

15th Annual Conference of the Metabolomics Society

METABOLOMICS 2019

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Unveiling non-cyclic flux modes in the plant TCA cycle by GC-MS-based ^{13}C -positional labelling



Prof. Dr. Danilo de Menezes Daloso

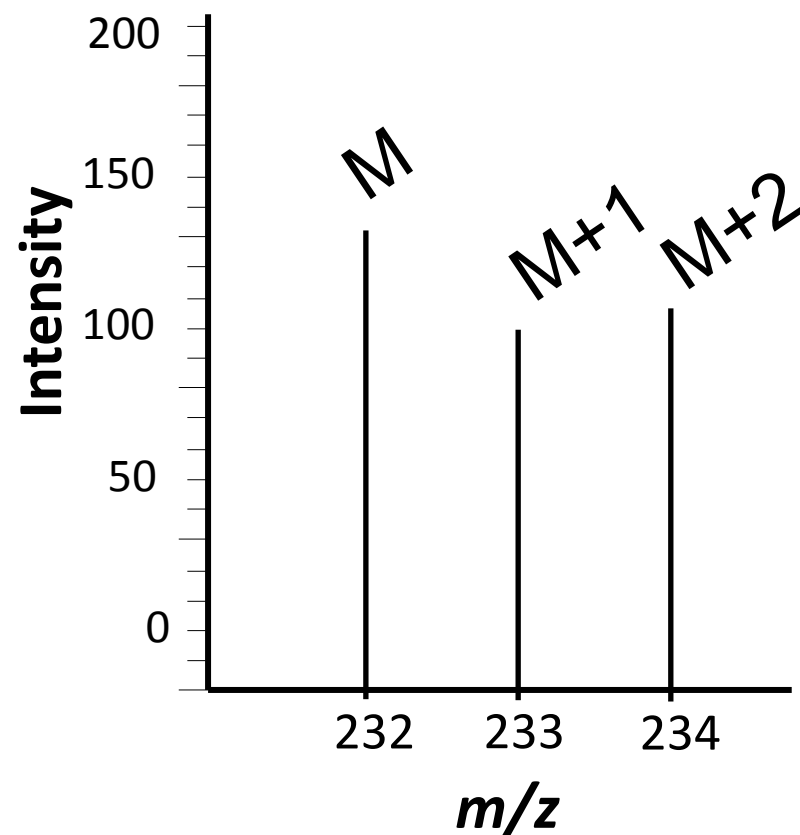
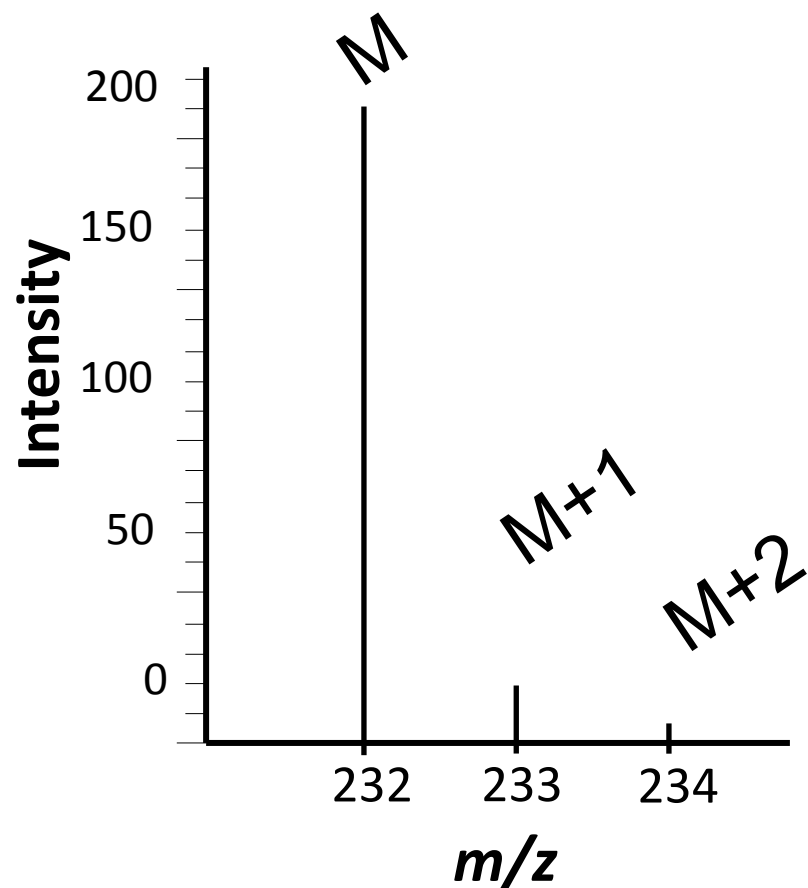
Federal University of Ceará, Fortaleza/CE, Brazil.

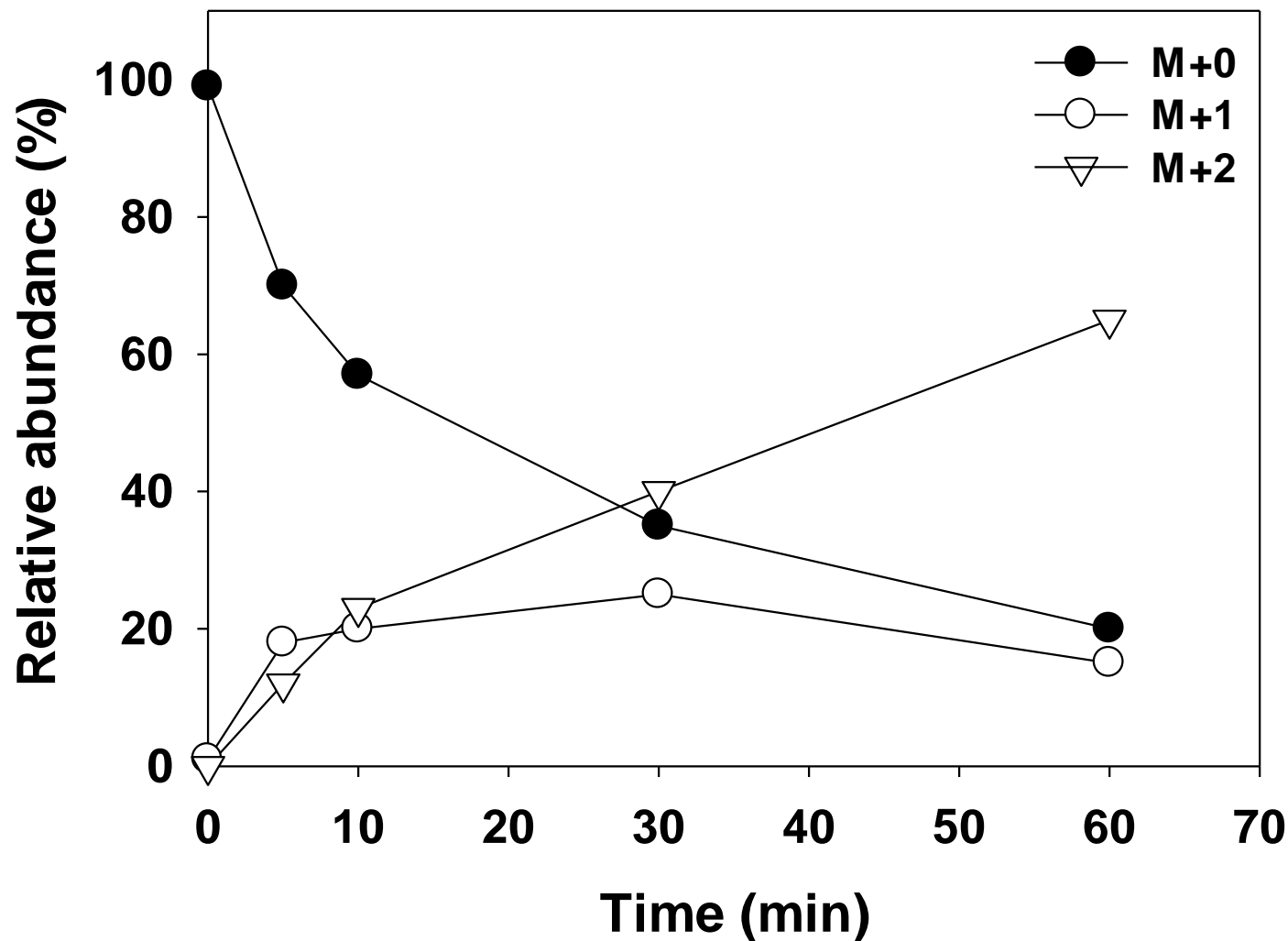
- **Metabolite profiling vs metabolic flux analysis (MFA)**
- **Positional isotope labelling**
 - NMR vs MS approaches
- **Non-cyclic modes in the plant TCA cycle and the origin of C for Glu synthesis**
- **The case of guard cells**
- **Take home messages**
- **Challenges and perspectives**
- **Acknowledgements**

Metabolite profiling vs MFA

○ ^{12}C
 ● ^{13}C

M ○○ – monoisotopic peak
M+1 ●○ ○● – incorporation of one ^{13}C
M+2 ●● – incorporation of two ^{13}C





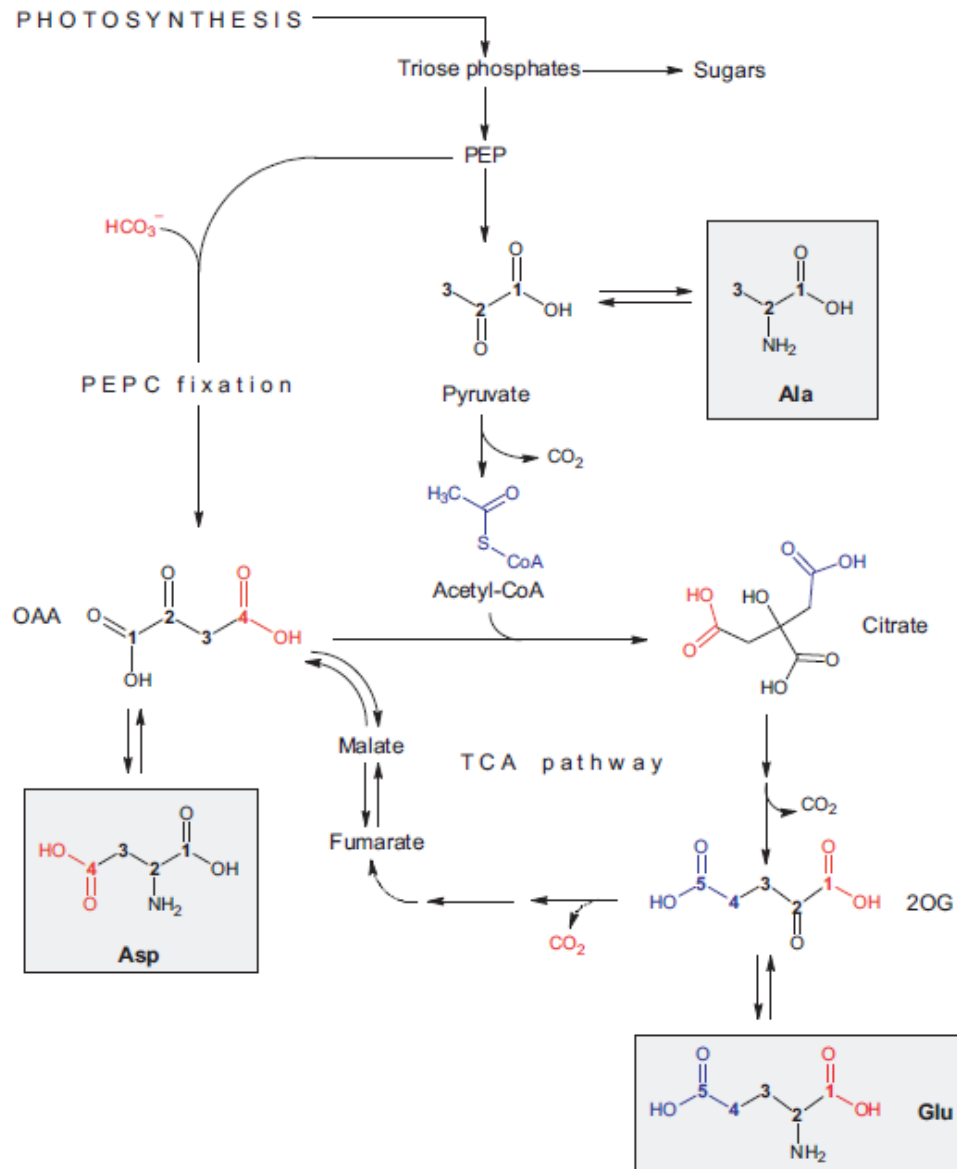
Metabolite profiling vs MFA

Ex: The level of Glu increase in plants transferred to light

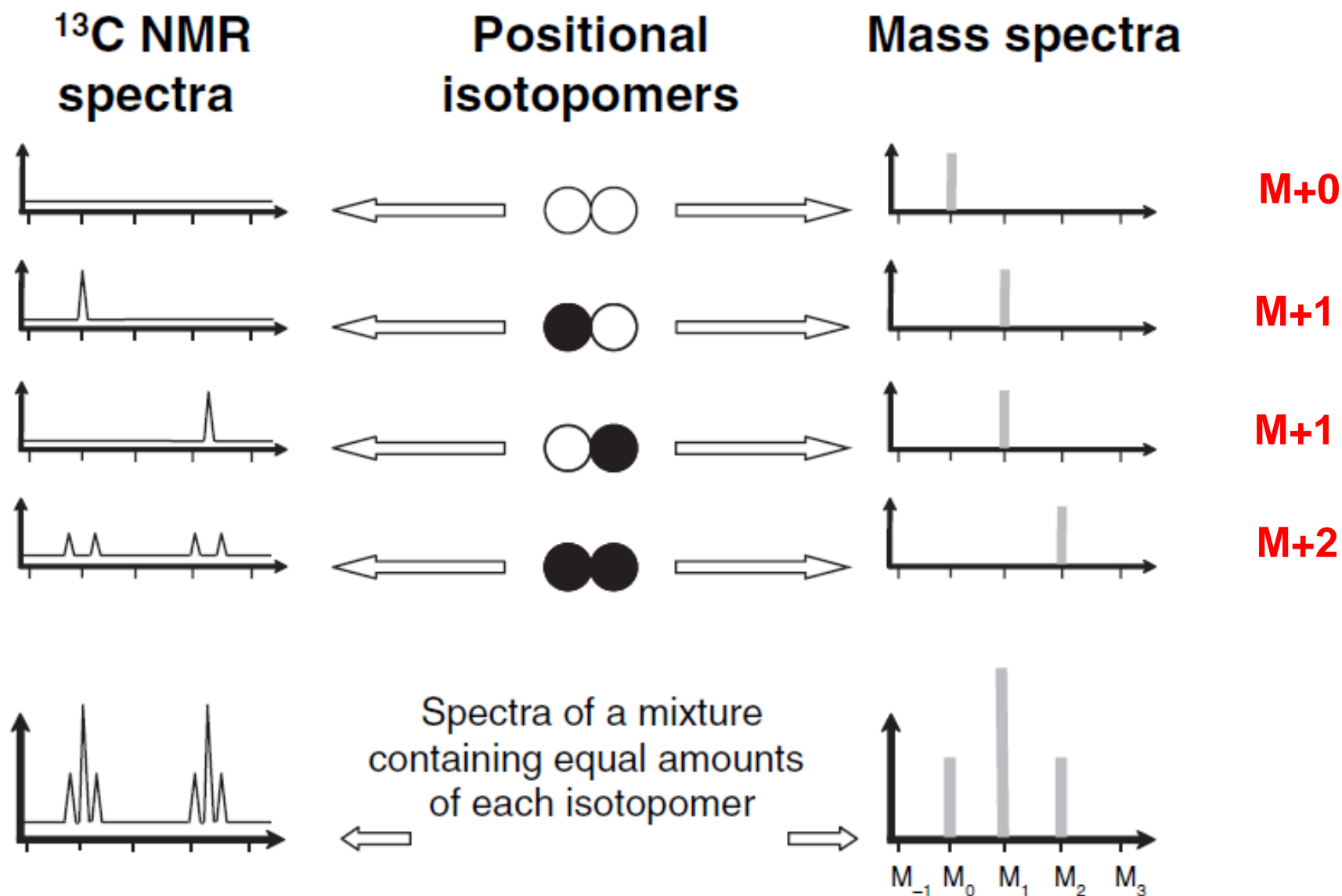
**Hypothesis:
Light**

Photosynthesis
Glutamate

Using positional isotope labelling to investigate the source of C for Glu synthesis

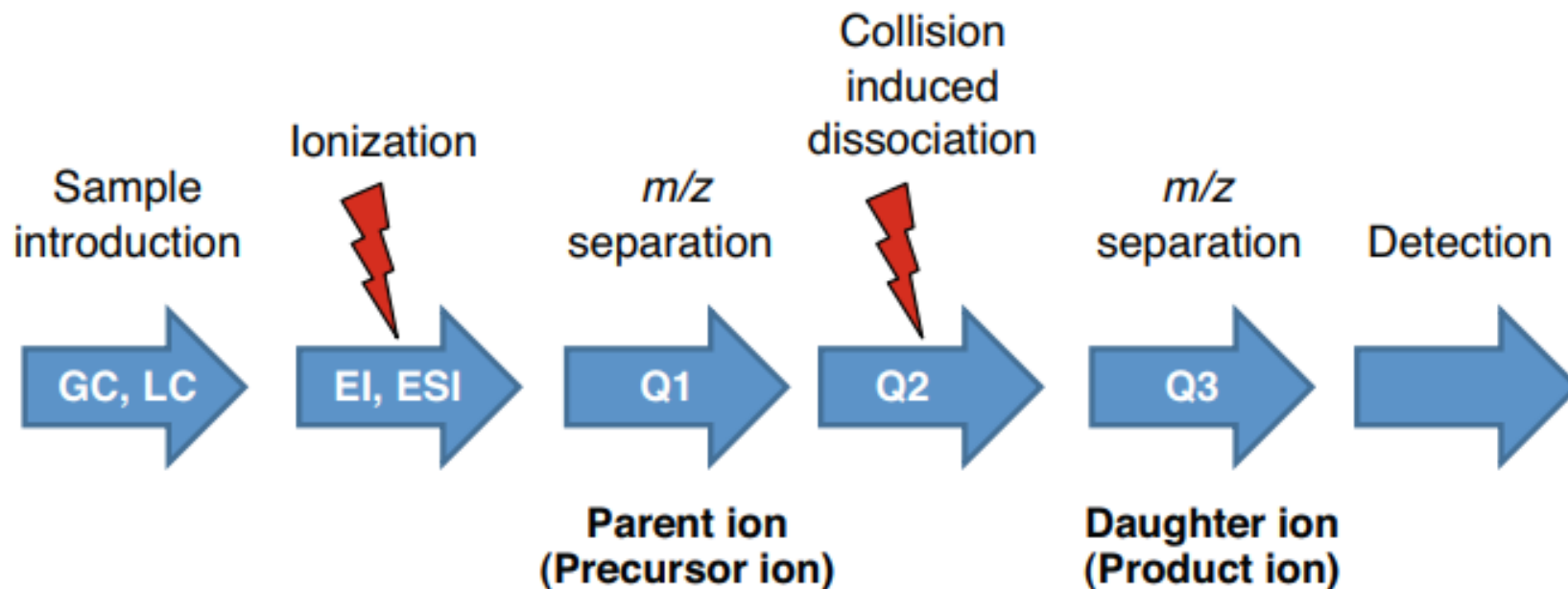


Positional isotope labelling – NMR vs MS



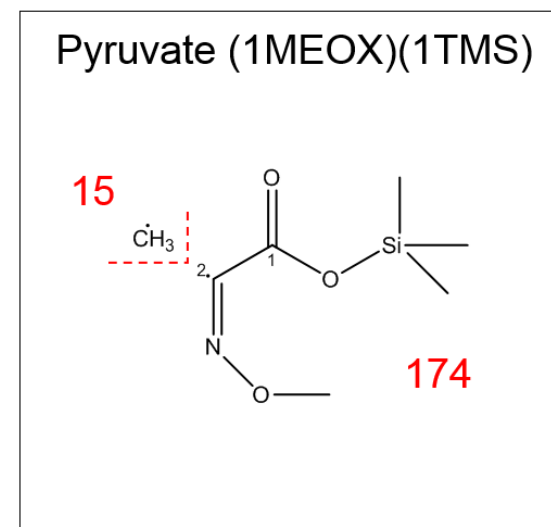
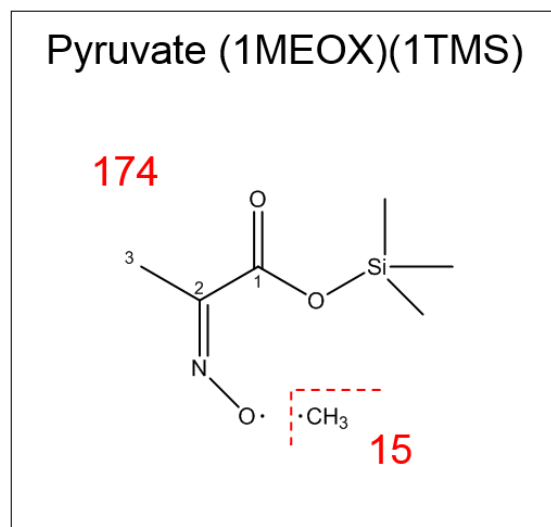
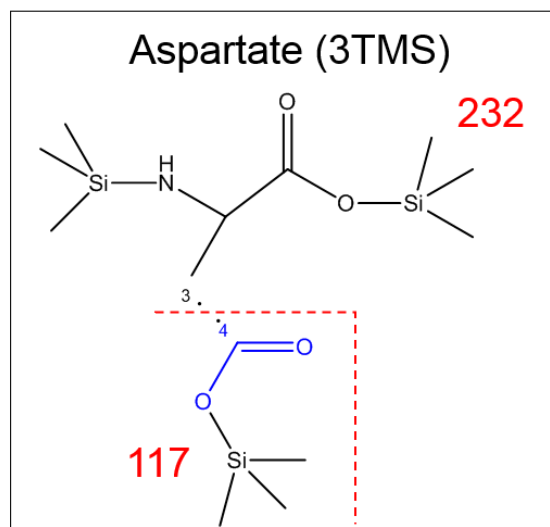
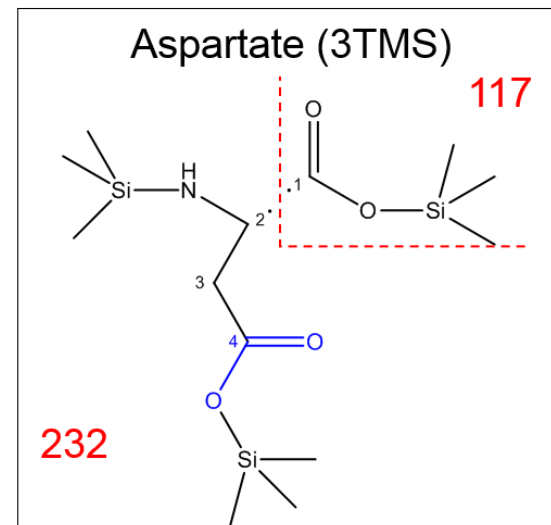
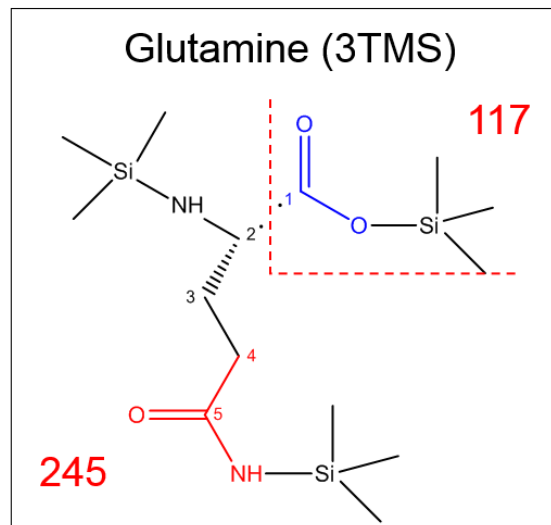
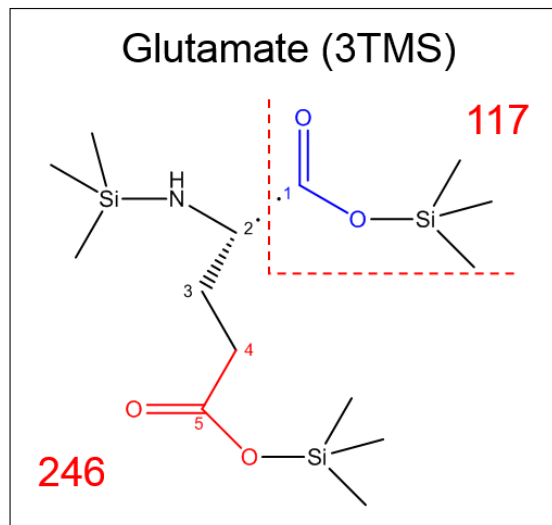
Positional isotope labelling via MS/MS

(a) Tandem mass spectrometry



Positional isotope labelling via GC-EI-MS

Which fragment should I use?





Research

Direct assessment of the metabolic origin of carbon atoms in glutamate from illuminated leaves using ^{13}C -NMR

Cyril Abadie¹, Jérémy Lothier², Edouard Boex-Fontvieille³, Adam Carroll¹ and Guillaume Tcherkez¹



Research

In vivo phosphoenolpyruvate carboxylase activity is controlled by CO_2 and O_2 mole fractions and represents a major flux at high photorespiration rates

Cyril Abadie and Guillaume Tcherkez

Origin of C for Glu synthesis

Labelling expected:

2 ¹³C from glycolysis

1 ¹³C from PEPc activity

2 ¹³C from the TCA cycle

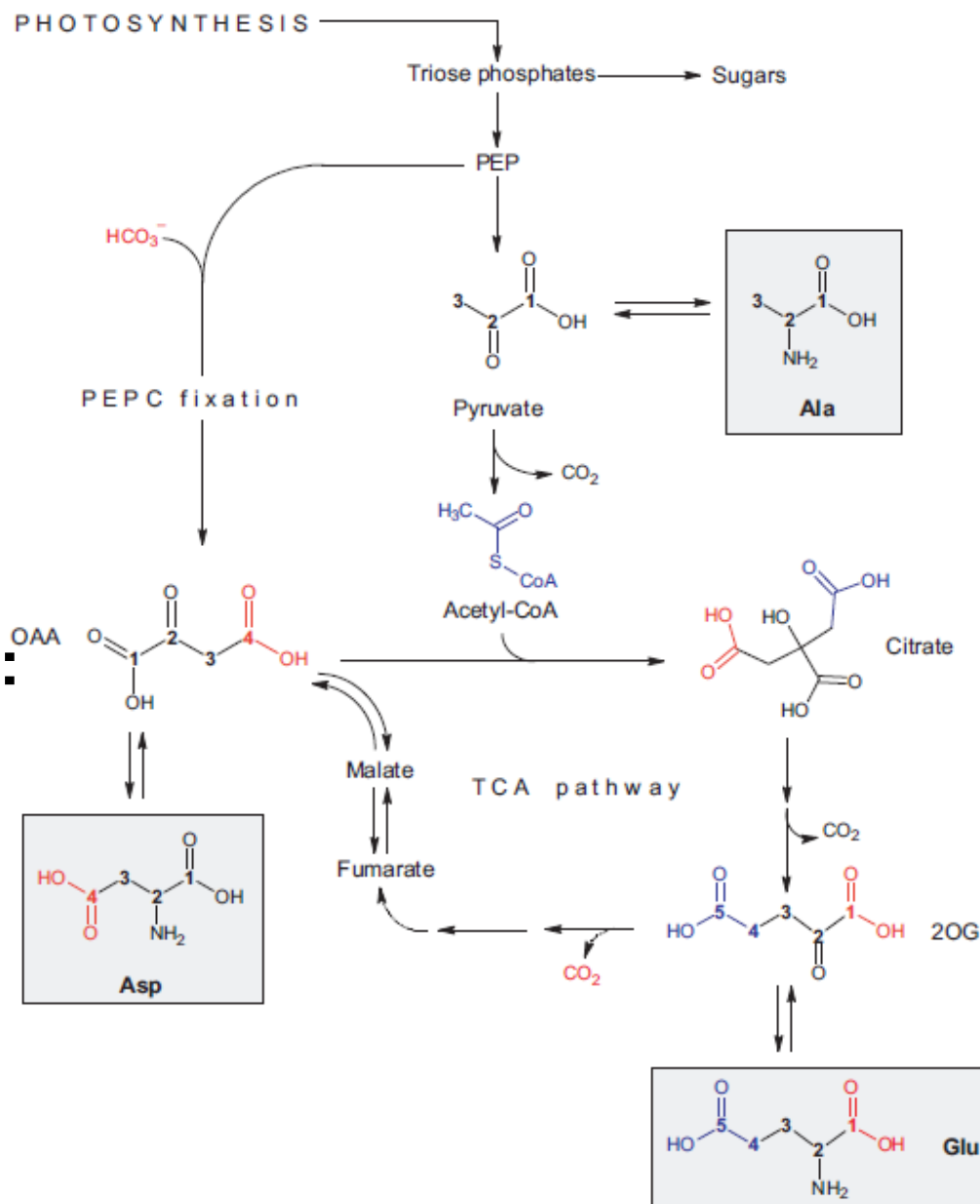
Labelling observed by NMR:

0 ¹³C from glycolysis

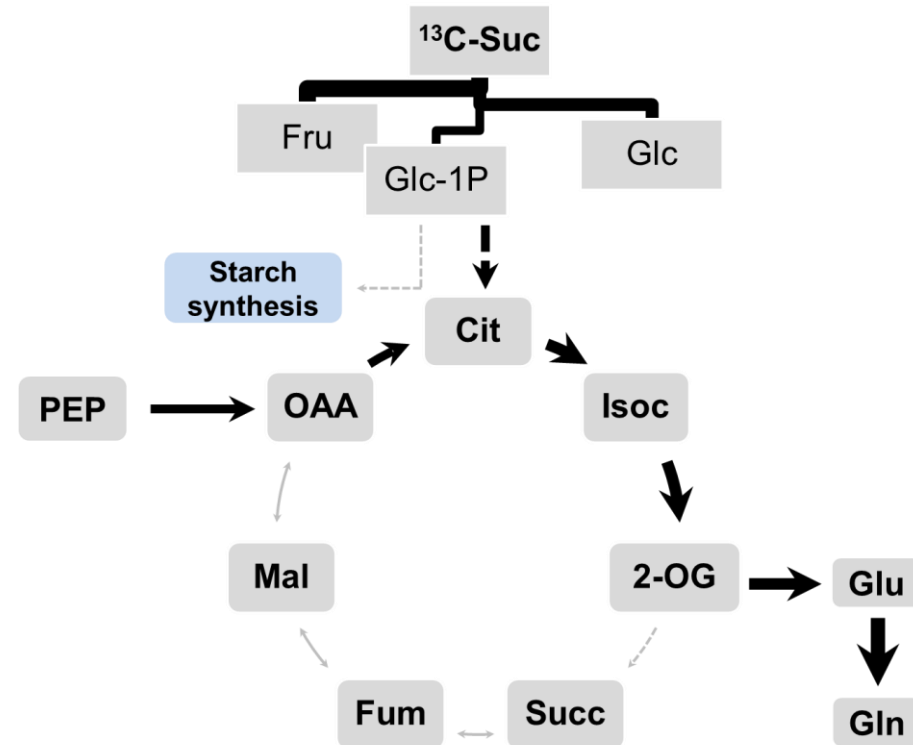
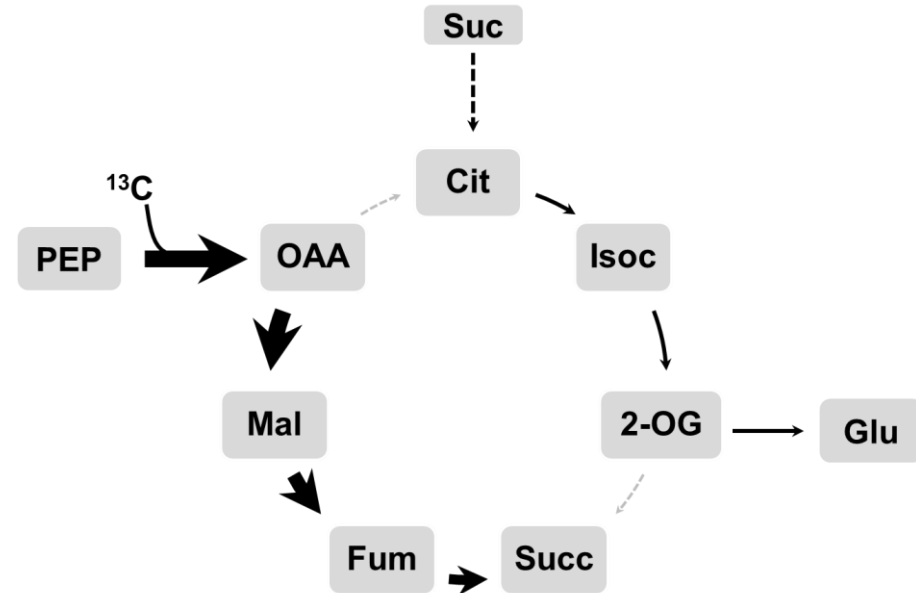
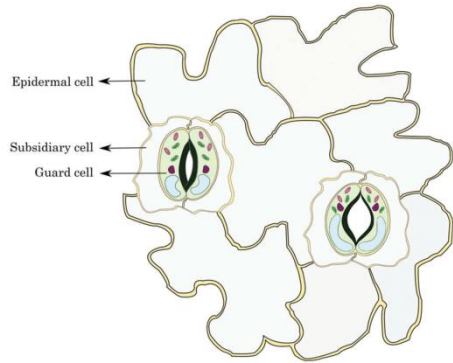
1 ¹³C from PEPc activity

0 ¹³C from the TCA cycle

Presence of ¹²C



Non-cyclic flux modes of guard cell TCA cycle



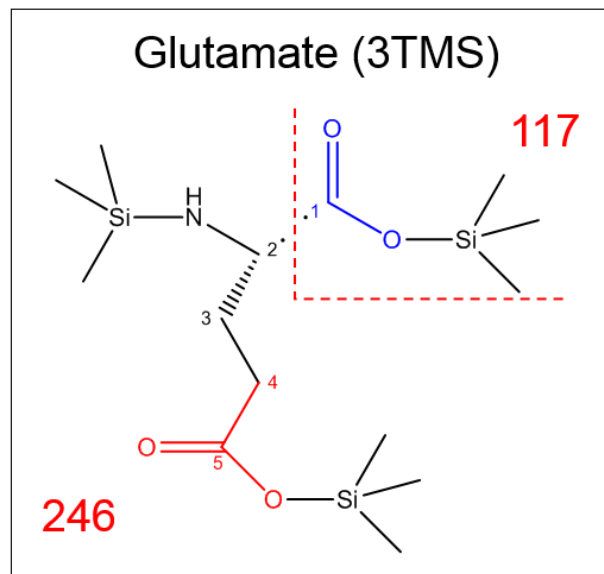
Daloso et al. (2016). New Phytologist 209

Daloso et al. (2015). Plant, Cell & Environment 38

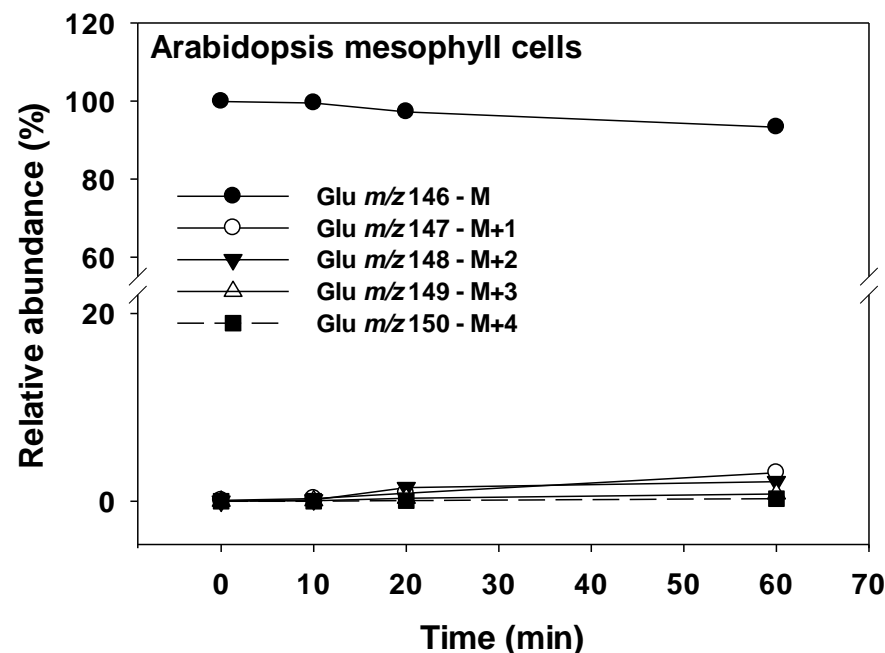
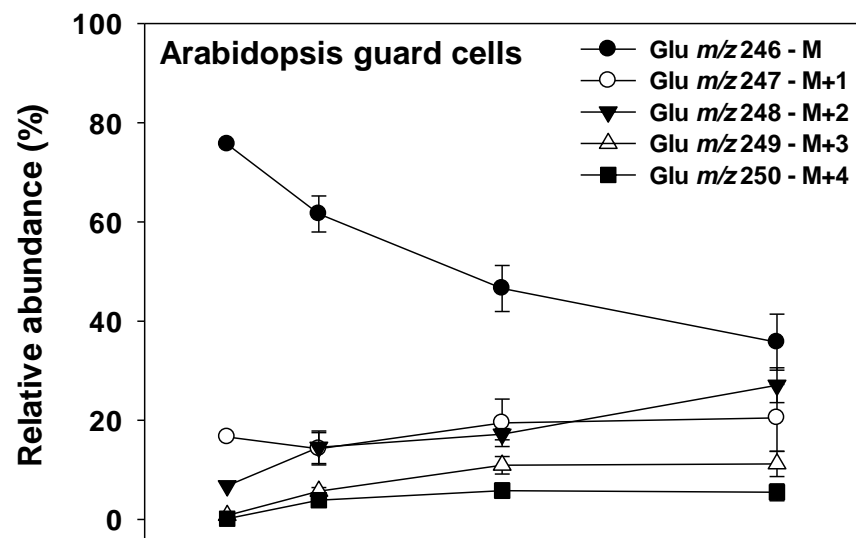
Robaina-Estévez et al. (2017). Scientific Reports 7

Medeiros et al. (2018). Plant Journal 94

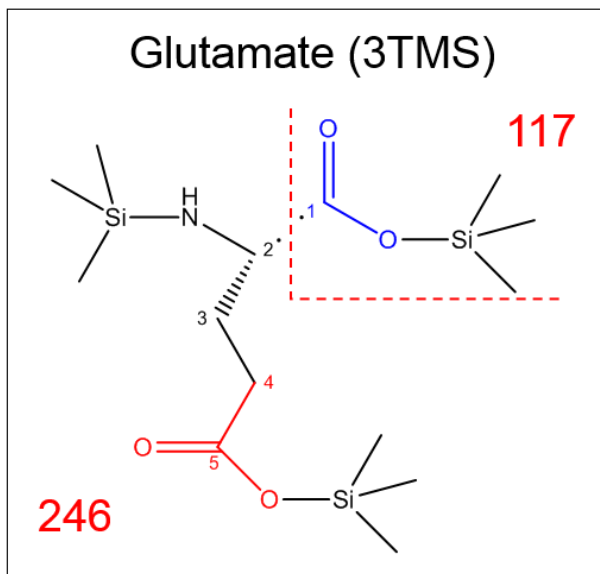
Positional isotope labelling in Glu m/z 246 - 146



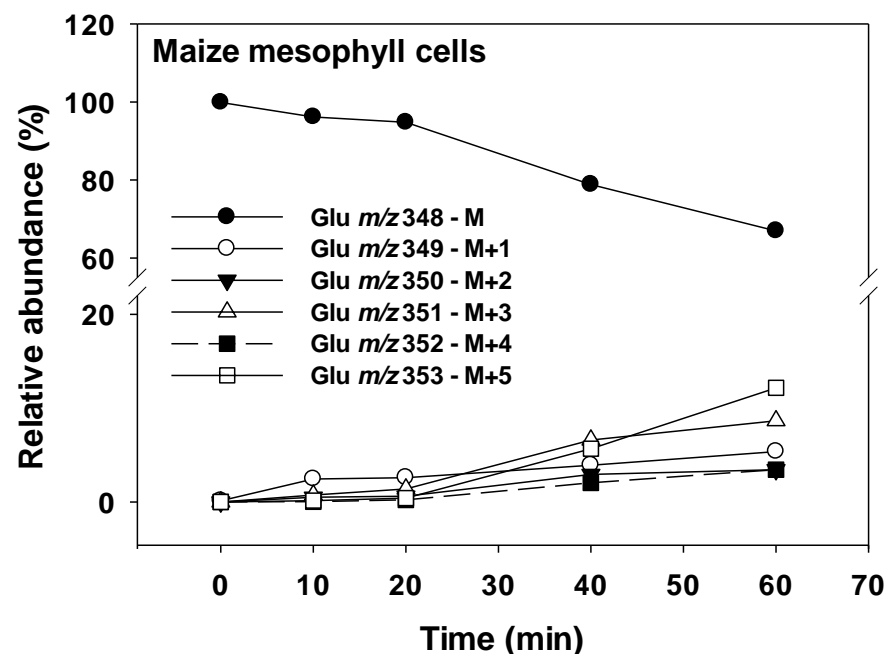
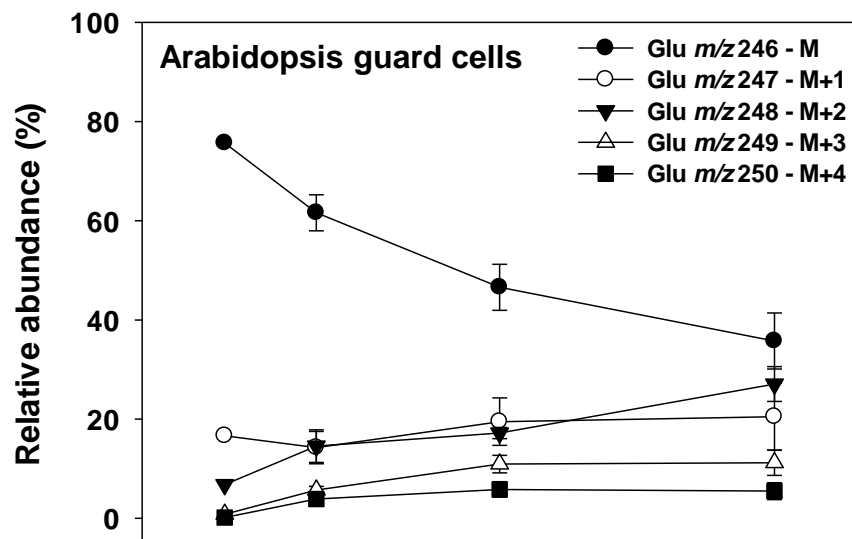
Labelled C:
 m/z 146 - 2,3,4,5
 m/z 246 - 2,3,4,5



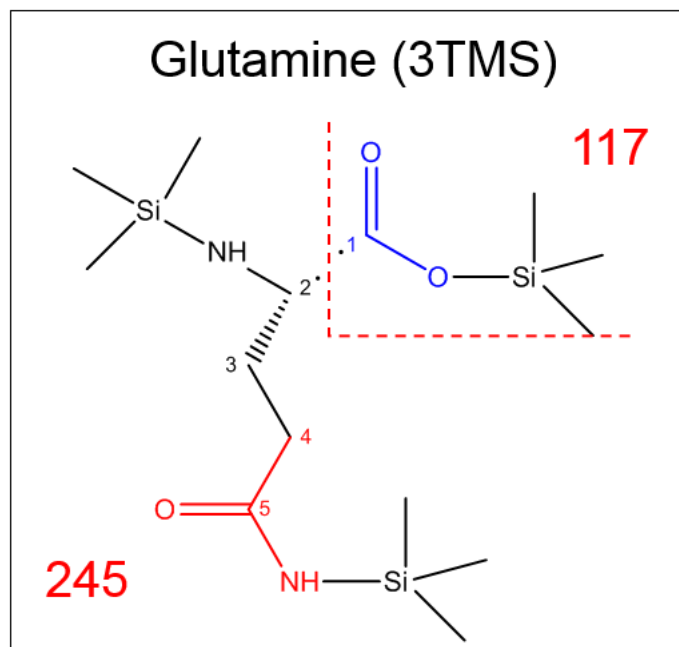
Positional isotope labelling in Glu m/z 246 - 353



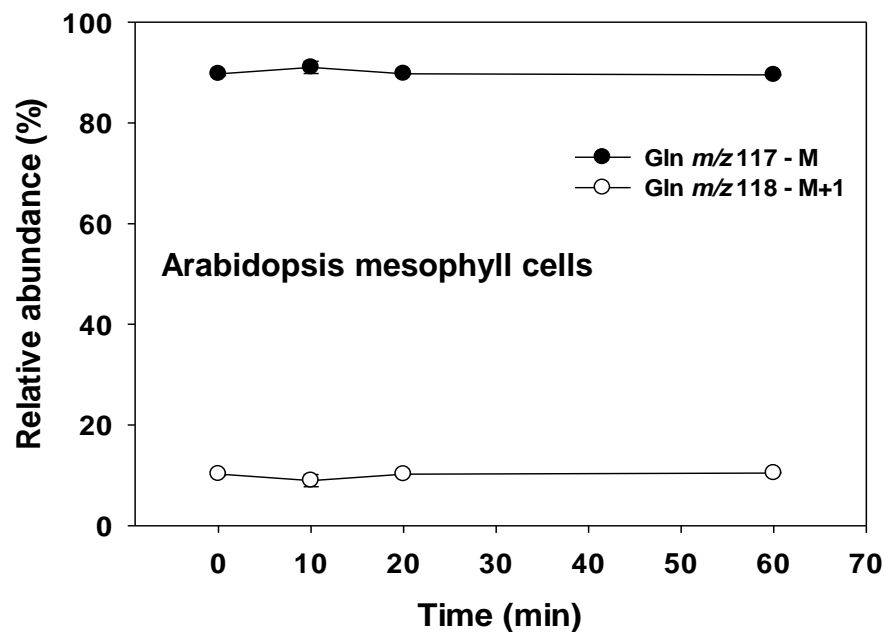
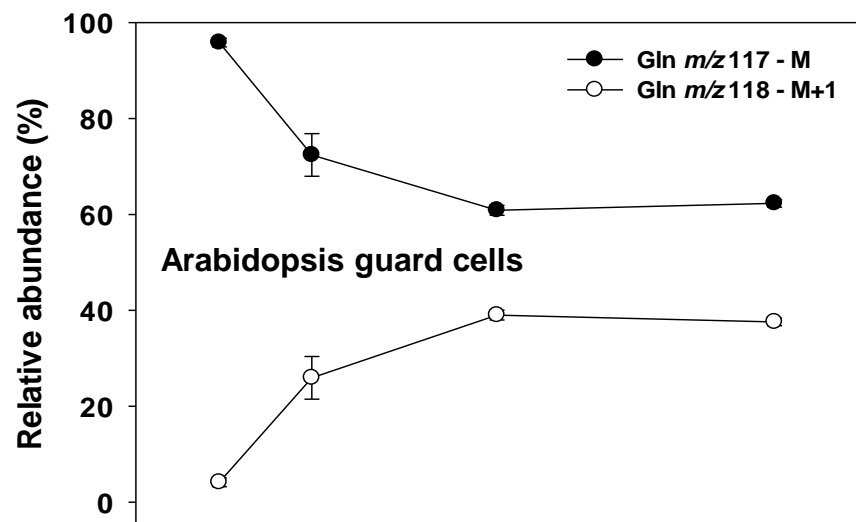
Labelled C:
 m/z 246 - 2,3,4,5
 m/z 353 - 1,2,3,4,5



Positional isotope labelling in Gln m/z 117

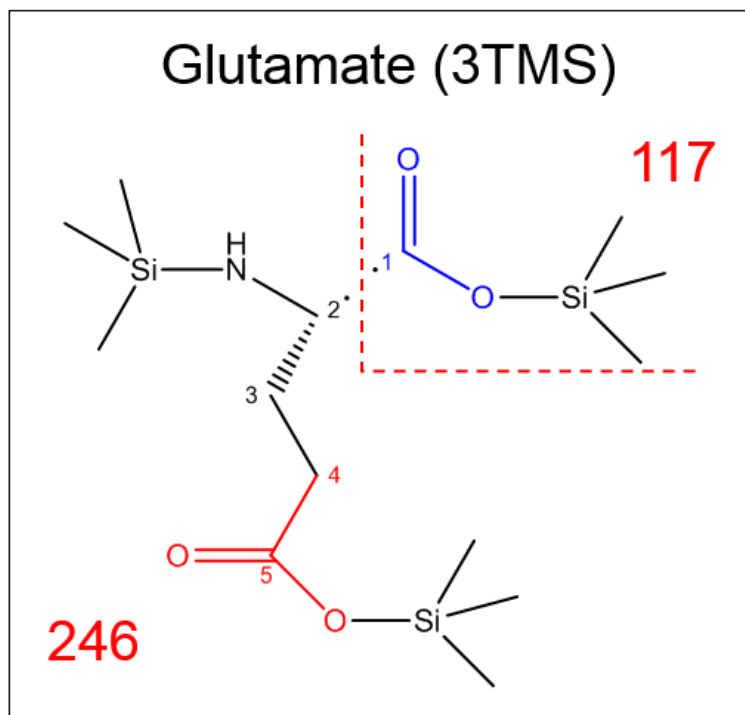


Labelled C:
 m/z 117 - 1



- **Do not use metabolite profiling data to discuss flux through the pathways**
- **The chosen labelled substrate will determine the fluxes that should be investigated**
- **Plant TCA cycle did not work as a cycle**
- **Positional isotope labelling is a useful approach to investigate ^{13}C redistribution throughout metabolic pathways**

- To create a complete map of GC-EI-MS fragmentaion



Glutamic acid 3TMS – GolmDB

m/z 117

m/z 128

m/z 156

m/z 246

m/z 348

m/z 363

- **To discriminate small fragments from noise and TMS derivatives**
 - Dry different volumes before derivatization
 - SIM method?
- **Measure enzyme activity *in vivo* using MS data**

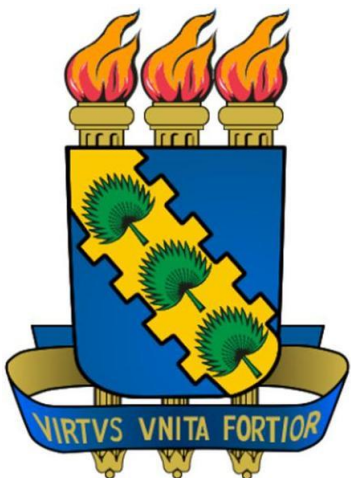


Research

In vivo phosphoenolpyruvate carboxylase activity is controlled by CO₂ and O₂ mole fractions and represents a major flux at high photorespiration rates

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Thanks for your attention