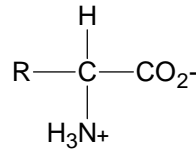


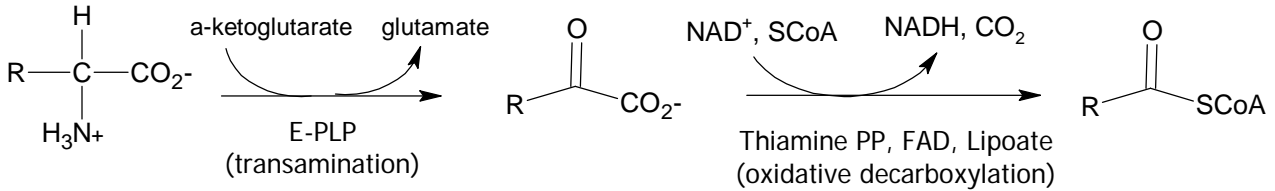
# Amino Acid Breakdown

## STEP I:

Amino Acids have the general structural motif of:



Convert the amino acid of interest to  $\text{R}-\text{C}(=\text{O})-\text{SCoA}$  using the following transformations:



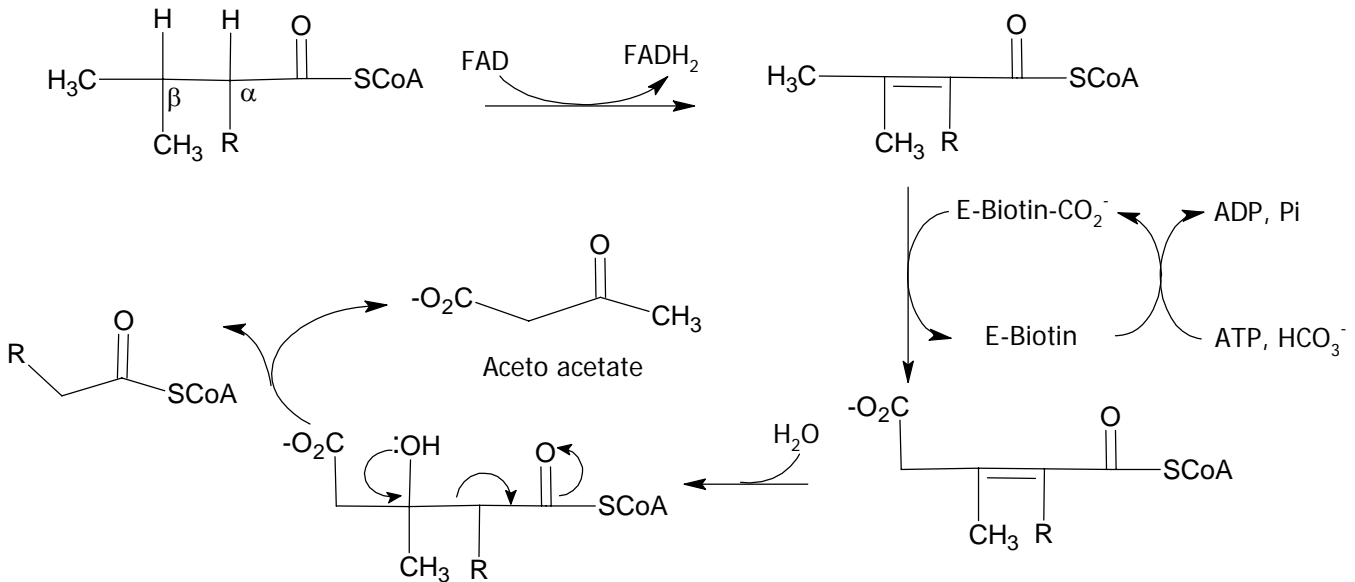
## STEP II:

Identify the LONGEST carbon chain backbone.

Examine the number of  $\beta$ -hydrogens on the longest carbon chain and follow the breakdown outline in the analysis below.

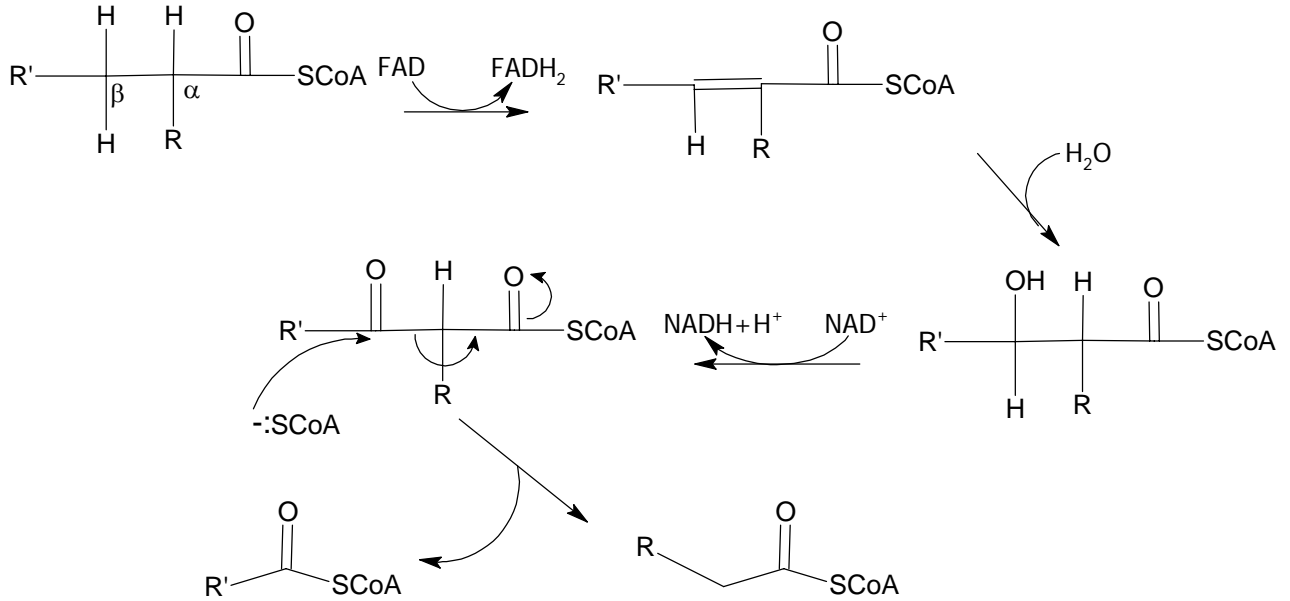
Note: Whenever there is a  $\beta$ -keto-acid present, spontaneous  $\beta$ -decarboxylation occurs.

## CASE I - one $\beta$ -hydrogen (analogous to Leucine breakdown)



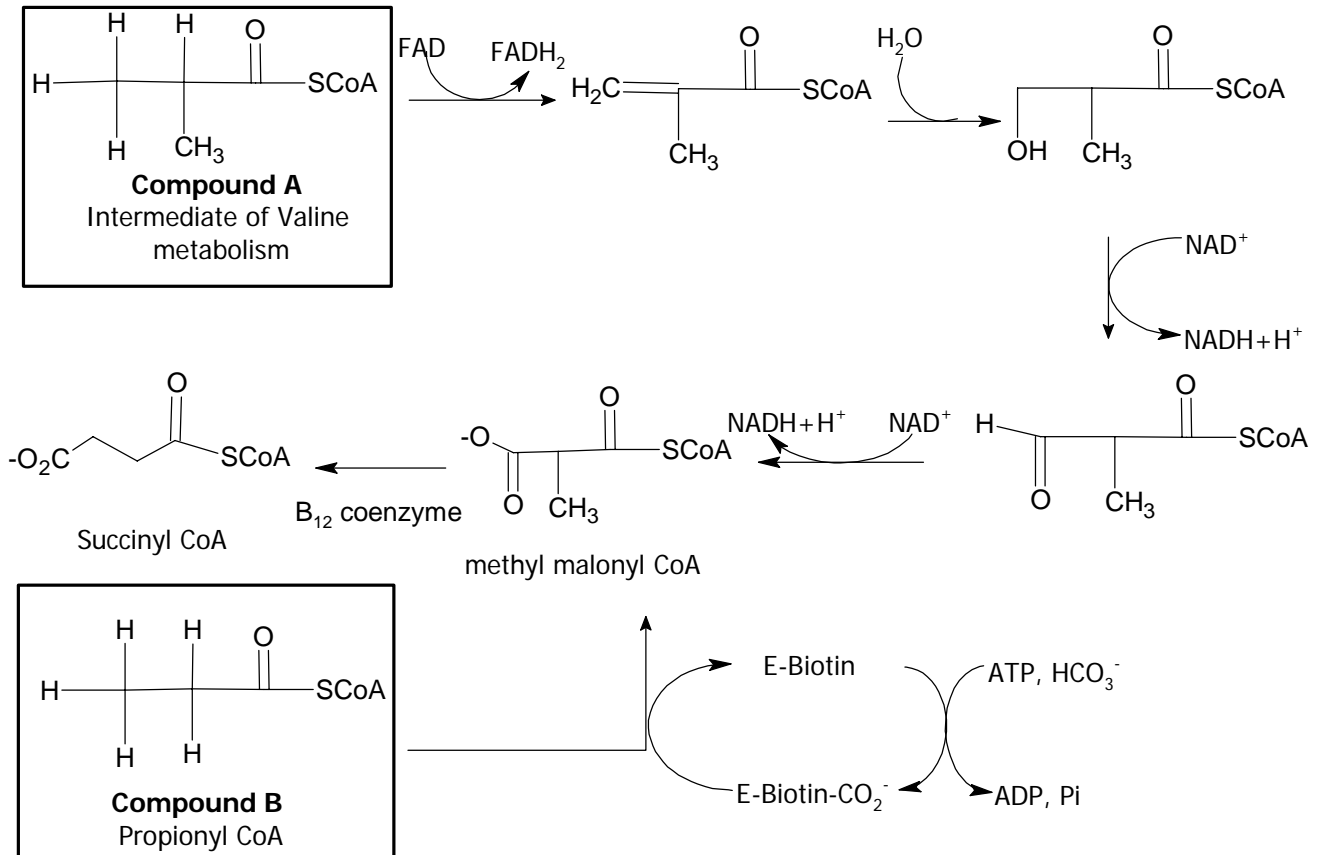
# Amino Acid Breakdown

**CASE II** - two  $\beta$ -hydrogens (analogous to Isoleucine and fatty acid breakdown)



**CASE III** - three  $\beta$ -hydrogens (analogous to valine breakdown)

Only two molecules fit this category.

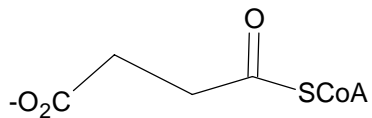


# Amino Acid Breakdown

## STEP III:

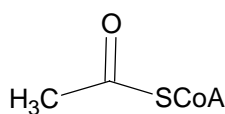
Repeat Step II until you obtain one of the following molecules.

**Glycogenic**

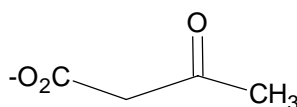


Succinyl CoA

**Ketogenic**



Acetyl CoA



Aceto acetate