

Decreased plasma citrulline

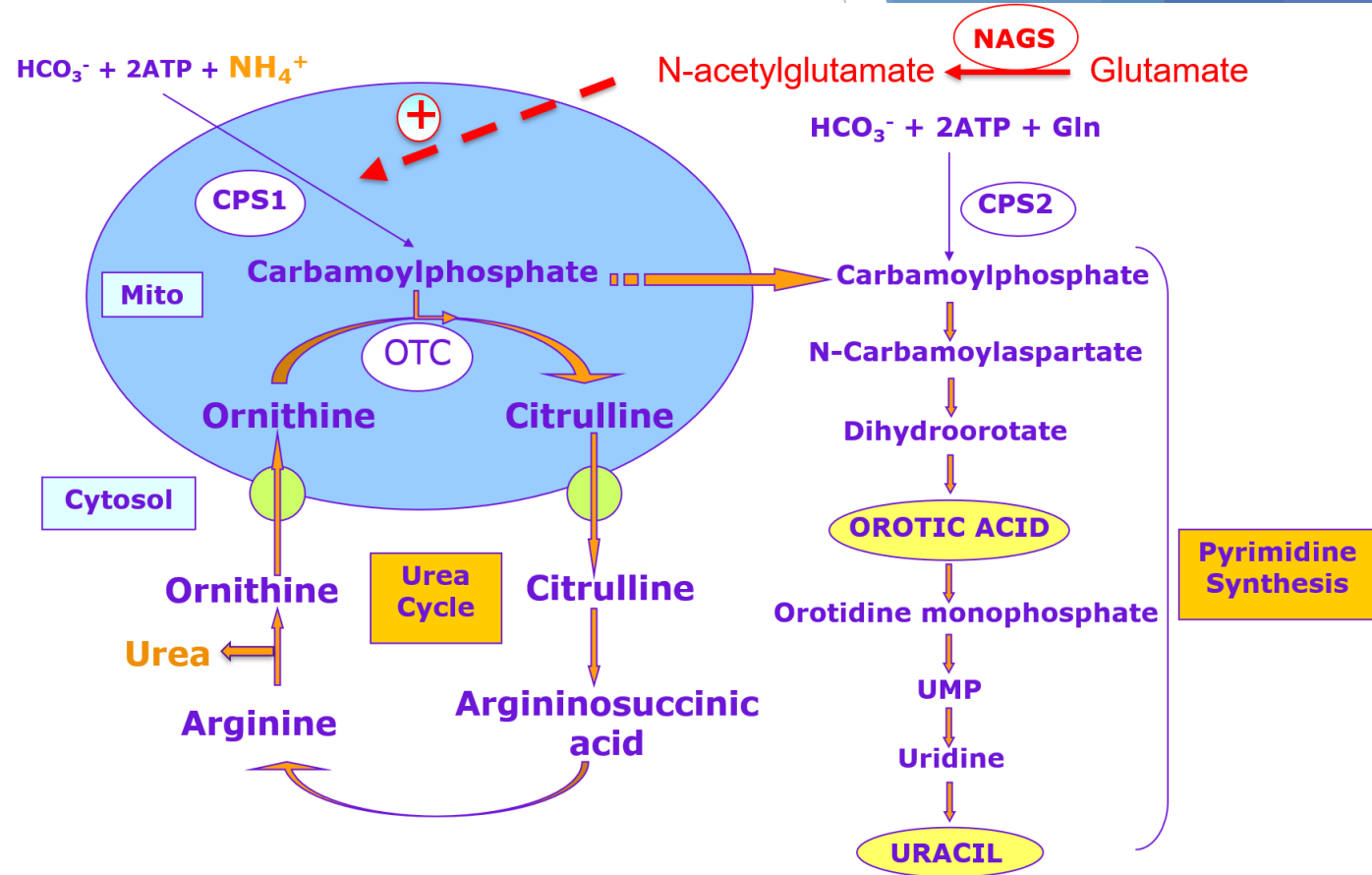
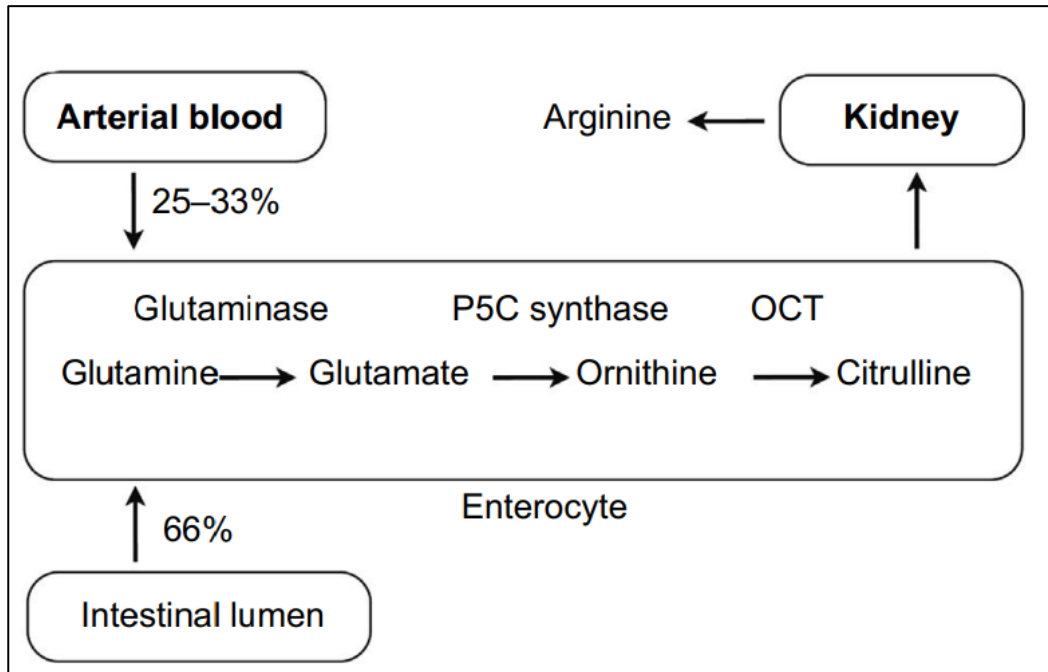
29th January 2024

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Decreased plasma citrulline - causes:

- ▶ Urea cycle defect (OTC / CPS1 / NAGS)
- ▶ Compromised small bowel function
- ▶ Generalised decrease in amino acids



Clinical Chemistry 69:6
661–668 (2023)

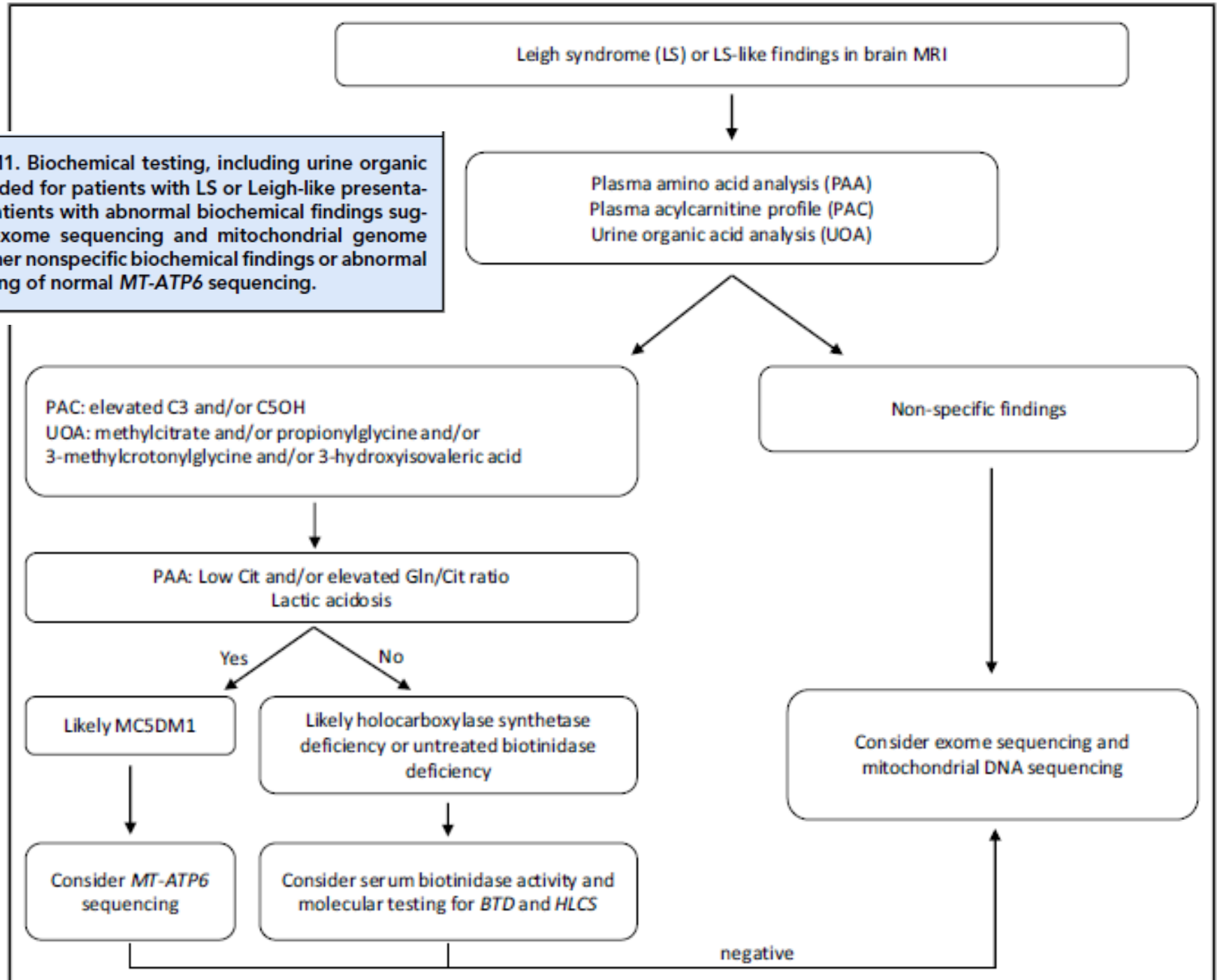
the Clinical Chemist

Genetic Metabolic Series

Low Plasma Citrulline Guiding the Diagnosis of a Mitochondrial Disorder

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Fig. 2. Recommended diagnostic algorithm for MC5DM1. Biochemical testing, including urine organic acid, plasma amino acid and PAC analyses, is recommended for patients with LS or Leigh-like presentation. Sequencing of *MT-ATP6* is recommended in the patients with abnormal biochemical findings suggestive of MC5DM1, while broader tests, including exome sequencing and mitochondrial genome sequencing, are recommended for patients who have either nonspecific biochemical findings or abnormal biochemical findings suggestive of MC5DM1 in the setting of normal *MT-ATP6* sequencing.



Plasma amino acids

Amino acids [Plasma]

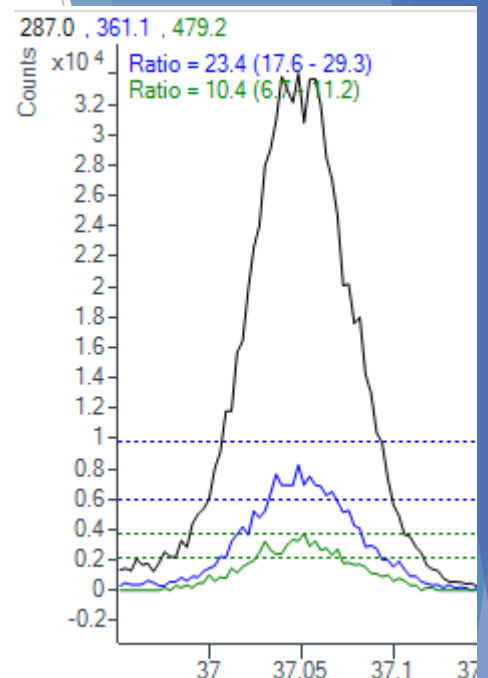
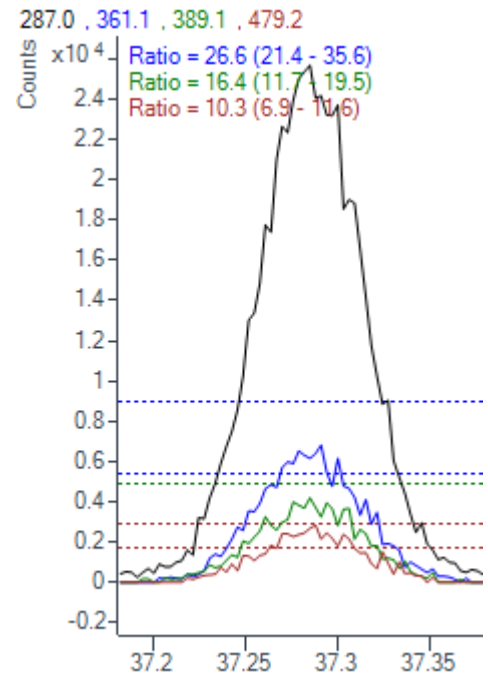
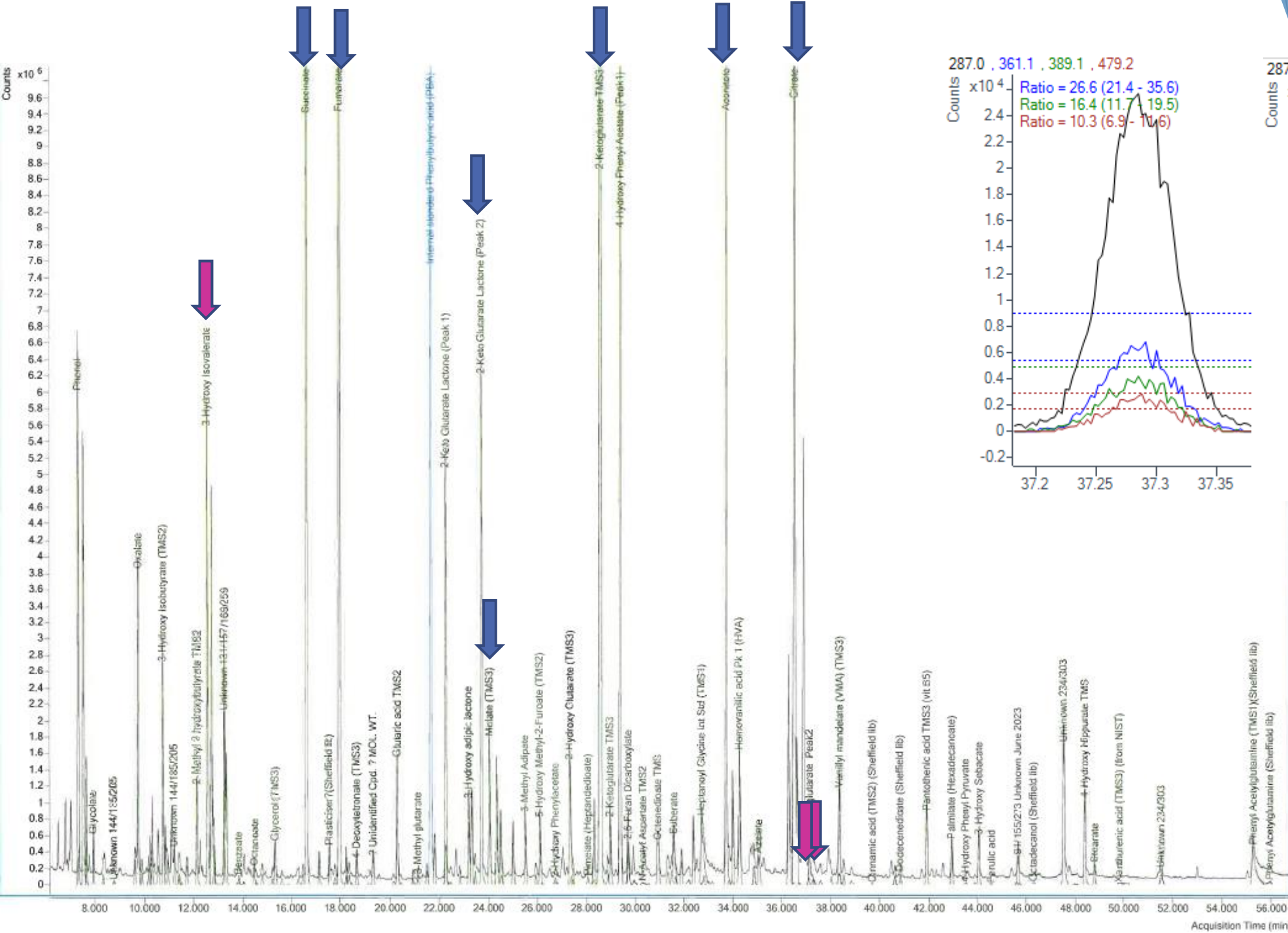
Taurine	43	umol/L	(28-212)	Valine	193	umol/L	(77-335)
Aspartic acid	8	umol/L	(4-52)	Cystine	27	umol/L	(10-80)
Hydroxyproline	20	umol/L	(6-75)	Methionine	18	umol/L	(11-43)
Threonine	181	umol/L	(47-228)	Isoleucine	110 H	umol/L	(27-101)
Serine	139	umol/L	(66-231)	Leucine	151	umol/L	(52-206)
Asparagine	44	umol/L	(23-80)	Tyrosine	47	umol/L	(32-125)
Glutamic acid	74	umol/L	(21-194)	Phenylalanine	42	umol/L	(35-103)
Glutamine	574	umol/L	(227-735)	Ornithine	41	umol/L	(24-144)
Proline	163	umol/L	(98-429)	Lysine	134	umol/L	(70-259)
Glycine	171	umol/L	(133-455)	Histidine	52	umol/L	(42-108)
Alanine	646 H	umol/L	(138-565)	Tryptophan	73	umol/L	(19-107)
Citrulline	6 L	umol/L	(7-48)	Arginine	28	umol/L	(12-145)

Urine organic acids

Organic acids [Urine]

Organic acids [Urine] Comment

Marked increases in TCA cycle intermediates (succinate, fumarate, malate and 2-ketoglutarate). Suggest further investigation to exclude a mitochondrial disorder in this patient.



Plasma acylcarnitines

Plasma acylcarnitine profile:

Tandem Mass Spectrometric Analysis of Acylcarnitines

Free Carnitine	38	umol/L	15-53
C3	4.29	umol/L	<1.30
C5-OH	0.14	umol/L	<0.06

Biotinidase

Biotinidase

Biotinidase	10.9	U/L	(5.0-14.9)
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Biotinidase activity analysed Spectrophotometrically.

Normal biotinidase activity detected.

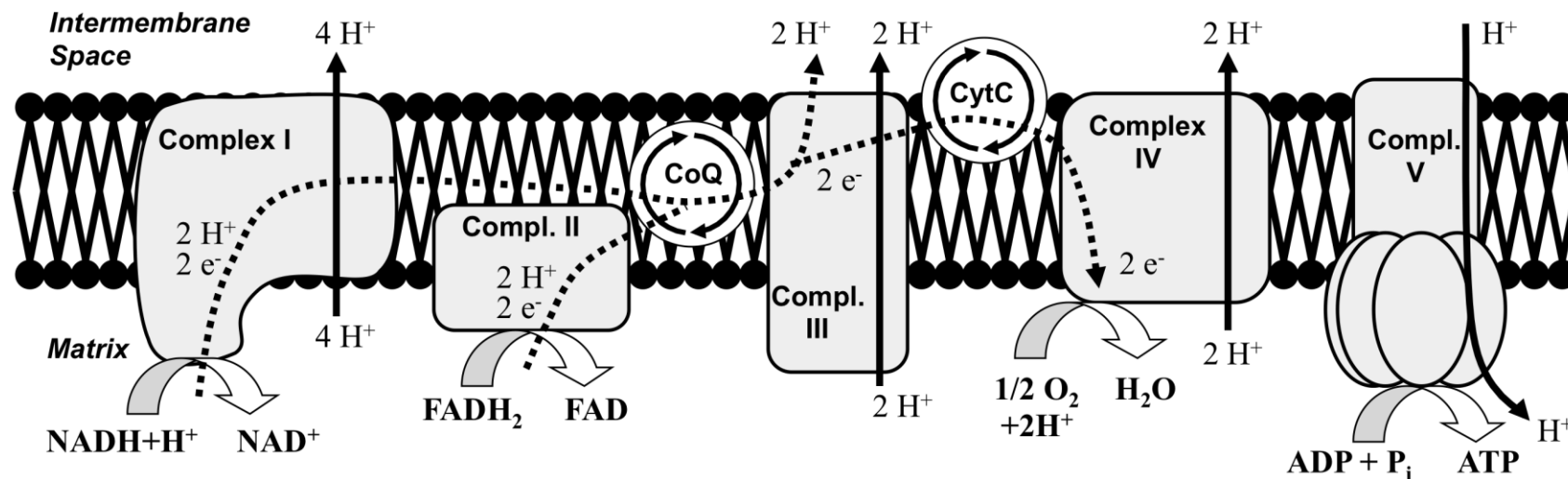
- ▶ NB: Doesn't exclude holocarboxylase synthetase deficiency

Summary of biochemistry MT-ATP6

	Plasma lactate (mmol/L)	Plasma citrulline (umol/L)	Plasma alanine (umol/L)	Organic acid increases	Acylcarnitine increases	Biotinidase
CASE 1	6.8	6	646	TCA cycle (3OH-IVA)	C3 & C5OH	Normal
CASE 2	4.8 - 5.3	4 & 0	793 & 739	3OH-IVA	C5OH	Normal
CASE 3	5.9	4 & 4	517 & 618	TCA cycle & 3OH-IVA	C3 & C5OH	Normal

MT-ATP6 & MC5DM1

- ▶ MT-ATP6 encodes subunit of Complex V of respiratory chain = ATP synthase
- ▶ MC5DM1: Mitochondrial complex V mitochondrial type 1 (MT-ATP6)
- ▶ MC5DM2: Mitochondrial complex V mitochondrial type 2 (MT-ATP8)
- ▶ Ganetsky et al 2019: 218 published cases of MT-ATP6 + 14 new cases
- ▶ m.8993T>G variant most prevalent (>100 cases)
- ▶ 18 other pathogenic variants described



MC5DM1 - Clinical presentation

- ▶ Developmental delay
- ▶ Epilepsy
- ▶ Cardiomyopathy
- ▶ Neuropathy
- ▶ Ataxia
- ▶ Retinitis pigmentosa
- ▶ Brain MRI abnormalities consistent with Leigh syndrome

- ▶ Variation in severity:
 - Infantile-onset Leigh syndrome → adult-onset progressive ataxia
- ▶ Dependent on genotype & heteroplasmy

Mechanism for biochemical abnormalities

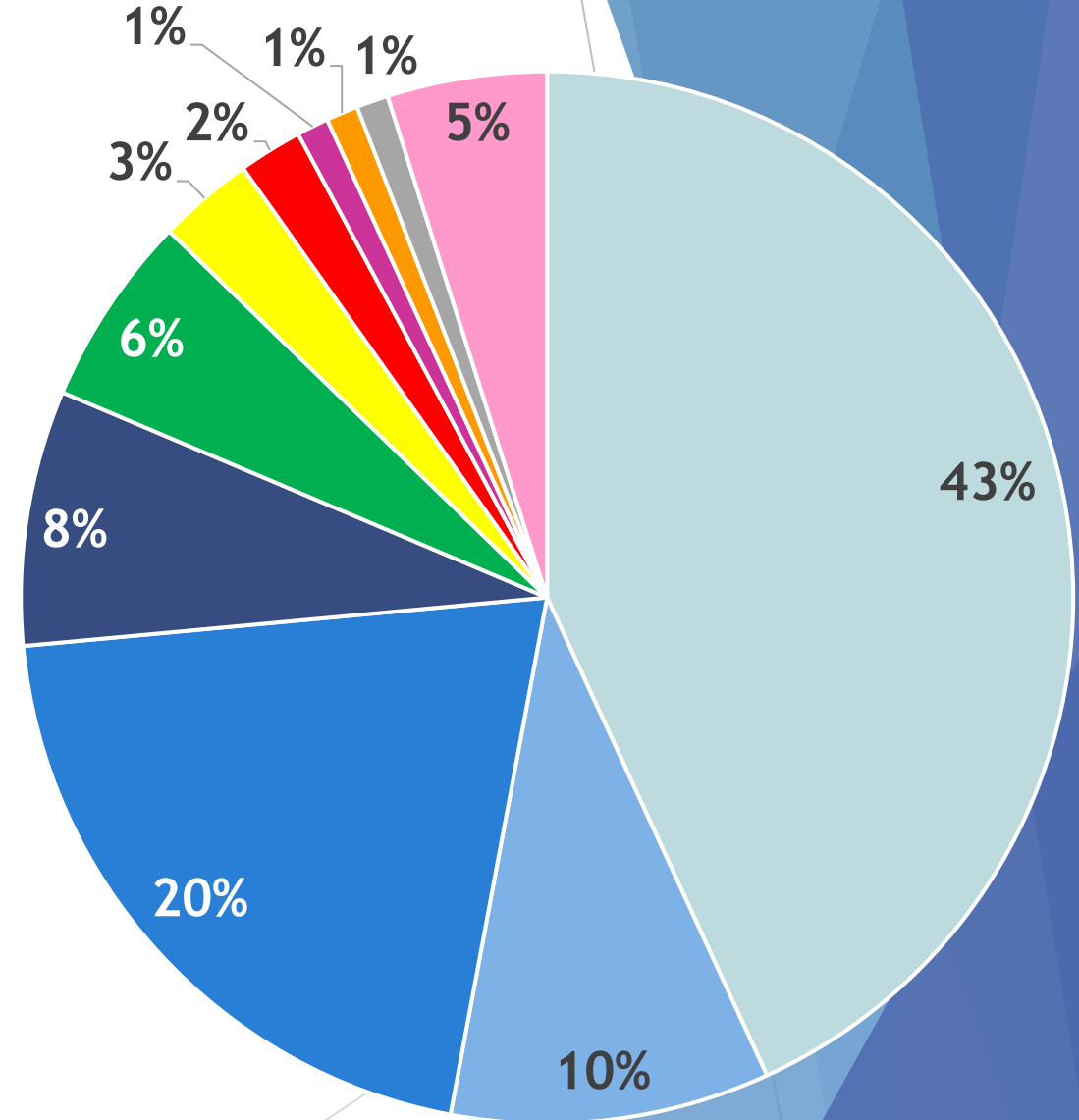
- ▶ Larson et al 2019: ? Secondary deficiencies in ATP-dependent enzymes:
 - ▶ CPS1
 - ▶ Δ_1 -pyrroline-5-carboxylate synthetase
 - ▶ Biotin-protein ligase
 - ▶ Propionyl-CoA carboxylase
 - ▶ 3-Methylcrotonyl carboxylase

Review of decreased citrulline results

- ▶ 3459 plasma amino acid profiles
- ▶ 102 citrulline (3%) <LLN of reference range (<7 $\mu\text{mol/L}$)
- ▶ 63 (62%) from HYPOGLYCAEMIA SCREENS

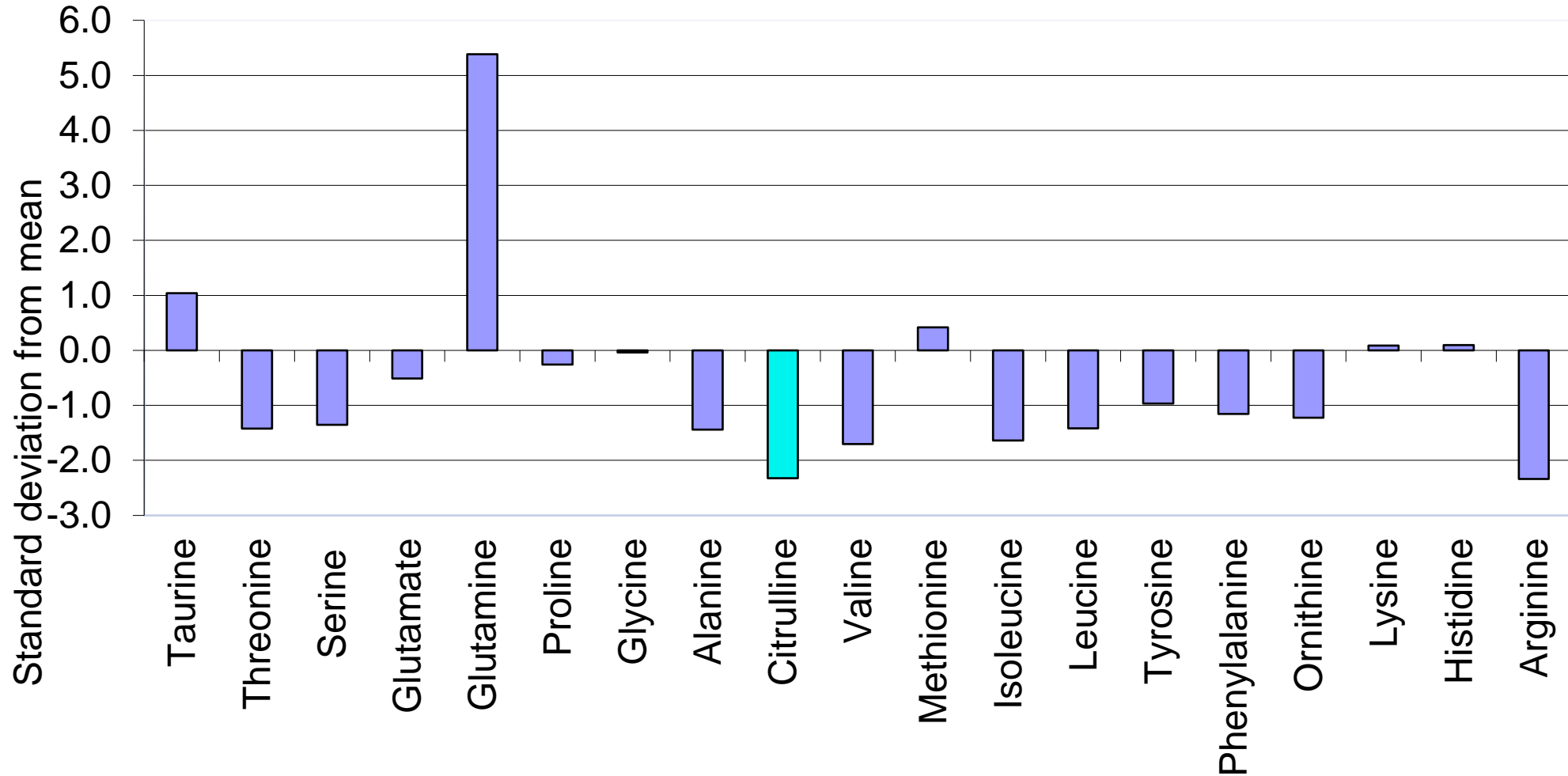
Plasma samples with decreased citrulline concentration

- Generalised decrease
- Increased BCAA
- Generalised decrease/increased BCAA
- HOG but not gen decrease / increased BCAA
- Compromised small bowel absorptive capacity
- MT-ATP6
- Deranged profile - contamination
- ALL - low Asn and Gln
- Increased ala, decreased cit - schizophrenia
- SCN3A mutation seizures
- Other



Use of SD plots to interpret low citrulline

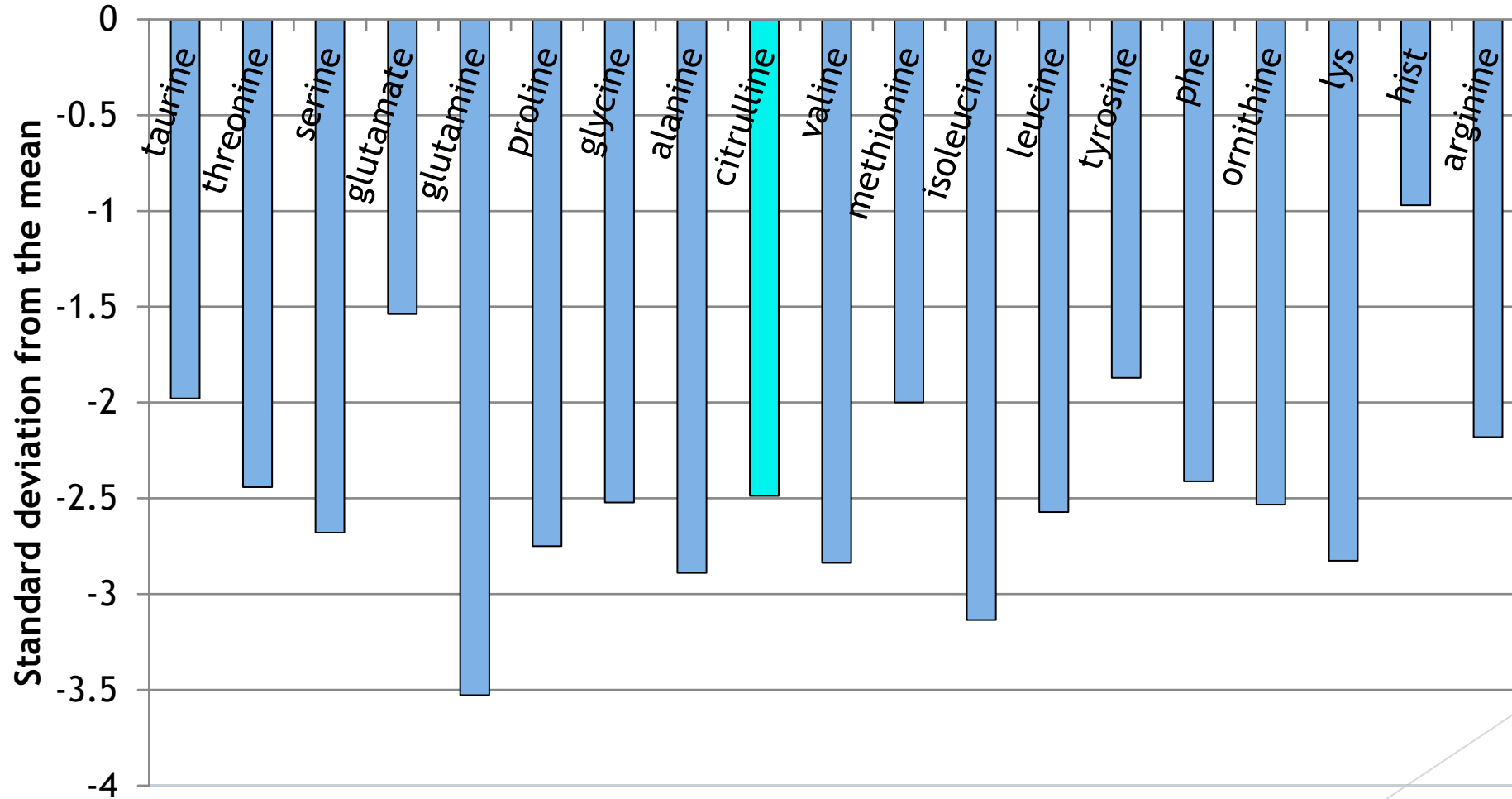
Citrulline
2 $\mu\text{mol/L}$



OTC
Deficiency

Citrulline
2 $\mu\text{mol/L}$

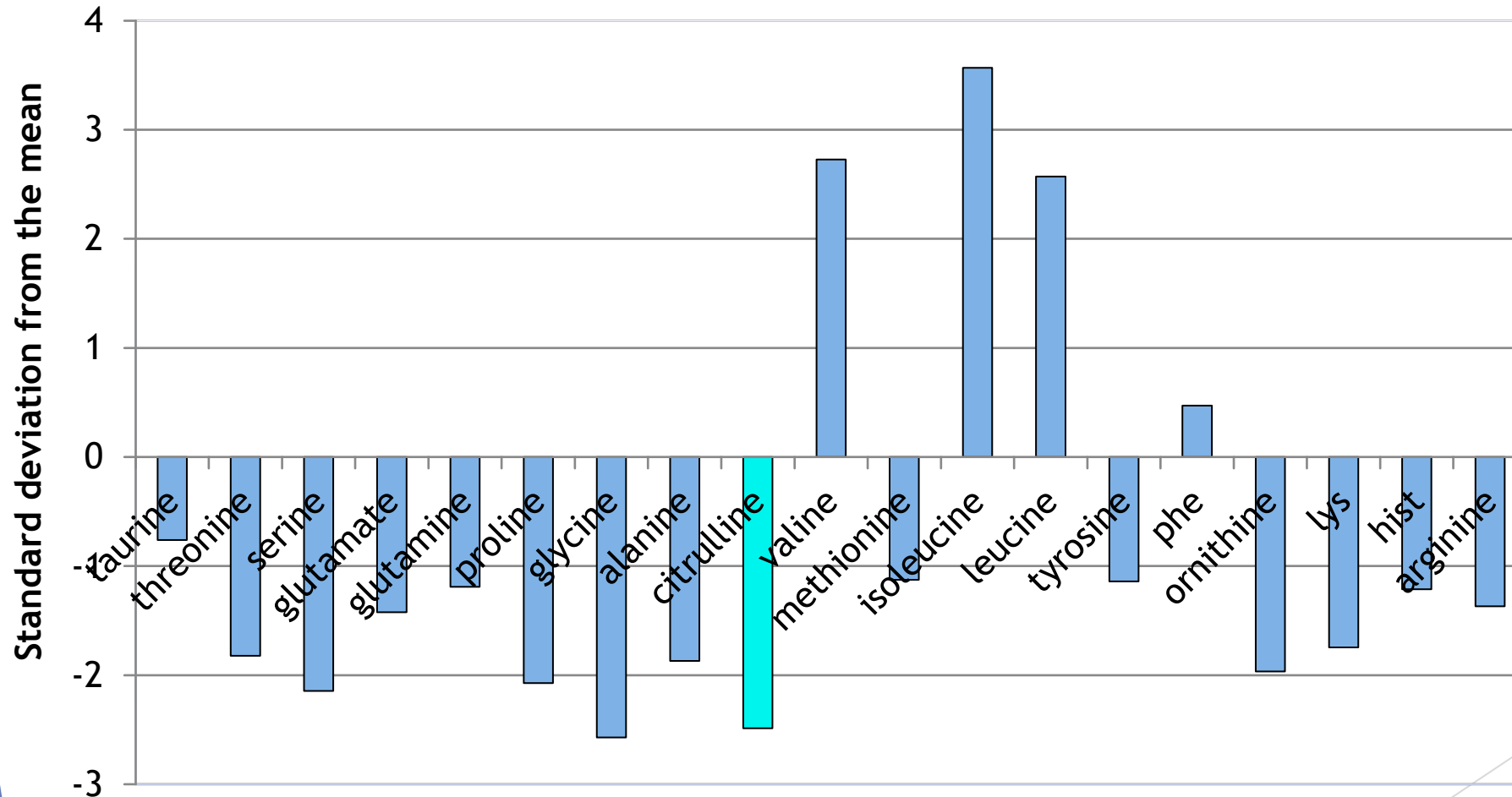
Use of SD plots to interpret low citrulline



Generalised decrease

Use of SD plots to interpret low citrulline

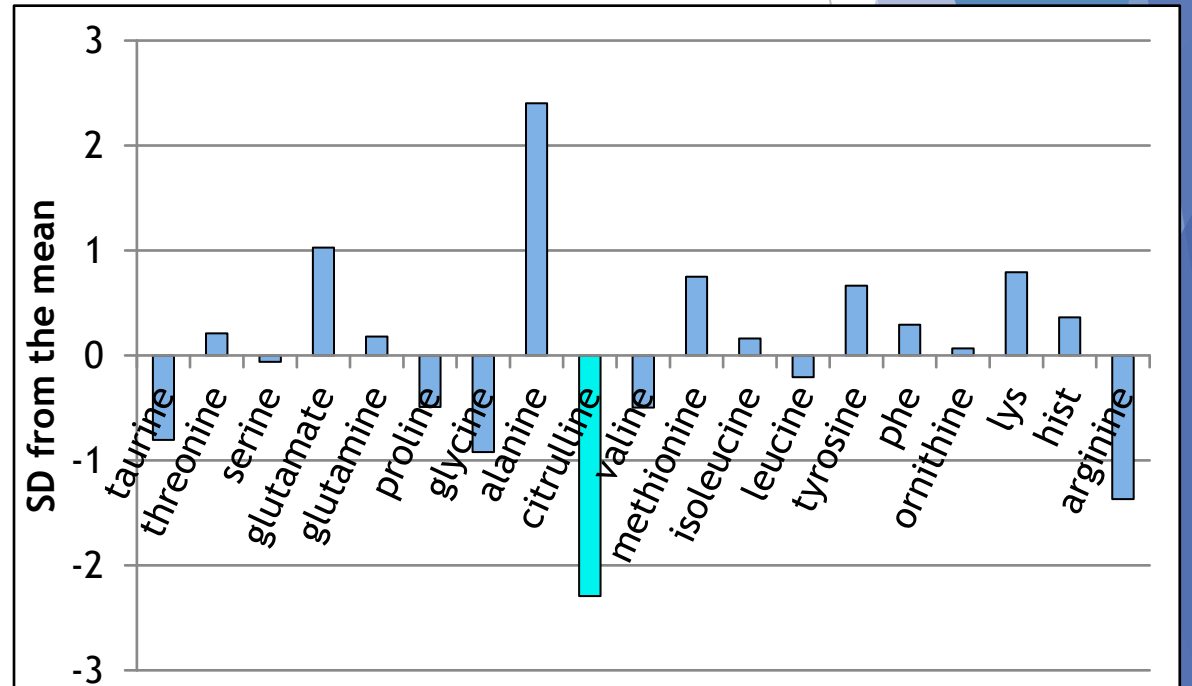
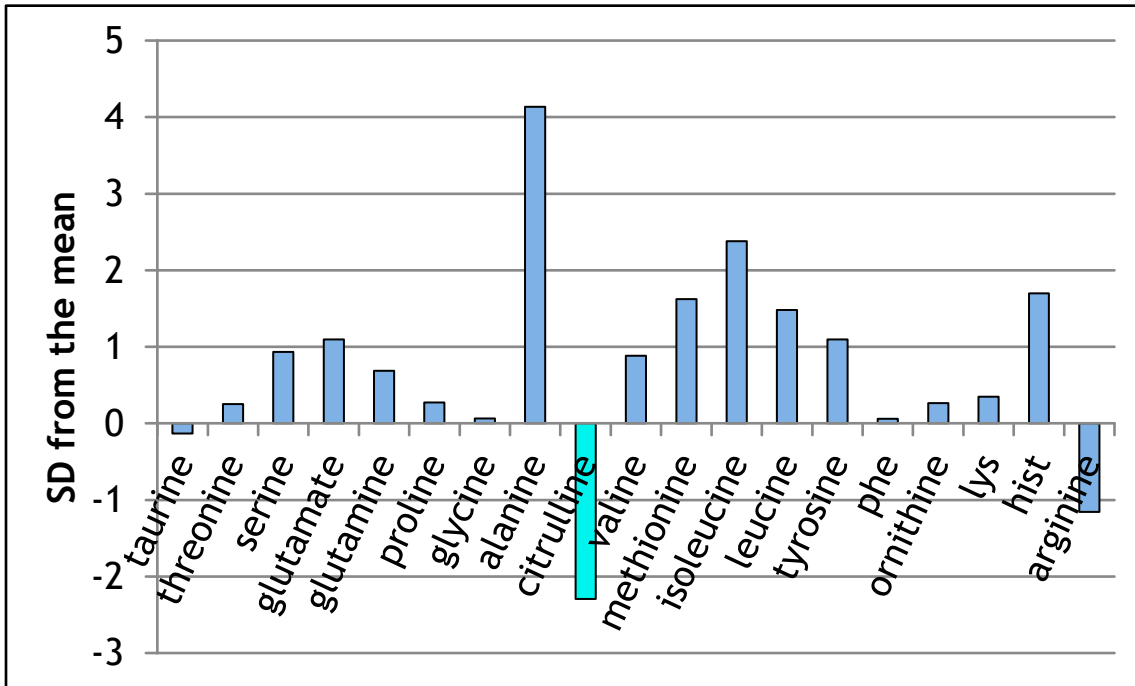
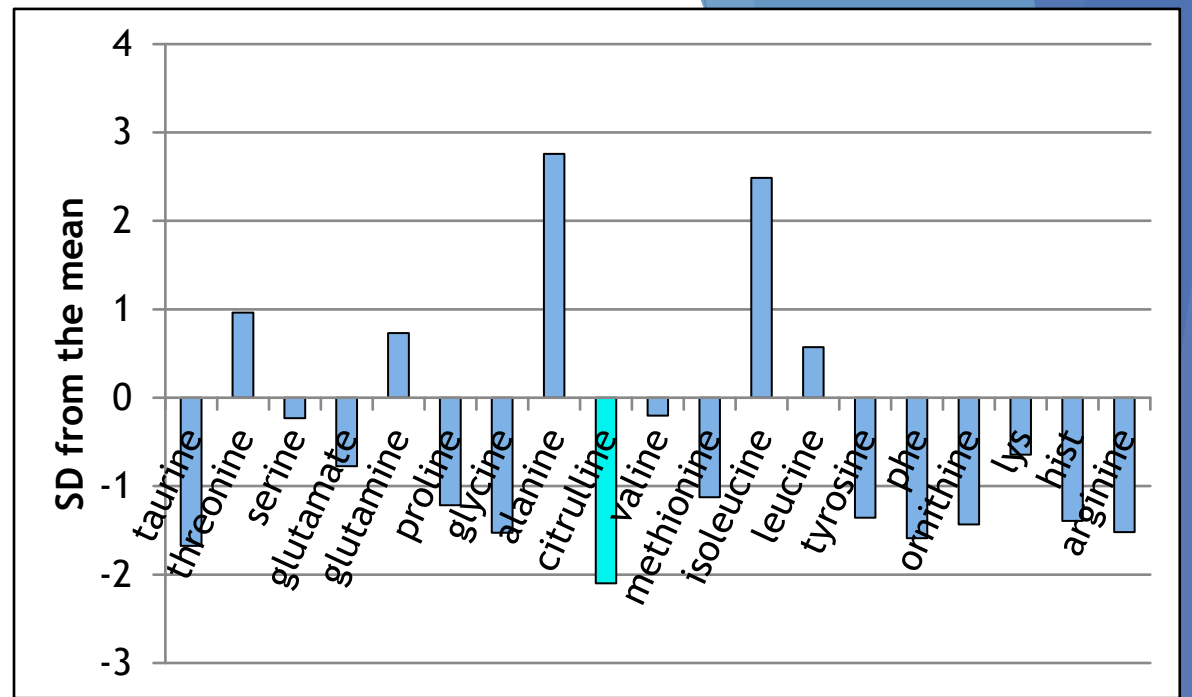
Citrulline
2 $\mu\text{mol/L}$



Generalised
decrease with
increased BCAA
(Ketotic)

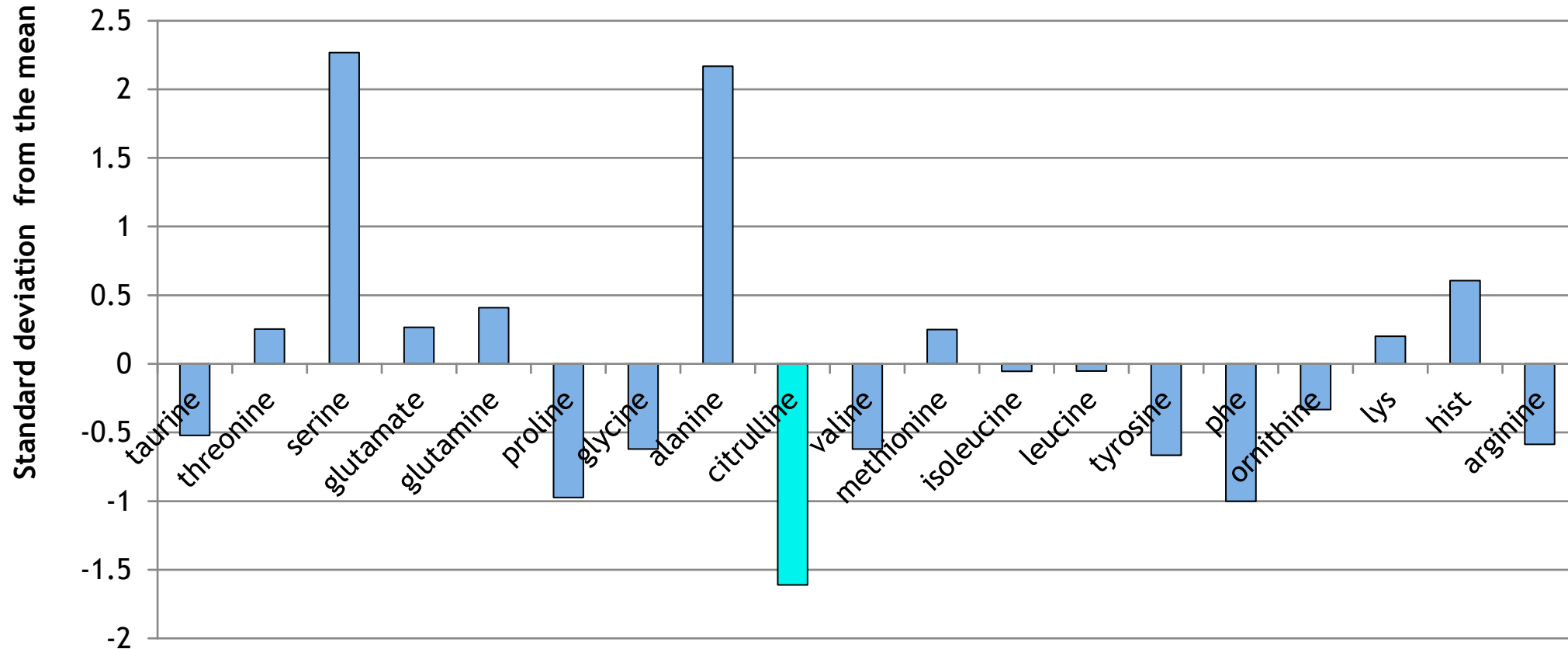
MT-ATP6 cases

- ▶ No generalised decrease
- ▶ Citrulline decreased
- ▶ Alanine increased 2° to lactic acidosis



Is decreased citrulline present in all mitochondrial / respiratory chain disorders?

**Complex 1 deficiency (NDUFS7):
CITRULLINE 11 μ mol/L (RR7-48)**



Summary

- ▶ Think MC5DM1 due to MT-ATP6 if you see:

Lactate	Plasma citrulline	Plasma alanine	Organic acid increases	Acylcarnitine increases	Biotinidase
↑	↓	↑	TCA +/- 3OH-IVA +/- Me.cit	C3 +/- C5OH	Normal

- ▶ SD Plots can be useful to highlight relative decreases (or increases)
- ▶ ? Other respiratory chain disorders / secondary mitochondrial dysfunction

References

- ▶ Peters et al Intravenous citrulline generation test to assess intestinal function in intensive care unit patients. *Clinical and Experimental Gastroenterology* 2017;10 75-81
- ▶ Ganetsky et al *MT-ATP6* Mitochondrial Disease Variants: Phenotypic and biochemical features analysis in 218 published cases and cohort of 14 new cases. *Hum Mutat* 2019; 40(5): 499-515.
- ▶ Capiou et al. Clinical Heterogeneity in *MT-ATP6* Pathogenic Variants: Same Genotype—Different Onset. *Cells* 2022; 11: 489.
- ▶ Wongkittichote et al. Low Plasma Citrulline Guiding the Diagnosis of a Mitochondrial Disorder. *Clin Chem* 2023; 69(6): 661-668
- ▶ Larson et al. Biochemical signatures mimicking multiple carboxylase deficiency in children with mutations in *MT-ATP6*. *Mitochondrion* 2019; 44: 58-64

Thank you!