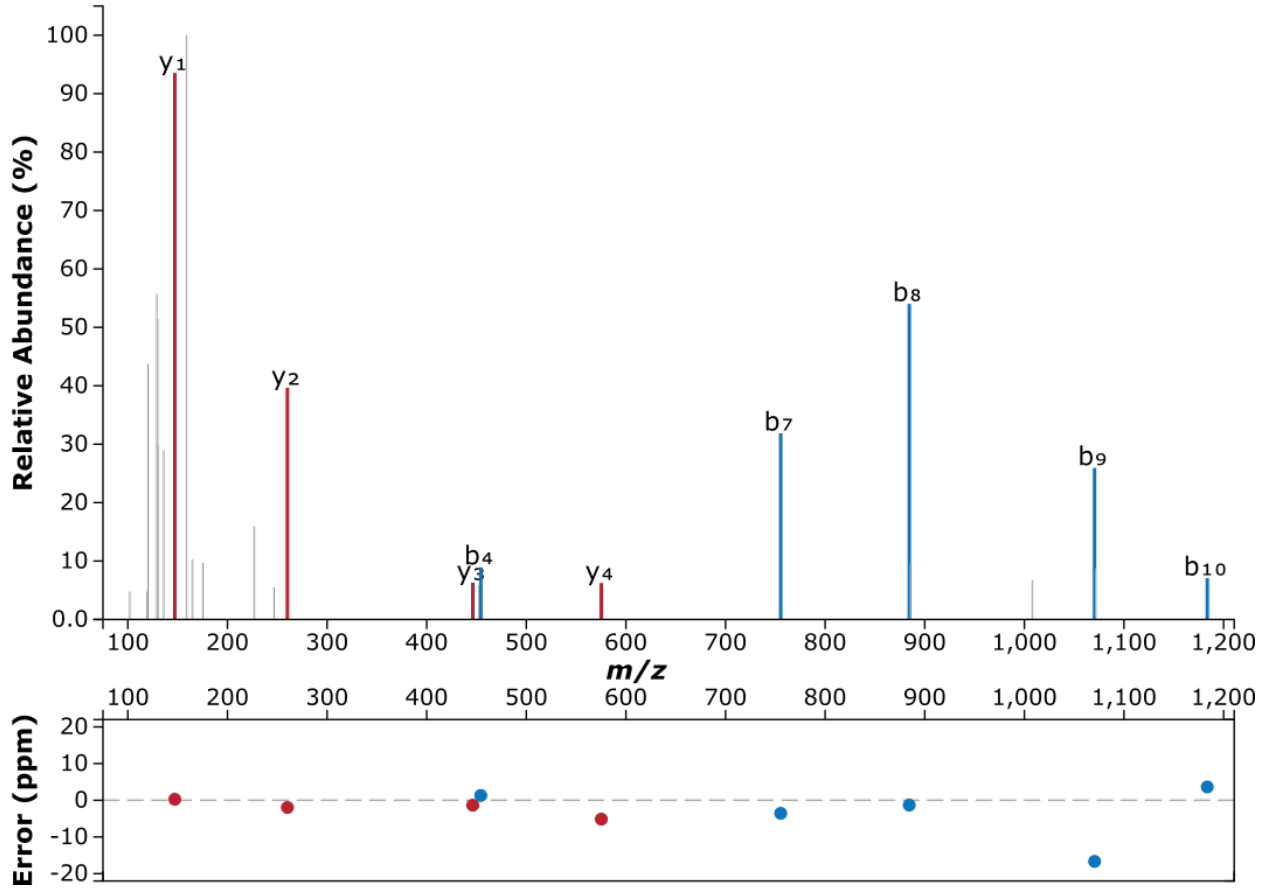
Precursor m/z: 443.2295 Charge: +3

Hypothetical exported protein (Q2LUP5)



A I A k^{But} E G D E W L K

Precursor m/z: 665.3561 Charge: +2
Acetyl-CoA synthetase (Q2LWR8)

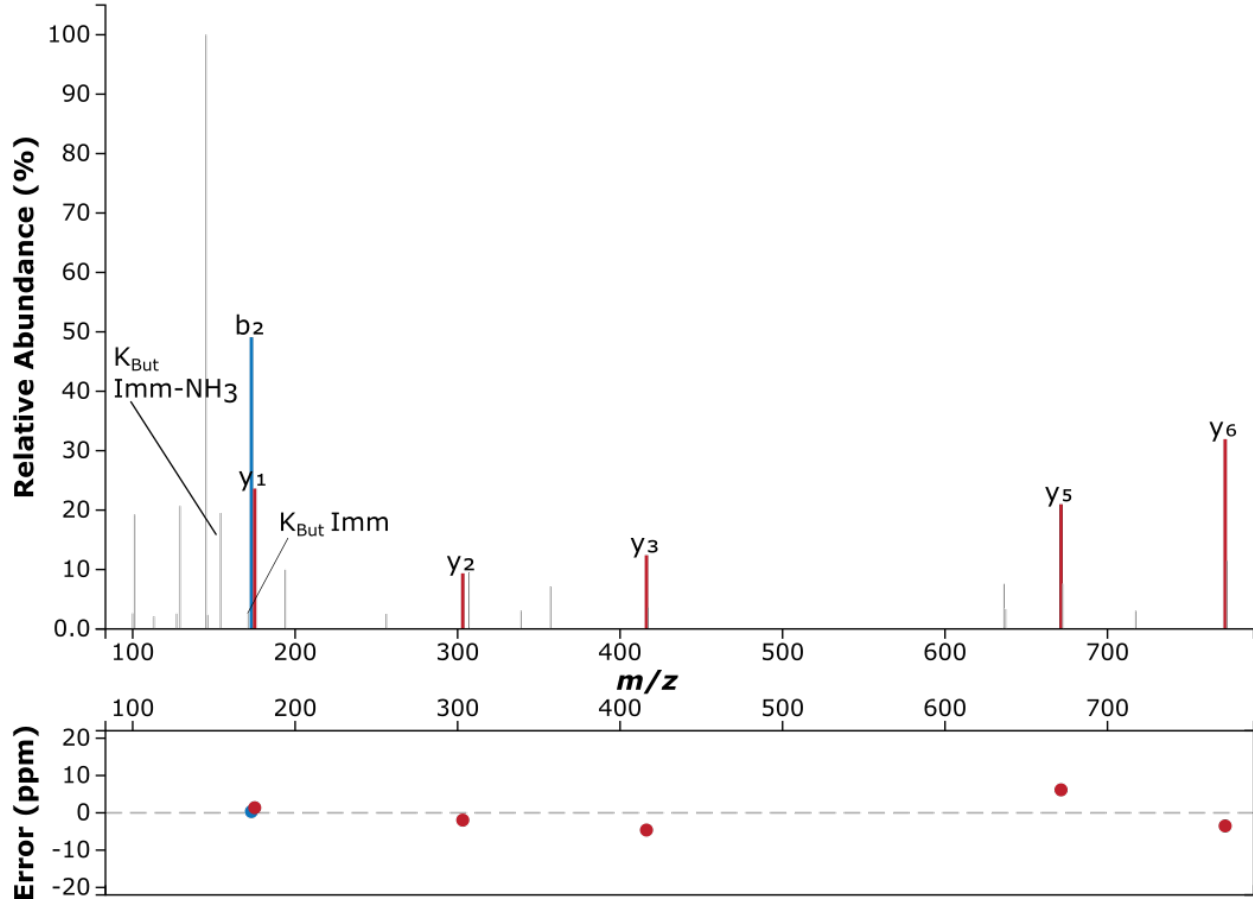


T A T G k I Q R
^{but}

Precursor m/z: 472.7798

Charge: +2

4-hydroxybenzoate-CoA ligase / benzoate-CoA ligase (Q2LRH0, Q2LRH7)



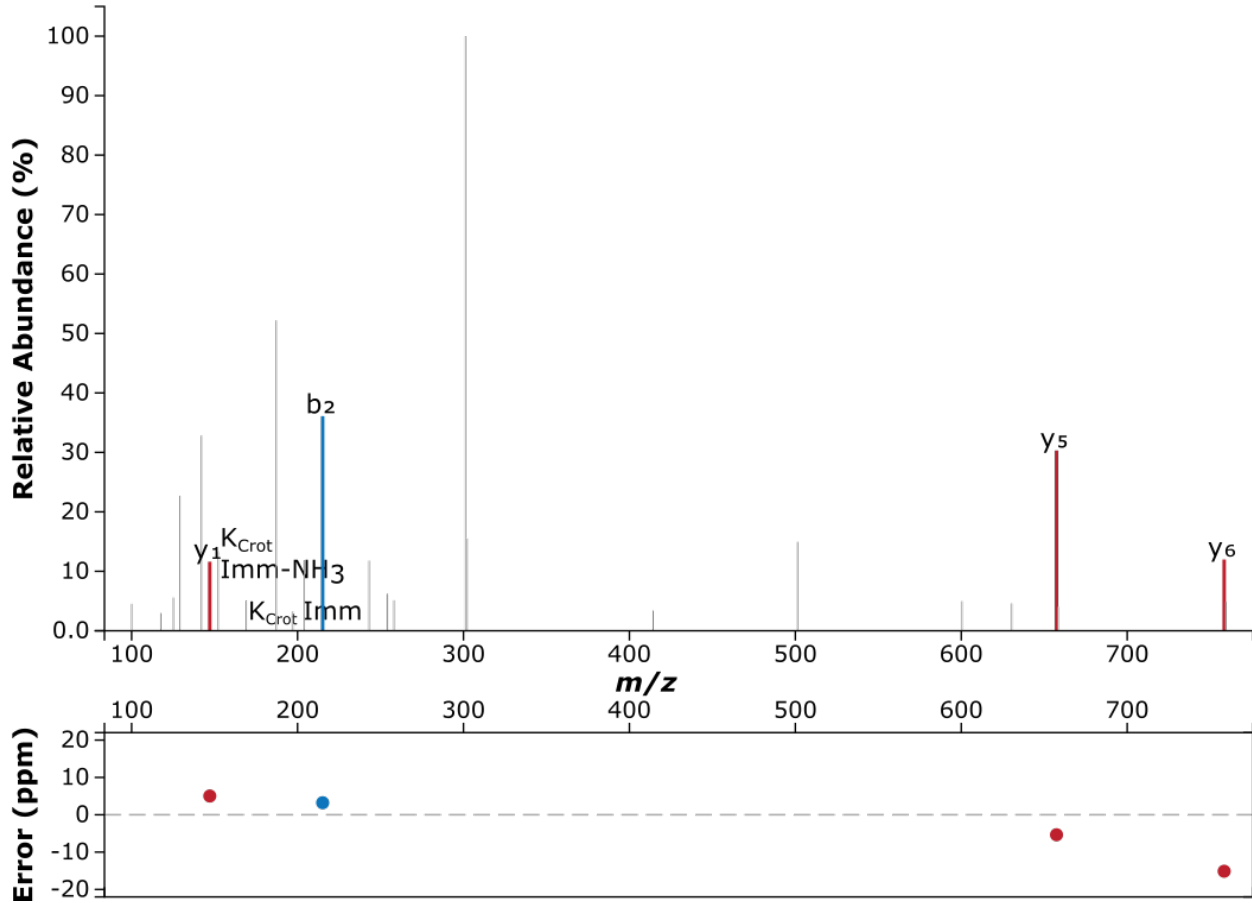
Crotonyl

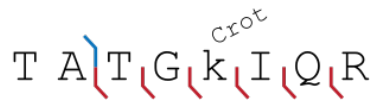


Precursor m/z: 436.2716

Charge: +2

Acetyl-CoA synthetase / acetyltransferase (GNAT) family protein (Q2LUQ3)

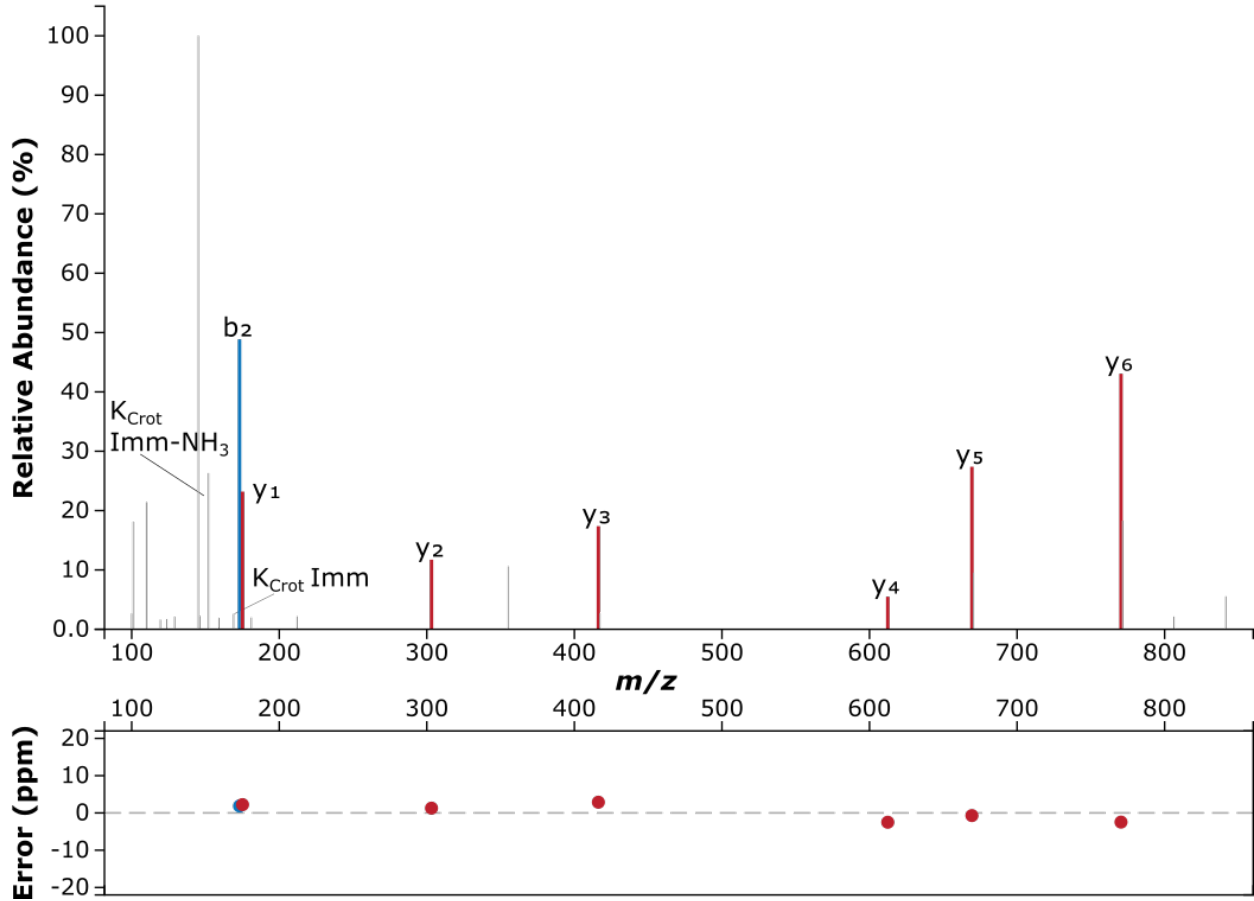




Precursor m/z: 471.7720

Charge: +2

4-hydroxybenzoate-CoA ligase / benzoate-CoA ligase (Q2LRH0, Q2LRH7)



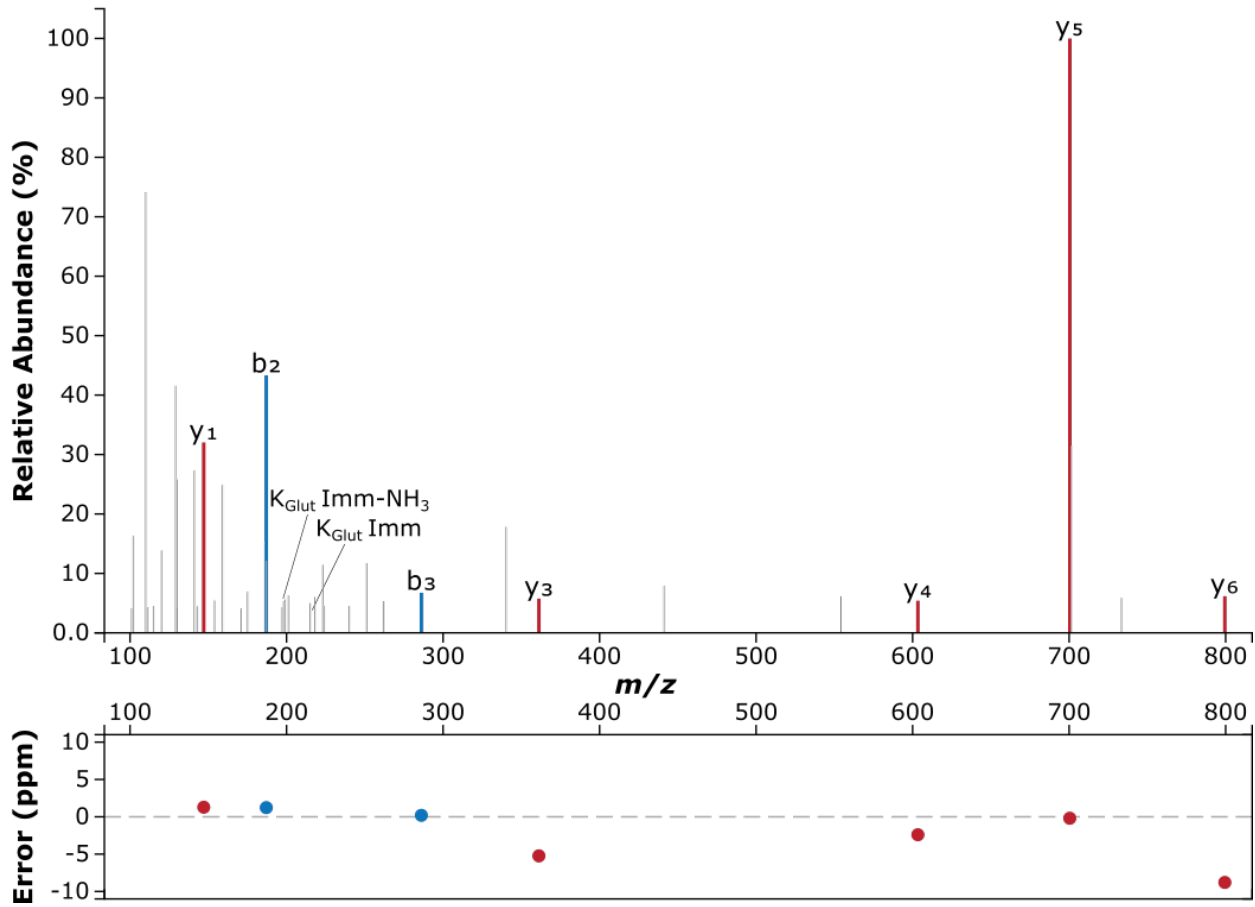
Glutaryl

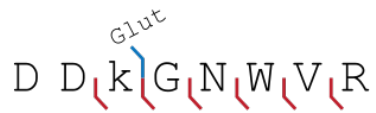


Precursor m/z: 493.2819

Charge: +2

6-hydroxycyclohex-1-ene-1-carboxyl-CoA dehydrogenase (Q2LXU3)

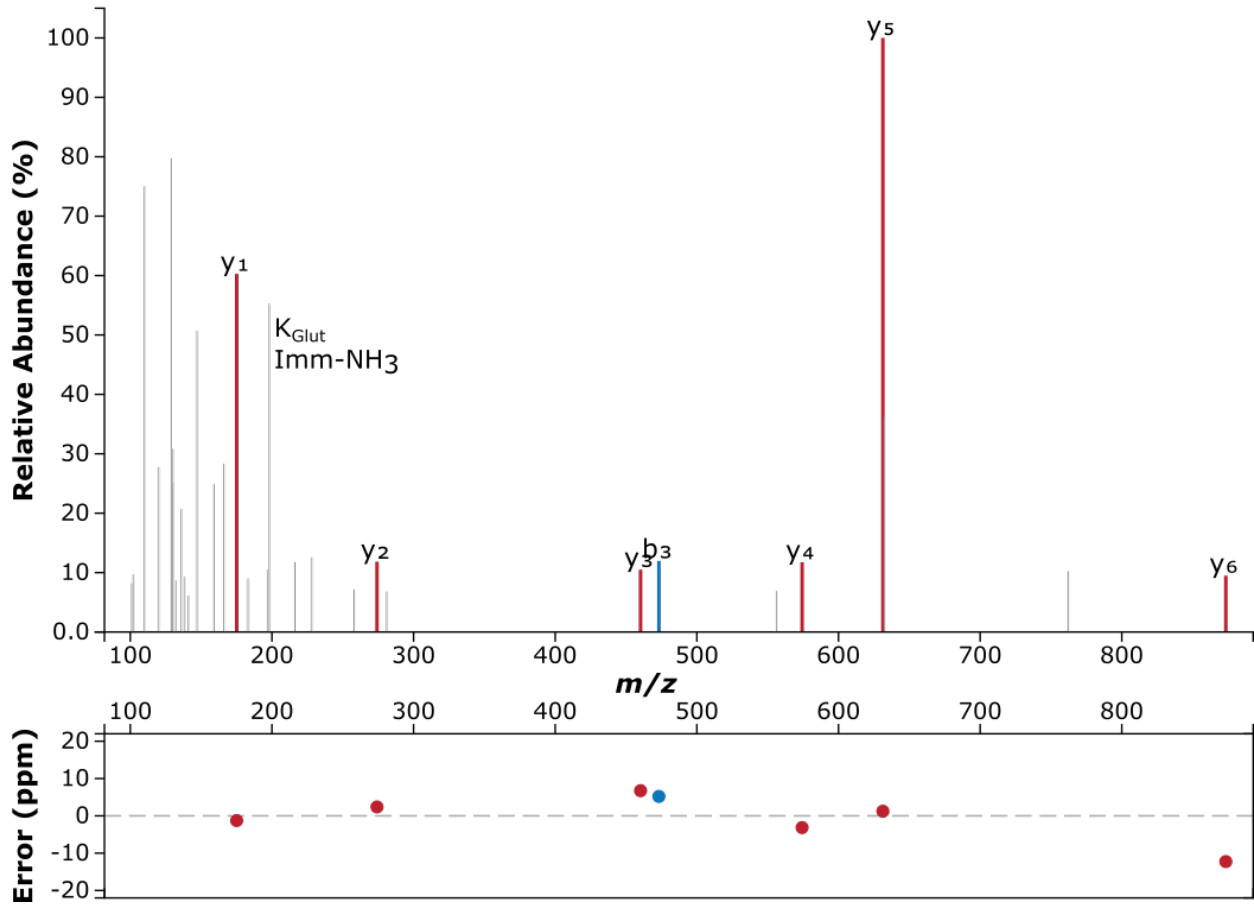




Precursor m/z: 552.2594

Charge: +2

6-oxocyclohex-1-ene-1-carboxyl-CoA hydrolase (Q2LXU2)

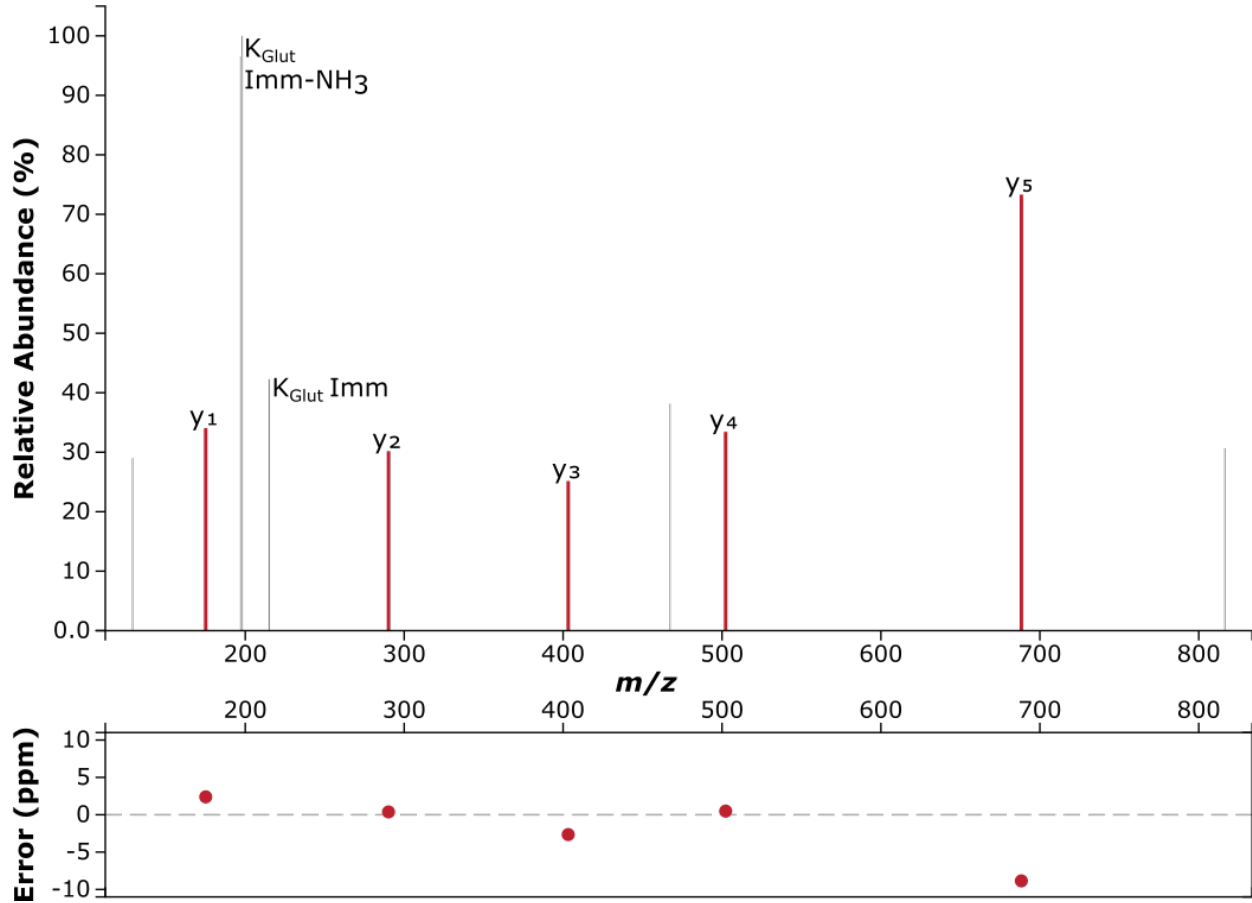


E L k ^{Glut} Q W V L D R

Precursor m/z: 650.8484

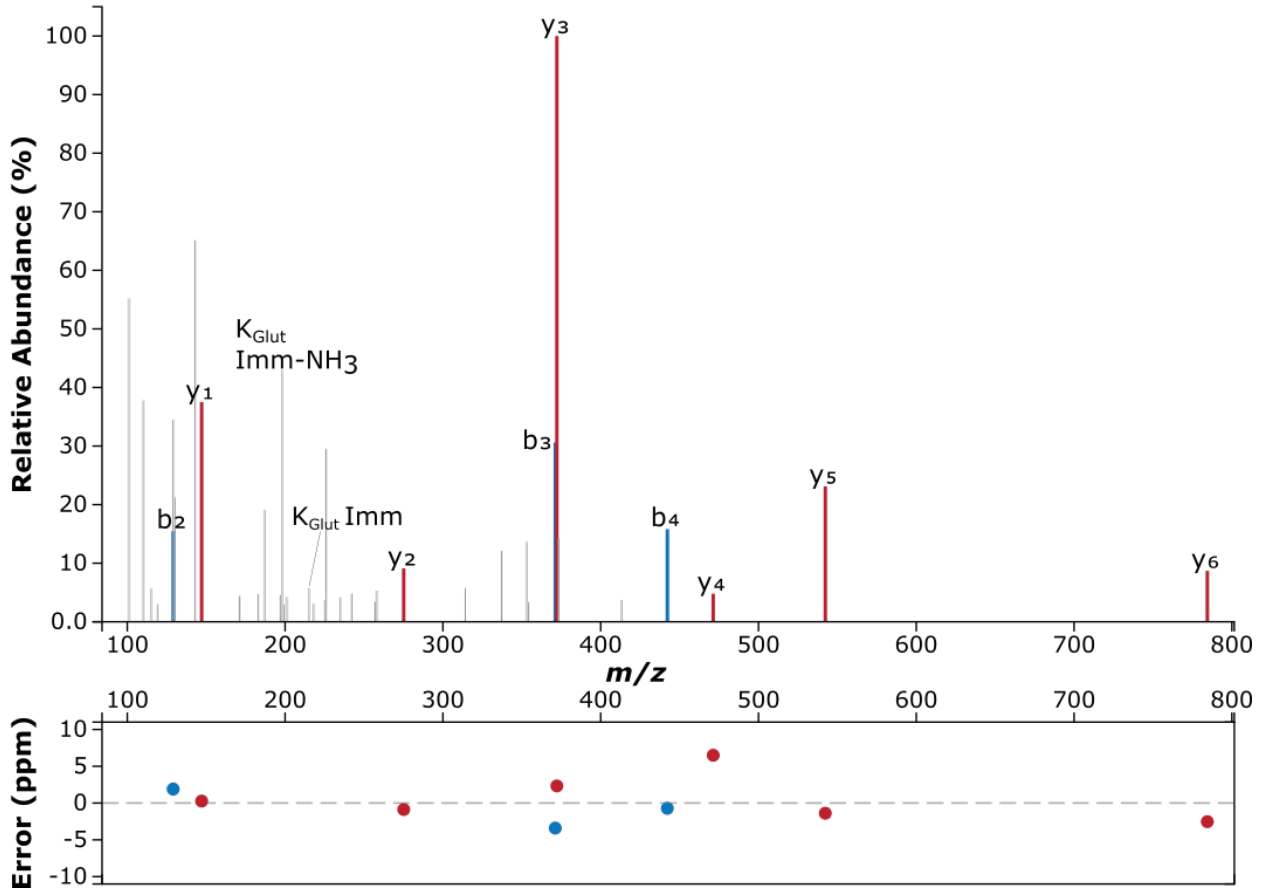
Charge: +2

4-hydroxybenzoate-CoA ligase / benzoate-CoA ligase (Q2LRH0, Q2LRH7)





Precursor m/z: 456.7611 Charge: +2
 Acetyl-CoA synthetase (Q2LWR8)

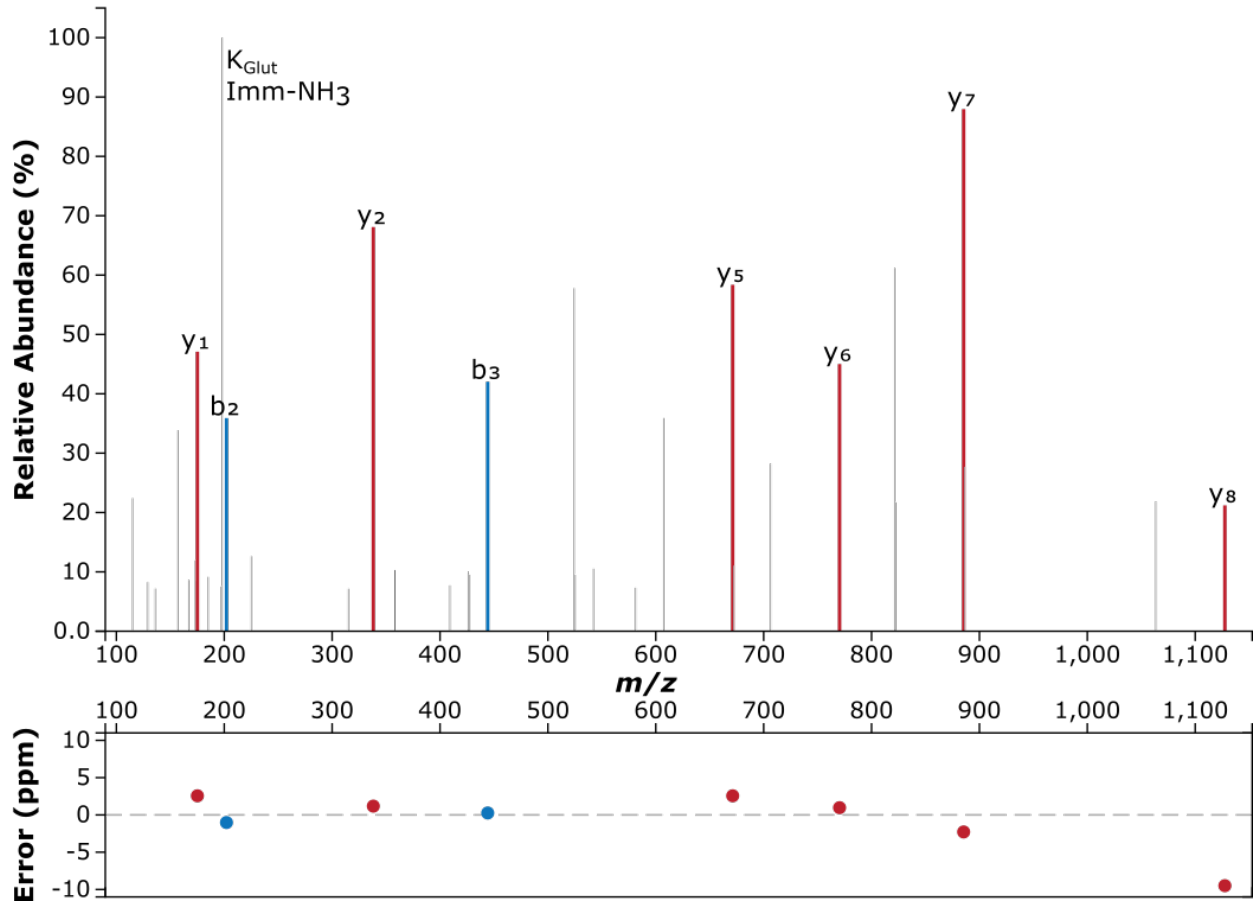


S N k^{Glut} D V m G E Y R

Precursor m/z: 664.7930

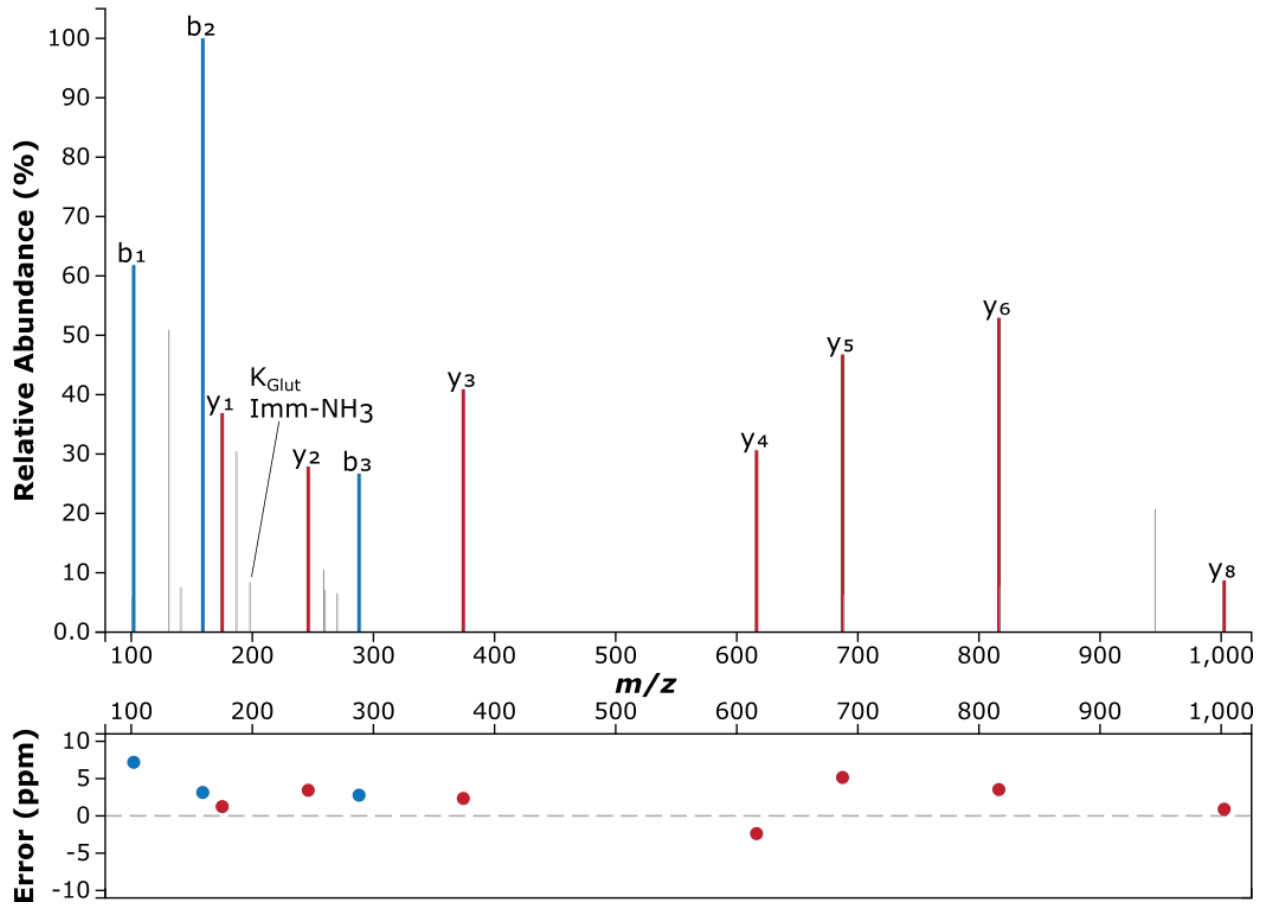
Charge: +2

6-hydroxycyclohex-1-ene-1-carboxyl-CoA dehydrogenase (Q2LXU3)





Precursor m/z : 552.2700 Charge: +2
 6-oxocyclohex-1-ene-1-carbonyl-CoA hydrolase (Q2LXU2)

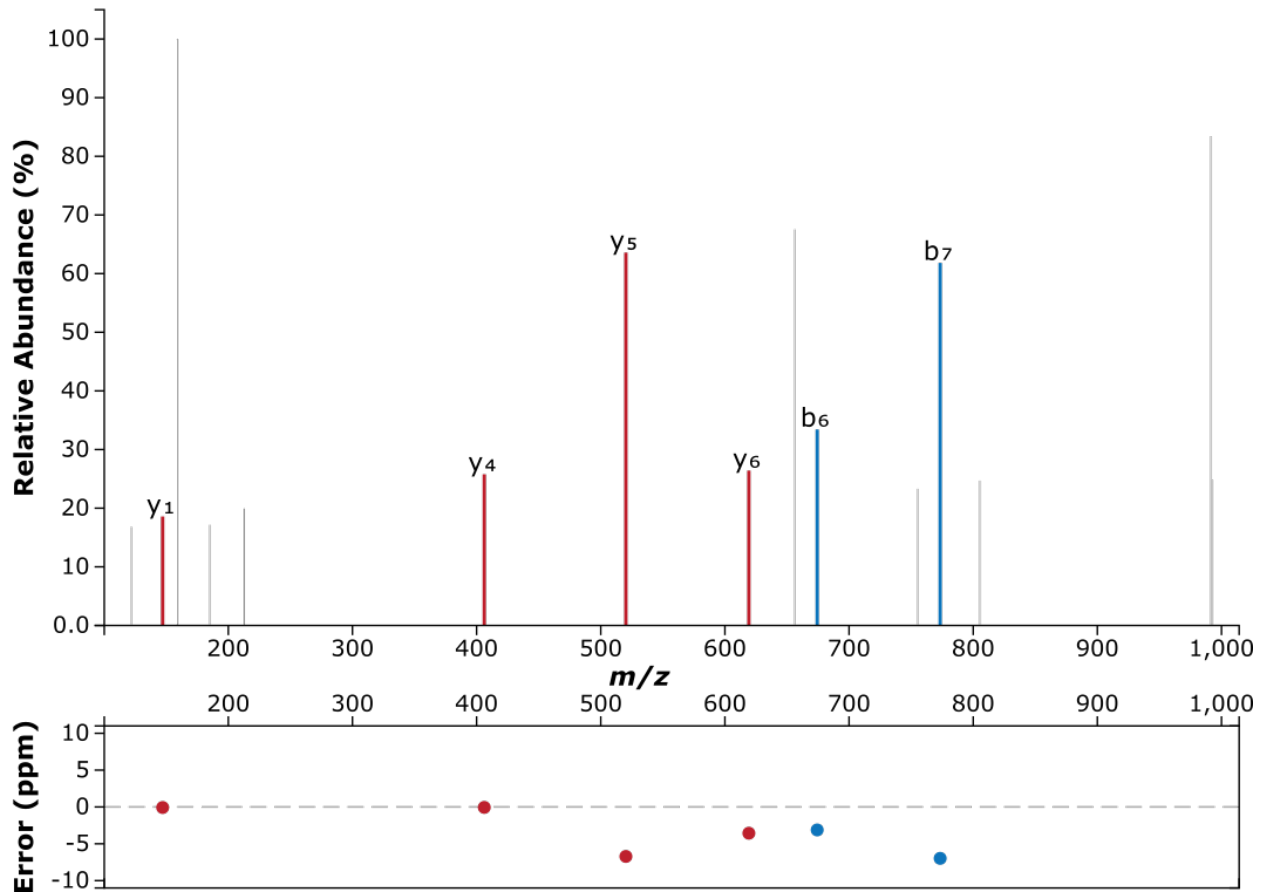


3-Hydroxybutyryl

3-OH Butyryl
D k T A A T V I G E W V N S G D K

Precursor m/z: 626.3144 Charge: +3

4-hydroxybenzoate-CoA ligase / benzoate-CoA ligase (Q2LRH7)

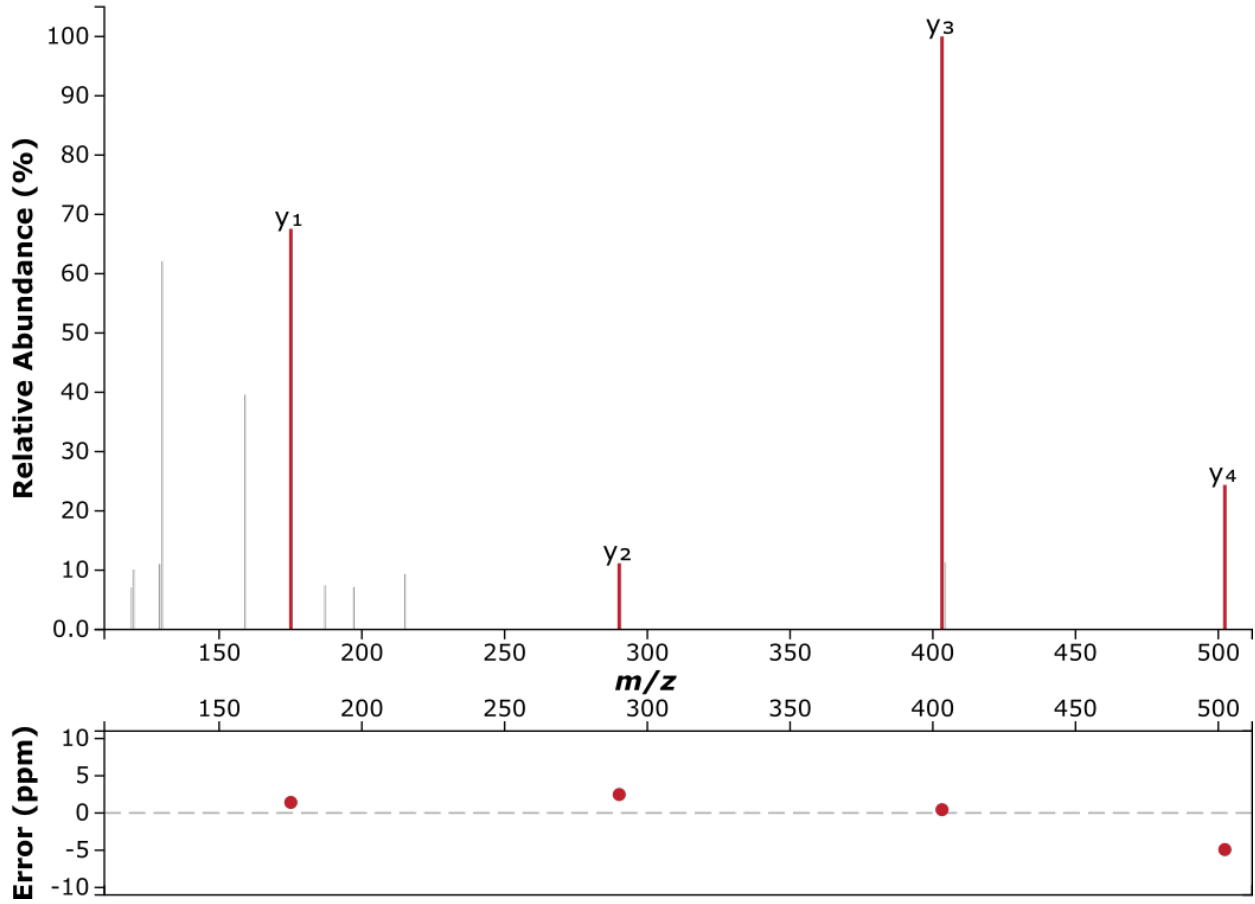


3-OH Butyryl
E L k Q W V L D R

Precursor m/z: 424.9031

Charge: +3

4-hydroxybenzoate-CoA ligase / benzoate-CoA ligase (Q2LRH0,Q2LRH7)

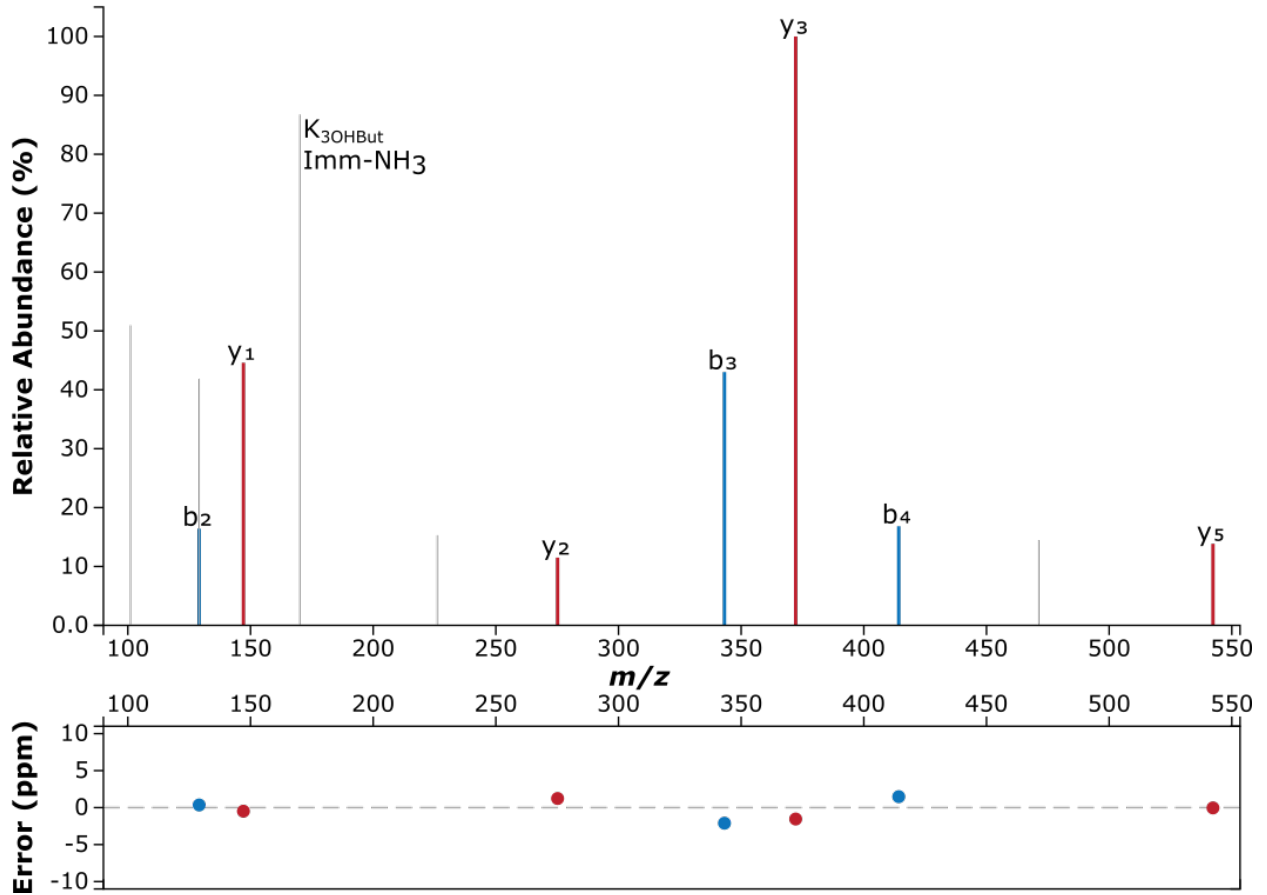




Precursor m/z: 442.7636

Charge: +2

Acetyl-CoA synthetase (Q2LWR8)

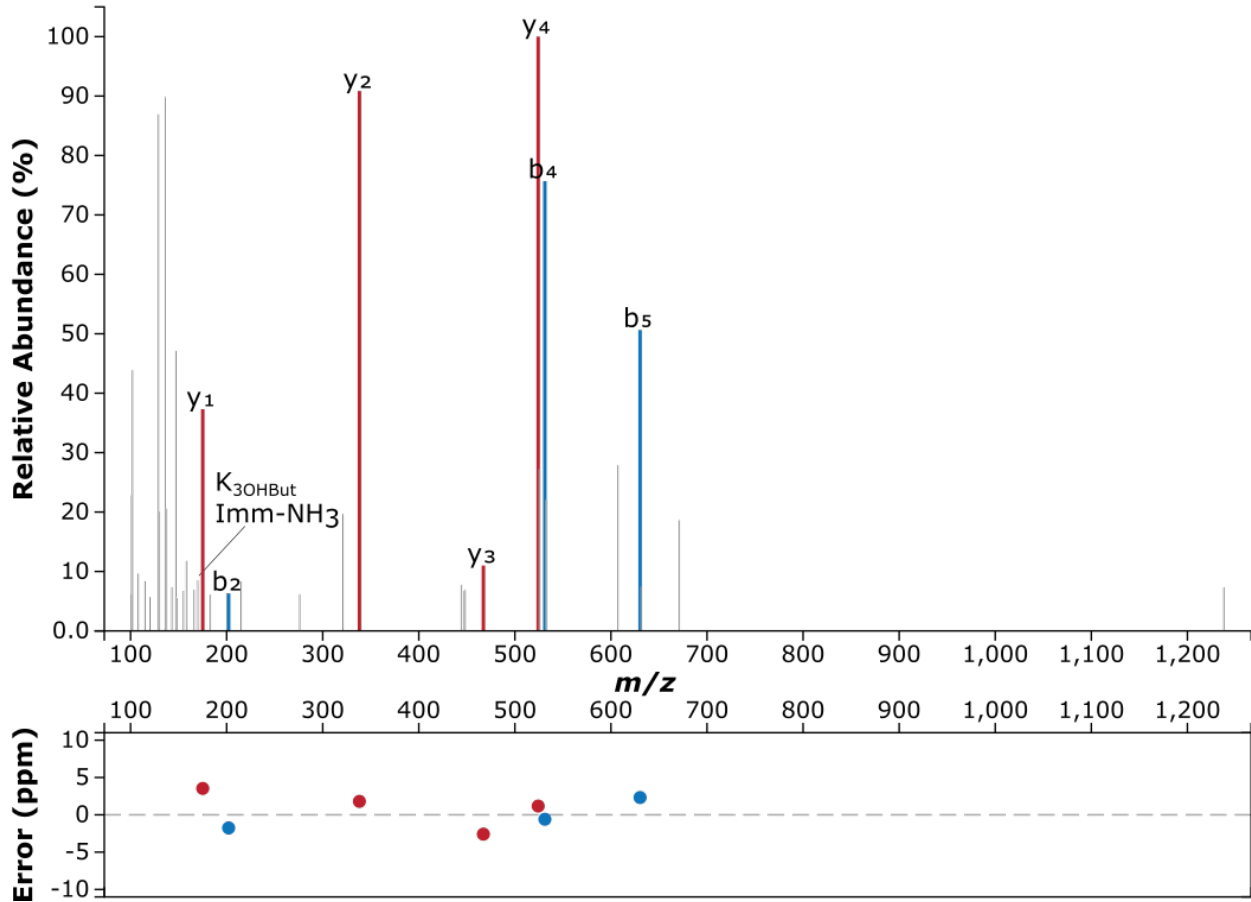


^{3-OH butyryl}
 S N k D V M G E Y R

Precursor m/z: 428.8678

Charge: +3

6-hydroxycyclohex-1-ene-1-carboxyl-CoA dehydrogenase (Q2LXU3)

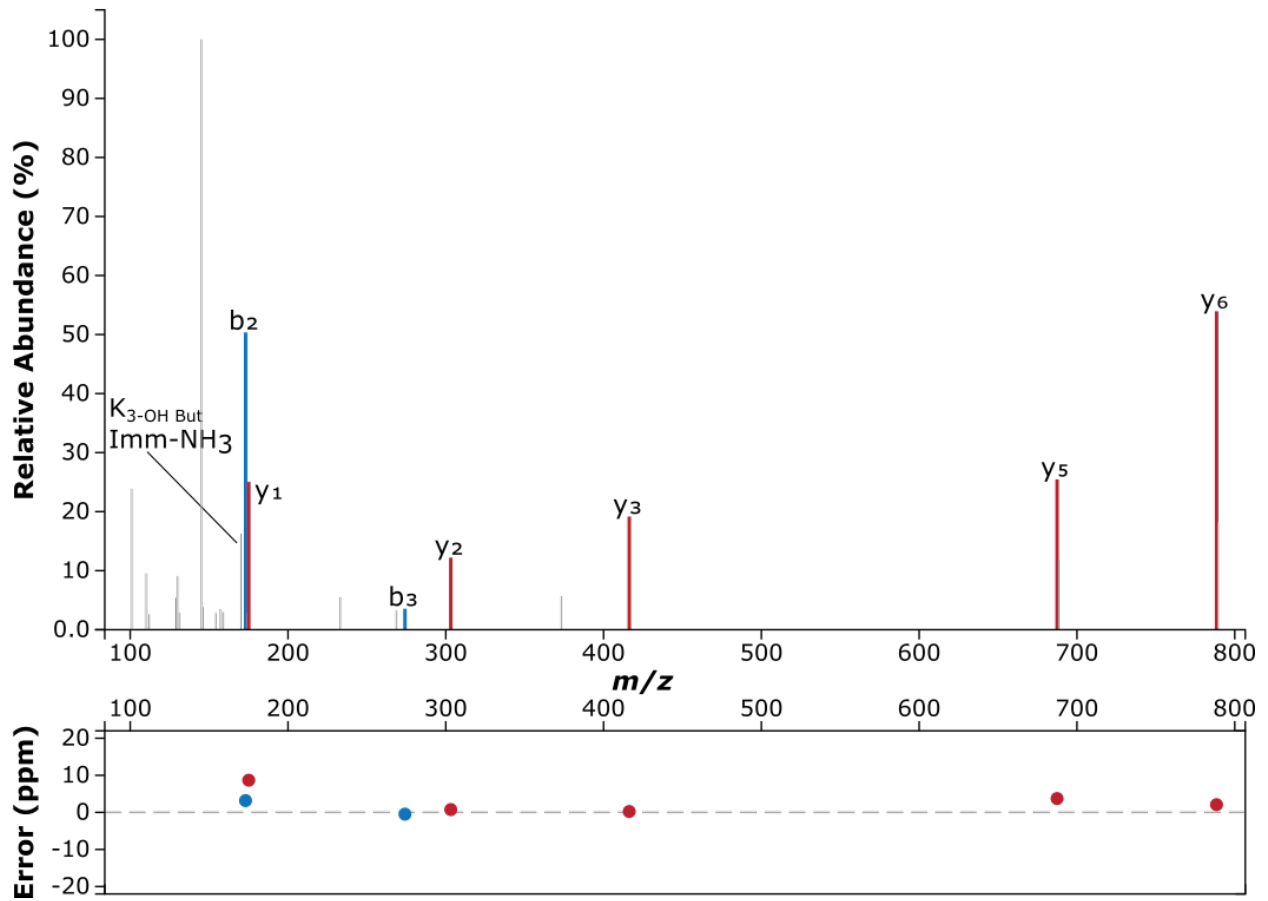


T A T G k I Q R
^{3-OH Butyryl}

Precursor m/z: 480.7773

Charge: +2

4-hydroxybenzoate-CoA ligase / benzoate-CoA ligase (Q2LRH0, Q2LRH7)

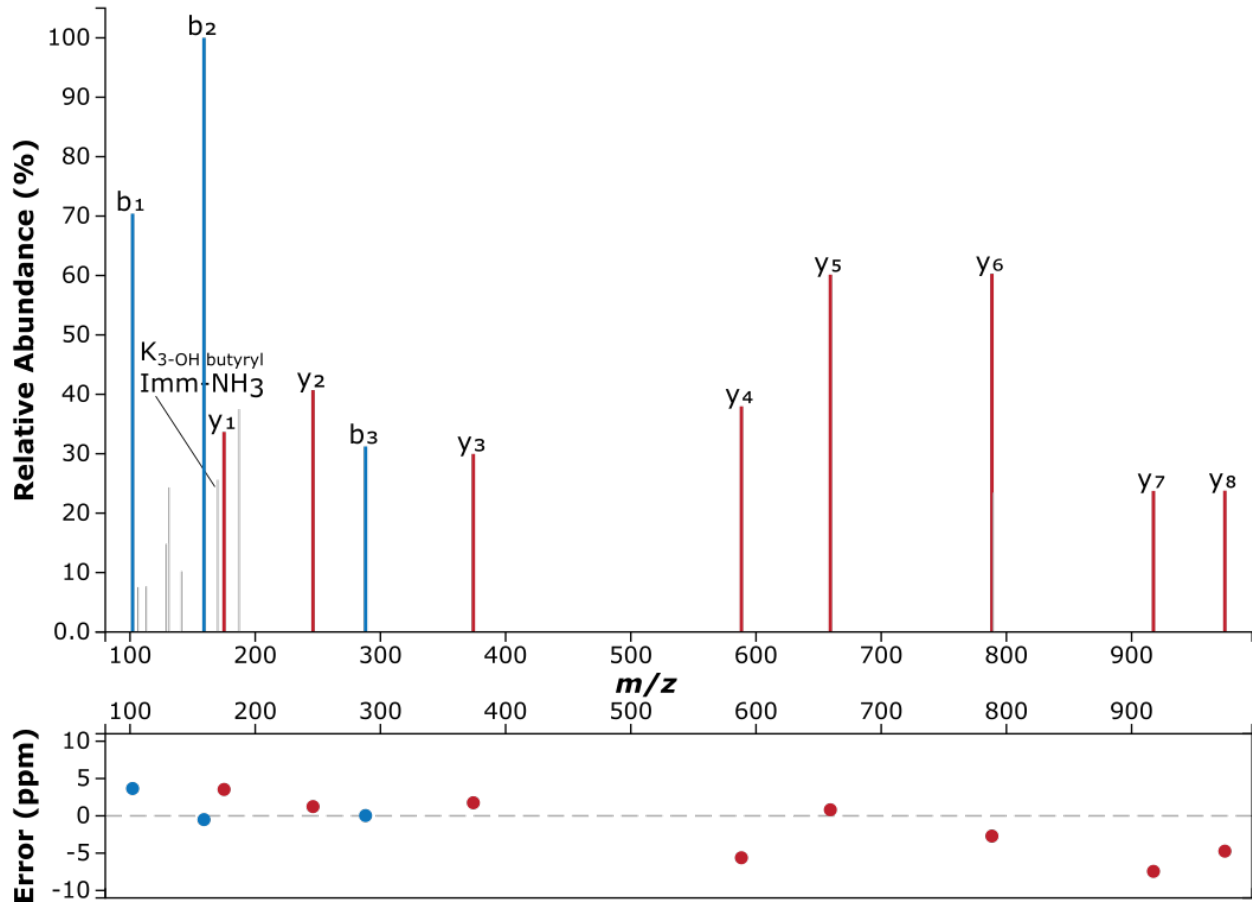




Precursor m/z: 538.2725

Charge: +2

6-oxocyclohex-1-ene-1-carbonyl-CoA hydrolase (Q2LXU2)



3-Hydroxypimelyl

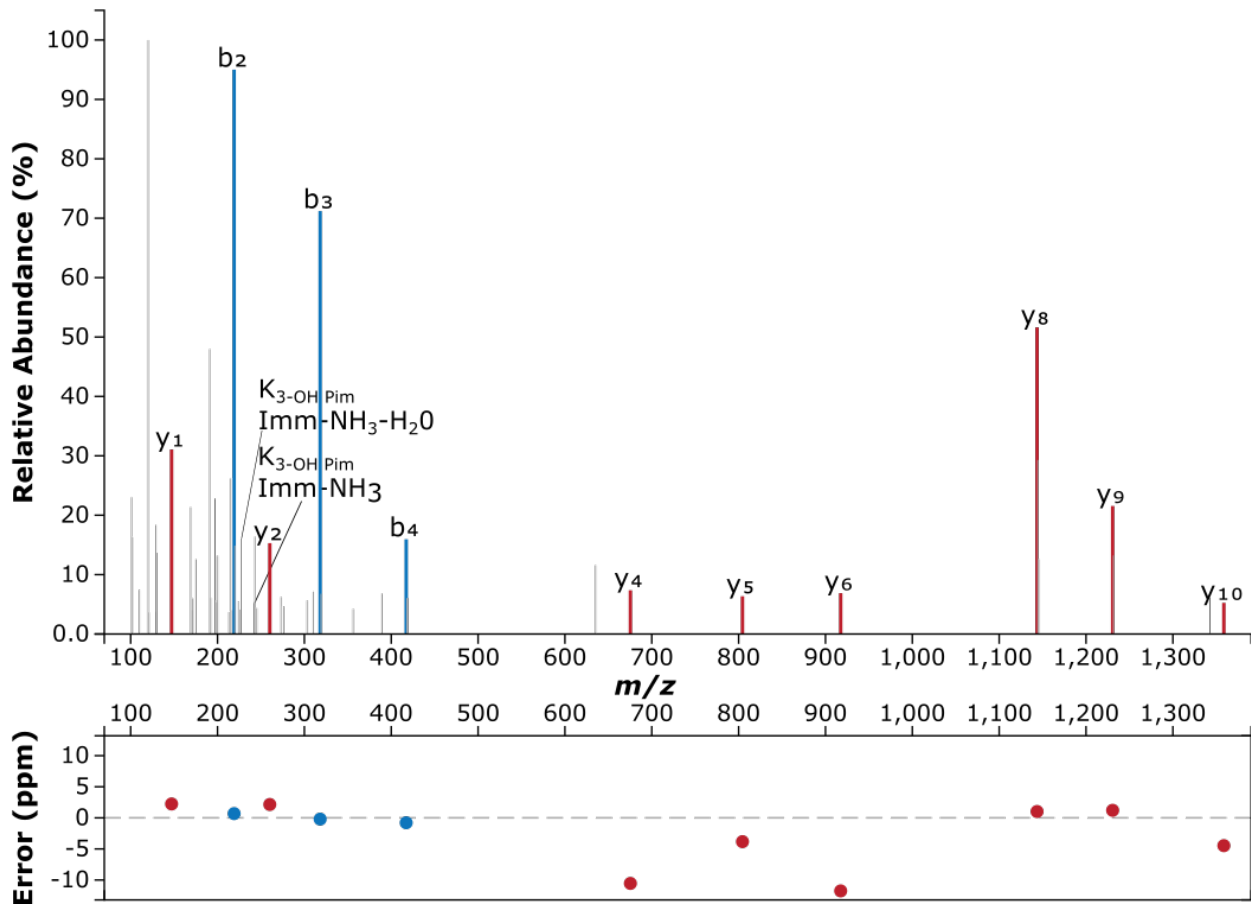
A F V V L N Q G F A Q S P E L E k E L K

3-OH Pimelyl

Precursor m/z: 802.4248

Charge: +3

4-hydroxybenzoate-CoA ligase / benzoate-CoA ligase (Q2LRH0)

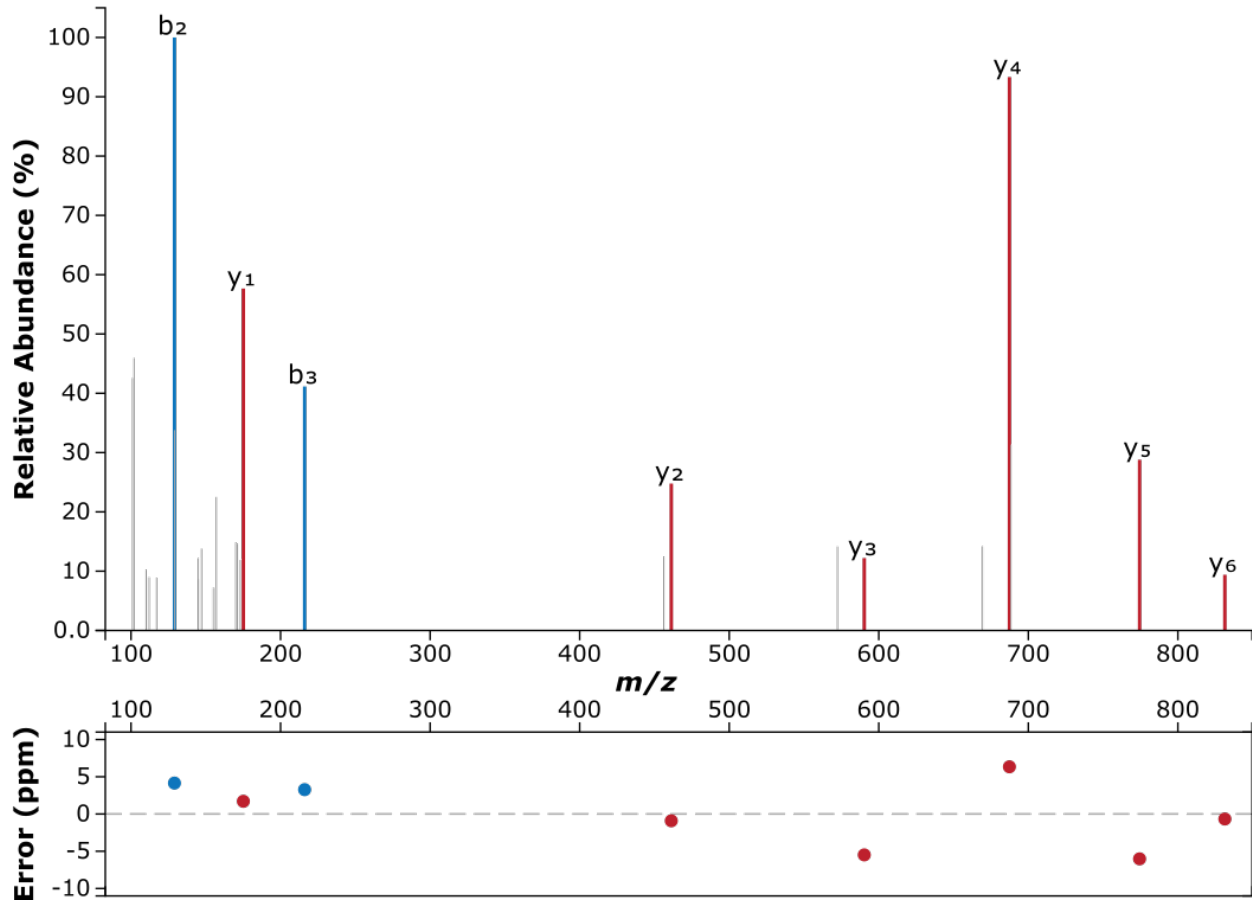


A₁G₁S₁P₁E₁k₁R₁^{3-OH pimelyl}

Precursor m/z: 451.7325

Charge: +2

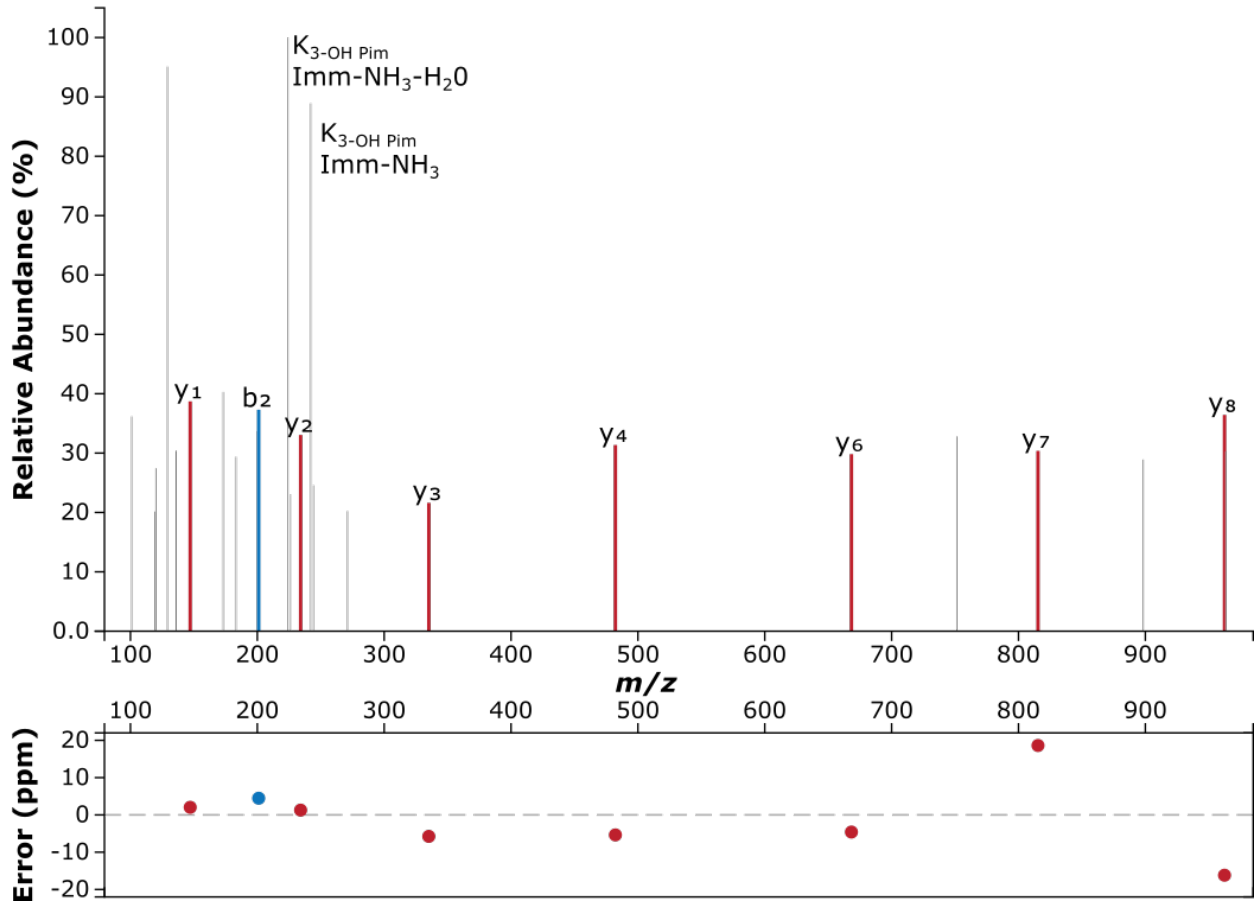
6-hydroxycyclohex-1-ene-1-carboxyl-CoA dehydrogenase (Q2LXU3)

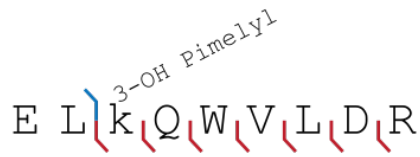


E A A Q k F m A D F T S K
^{3-OH Pimelyl}
^{ox}

Precursor m/z: 824.3822 Charge: +2

Putative enoyl-CoA hydratase (Q2LXU6)

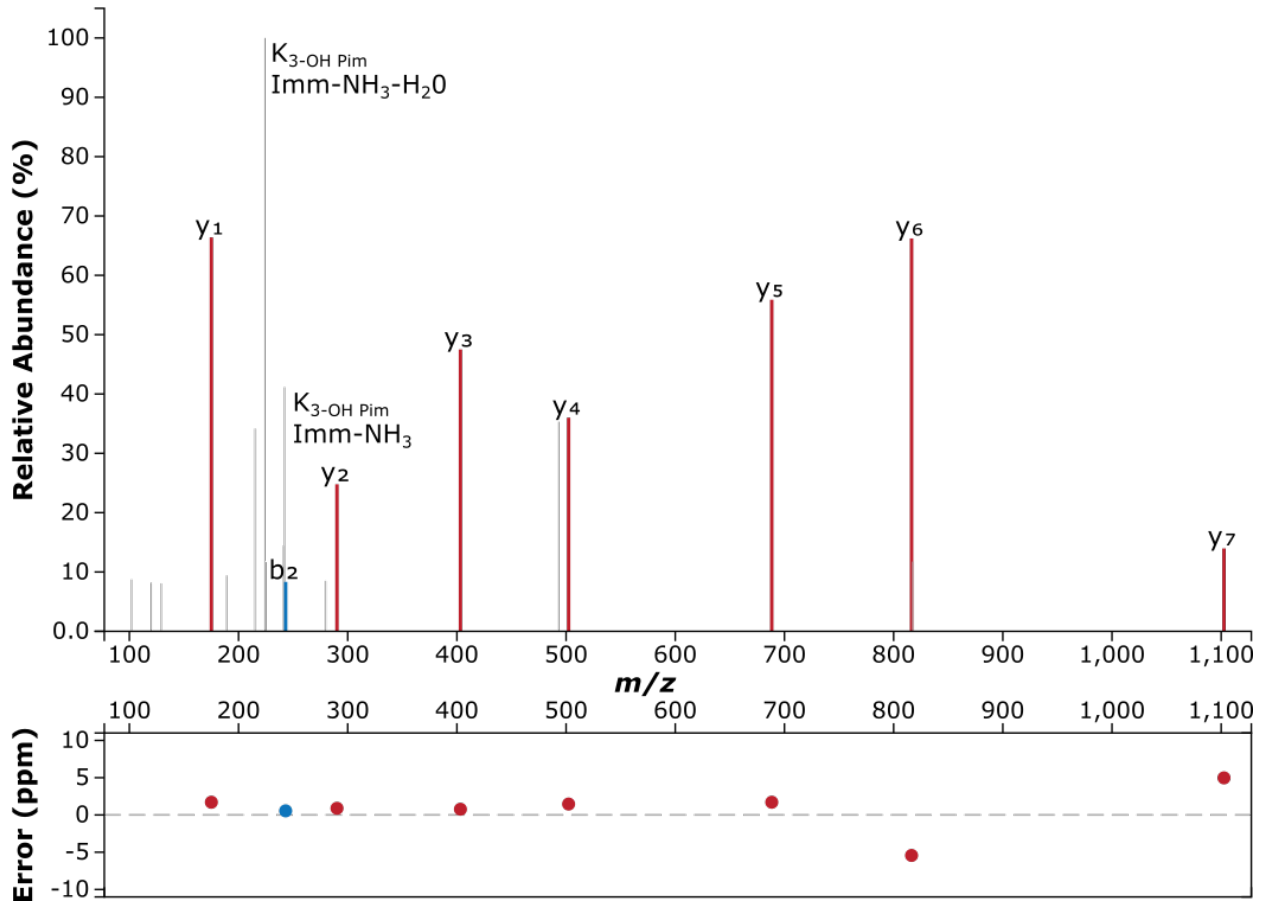




Precursor m/z: 672.8615

Charge: +2

4-hydroxybenzoate-CoA ligase / benzoate-CoA ligase (Q2LRH0, Q2LRH7)

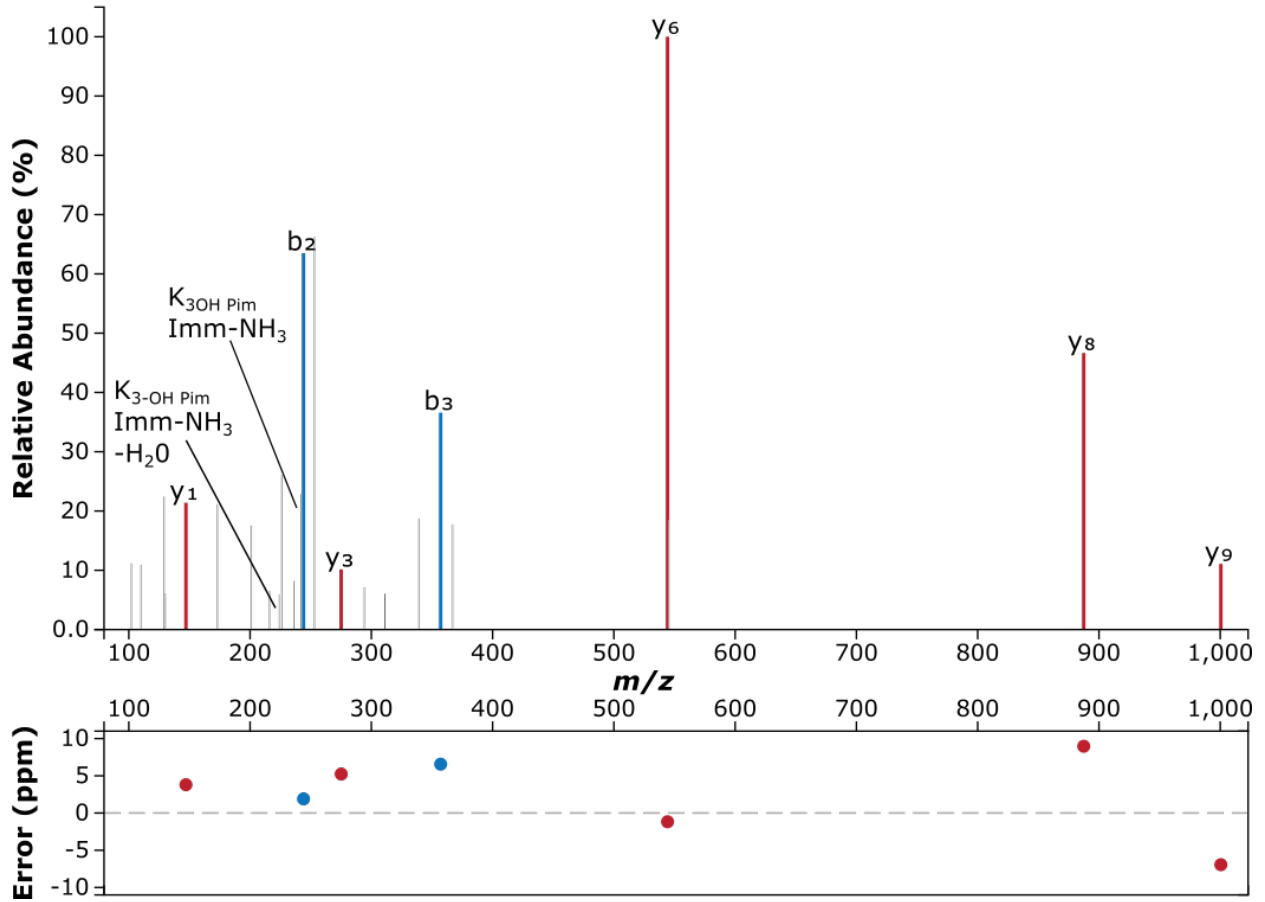


E N L L G k P G D G A K
^{3-OH Pimelyl}

Precursor m/z: 678.8539

Charge: +2

Acyl-CoA dehydrogenase, short-chain specific (Q2LUY8)

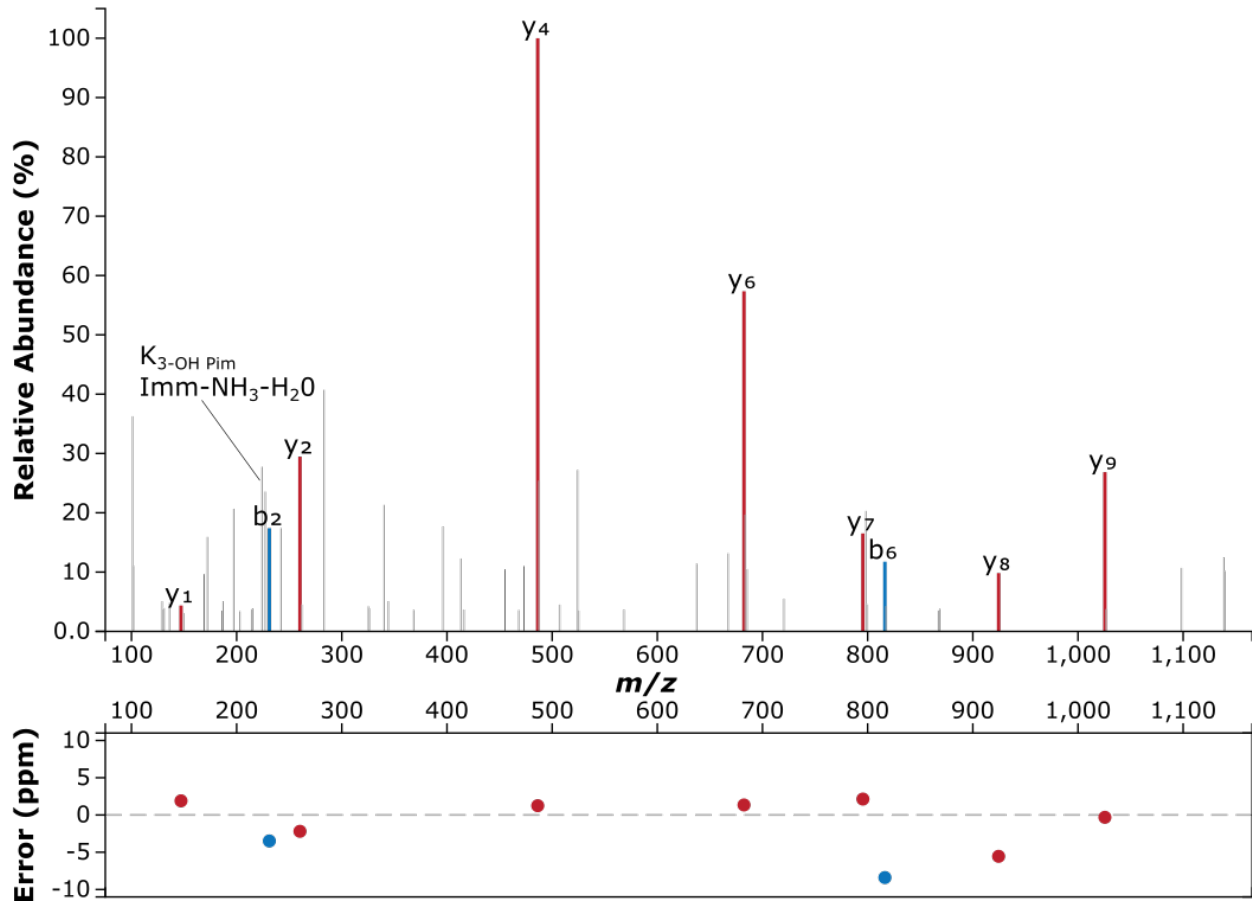


^{3-OH Pimelyl}
 E T G E k I P G Q L Q L T E I P V P E L K

Precursor m/z: 826.4510

Charge: +3

6-hydroxycyclohex-1-ene-1-carboxyl-CoA dehydrogenase (Q2LXU3)

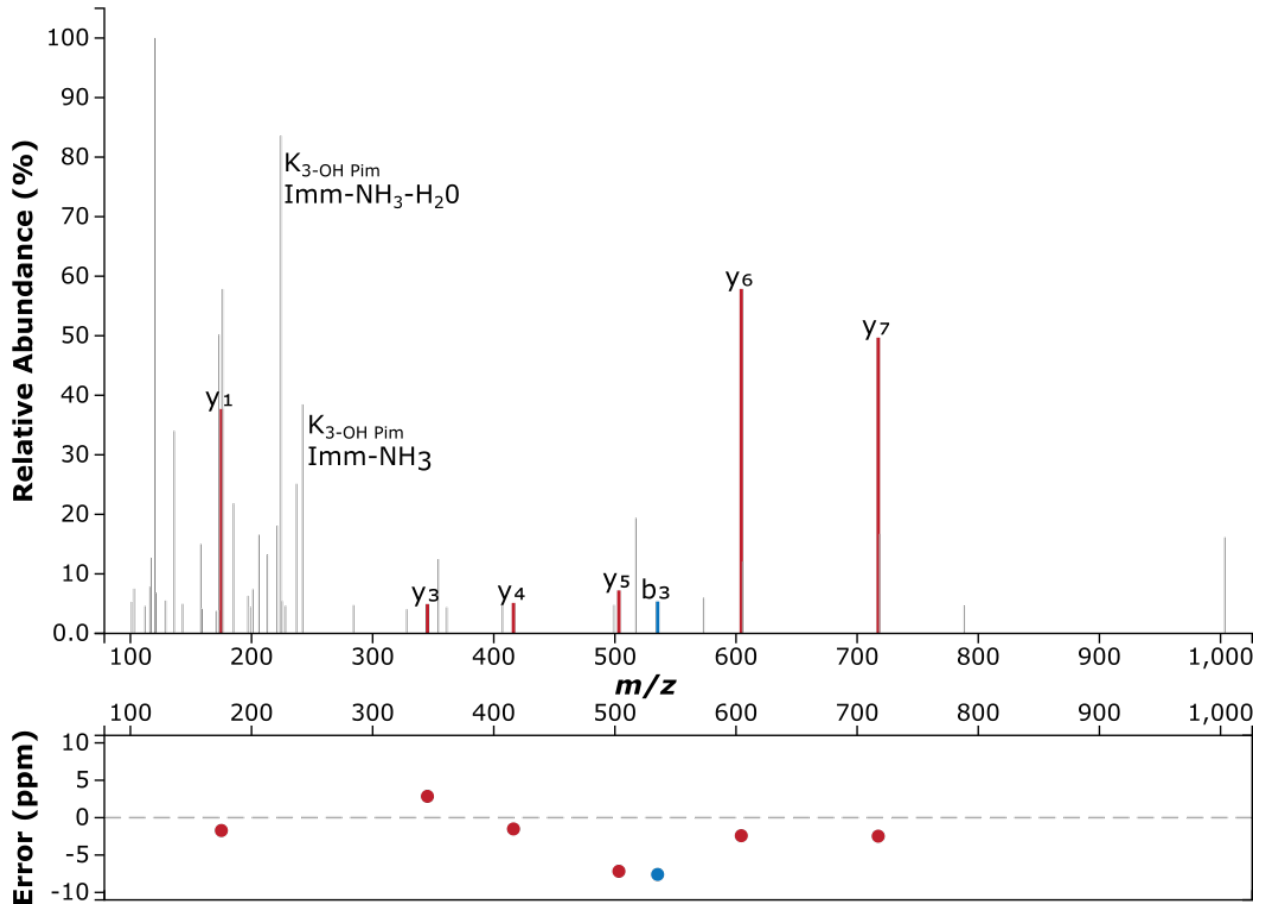


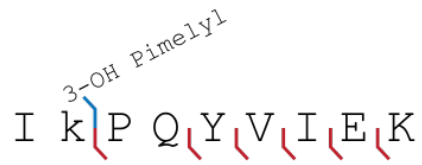


Precursor m/z: 626.3508

Charge: +2

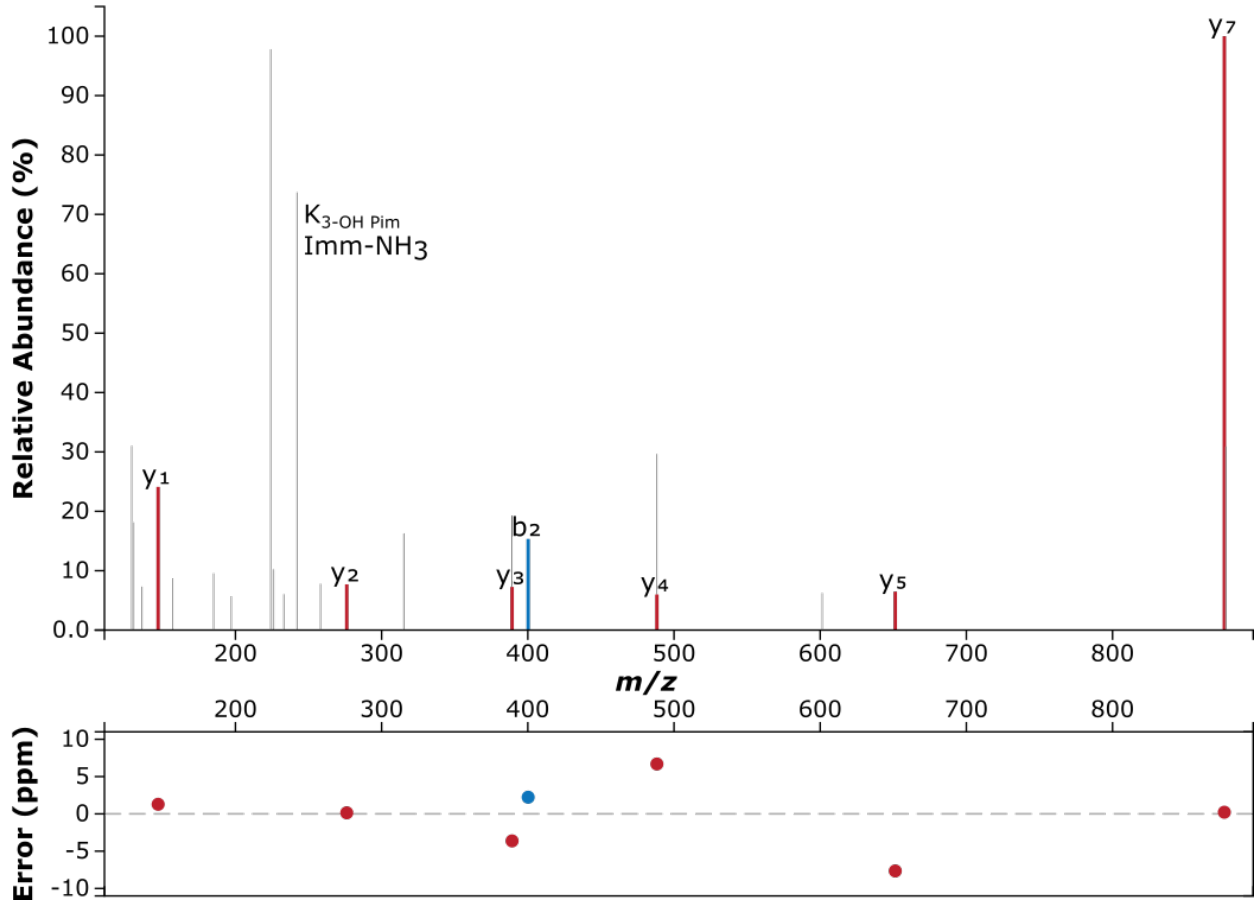
Inosine-5'-monophosphate dehydrogenase (Q2LTT4)





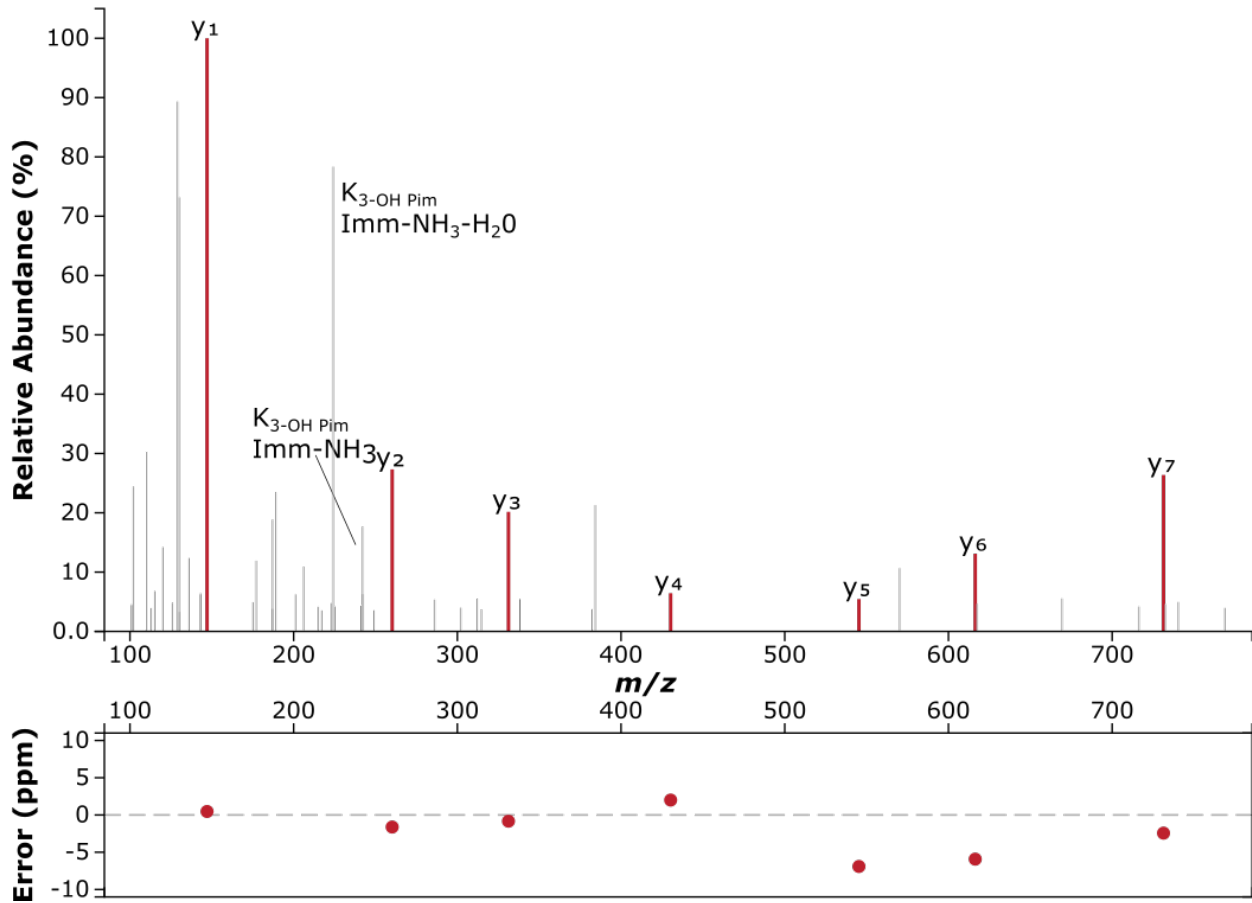
Precursor m/z: 638.3634

Charge: +2



3-OH Pimelyl
k D A D V A L K

Precursor m/z: 509.2768 Charge: +2
Acetyl-CoA synthetase (Q2LWR8)



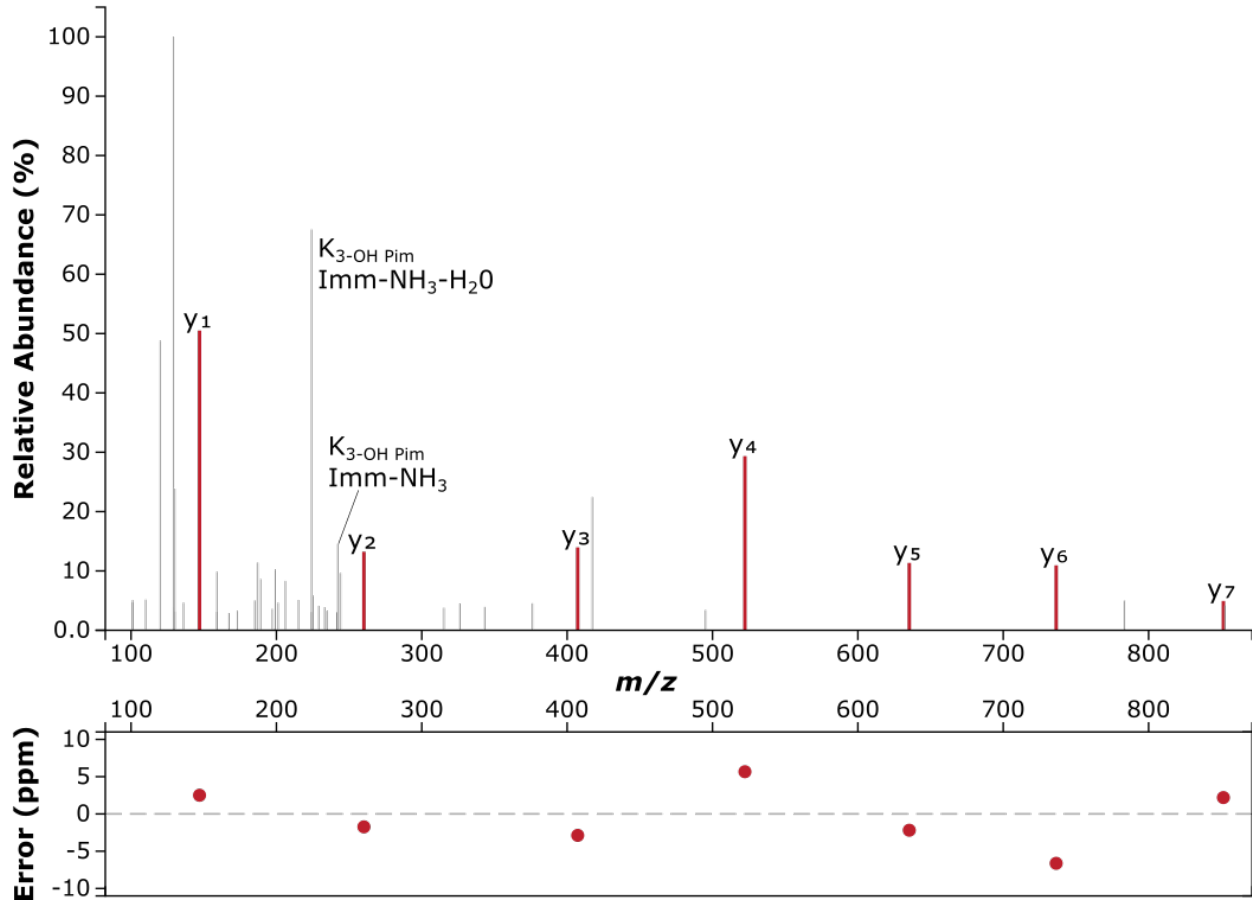
3-OH Pimelyl

k I T G K D T I D F I K

Precursor m/z: 512.9555

Charge: +3

6-oxocyclohex-1-ene-1-carbonyl-CoA hydrolase (Q2LXU2)



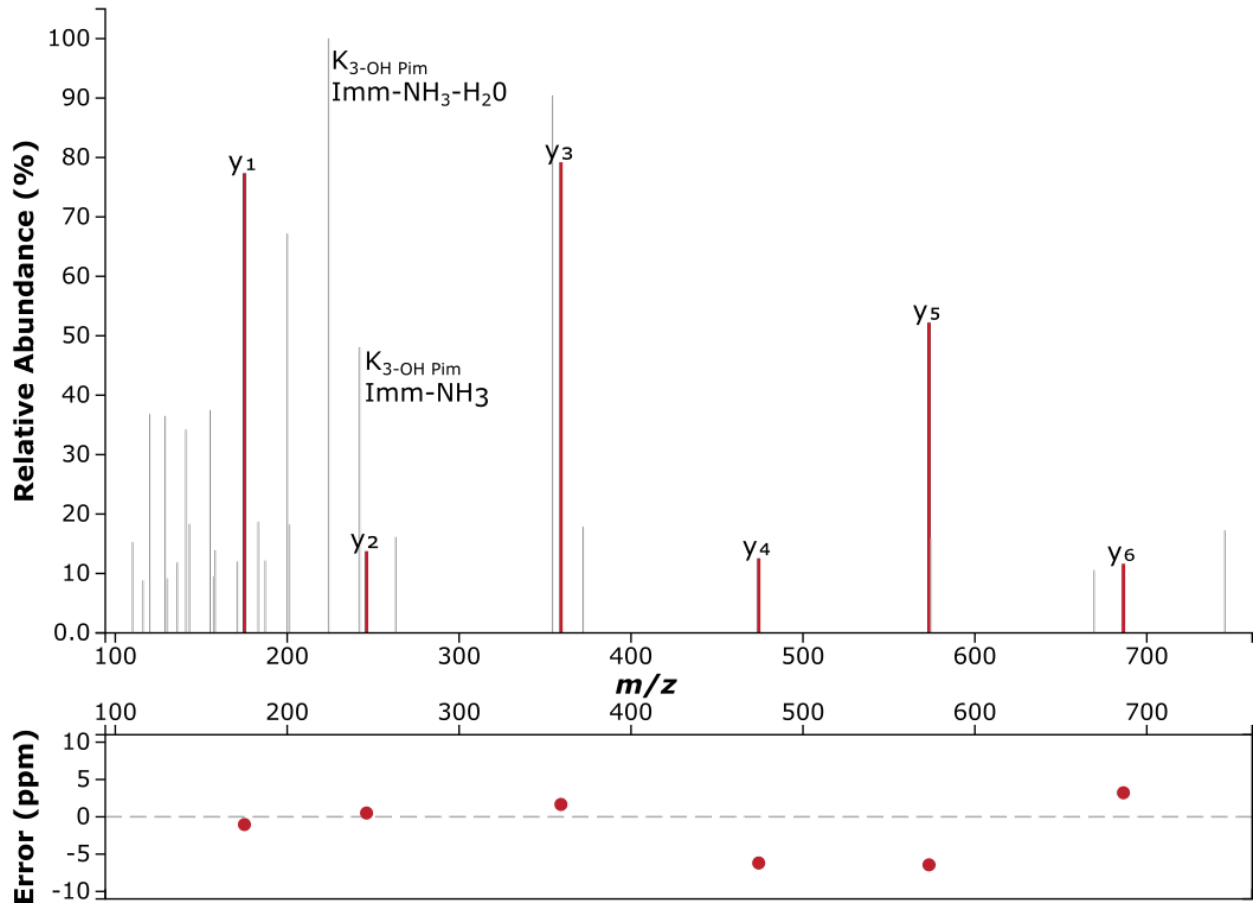
3-OH Pimelyl

k I V D I A R

Precursor m/z: 486.7898

Charge: +2

Aspartate aminotransferase (Q2LWK1)

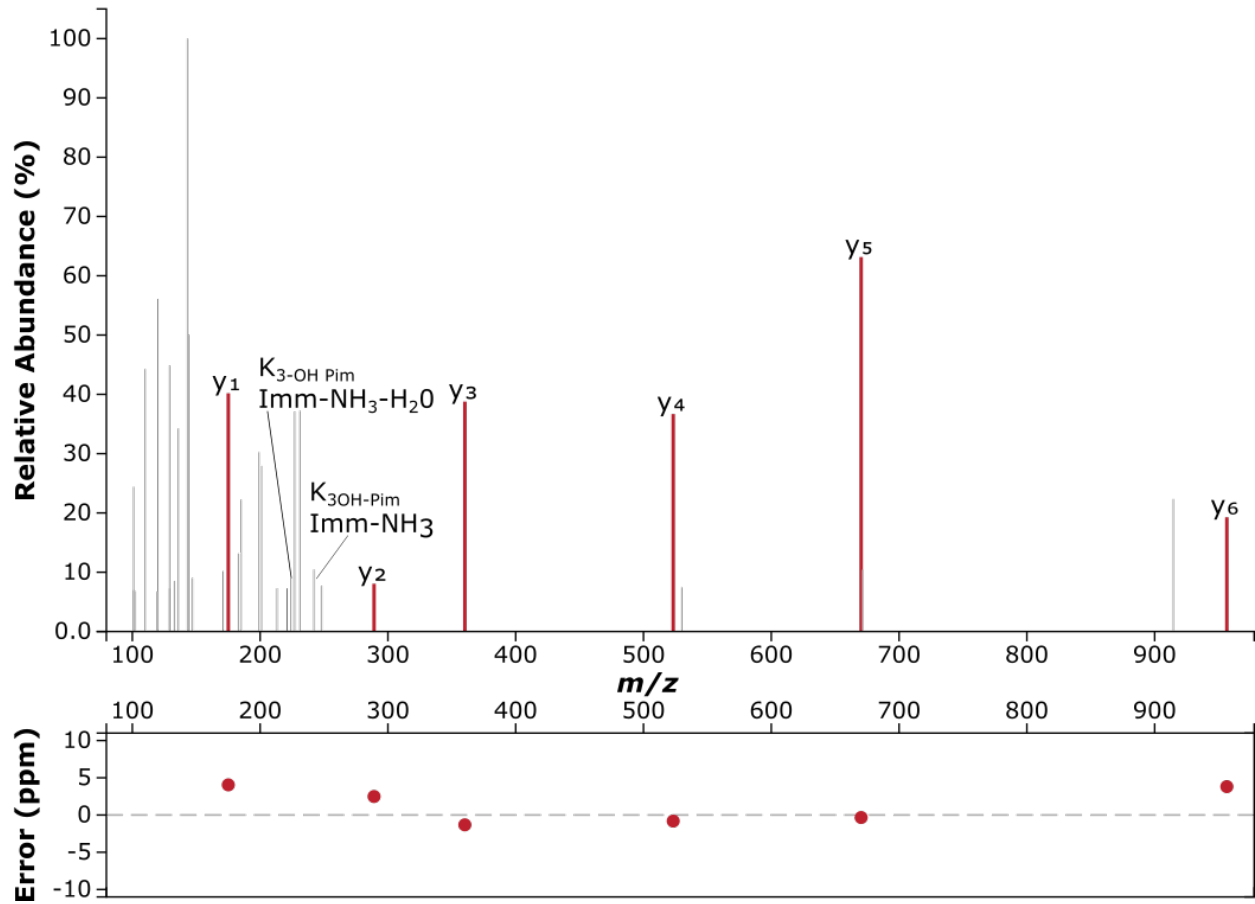


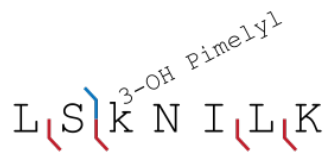
L P E Q H ^{3-OH Pimelyl} k F Y A N R

Precursor m/z: 520.9317

Charge: +3

Glutaconyl-CoA decarboxylase A subunit (Q2LUY7)

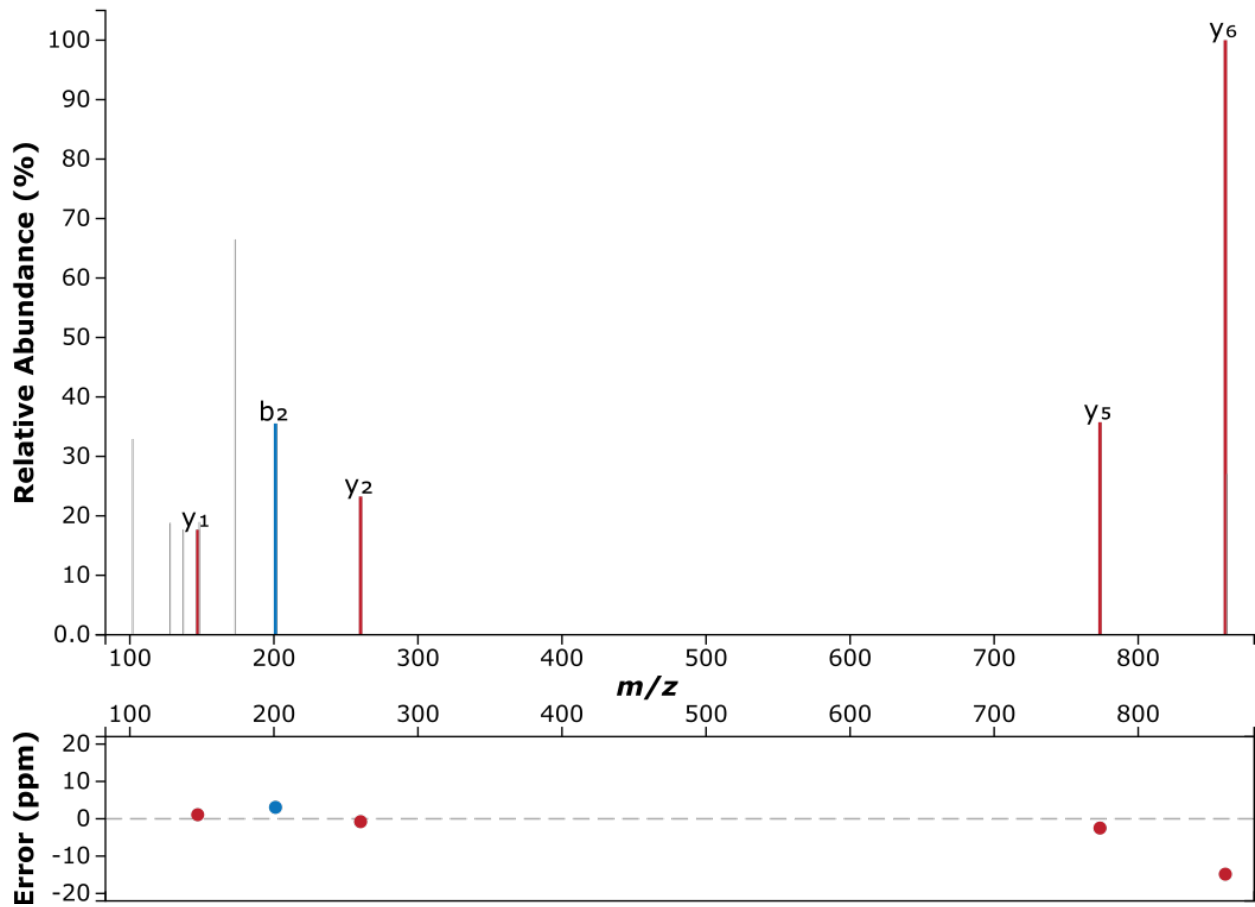




Precursor m/z: 487.3000

Charge: +2

Acetyl-CoA synthetase beta subunit (Q2LVF9)

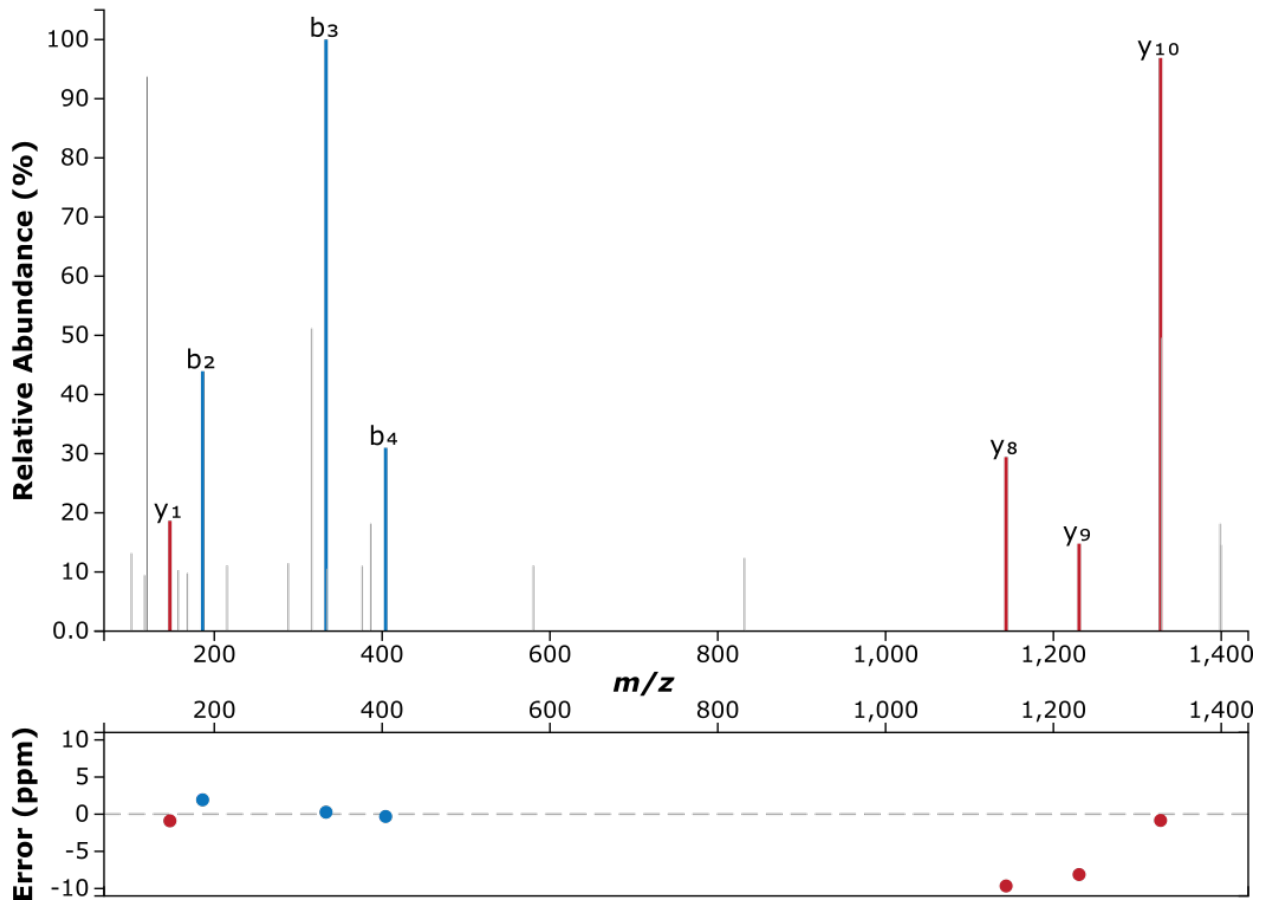


Q G F A P S P E L E k E L K ^{3-OH Pimelyl}

Precursor m/z: 865.9460

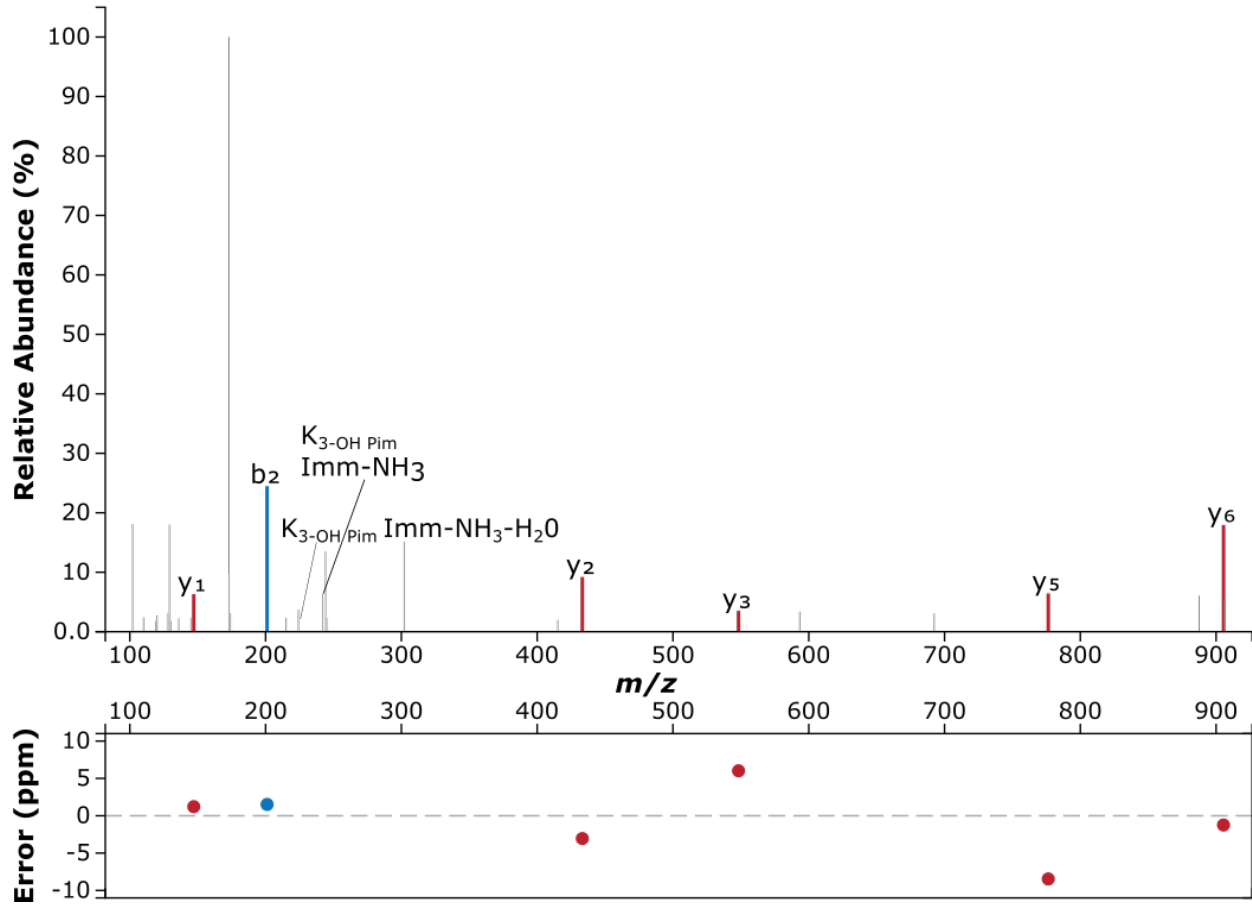
Charge: +2

4-hydroxybenzoate-CoA ligase / benzoate-CoA ligase (Q2LRH7)





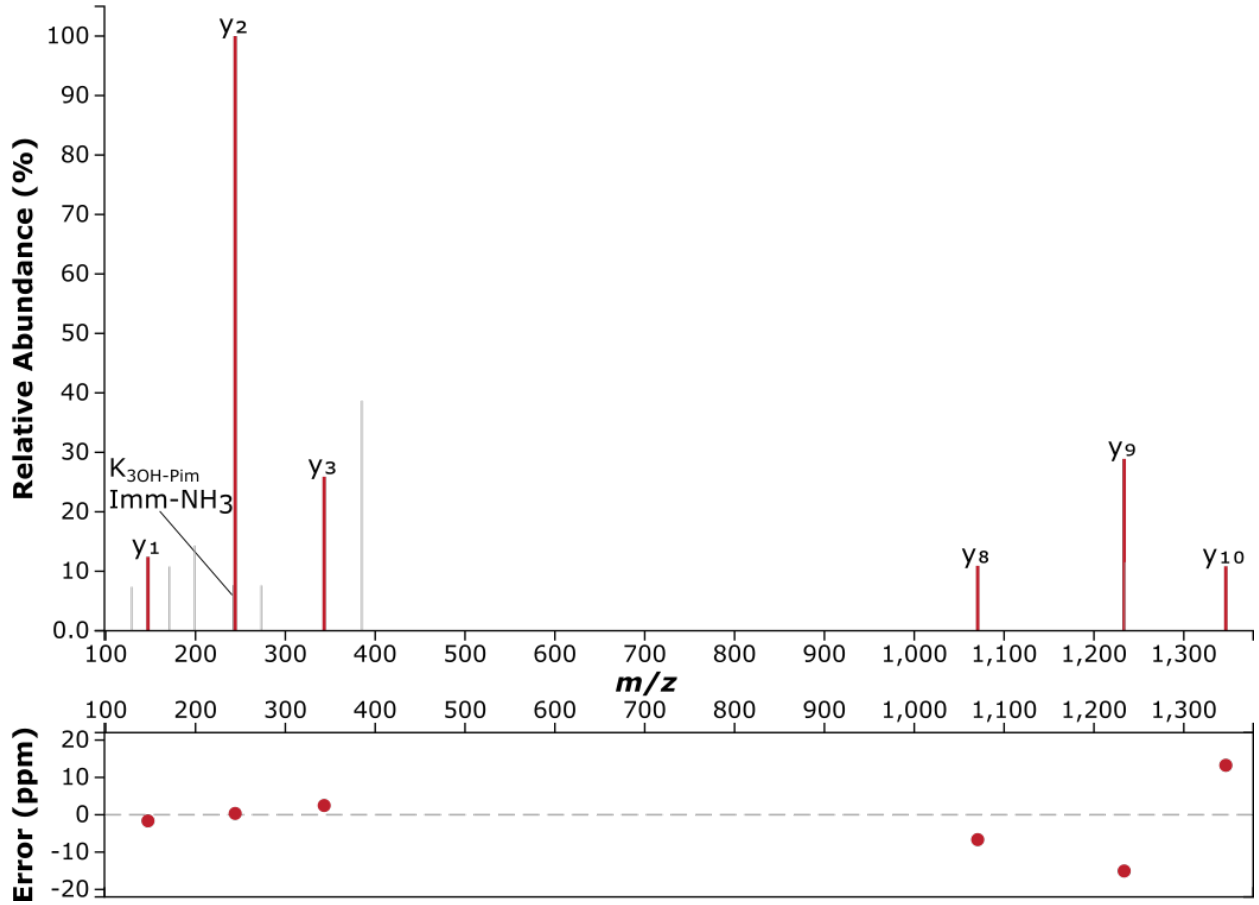
Precursor m/z: 553.2848 Charge: +2
 Phosphoglycerate mutase (Q2LTT8)



3-OH Pimelyl

S S P T G E I Y L D N V k V P K

Precursor m/z: 952.9962 Charge: +2
Acyl-CoA dehydrogenase, short-chain specific (Q2LUY8)

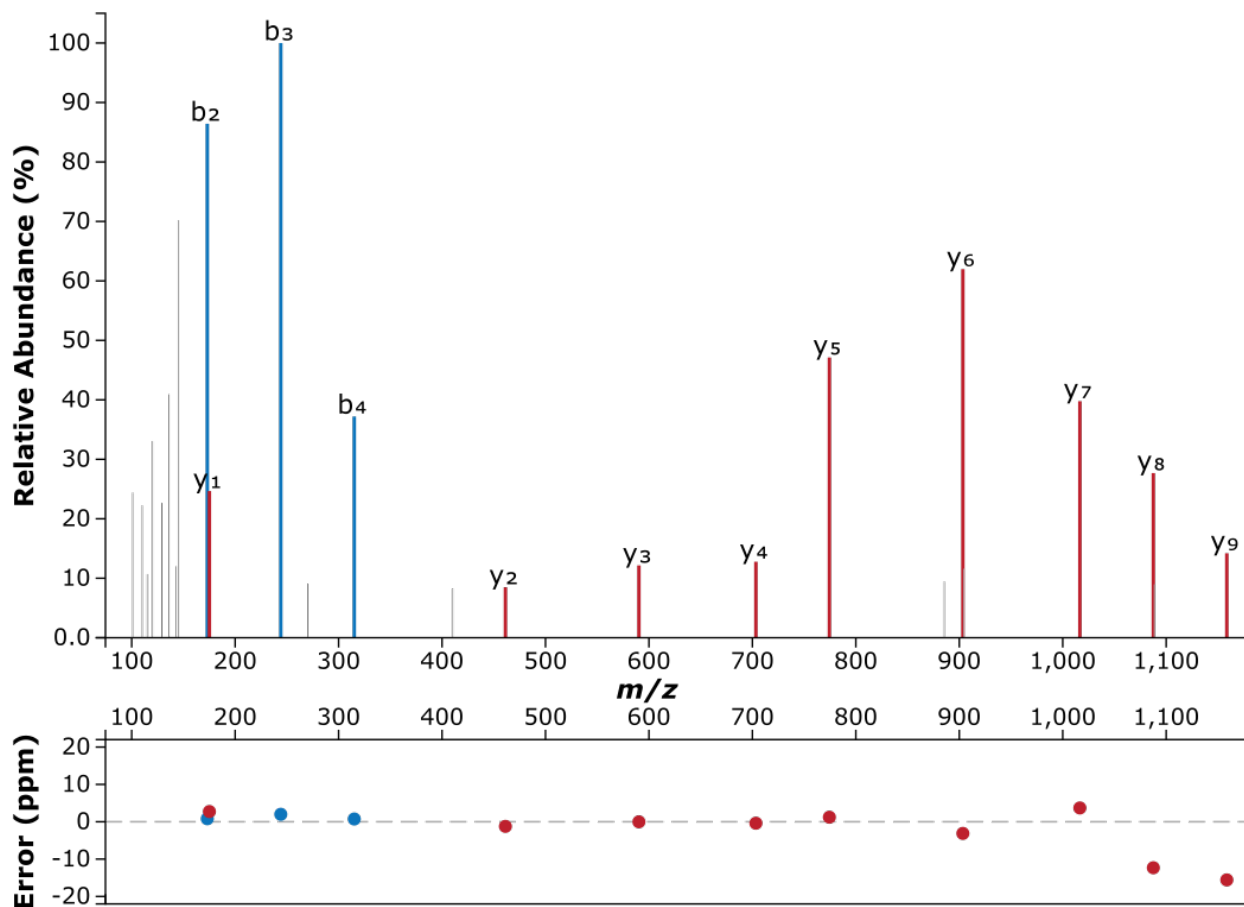


T A A A L E A L E k R ^{3-OH Pimelyl}

Precursor m/z: 665.8643

Charge: +2

Polyribonucleotide nucleotidyltransferase (Q2LWT4)

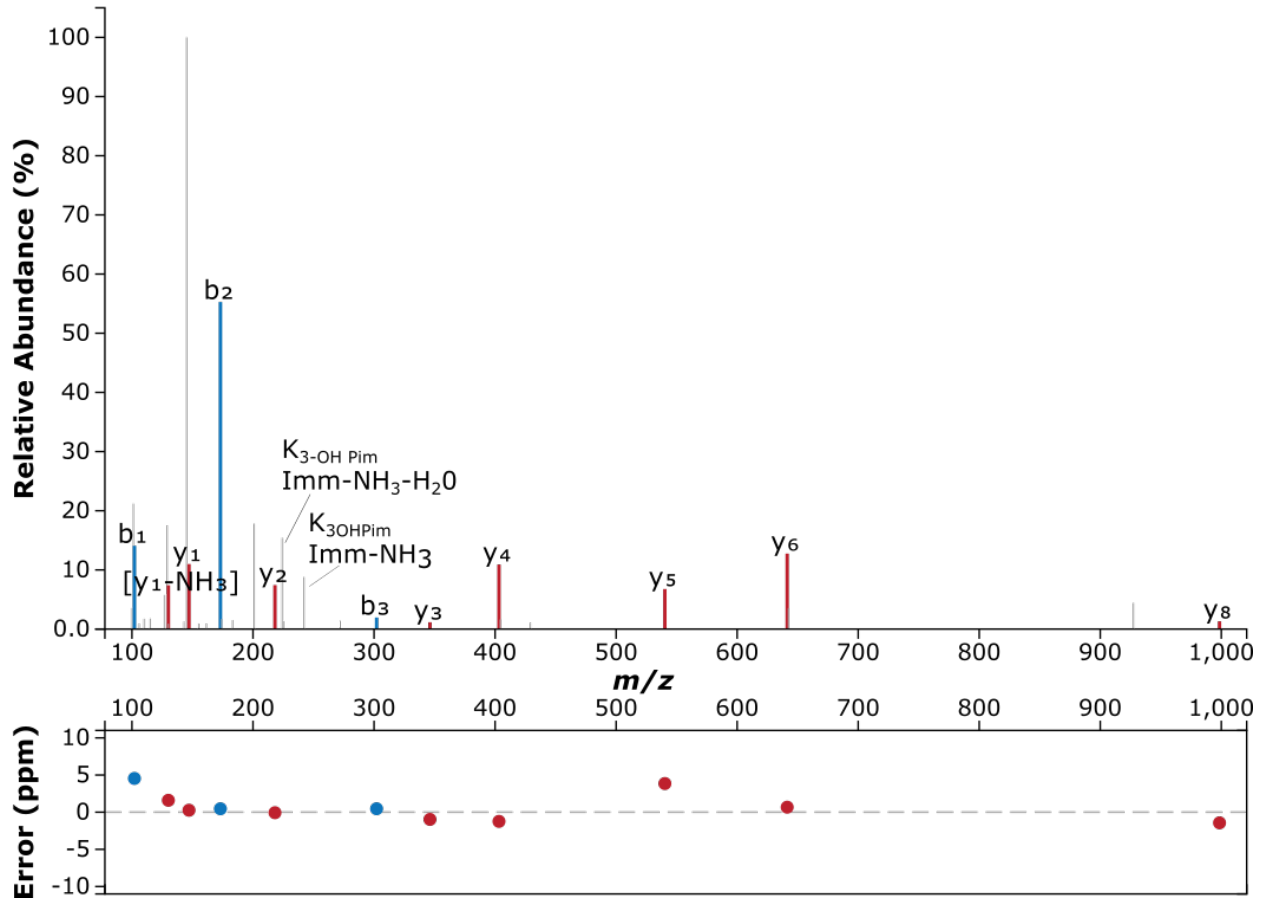


T A E A A k T H G Q A K
^{3-OH pimelyl}

Precursor m/z: 457.5685

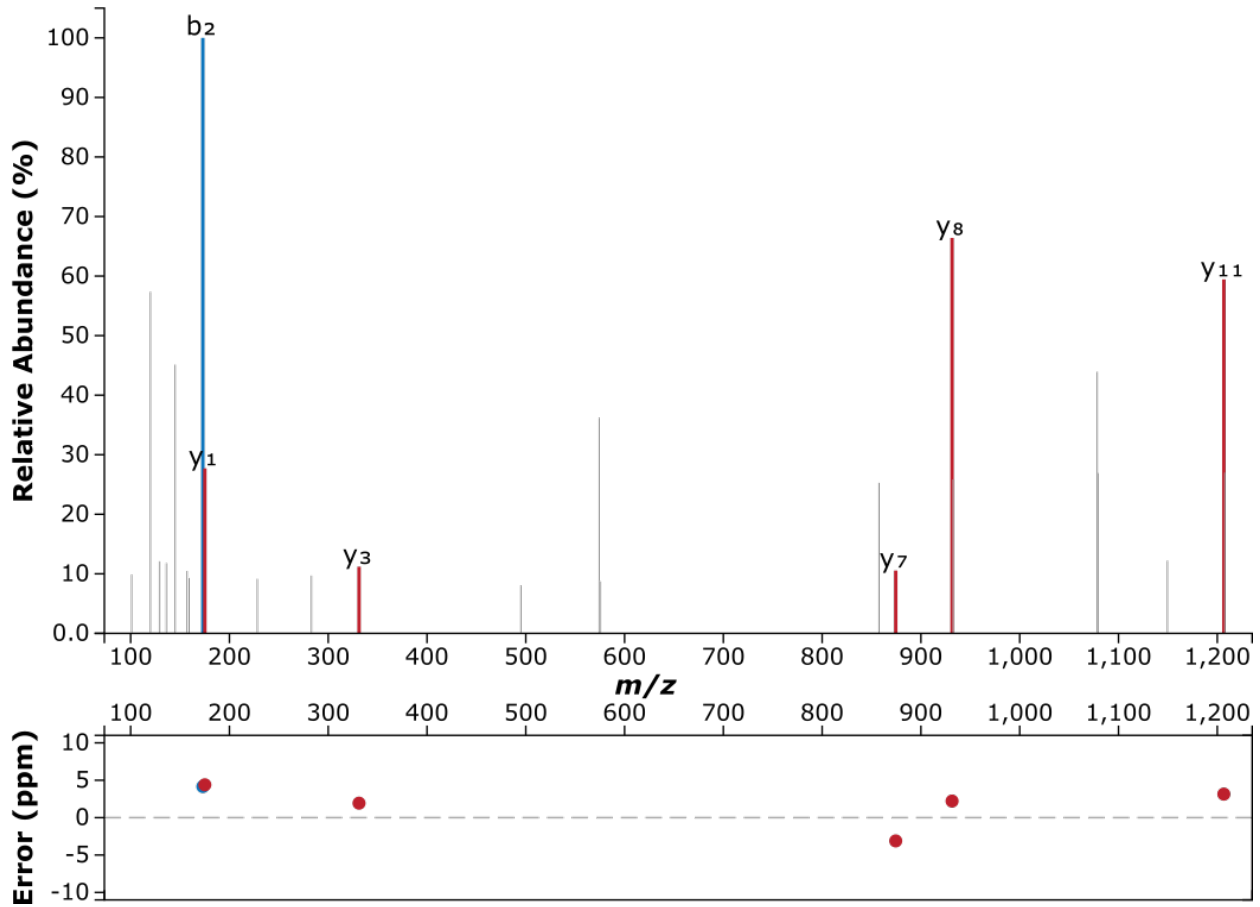
Charge: +3

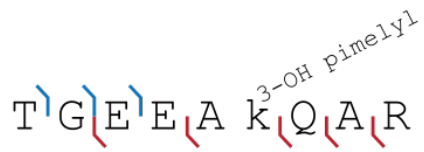
Putative K(+)-stimulated pyrophosphate-energized sodium pump (Q2LUL1)



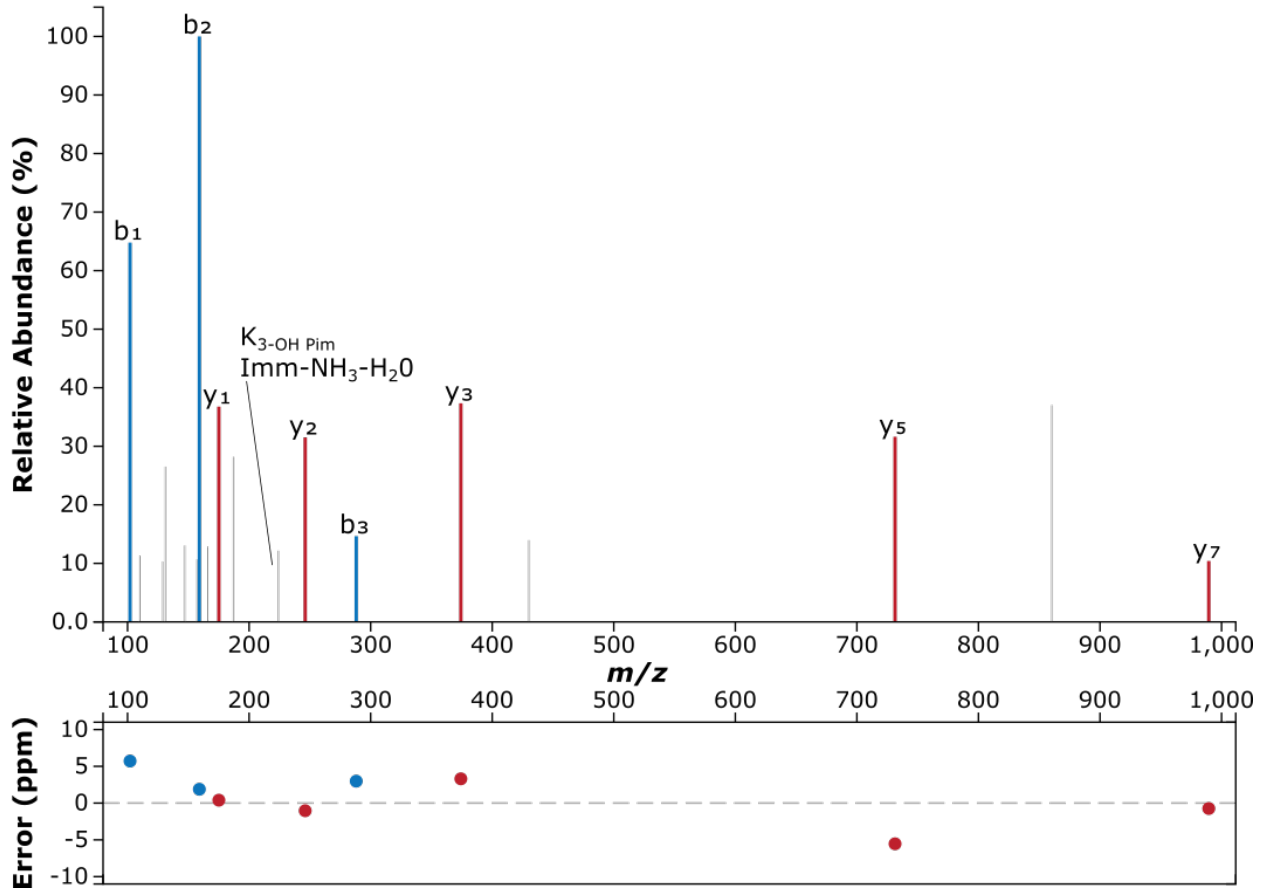
T A V G A F G G S L k G V R ^{3-OH Pimelyl}

Precursor m/z: 739.4041 Charge: +2
 Acetyl-CoA acetyltransferase (Q2LWQ5)





Precursor m/z: 574.2831 Charge: +2
 6-oxocyclohex-1-ene-1-carbonyl-CoA hydrolase (Q2LXU2)



V T A E k D P A N L K
^{3-OH Pimelyl}

Precursor m/z: 672.3563

Charge: +2

Glyceraldehyde-3-phosphate dehydrogenase (Q2LWW2)

