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# Opportunities to Redefine Carbon Partitioning in Soybeans



DONALD DANFORTH  
PLANT SCIENCE CENTER

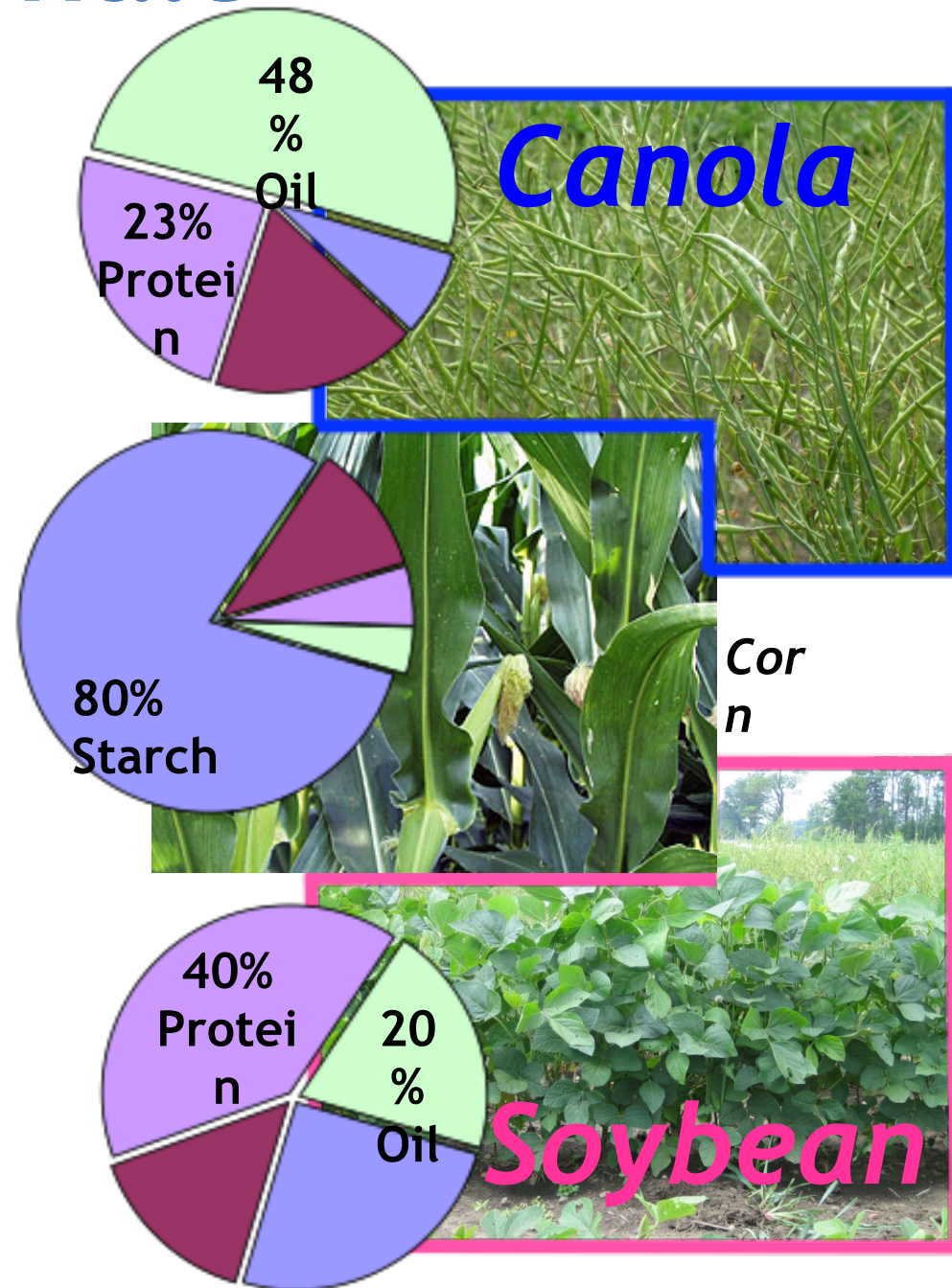
DISCOVERY | COMMUNITY | IMPACT

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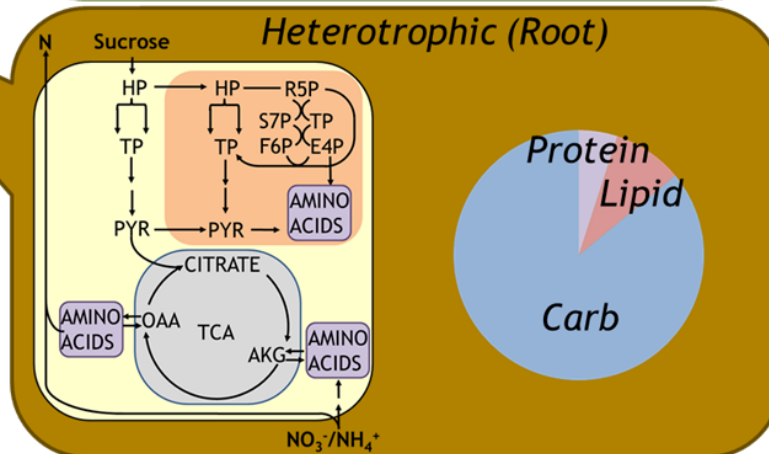
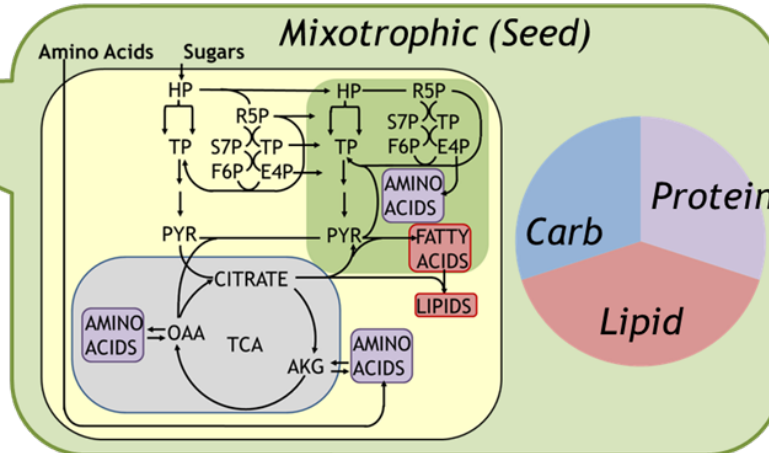
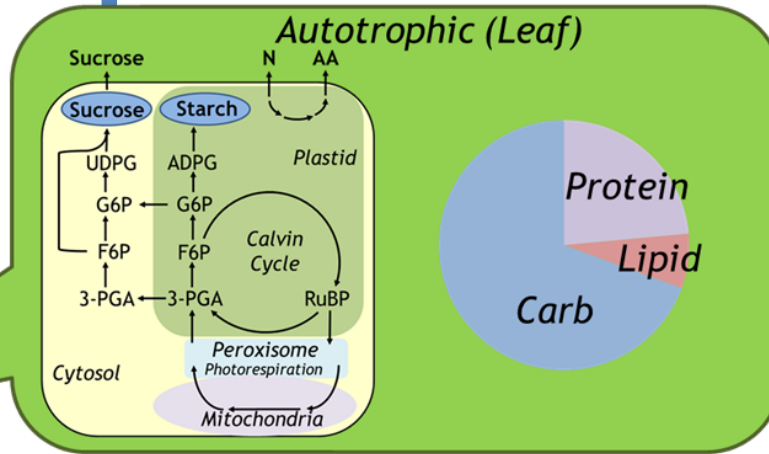
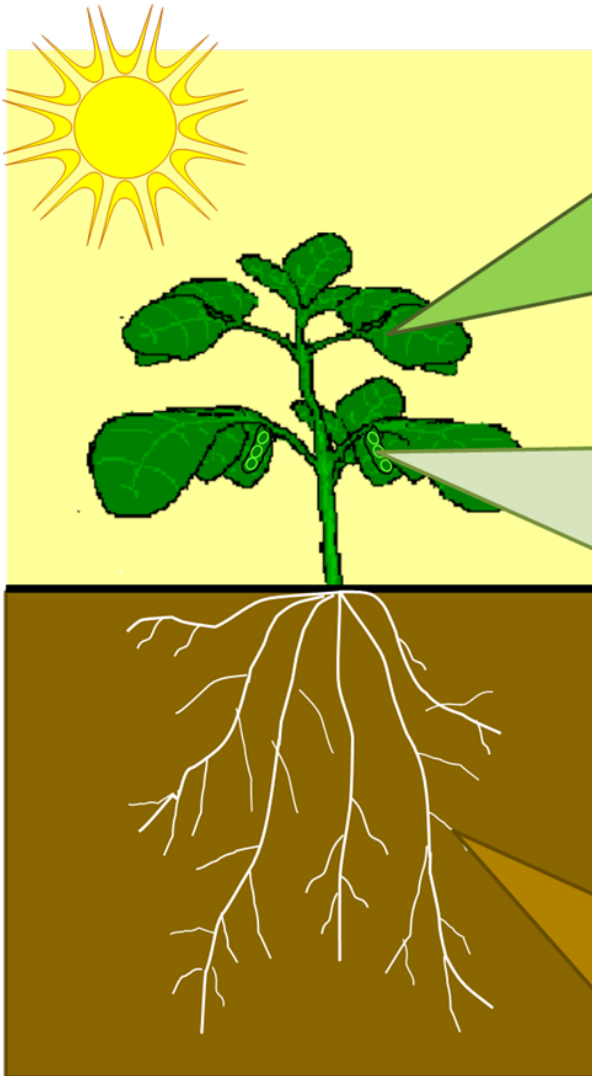


# Rationale

- Crop composition varies
- Storage reserves are generated by central carbon metabolism
- Carbon and energy partitioning varies among species
- “Engineering” benefits (petroleum replacements & improved nutrition)



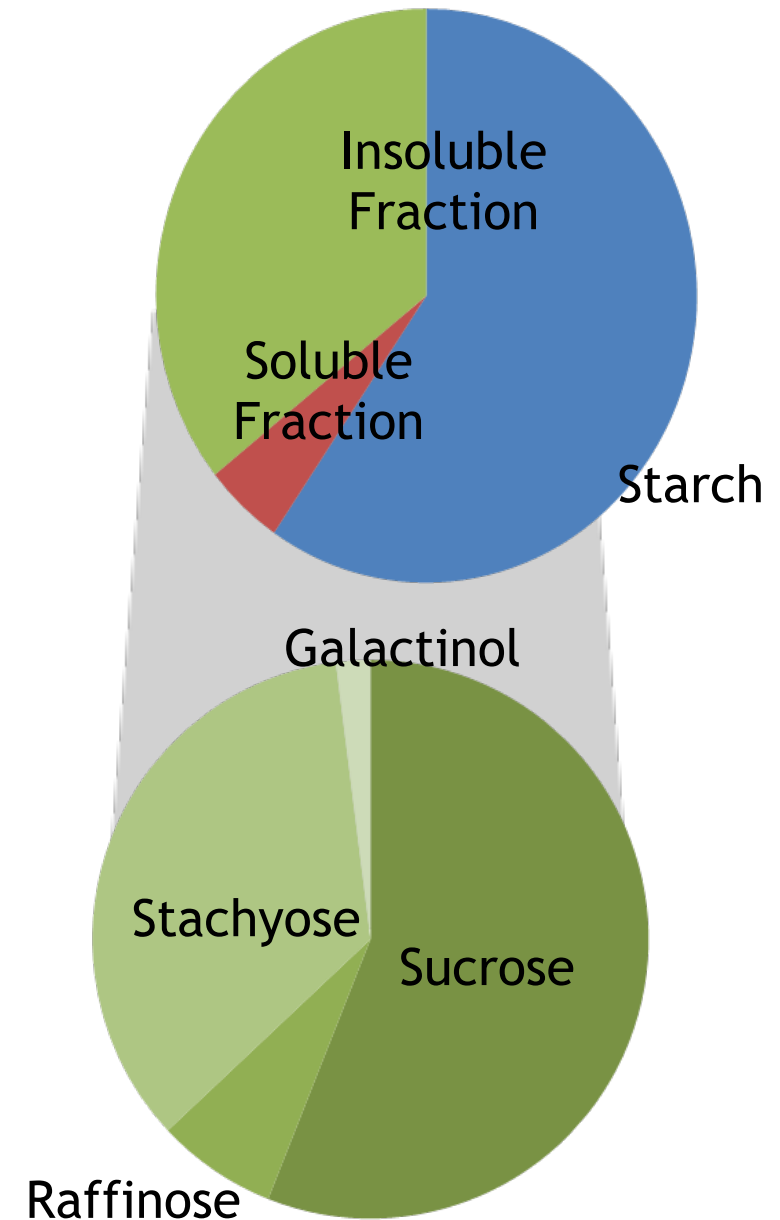
# Cells Operate Differently



- Different compositions
- Different metabolism
- Many of the same enzymes
- Enzyme operation is specific to

# Carbohydrate Profile

- Cell wall/insoluble fiber → 20-25%
- Starch (decreased at maturation) → <4%
- Soluble carbohydrates → ~10-15%
  - sucrose/galactinol
  - oligosaccharides: raffinose, stachyose



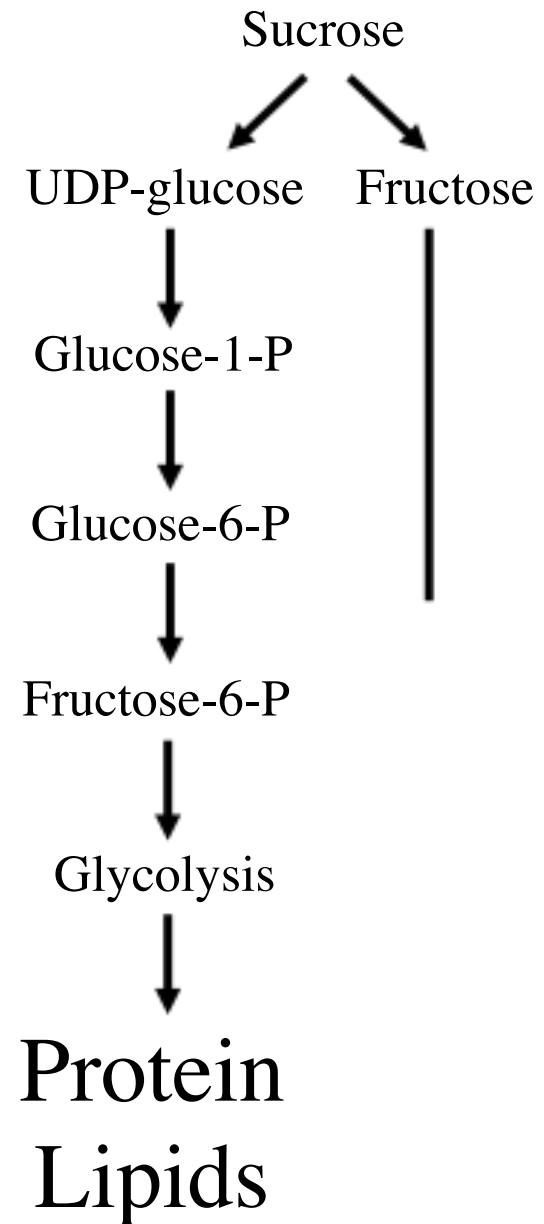
# Reducing Oligosaccharides

(Bilyeu, Dierking, Wiebold, Fehr, Schillinger, Obendorf  
some work sponsored by USB)

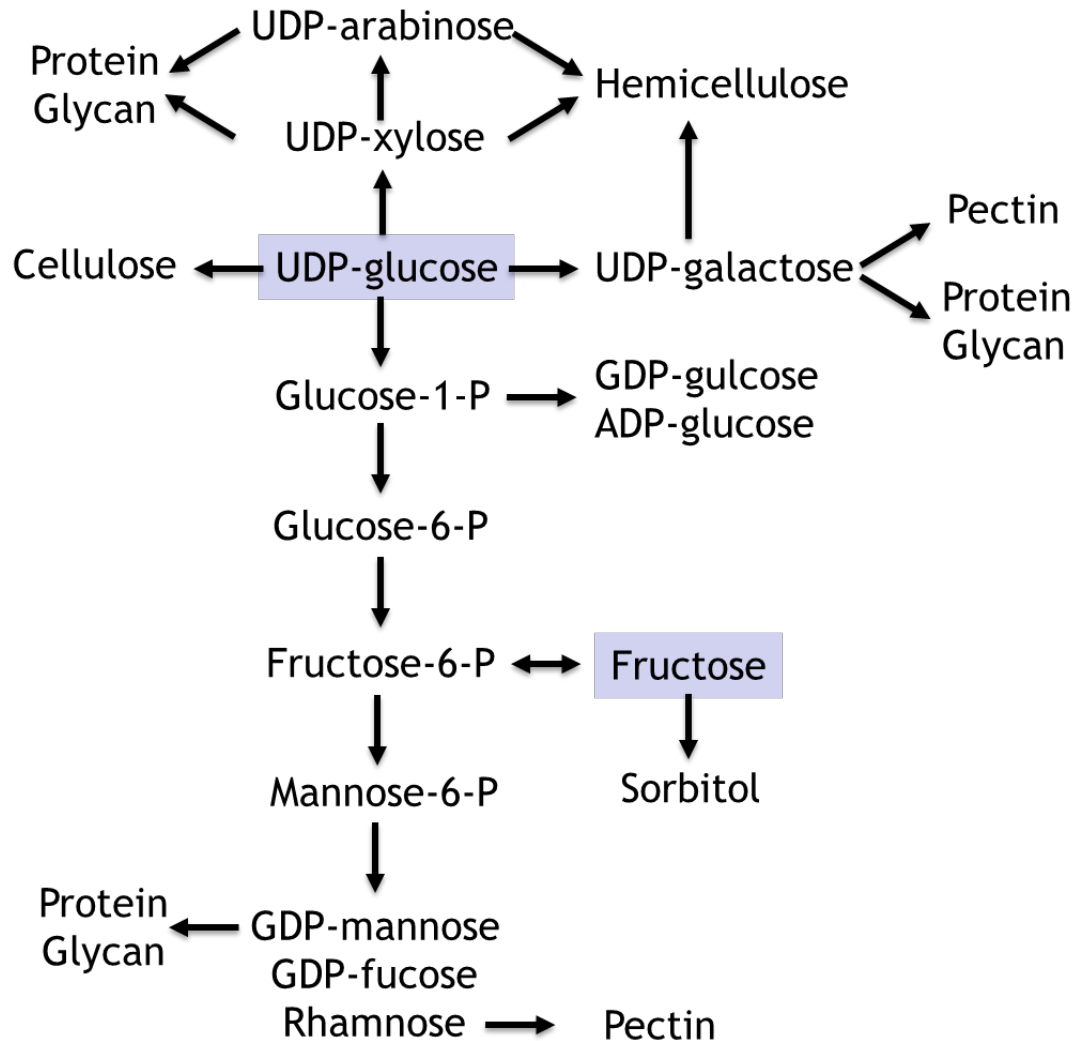
- Impact metabolizable energy negatively
- Considered as anti-nutritional factors
- Soluble carbohydrates not critical for desiccation tolerance (no reduction in germination/emergence) (Obendorf 1997; Schillinger et al Patent 8471107, Neus et al 2005)
- Variant alleles with altered genes (e.g. RS2) produce less oligosaccharides,
  - more sucrose (up to 85% of soluble carbohydrate)  
(Hageley, Palmquist and Bilyeu 2013)
  - higher metabolizable energy in soy meal

# What *FURTHER* can be done?

- Can insoluble components be reduced?
  - UDP-glucose metabolic partitioning
  - sucrose is imported in seeds
- Can sucrose be converted to something of higher value?
  - protein?
  - oil?
  - transcription factors such as *Wrinkled 1*



# Wall Biosynthesis from Glycosides including UDP-glucose & Fructose



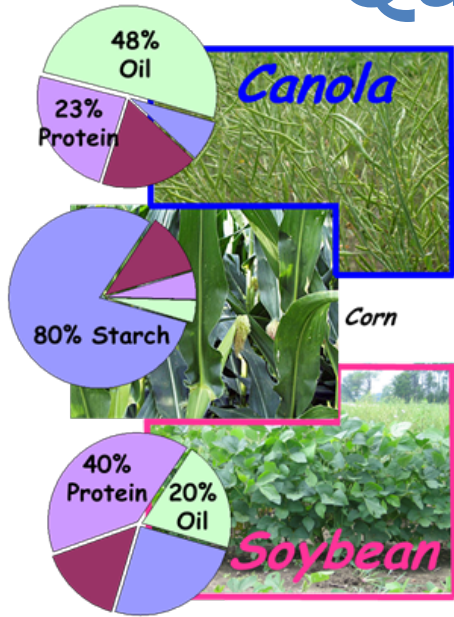
- Reduced glycosyltransferase mutants indicate 20% reduction in arabinose content (Egelund et al 2007)
- Structural and protective functional features of cell wall limit complete removal
- Cell wall biosynthesis is still unknown, though composition can be highly variable
- Removing recalcitrant wall

# Storage Reserve Generation

- Soybeans require a lot of nitrogen for protein biosynthesis
- Balance of amino acids with sugars
  - Amino acid transporters impact seed protein levels (Sanders et al. 2009)
- Protein and Oil and Protein and Yield are inversely related → different inputs from maternal plant
- Mechanisms determining differential partitioning of seed reserves remains unknown



# Quantify Cellular Metabolism

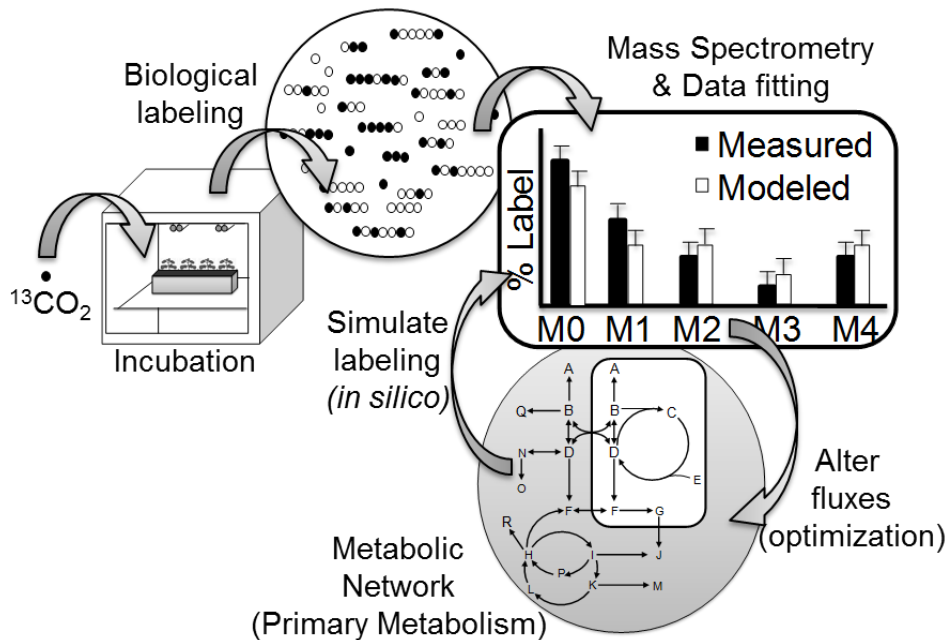


Plant seed composition varies: oil, protein and carbohydrate

Engineering nutrition & chemical feed stocks



Different metabolic functions:  
Seeds = storage or germination  
Leaves = sink then photosynthesis



Isotopic labeling & computational metabolic flux analysis: central carbon, lipid and protein metabolism

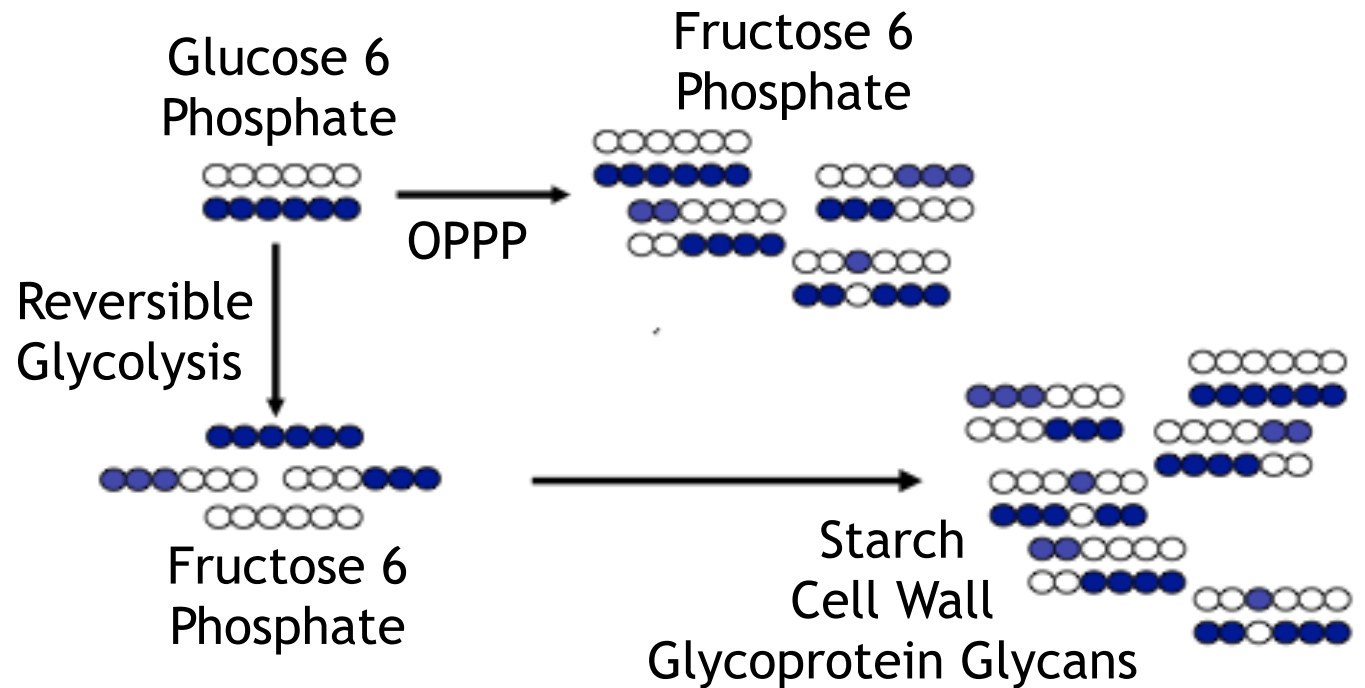
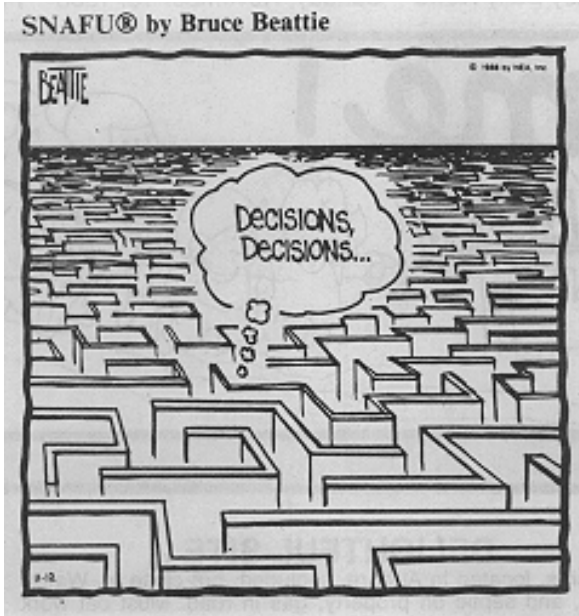
# Questions/Aims

- What are *ROUTES* & *RATES* of carbon flow?
- What are the *SOURCES* of energy & redox for storage reserve production?
- What is the impact of environmental or transgenic perturbations

# Approaches

- Measure uptake/export rates & composition
- Balance energy & cofactor supply & demand
- Perform isotopic labeling experiments and measure metabolite enrichment
- Develop flux maps with computational tools

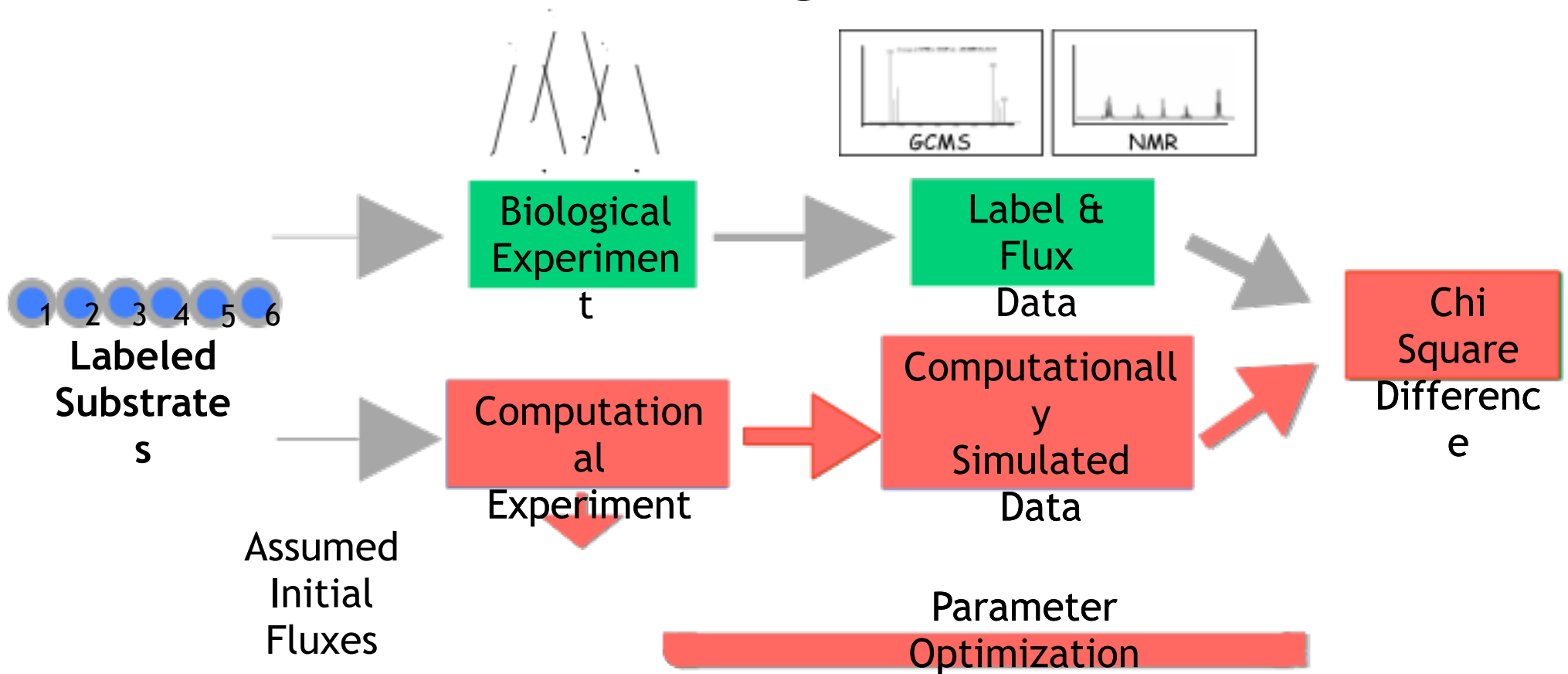
# Isotopic Tracers



**Flux analysis allows quantification of different pathways**

**Fluxes determine the balances of redox, energy and elements**

# Determining Flux Values



Model calculates labeling from flux values

Compare measured and simulated labeling patterns and minimize differences to obtain best fit flux map

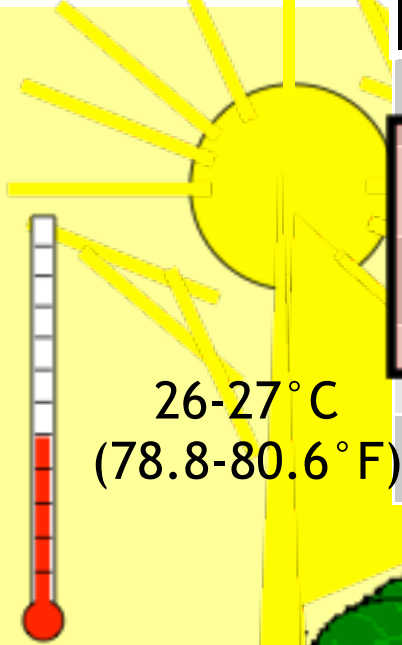
*Inspired by Wiechert 2001*

Forward problem  
Inverse problem



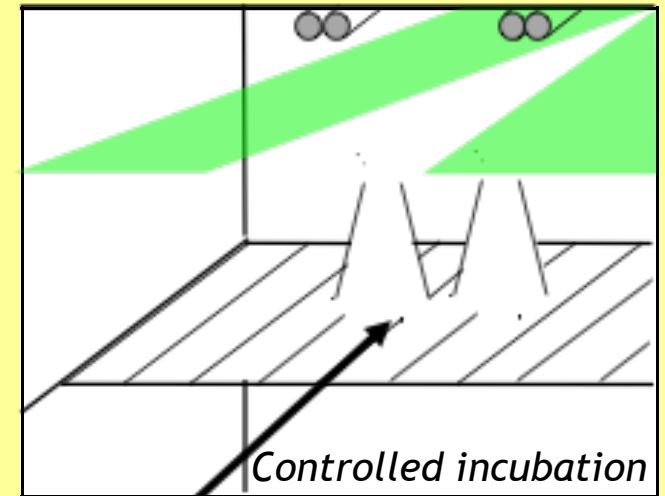
# Environmental Variables & Culturing Conditions

C:N	Sucrose	Glucose	Glutamine	Asparagine
6	90	45	140	50
13	140	70	70	25
21	150	75	45	16
37	170	85	25	12
91	180	90	11	4



26-27°C  
(78.8-80.6°F)

30  $\mu$ E  
light

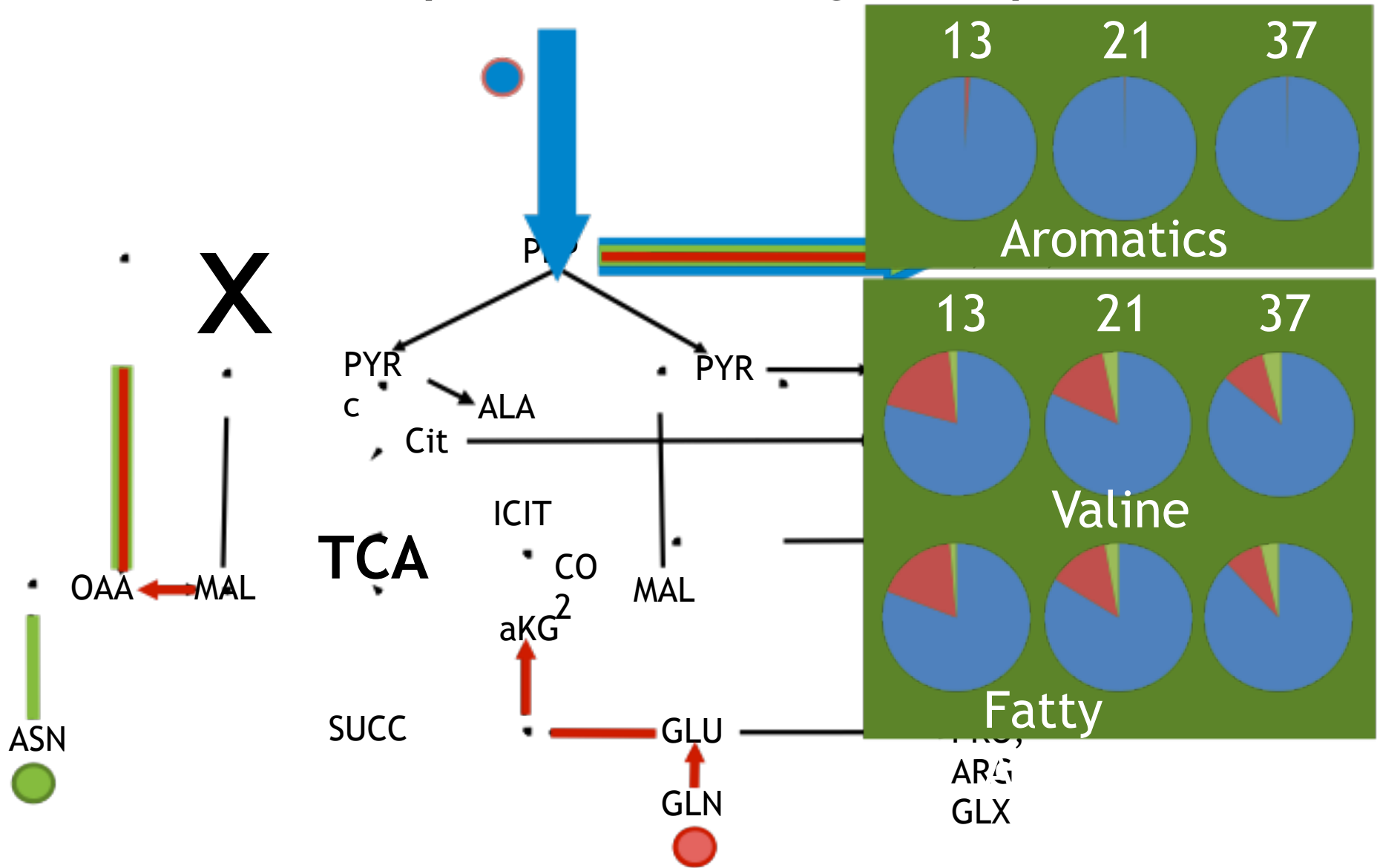


Starting Fresh  
Weight: 25-40mg

14 day culture experiments

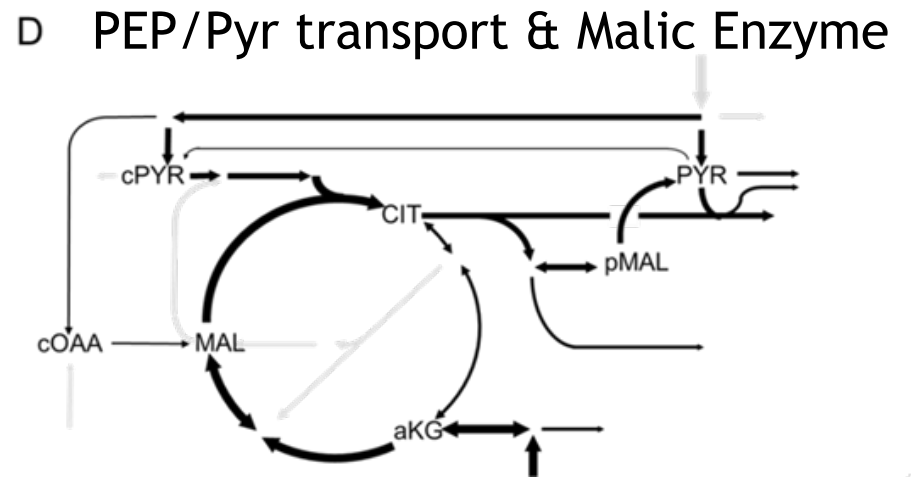
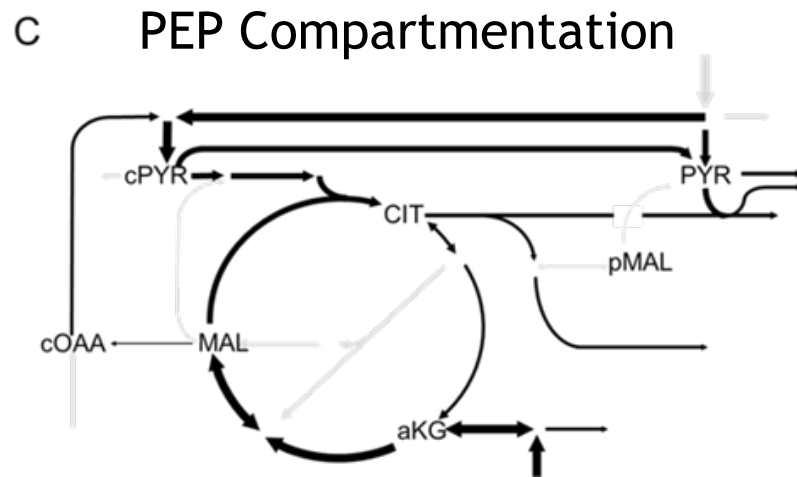
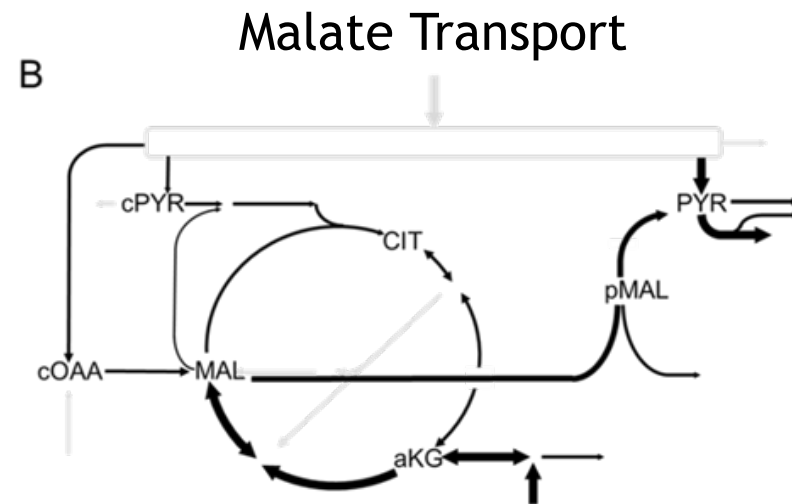
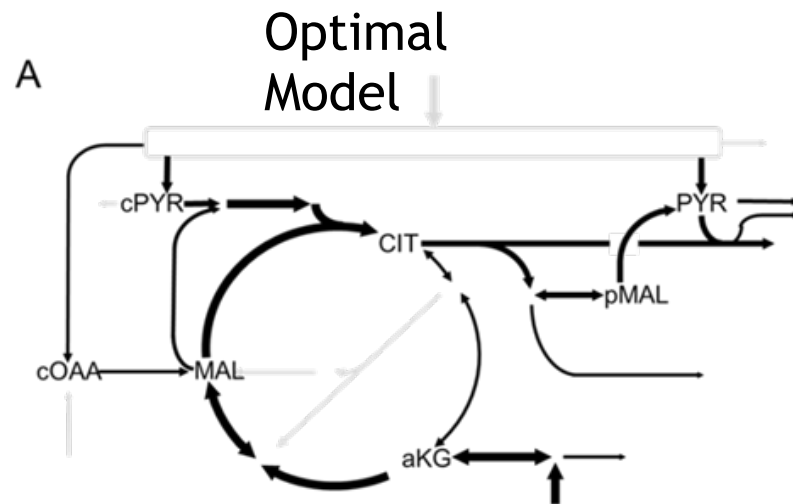
*Allen and Young 2013, Plant Phys.*  
*Allen et al. 2009, Plant J.*

# PEPCK is not required for labeling descriptions

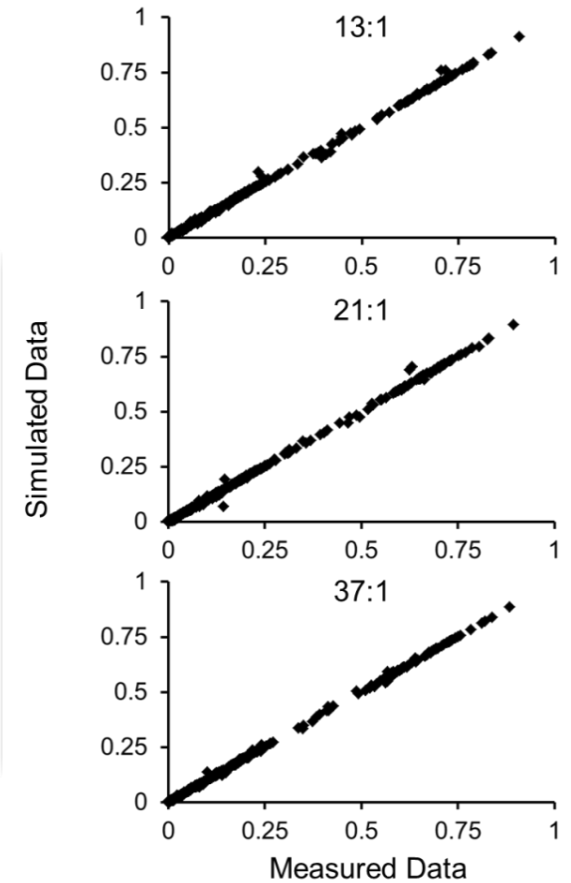
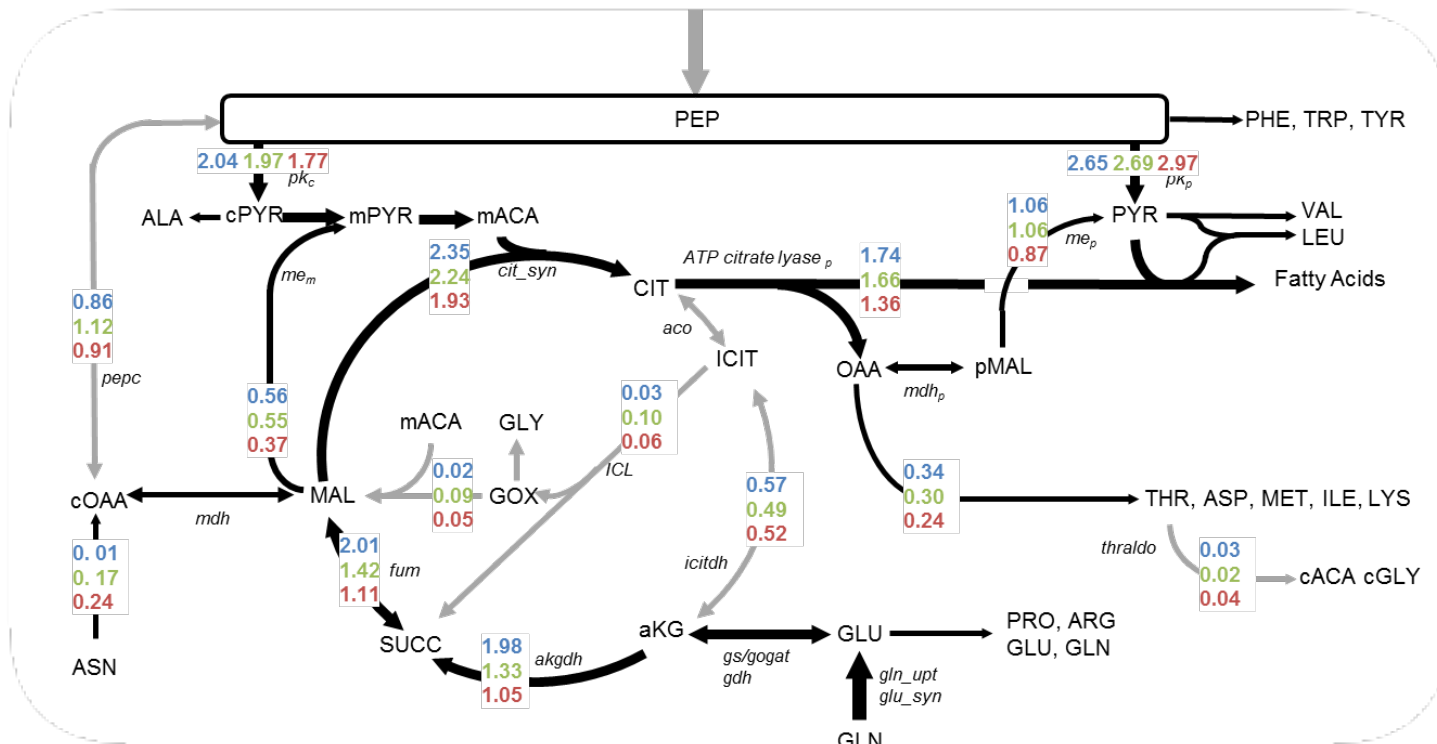


- PEPCK activity frequently unknown
- Modeling results: PEPCK activity is not necessary for fit

# Models Considered



# Fluxes vary for C:N ratios

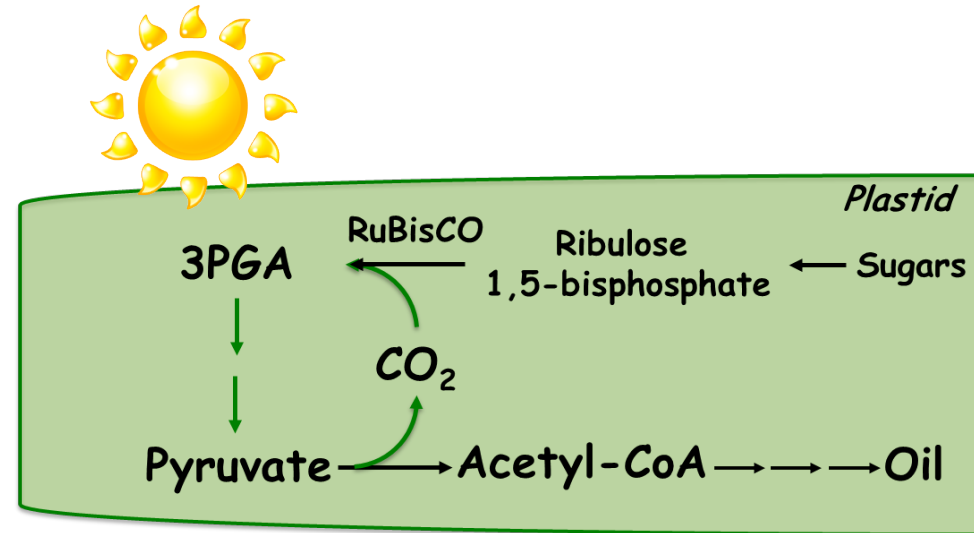




# Metabolism in Green Oilseeds

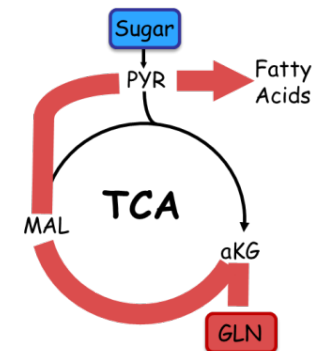
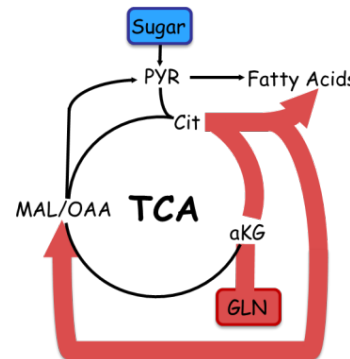
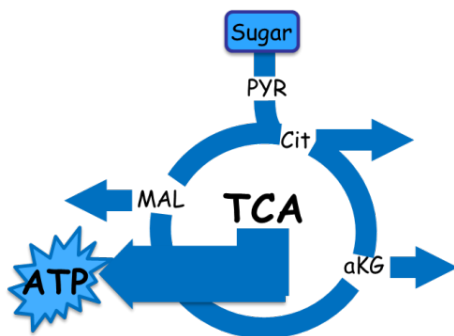
- Hypothesized Roles for Light:

- Prevent hypoxia
- Produce NADPH, ATP
- Improve carbon use efficiency



- Altered Tricarboxylic Acid Cycles:

- Cyclic vs. non-cyclic
- Citrate export for cytosolic acetyl-CoA
- Amino acids supplied maternally

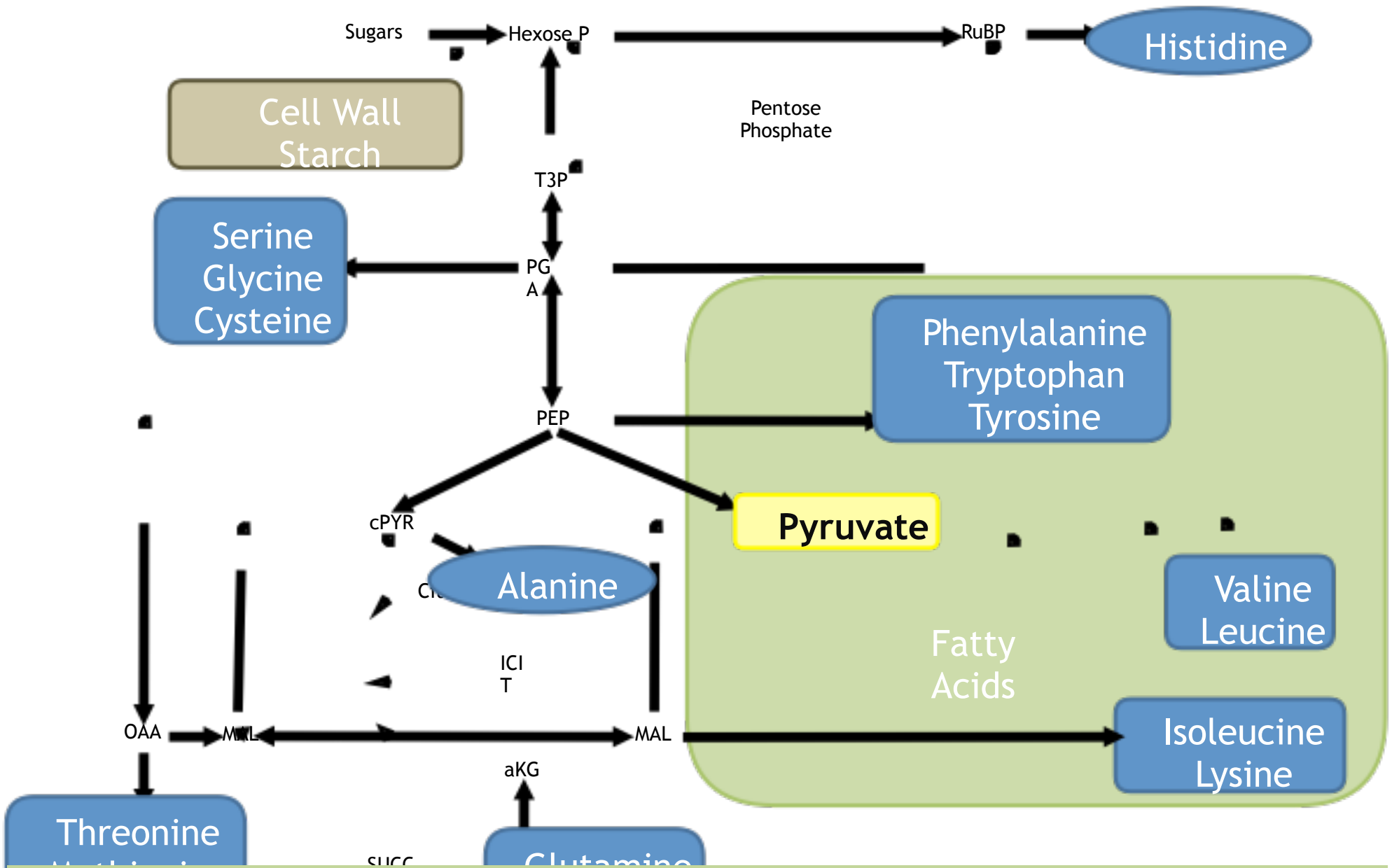


# 14C Labeling Experiments (Balancing Carbon)

C:N Ratio	Biomass growth (mg day <sup>-1</sup> embryo <sup>-1</sup> )	Protein (DPM)	Oil (DPM)
13	5.0 ± 1.3	27809 ± 2684	9279 ± 513
21	4.8 ± 0.7	24324 ± 4217	10523 ± 717
37	5.1 ± 0.4	20860 ± 1070	14163 ± 1273

- Oil production changes inversely with protein on a carbon basis
- Competition for carbon

# Pyruvate Carbon Used for Oil or Protein



If amino acid production is coordinated, a small change in oil will have more dramatic impact on protein levels

# Summary

- Oligosaccharides have been successfully diverted to sucrose
- To move sucrose to protein or oil will require explorations in central metabolism
- Can the insoluble fraction within a seed be reduced?
- Protein and oil are inversely tied, possibly through the node of pyruvate



# Acknowledgements

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