



Dr Test Doctor (Test Doctor) Test Clinic. 123 Test Street, Test Suburb Victoria 3125

Lab ID
Patient ID P000060
Ext ID 26133-0053

Test Patient

Sex: Female • 56yrs • 01-Jan-70
123 Home Street, Test Suburb VIC 3125

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13-May-26

Clinical Notes .

NutriSTAT

Specimen type - Blood-Serum

Specimen type - Blood-Na Citrate

Specimen type - Blood-K2EDTA

Specimen type - Blood-Trace Metal

Specimen type - Blood-Red Cell

Specimen type - Blood-Whole

Specimen type - Blood-EDTA

Specimen type - Urine, Spot

Collected

20-Apr-26

20-Apr-26

20-Apr-26

20-Apr-26

20-Apr-26

20-Apr-26

20-Apr-26

20-Apr-26

RENAL FUNCTION TESTS (UEC)

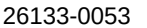
TEST	RESULT	H/L		REFERENCE	UNITS
Sodium	135			(135-145)	mmol/L
Potassium	5.2			(3.5-5.2)	mmol/L
Chloride	101			(95-110)	mmol/L
Bicarbonate	25			(22-32)	mmol/L
Anion Gap	14			(8-16)	mmol/L
Urea Nitrogen (BUN)	3.70			(3.50-8.00)	mmol/L
Creatinine	58			(45-90)	umol/L
Estimated GFR	>90			(>90)	mL/min/1.73m^2

LIVER FUNCTION TESTS (LFT)

[illegible]

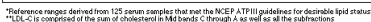
CVD INFLAMMATORY MARKERS

TEST	RESULT	H/L		REFERENCE	UNITS
Glucose	5.0			(3.0-5.4)	mmol/L
Homocysteine	18.0	H		(5.0-15.0)	umol/L
High Sensitivity C-Reactive Protein	5.5	H		(<5.0)	mg/L
Fibrinogen	4.2			(2.0-4.5)	g/L



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Type A - Normal



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[illegible]

Condition/Symptom	IRON	TRANSFERRIN SATURATION	FERRITIN	TRIAL OF ORAL IRON
Iron Deficiency	Decreased	Decreased	Decreased	Haemoglobin normalises
Iron Deficiency & Acute Phase Response	Decreased	Normal or Decreased	"Normal" <100ug/L	Partial Response
Acute Phase Response	Decreased	Decreased	Increased	No Response
Iron Overload	Increased	Increased	Increased	Not Appropriate

[illegible]



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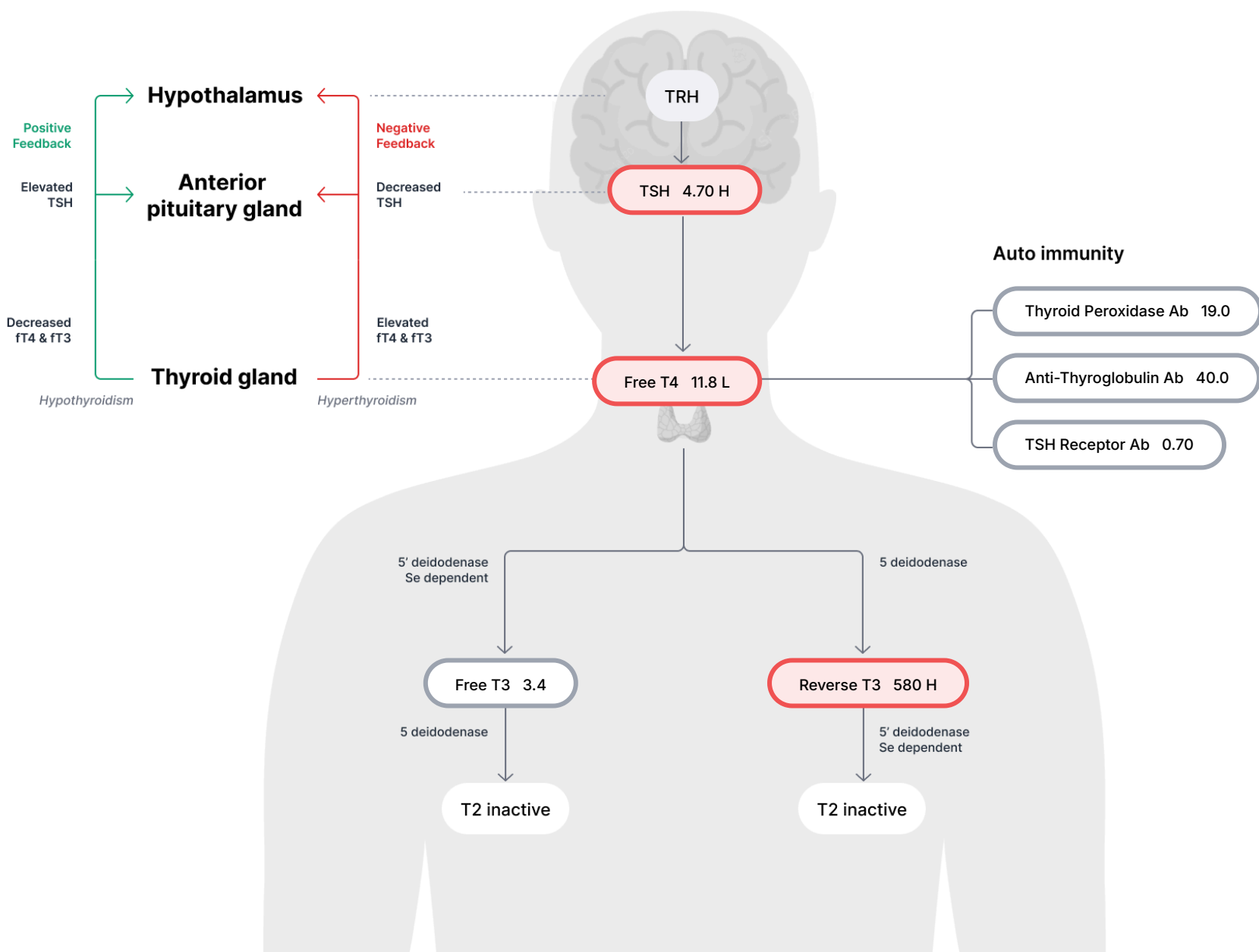
Test Patient

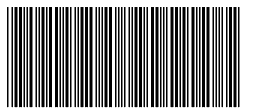
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Legend Not Tested Within Range Out of Range L = Low, LL = Critically Low H = High, HH = Critically High Regulator Enzyme

Thyroid System

[illegible]



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RED CELL MINERALS

TEST	RESULT	H/L		REFERENCE	UNITS
Chromium	1.09			(1.00-2.00)	ug/L
Cobalt	0.22			(0.13-1.70)	ug/L
Copper	1.0	H		(0.5-0.8)	ug/L
Iodine	87.00			(20.00-160.00)	ug/L
Magnesium	41.30			(39.00-58.00)	mg/L
Manganese	10.40			(9.00-33.00)	ug/L
Molybdenum	0.58	L		(0.60-2.00)	ug/L
Selenium	204.00			(190.00-500.00)	ug/L
Vanadium	0.25			(0.10-0.50)	ug/L
Zinc	9.7			(8.6-14.5)	ug/L

WHOLE BLOOD METALS

TEST	RESULT	H/L		REFERENCE	UNITS
Aluminium	24.00			(<30.00)	ug/L
Antimony	0.20			(<5.00)	ug/L
Arsenic	39.00	H		(<12.00)	ug/L
Beryllium	<DL			(<4.00)	ug/L
Bismuth	<DL			(<1.00)	ug/L
Cadmium	0.72			(<1.00)	ug/L
Lead	3.90			(<50.00)	ug/L
Mercury	3.8			(<10.1)	ug/L
Nickel	0.68			(<2.00)	ug/L
Platinum	<DL			(<0.40)	ug/L
Silver	0.04			(<2.00)	ug/L
Thallium	<DL			(<1.00)	ug/L
Tin	0.22			(<1.30)	ug/L
Uranium	<DL			(<0.10)	ug/L
Zirconium	0.75			(<1.32)	ug/L

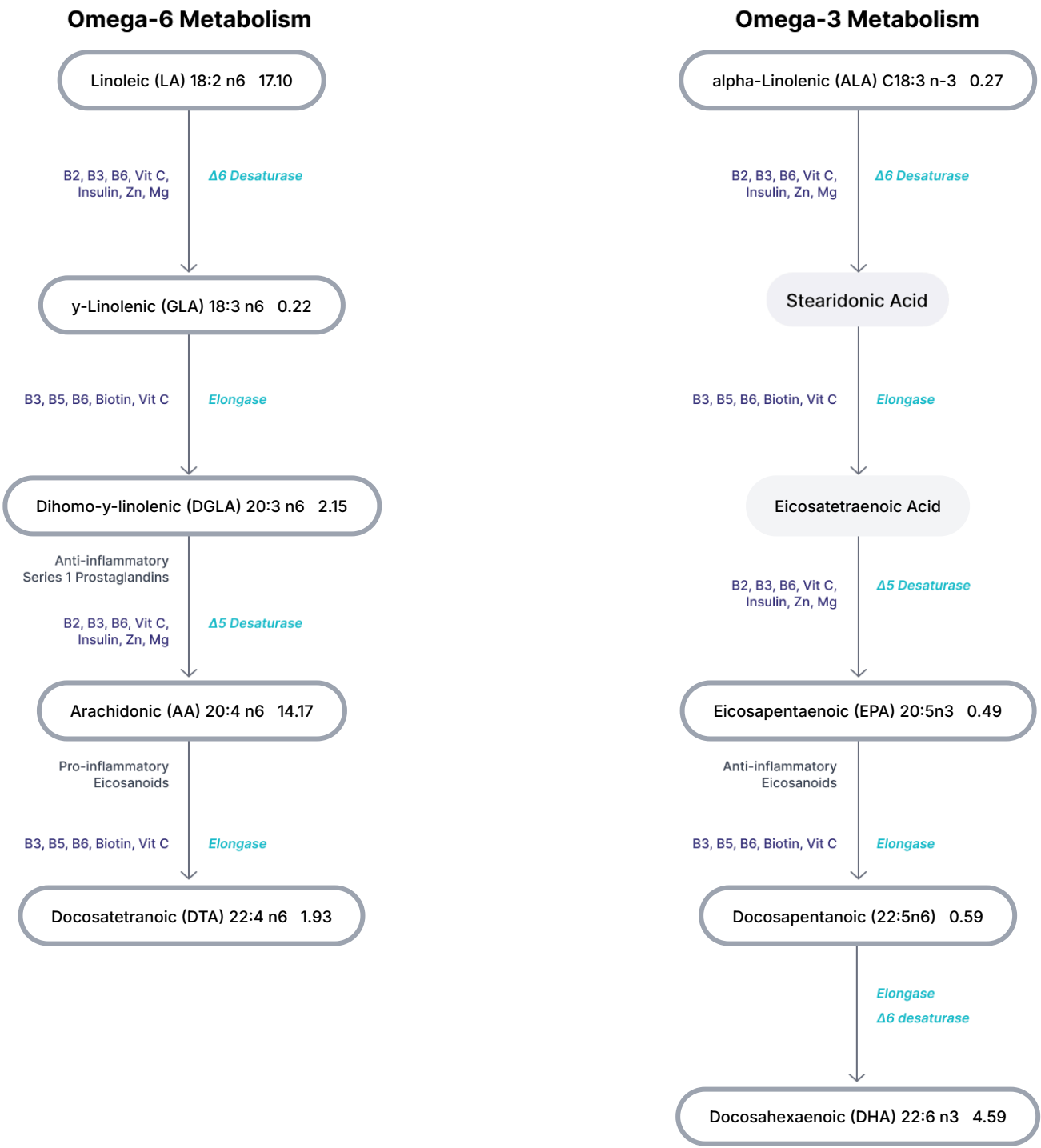
COPPER/ZINC BALANCE

TEST	RESULT	H/L		REFERENCE	UNITS
Copper	23.0	H		(13.0-22.0)	umol/L
Zinc	13.1			(9.0-19.0)	umol/L
Copper/Zinc Ratio	1.76	H		(0.54-1.68)	ratio
Caeruloplasmin	0.24			(0.20-0.60)	g/L
% Free Copper	51	H		(5-25)	%

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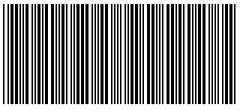
Essential Fatty Acids Pathway



Analyte with molecular structure shorthand

Linoleic acid (18:2 n-6)
 γ-Linolenic acid (18:3 n-6)
 Dihomo-γ-linolenic acid (20:3 n-6)
 Arachidonic acid (20:4 n-6)
 Docosatetraenoic acid (22:4 n-6)

- α-Linolenic acid (18:3 n-3)
- Eicosatetraenoic acid (20:4 n-3)
- Eicosapentaenoic acid (20:5 n-3)
- Docosapentaenoic acid (22:5 n-3)
- Docosahexaenoic acid (22:6 n-3)



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RED CELL FATTY ACIDS SUMMARY

TEST	RESULT	H/L	REFERENCE	UNITS
Saturated Fats, Total	37.38		(29.89-42.10)	%
Monounsaturated Fats, Total	18.57		(15.65-31.82)	%
Omega 3 Fatty Acids, Total	7.00		(2.57-15.15)	%
Omega 6 Fatty Acids, Total	36.36		(24.85-44.15)	%
Omega 3/Omega 6 Ratio	0.19		(0.07-5.30)	ratio
AA/EPA Ratio	28.92		(1.10-30.00)	ratio
Omega 3 Index	5.12		(>4.00)	%
Delta 6 Desaturase Activity (LA/DGLA)	7.95		(6.00-25.00)	ratio

OMEGA 3 FATTY ACIDS

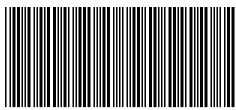
TEST	RESULT	H/L		REFERENCE	UNITS
alpha-Linolenic (ALA) C18:3 n-3	0.27			(0.10-1.90)	%
Eicosapentaenoic (EPA) 20:5n3	0.49			(0.14-6.92)	%
Docosapentaenoic (DPA) 22:5 n3	1.65			(0.53-2.81)	%
Docosaheptaenoic (DHA) 22:6 n3	4.59			(1.00-6.50)	%
Total Omega 3 Fatty Acids	7.00			(2.57-15.15)	%

OMEGA 6 FATTY ACIDS

[illegible]

MONOUNSATURATED FATTY ACIDS

[illegible]



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ESSENTIAL AMINO ACIDS

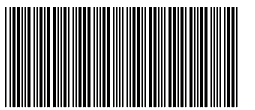
SERVICE		RESULT	H/L						REFERENCE	UNITS
1	Histidine	40.8			(39.0-123.0)	nmol/ml				
2	Isoleucine	46.0			(36.0-107.0)	nmol/ml				
3	Leucine	75.3			(68.0-254.0)	nmol/ml				
4	Lysine	65.0	L		(71.0-434.0)	nmol/ml				
5	Methionine	6.3			(4.0-44.0)	nmol/ml				
6	Phenylalanine	40.6			(35.0-80.0)	nmol/ml				
7	Threonine	63.8			(59.0-231.0)	nmol/ml				
8	Tryptophan	37.4			(29.0-77.0)	nmol/ml				
9	Valine	138.0			(119.0-336.0)	nmol/ml				
10	Total Branched Chain AAs	259			(223-697)	nmol/ml				

CONDITIONALLY ESSENTIAL AMINO ACIDS

[illegible]

NON-ESSENTIAL AMINO ACIDS

[illegible]



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INTERMEDIARY & FUNCTIONAL METABOLITES

(Functional markers of B-Vitamin status, Urea Cycle, & one-carbon metabolism)

[illegible]

DIETARY PEPTIDE RELATED MARKERS

SERVICE		RESULT	H/L		REFERENCE	UNITS
33	1-Methyl Histidine	4.76			(0.00-28.00)	nmol/ml
34	3-Methyl Histidine	12.9			(1.7-47.1)	nmol/ml
35	beta-Alanine	1.4			(0.0-5.0)	nmol/ml
36	Anserine	3.8			(0.0-43.0)	nmol/ml
37	Hydroxyproline	6.2			(3.0-29.0)	nmol/ml
38	Hydroxylysine	0.4	L		(0.5-3.2)	nmol/ml
39	Carnosine	0.56			(0.00-1.00)	nmol/ml

AMINO ACID FUNCTIONAL RATIOS

SERVICE		RESULT	H/L		REFERENCE	UNITS
40	Phenylalanine/Tyrosine	1.2			(0.0-2.0)	ratio
41	Glutamate/Glutamine	1.03	H		(0.10-0.40)	ratio
42	Hydroxyproline/Proline	0.1			(0.0-0.3)	ratio
43	a-Aminobutyrate/Leucine	0.2			(0.0-0.6)	ratio
44	Tryptophan/LNAAs	0.11			(0.08-0.40)	ratio



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Description: Hydroxylysine is formed by post-translational hydroxylation of lysine in collagen and elastin by lysyl hydroxylase (procollagen-lysine 2-oxoglutarate 5-dioxygenase), a vitamin C- and iron-dependent enzyme.

Clinical Significance: Low plasma hydroxylysine indicates impaired collagen cross-linking, which may be due to vitamin C deficiency, iron deficiency, or insufficient lysine or collagen precursor availability. It is associated with reduced collagen structural integrity, poor wound healing, joint laxity, and vascular fragility.

Suggested Treatment: Ensure adequate vitamin C (as ascorbate), iron, and lysine intake to support lysyl hydroxylase activity. Collagen hydrolysate or bone broth supplementation may provide hydroxylysine directly. Address any deficiencies identified in related nutrients.

GLUTAMATE/GLUTAMINE RATIO ELEVATED:

Description: The Glutamate/Glutamine ratio reflects glutamate-glutamine cycling and neurotransmitter balance, with glutamine synthetase converting glutamate to glutamine and glutaminase reversing this reaction.

Clinical Significance: An Elevated Glutamate/Glutamine ratio indicates excess excitatory tone relative to inhibitory buffering capacity. It may be associated with anxiety, neurological irritability, excitotoxicity risk, mitochondrial dysfunction, or impaired glutamine synthetase activity. Elevated dietary glutamate (MSG) or impaired glial recycling may also elevate this ratio.

Suggested Treatment: Address excitatory/inhibitory neurotransmitter balance by supporting GABA synthesis (B6, magnesium, taurine). Reduce dietary glutamate/MSG exposure. Evaluate mitochondrial function. Magnesium supplementation helps modulate NMDA receptor sensitivity to glutamate.



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MICROBIAL OVERGROWTH

3,4-Dihydroxybenzoic Acid (3,4-DHBA)
3-Hydroxyphenylacetic Acid (3-HPAA)
4-Hydroxybenzoic Acid (4-HBA)
Benzoic Acid
Furancarbonylglycine

YEAST/FUNGAL METABOLITES

5-Hydroxymethyl-2-furoic Acid
Arabinose

ENVIRONMENTAL EXPOSURES

3-Methylhippuric Acid
4-Hydroxybenzoic Acid (4-HBA)
5-Methylcytosine
Benzoic Acid
Quinolinic Acid
t,t-Muconic Acid

MITOCHONDRIA/ENERGY

2-Oxoglutaric Acid
Isocitric Acid
Mevalonolactone

FATTY ACID / KETONE METABOLISM

Adipic Acid

CARBOHYDRATE / GLYCAEMIC METABOLISM

N/A

NEUROTRANSMITTER METABOLISM

Kynurenic Acid
Quinolinic Acid
Quinolinic Acid/5-HIAA Ratio
Xanthurenic Acid

AMINO ACIDS METABOLISM

2-Oxisocaproic Acid
Guanidinoacetic Acid
N-Acetylphenylalanine
Phenylpyruvic Acid

VITAMINS / NUTRITIONAL MARKERS

Kynurenic Acid
Quinolinic Acid
Xanthurenic Acid

DETOXIFICATION / GLUTATHIONE FUNCTION

5-Hydroxymethyl-2-furoic Acid

OXIDATIVE DAMAGE / INFLAMMATION

8-hydroxy-deoxyguanosine
Leukotriene E4
Quinolinic Acid

OXALATE METABOLISM

N/A

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ORGANIC ACIDS INTRODUCTION

Organic acids are small carbon-based molecules produced as by-products of everyday metabolic processes in the body. They are measurable in urine and serve as functional windows into the biochemical pathways that govern energy production, detoxification, immune regulation, neurotransmitter synthesis, and nutritional status. Organic acid testing offers a uniquely deep and clinically meaningful picture of a patient's metabolic health.

The Organic Acids Test (OAT) is a comprehensive urine-based analysis that examines over 120 individual metabolic markers across the following key domains:

Microbial Overgrowth - Identifies markers associated with bacterial dysbiosis and overgrowth of potentially pathogenic organisms in the gastrointestinal tract, including phenolic and aromatic compounds produced by microbial fermentation of dietary substrates.

Yeast & Fungal Metabolites - Screens for metabolites associated with yeast and fungal overgrowth, including Candida-related markers such as arabinitol, tartaric acid, and tricarballic acid, which may interfere with mitochondrial function and nutrient absorption.

Mitochondrial & Energy Metabolism - Evaluates the functional integrity of the citric acid (TCA) cycle through measurement of key intermediates including citric acid, cis-aconitic acid, isocitric acid, and alpha-ketoglutaric acid.

Fatty Acid Oxidation & Ketone Metabolism - Assesses the efficiency of both mitochondrial beta-oxidation and the alternative omega-oxidation pathway through dicarboxylic acid markers, providing insight into fatty acid handling, carnitine sufficiency, and metabolic flexibility.

Amino Acid & Branched-Chain Metabolism - Examines the catabolism of essential and branched-chain amino acids, identifying impairments in leucine, isoleucine, valine, methionine, threonine, and tyrosine pathways that may contribute to fatigue, mood dysregulation, and neurological symptoms.

Neurotransmitter Metabolism -Profiles the major catecholamine and indole pathways, including dopamine, norepinephrine, and serotonin metabolites, as well as the tryptophan-kynurenine pathway, offering functional insight into mood, cognition, stress response, and sleep regulation.

Vitamins & Nutritional Cofactor Markers - Provides functional assessment of B-vitamin status (B1, B2, B3, B5, B6, B12, folate, biotin), glutathione and antioxidant capacity, and key cofactors required for enzymatic activity across multiple metabolic pathways.

Detoxification & Glutathione Function - Evaluates markers of phase I and phase II hepatic detoxification, glutathione cycling via pyroglutamic acid, and NAC status, reflecting the body's capacity to manage oxidative burden and chemical exposure.

Oxidative Stress & Inflammation - Measures markers of DNA oxidative damage and inflammatory mediators, including 8-hydroxy-deoxyguanosine and leukotriene E4, providing a functional picture of systemic oxidative load.

Oxalate Metabolism - Screens for markers of excess oxalate production or absorption, which may be associated with gut dysbiosis, dietary factors, or impaired detoxification.

Environmental & Xenobiotic Exposures - Identifies urinary metabolites of common environmental chemicals including xylene, benzene, toluene, phthalates, parabens, and endocrine-disrupting compounds such as Bisphenol A (BPA) and Bisphenol S (BPS), providing objective evidence of toxic load.

Please Note: We acknowledge that certain compounds appear in multiple sections of the report. This is intentional, as these analytes have relevance across different clinical pathways. Their inclusion in multiple categories supports more accurate pattern recognition and enhances the interpretive value of Organic Acids testing.



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Food Source Fats

- Fatty Acids → Fatty Acyl-CoA → Adipic Acid 4.30 H → Suberic Acid 1.33 → β-Oxidation → Acetyl-CoA
- Omega Oxidation: Fatty Acids → Ethylmalonic Acid 4.10 → Methylsuccinic Acid 1.65
- Vitamins/Minerals: B2, B3, Mg, L-carnitine; Mg, Mn

Food Source Carbohydrates

- Glucose (OA) 0.98 → Glycolysis → Pyruvic Acid 1.60 → Acetyl-CoA
- Vitamins/Minerals: B1, B2, B3, B6, Mn, Mg, K

Food Source Proteins

- Amino Acids → Alanine → Gluconeogenesis → Glucose (OA) 0.98
- Vitamins/Minerals: B6, Zn, GSH; B1, B2, B3, B5

Citric Acid Cycle

- Acetyl-CoA + Oxaloacetic Acid → Citric Acid 280
- Citric Acid → cis-Aconitic Acid 12.70
- cis-Aconitic Acid → Isocitric Acid 88.00 H
- Isocitric Acid → 2-Oxoglutaric Acid 65.00 H
- 2-Oxoglutaric Acid → Succinyl-CoA
- Succinyl-CoA → Succinic Acid 12.00
- Succinic Acid → Fumaric Acid 4.30
- Fumaric Acid → Malic Acid 1.50
- Malic Acid → Oxaloacetic Acid
- Vitamins/Minerals: B3; B2, B3, Fe, Zn, Antioxidants; B2, B3, Cu, Fe, Zn, Resveratrol, CoQ10, Antioxidants; Mg; B1, B2, B3, r-lipoc acid, Ca

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(Phenylalanine, Tyrosine)

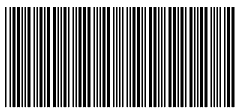
TEST	RESULT	H/L	REFERENCE	UNITS
Inhibitory (Serotonin)				
101	5-Hydroxyindoleacetic Acid (5-HIAA)	1.14	(<4.30)	mmol/molCR
Excitatory (Dopamine)				
102	3,4-Dihydroxyphenylacetic Acid (DOPAC)	0.15	(0.08-3.50)	mmol/molCR
103	3-Methyl-4-hydroxyphenylglycol (MHPG)	0.18	(0.05-0.50)	mmol/molCR
104	Homovanillic Acid (HVA)	0.62	(0.10-5.30)	mmol/molCR
105	Vanillylmandelic Acid (VMA)	1.13	(0.40-3.60)	mmol/molCR
106	HVA/DOPAC Ratio	4.1	(<10.0)	ratio
107	HVA/VMA Ratio	0.5	(<2.0)	ratio

[illegible]

Enzyme

The diagram illustrates the metabolic pathways of catecholamines. It begins with Phenylalanine, which can be converted to PEA (Phenylethylamine) via Vitamin B6. Phenylalanine is also converted to Tyrosine, a process requiring MTHF, BH4, Fe, and Vitamins B3, C, and D. Tyrosine is then converted to L-DOPA, also requiring MTHF, BH4, Fe, and Vitamins B3, C, and D. L-DOPA is converted to Dopamine via Vitamin B6. Dopamine can be converted to 3,4-Dihydroxyphenylacetic Acid (DOPAC) via MAO (Monoamine oxidase), which then leads to Homovanillic Acid (HVA). Dopamine is also converted to Noradrenalin via MTHF, BH4, Fe, and Vitamins B3, C, and D. Noradrenalin can be converted to Normetanephrine via COMT (Catechol-O-methyltransferase), which then leads to MHPG (3-Methoxyphenylethanolamine). Noradrenalin is also converted to DHPG (3,4-Dihydroxyphenylethanolamine) via MAO, which then leads to MHPG. Adrenalin is converted to Metanephrine via COMT, which then leads to MHPG. MHPG is converted to Vanillylmandelic Acid (VMA) via Mg. The final products are DOPAC (0.15), HVA (0.62), and VMA (1.13).

```
graph TD
    Phenylalanine -- "Vit B6" --> PEA
    Phenylalanine -- "MTHF, BH4, Fe, Vits B3, C, D" --> Tyrosine
    Tyrosine -- "Vit B6" --> Tyramine
    Tyrosine -- "MTHF, BH4, Fe, Vits B3, C, D" --> LDOPA[L-DOPA]
    LDOPA -- "Vit B6" --> Dopamine
    Dopamine -- "Cu, Vit B2, MAO" --> DOPAC["3,4-Dihydroxyphenylacetic Acid (DOPAC) 0.15"]
    DOPAC --> HVA["Homovanillic Acid (HVA) 0.62"]
    Dopamine -- "MTHF, BH4, Fe, Vits B3, C, D" --> Noradrenalin
    Noradrenalin -- "Mg, SAmE, COMT" --> Normetanephrine
    Normetanephrine -- "Cu, Vit B2, MAO" --> MHPG
    Noradrenalin -- "MAO" --> DHPG
    DHPG -- "Mg, SAmE, COMT" --> MHPG
    Adrenalin -- "Mg, SAmE, COMT" --> Metanephrine
    Metanephrine -- "Cu, Vit B2, MAO" --> MHPG
    MHPG -- "Mg" --> VMA["Vanillylmandelic Acid (VMA) 1.13"]
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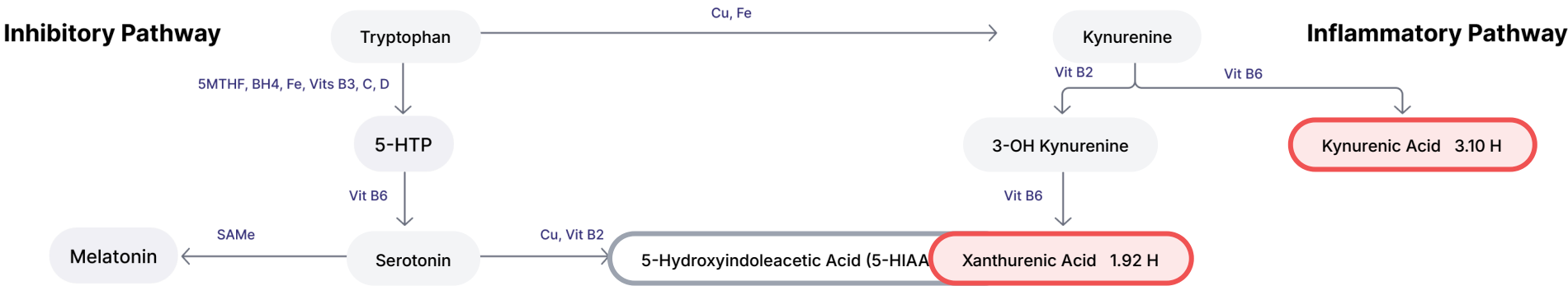
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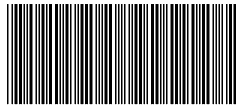
TRYPTOPHAN/KYNURENINE PATHWAY

(Tryptophan, B6, Antioxidants)

TEST	RESULT	H/L		REFERENCE	UNITS
Inhibitory					
109 5-Hydroxyindoleacetic Acid (5-HIAA)	1.14			(<4.30)	mmol/molCR
Inflammatory					
110 Kynurenic Acid	3.10	H		(<2.20)	mmol/molCR
111 Quinolinic Acid	11.80	H		(<9.10)	mmol/molCR
112 Picolinic Acid	4.20			(<10.28)	mmol/molCR
113 Xanthurenic Acid	1.92	H		(<0.96)	mmol/molCR
114 Kynurenic/Quinolinic Ratio	0.3			(<2.0)	ratio
115 Quinolinic Acid/5-HIAA Ratio	10.4	H		(<5.0)	ratio

Legend Not Tested Within Range Out of Range L = Low, LL = Critically Low H = High, HH = Critically High Regulator Enzyme





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DETOXIFICATION & GLUTATHIONE FUNCTION

(Arg, NAC, Meth, Mg, Antioxidants)

TEST	RESULT	H/L	REFERENCE	UNITS
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Ammonia Metabolism

116	Orotic Acid	1.95		(<3.20)	mmol/molCR
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Glutathione Metabolism

117	Pyroglutamic Acid	28.80		(4.50-33.00)	mmol/molCR
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118	N-Acetylcysteine (NAC)	0.05		(0.02-0.28)	mmol/molCR
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119	Glucaric Acid	8.40		(<11.00)	mmol/molCR
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Phase I / Xenobiotic Markers

120	2-Hydroxyhippuric Acid	0.80		(<1.50)	mmol/molCR
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Index	Compound Name	Score	Label	Bar Chart	Value	Unit
121	5-Hydroxymethyl-2-furoic Acid	18.50	H		(<10.00)	mmol/molCR

122	Furan-2,5-dicarboxylic Acid	12.90	●	(<15.00)	mmol/molCR
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OXIDATIVE STRESS & INFLAMMATION

(Vitamin C, Other Antioxidants)

TEST	RESULT	H/L	REFERENCE	UNITS
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Index	Metabolite	Value	Unit	Color Scale	Reference	Unit
123	8-hydroxy-deoxyguanosine	3.80	H		(<2.70)	umol/molCR

124	Leukotriene E4	195.00	H		(<100.00)	pg/mgCR
-----	----------------	--------	---	---	-----------	---------

125	Quinolinic Acid	11.80	H						●		(<9.10)	mmol/molCR
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126	Pyroglutamic Acid	28.80		(4.50-33.00)	mmol/molCR
-----	-------------------	-------	---	--------------	------------

127	Ascorbic Acid (Vit C)	59.00		(0.50-200.00)	mmol/molCR
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128	Quercetin	0.68		(0.01-2.00)	mmol/molCR
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OXALATE METABOLISM

TEST	RESULT	H/L	REFERENCE	UNITS
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129	Glycolic Acid	20.80		(<67.00)	mmol/molCR
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Index	Compound	Value	Color Scale	Unit
130	Glyceric Acid	3.80		(<6.00) mmol/molCR

131	Oxalic Acid	42.00		(<78.00)	mmol/molCR
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NUTRITION GUIDE			
SERVICE	Nutritional Need	DOSE	UNITS

Amino Acids

5-HTP	HIGH	80.0	mg
Acetyl-L-Carnitine	Adequate	0.0	mg
L-Arginine	Adequate	0.0	mg
Aspartic Acid	Adequate	0.0	mg
Glutamine	Adequate	0.0	mg
Glycine	Adequate	0.0	mg
Lysine	Adequate	0.0	mg
Methionine	Adequate	0.0	mg
Ornithine	Adequate	0.0	mg
Phenylalanine	Adequate	0.0	mg
Serine	Adequate	0.0	mg
Taurine	Adequate	0.0	mg
Tryptophan	Adequate	0.0	mg
Tyrosine	Adequate	0.0	mg

Probiotics

● D-Lactate-free probiotics	Mild	20.0	billion CFU
● Lactobacillus	HIGH	40.0	billion CFU
● Probiotics (Multistrain)	Mild	20.0	billion CFU

Other

● Malic Acid	Adequate	0.0	mg
● EPA/DHA	Adequate	0.0	mg
● Folinic Acid	Adequate	0.0	ug



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About the Nutrition Guide & Supplement Schedule

The Nutrition Guide presented in this report represents a personalised supplementation schedule generated directly from the organic acid findings of this individual patient. It is not a generic recommendation it is dynamically calculated for each patient based on the specific metabolic pattern identified across all measured domains.

The schedule is generated using a proprietary weighted algorithm developed by our clinical and scientific team. The algorithm analyses the full pattern of organic acid results not individual markers in isolation and assigns a weighted score to each finding based on its functional significance, its relationship to nutrient-dependent enzymatic pathways, and its interaction with other metabolic markers present in the same report. The outcome of this calculation determines both the nutritional priority level (Adequate, Mild, Moderate, or High) and the suggested starting dose for each nutrient, amino acid, probiotic, or cofactor listed.

Important Disclaimer

The Nutrition Guide and supplementation suggestions in this report are generated for clinical decision-support purposes only and are not intended as standalone medical advice or a prescription.

The proprietary algorithm used to generate this schedule is based on established relationships between organic acid biomarkers and nutritional cofactor requirements. However, it operates solely on biochemical data and cannot account for a patient's complete medical history, current medications, existing diagnoses, organ function, pregnancy or breastfeeding status, or any other health condition that may influence the safety or appropriateness of the suggested nutrients and doses.

The final therapeutic decision including whether to implement, modify, or withhold any recommendation in this report rests entirely with the treating practitioner. It is the practitioner's responsibility to integrate these findings with the full clinical picture before making any therapeutic recommendation.

This report does not diagnose, treat, cure, or prevent any disease or medical condition.

Nutritional Guide Practitioner Notes



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BENZOIC ACID ELEVATED (URINE):

Consider: treatment for dysbiosis and diet changes, mucosal support, pre and probiotics

Consider: Treatment for dysbiosis and diet changes, mucosal support, pre and probiotics.

Evaluate catecholamine pathway function and oxidative stress markers. Support antioxidant status with N-acetylcysteine, glutathione precursors, and vitamin C. Address gut dysbiosis if implicated.

Consider targeted antimicrobial or herbal antimicrobial therapy (e.g., oregano oil, berberine). Implement gut restoration protocol including dietary modification, probiotics, and intestinal barrier support (L-glutamine, zinc carnosine). Retest after intervention.



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FURANCARBONYLGLYCINE ELEVATED:

Furancarboxylglycine (FCG) is a glycine-conjugated metabolite formed during the hepatic detoxification of furan-derived compounds, which are commonly generated during the thermal processing of foods and from certain environmental exposures.

Elevated urinary furancarboxylglycine may reflect increased exposure to furan-containing compounds, with subsequent metabolism via glycine conjugation pathways and renal excretion. This is most commonly associated with dietary intake of heat-processed foods, although environmental sources may also contribute.

As a phase II detoxification product, this marker primarily reflects exposure and metabolic processing rather than intrinsic metabolic dysfunction, and is therefore non-specific when interpreted in isolation.

Suggested Treatment Considerations:

Consider review of dietary intake, particularly consumption of highly processed or heat-treated foods. Assessment of environmental exposures may be appropriate where clinically indicated. No specific medical intervention is typically required beyond addressing identifiable sources. Interpretation should be guided by the overall clinical context and associated findings.

Yeast Dysbiosis Comment

ARABINOSE ELEVATED:

Arabinose is a pentose sugar that may be detected in urine as a product of dietary intake and gastrointestinal microbial metabolism. It is not a primary endogenous human metabolite and is typically present at low concentrations.

Elevated urinary arabinose may be associated with increased gastrointestinal microbial production of arabinose, which has been described in the context of altered gut microbial activity, including possible overrepresentation of certain yeast or bacterial species. However, this finding is non-specific and may also be influenced by dietary sources and intestinal metabolism.

As such, arabinose should not be used in isolation as a diagnostic marker of a specific organism, and is best interpreted alongside other markers of microbial metabolism and gastrointestinal function.

Suggested Treatment Considerations:

Consider correlation with other markers of gastrointestinal microbial activity and clinical features suggestive of altered gut function. Review dietary intake and factors that may influence microbial balance. Further evaluation of gastrointestinal health may be appropriate where clinically indicated. Management should be guided by the overall clinical context and associated laboratory findings rather than the isolated elevation of this marker.

5-HYDROXYMETHYL-2-FUROIC ACID ELEVATED:

5-Hydroxymethyl-2-furoic acid (5-HMFA) is a urinary metabolite derived from the metabolism of 5-hydroxymethylfurfural (HMF), a compound formed during the thermal processing of carbohydrate-containing foods, particularly under high heat conditions (e.g. baking, roasting, or caramelisation).

Elevated urinary 5-HMFA typically reflects increased dietary exposure to heat-processed foods, with subsequent hepatic metabolism and excretion. As such, this marker is primarily considered an indicator of dietary intake and exposure, rather than endogenous metabolic dysfunction.

In some contexts, elevated levels may also reflect increased formation of furan-derived compounds associated with carbohydrate degradation during cooking. This finding is non-specific and should be interpreted in conjunction with dietary history and other relevant exposure markers.

Suggested Treatment Considerations:

Review dietary intake, particularly consumption of highly processed or heat-treated foods. Where appropriate, consider reducing intake of foods subjected to high-temperature processing. No specific medical intervention is typically required beyond addressing identifiable dietary sources. Interpretation should be guided by the overall clinical context and associated findings.

Mitochondrial/Energy Metabolism Comment

ISOCITRIC ACID ELEVATED (URINE):



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Elevated urinary isocitric acid suggests altered downstream TCA cycle efficiency or compensatory metabolic flux.

Clinically, symptoms are non-specific.

From an organic acid pattern perspective, interpretation alongside alpha-ketoglutaric acid assists in determining whether downstream enzymatic congestion is present.

From a functional medicine perspective, this finding should be interpreted as part of an overall mitochondrial pattern.

A high level is suggestive of inhibition to the enzyme by Aluminum.

Supplementation Recommendations: Cofactors needed to increase the breakdown of isocitrate to alpha-ketoglutarate are: Vit B3, (NAD), Mg, Mn.

2-OXOGLUTARIC ACID ELEVATED (URINE)

Elevated urinary alpha-ketoglutaric acid suggests a bottleneck within the TCA cycle and altered nitrogen handling.

Clinically, elevated alpha-ketoglutaric acid may be associated with fatigue, cognitive symptoms, or reduced exercise tolerance.

From an organic acid pattern perspective, elevation often occurs alongside elevated succinic acid, reflecting downstream congestion and potential oxidative stress.

From a functional medicine perspective, this finding should be interpreted in the context of mitochondrial efficiency, redox balance, and amino-acid metabolism.

Elevations can be seen with nutrient cofactor deficiencies needed for the enzymatic conversion of α ketoglutarate such as vitamin B3, zinc, magnesium, manganese.

MEVALONOLACTONE ELEVATED:

Mevalonolactone is the lactone form of mevalonic acid, an intermediate in the mevalonate pathway, which is essential for the synthesis of cholesterol, isoprenoids, and other biologically important molecules. This pathway is regulated by HMG-CoA reductase, the rate-limiting enzyme in cholesterol biosynthesis.

Elevated urinary mevalonolactone may reflect increased activity of the mevalonate pathway, potentially indicating upregulated cholesterol and isoprenoid synthesis. This pattern may be observed in the context of metabolic dysregulation, increased anabolic demand, or altered lipid metabolism. In some cases, elevations may also be associated with inflammatory states, as the mevalonate pathway is involved in the production of intermediates important for immune cell function.

While increased levels may provide insight into pathway activity, this finding is non-specific and should be interpreted in conjunction with lipid markers, inflammatory indicators, and overall metabolic context.

Suggested Treatment Considerations:

Consider correlation with lipid profile and markers of metabolic and inflammatory status. Review dietary and metabolic factors influencing cholesterol synthesis where clinically indicated. Further evaluation may be warranted in cases of persistent or significant elevation. Management should be guided by the overall clinical context and associated laboratory findings.

3-METHYLGLUTACONIC ACID ELEVATED:

3-Methylglutaconic acid (3-MGC) is a dicarboxylic acid intermediate in the leucine catabolism pathway. Under normal conditions, 3-methylglutaconyl-CoA is hydrated by the enzyme 3-methylglutaconyl-CoA hydratase to form HMG-CoA, which enters ketogenesis. When this step is impaired — due to genetic enzyme deficiency or acquired mitochondrial dysfunction — 3-MGC accumulates and is excreted in urine. Beyond leucine metabolism, 3-MGC has been established as a broader non-specific marker of mitochondrial inner membrane dysfunction, elevated across a range of mitochondrial disorders including Barth syndrome (TAZ gene mutation), DNAJC19-related cardiomyopathy, and SERAC1-related MEGDEL syndrome. In the functional medicine context, mild to moderate elevation most commonly reflects acquired mitochondrial stress and nutritional cofactor insufficiency (CoQ10, B2, carnitine, magnesium) rather than primary genetic deficiency.

Clinical Significance:

Elevated urinary 3-MGC is a meaningful indicator of mitochondrial dysfunction and warrants assessment of the broader mitochondrial marker pattern. Isolated mild elevation in a clinically well adult most likely reflects functional mitochondrial stress, cofactor depletion, or



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reactive oxygen species under conditions of elevated oxidative stress. KMBA therefore signals potential disruption at the critical junction between methionine metabolism, SAMe synthesis, transsulfuration, and glutathione production.

Clinical Significance:

Elevated KMBA indicates impaired methionine catabolism and/or transsulfuration pathway dysfunction with downstream consequences for methylation and glutathione synthesis. Reduced SAMe production impairs methylation reactions essential for neurotransmitter synthesis (dopamine, serotonin, norepinephrine), DNA and histone methylation, phospholipid metabolism, and immune regulation. Impaired transsulfuration reduces cysteine availability and consequently limits glutathione production compromising phase II detoxification, antioxidant defence, and the body's capacity to manage chemical and oxidative burden. If remethylation is concurrently impaired, homocysteine may accumulate, representing an independent cardiovascular and neurological risk factor. Clinically, this pattern is associated with fatigue, cognitive impairment and brain fog, poor detoxification, chemical sensitivity, mood dysregulation, and susceptibility to inflammatory and autoimmune conditions. The combination of elevated KMBA, elevated xanthurenic acid, and low pyroglutamic acid is a strong functional indicator of B6 insufficiency as the primary driver. Plasma homocysteine and serum B12, folate, and B6 levels should be assessed to directly characterise methylation pathway status.

PHENYLPYRUVIC ACID ELEVATED:

Phenylpyruvic acid is a keto-acid formed during the catabolism of phenylalanine, typically generated when phenylalanine is transaminated rather than converted via its primary metabolic pathway to tyrosine.

Elevated urinary phenylpyruvic acid may reflect impaired phenylalanine metabolism, most notably due to reduced activity of phenylalanine hydroxylase, as seen in phenylketonuria (PKU). In such cases, accumulation of phenylalanine leads to increased production of alternative metabolites, including phenylpyruvate.

Milder elevations may be observed in the context of increased phenylalanine load, altered amino acid metabolism, or metabolic stress, although these findings are non-specific and should be interpreted cautiously.

Given its clinical relevance, particularly in inherited metabolic disorders, this marker should be interpreted in conjunction with plasma amino acid profiles and other phenylalanine-related metabolites.

Suggested Treatment Considerations:

Consider correlation with plasma phenylalanine levels and clinical history. In cases of significant or persistent elevation, further evaluation for disorders of phenylalanine metabolism, including phenylketonuria, may be warranted. Referral for specialist metabolic assessment may be appropriate. Management should be guided by the underlying clinical diagnosis and overall context.

N-ACETYLPHENYLALANINE ELEVATED:

N-acetylphenylalanine is an acetylated derivative of the amino acid phenylalanine, formed through N-acetylation pathways as part of amino acid metabolism and phase II detoxification processes. It represents a minor urinary metabolite reflecting aromatic amino acid metabolism and conjugation activity.

Elevated urinary N-acetylphenylalanine may reflect increased flux through phenylalanine metabolism and/or enhanced N-acetylation activity, which may occur in the context of increased substrate availability, altered amino acid metabolism, or metabolic demand. In addition, elevations may be influenced by gastrointestinal microbial metabolism of aromatic compounds, contributing to altered phenylalanine handling.

This marker is non-specific and is best interpreted alongside other metabolites within the phenylalanine/tyrosine pathway and markers of microbial activity to assess potential patterns of metabolic or gastrointestinal involvement.

Suggested Treatment Considerations:

Consider correlation with other markers of aromatic amino acid metabolism and gastrointestinal function. Review dietary protein intake and assess for factors influencing microbial metabolism where clinically indicated. Further evaluation should be guided by the overall clinical context and associated laboratory findings rather than the isolated marker elevation.

GUANIDINOACETIC ACID ELEVATED:

Guanidinoacetic acid (GAA) is an intermediate in creatine biosynthesis, formed from arginine and glycine via arginine:glycine amidinotransferase (AGAT) and subsequently converted to creatine by guanidinoacetate methyltransferase (GAMT) using S-



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Detoxification / Oxidative Damage Comment:

8-HYDROXY-2-DEOXYGUANOSINE ELEVATED (URINE):

8-OHdG is a marker of oxidative DNA damage. Elevation reflects increased oxidative stress and impaired antioxidant defence.

Clinically, elevations may be associated with fatigue, accelerated ageing features, inflammatory symptoms. Interpretation should consider dietary intake, medications, and the broader metabolic context.

From a functional medicine perspective, management focuses on reducing oxidative load, enhancing antioxidant capacity, and addressing environmental or metabolic stressors, addressing underlying contributors rather than isolated suppression of the marker.

Consider: Supplementation with antioxidants such as vitamin C, E, N-acetyl cysteine, lipoate.

LEUKOTRIENE E4 ELEVATED:

Leukotriene E4 (LTE4) is a stable urinary metabolite of the cysteinyl leukotrienes (LTC4, LTD4, and LTE4), which are lipid mediators derived from arachidonic acid via the 5-lipoxygenase pathway. These mediators play a key role in inflammatory and immune responses, particularly in MCAS, bronchoconstriction, vascular permeability, and recruitment of inflammatory cells.

Elevated urinary LTE4 reflects increased systemic leukotriene production and is considered a non-invasive marker of inflammatory pathway activation, particularly within allergic and respiratory conditions. Increased levels may be observed in association with asthma, allergic disease, mast cell activation, and other inflammatory states, reflecting upregulation of leukotriene-mediated signalling.

As LTE4 represents a downstream metabolite, it provides an indication of overall leukotriene pathway activity rather than a specific site of inflammation, and should be interpreted in conjunction with clinical findings and other markers of inflammatory or immune activity.

Suggested Treatment Considerations:

Consider correlation with clinical features of inflammatory or allergic disease, including respiratory symptoms or hypersensitivity patterns. Evaluation of factors contributing to inflammatory activation, including environmental exposures and immune triggers, may be appropriate. Management should be directed toward the underlying inflammatory process, with further investigation guided by the overall clinical context and associated laboratory findings.

Environmental/Xenobiotic Exposure Comment:

ENVIRONMENTAL POLLUTANTS PROFILE:

The reported markers in the Environmental Pollutants Profile commonly originate from industrial/manufacturing products or their associated byproducts. Exposures are often occupationally-related and typically through either inhalation or topical exposure.

Metabolism of these products occurs via the liver detoxification pathways leading to excretion into the urine. Chronic exposures may also lead to build up of these products in fatty tissue deposits.

BENZOIC ACID ELEVATED (URINE):

Elevated urinary benzoic acid suggests increased exposure to benzoate-containing foods, preservatives, or altered gut microbial metabolism. Benzoic acid is detoxified primarily through glycine conjugation to form hippuric acid.

Clinically, elevated benzoic acid may be associated with headaches, fatigue, gastrointestinal discomfort, or chemical sensitivity, although symptoms are often non-specific.

From an organic acid pattern perspective, elevated benzoic acid may be observed alongside elevated hippuric acid, indicating increased benzoate load or increased demand on glycine-dependent conjugation pathways.

From a functional medicine perspective, elevated benzoic acid should be interpreted in the context of dietary intake, glycine availability, gut microbial activity, and overall detoxification capacity rather than as an isolated finding.

Treatment: Limiting exposure to toluene. Supportive supplements such as glycine and N-acetyl cysteine can support natural detoxification.

4-HYDROXYBENZOIC ACID ELEVATED (URINE):

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Elevated urinary 4-hydroxybenzoic acid suggests increased exposure to parabens or altered metabolism of aromatic compounds. Parabens are commonly used as preservatives in cosmetics, personal care products, and some pharmaceuticals.

Clinically, elevated 4-hydroxybenzoic acid may be associated with non-specific symptoms such as fatigue, headaches, or endocrine-related concerns, although many individuals remain asymptomatic.

From an organic acid pattern perspective, elevated 4-hydroxybenzoic acid may be seen alongside other aromatic or phenolic metabolites, reflecting cumulative preservative or polyphenol exposure.

From a functional medicine perspective, this finding should be interpreted in the context of environmental and personal care exposures, hepatic conjugation capacity, and cumulative endocrine-disrupting burden, with emphasis on reducing ongoing exposure.

Treatment: Treatments focus on improving gut health through dietary changes like increasing plant-based foods, fermented foods, and prebiotics while reducing artificial sweeteners and potentially eliminating paraben-containing products.

t,t-MUCONIC ACID ELEVATED (URINE):

Elevated urinary t,t-muconic acid suggests increased exposure to benzene, as this metabolite represents a recognised biomarker of benzene metabolism and detoxification.

Clinically, elevated t,t-muconic acid may be associated with fatigue, headaches, dizziness, or non-specific neurological symptoms, although clinical effects are highly dependent on exposure magnitude and duration.

From an organic acid pattern perspective, elevated t,t-muconic acid may be observed alongside other aromatic or solvent-related metabolites, reflecting increased aromatic hydrocarbon burden and hepatic detoxification demand.

From a functional medicine perspective, this finding should be interpreted in the context of environmental and occupational exposure sources (e.g. fuel vapours, tobacco smoke, solvents), indoor air quality, and overall detoxification capacity, with primary emphasis on exposure identification and reduction.

Treatment: Treatment usually involves removing/limiting exposure; and therefore avoiding chronic exposure which can lead to severe consequences.

QUINOLINIC ACID ELEVATED (URINE):

Elevated urinary quinolinic acid suggests increased flux through the kynurenine pathway of tryptophan metabolism, often associated with immune activation or inflammatory signalling. Quinolinic acid is a neuroactive metabolite with excitatory properties.

Clinically, elevated quinolinic acid may be associated with neurocognitive symptoms such as brain fog, mood disturbance, irritability, sleep disruption, or heightened pain sensitivity, although symptom expression varies.

From an organic acid pattern perspective, elevated quinolinic acid is often observed alongside alterations in other tryptophan metabolites, indicating inflammatory diversion of tryptophan metabolism away from serotonin and melatonin pathways.

From a functional medicine perspective, this finding should be interpreted in the context of immune activation, inflammatory burden, and overall balance of tryptophan metabolism rather than as a primary neurological disorder.

Treatment: Treatments for high quinolinic acid focus on reducing its production and counteracting its neurotoxic effects, including dietary changes like increasing vitamin B6 and consuming antioxidants (e.g., sulforaphane, green tea polyphenols, curcumin), and supplements (melatonin, selenium, theanine).

3-METHYLHIPPURIC ACID ELEVATED (URINE):

Elevated urinary 3-methylhippuric acid suggests increased exposure to xylene, as this metabolite represents glycine conjugation of xylene-derived methylbenzoic acids. Urinary methylhippuric acids are well-established biomarkers of xylene exposure.

Clinically, elevated 3-methylhippuric acid may be associated with headaches, dizziness, fatigue, mucosal irritation, or central nervous system symptoms, depending on exposure magnitude and duration.

From an organic acid pattern perspective, elevated 3-methylhippuric acid may be observed alongside increased hippuric acid or other aromatic conjugates, reflecting increased aromatic solvent burden and glycine-dependent detoxification demand.

From a functional medicine perspective, this finding should be interpreted in the context of occupational, environmental, or household solvent exposure (e.g. paints, fuels, adhesives), alongside assessment of hepatic conjugation capacity and glycine availability, with



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emphasis on exposure identification and reduction.

Treatment: Treatment options include limiting exposure to xylenes and supportive supplements such as glycine and N-acetyl cysteine can support natural detoxification.

5-METHYLCYTOSINE ELEVATED:

Urinary 5-methylcytosine (5-MC) is a modified nucleoside derived from DNA methylation and reflects global DNA methylation turnover and nucleic acid metabolism. Elevated levels may indicate increased DNA turnover, enhanced methylation flux, or increased degradation of methylated cytosine residues.

This pattern may be observed in the context of increased cellular turnover, oxidative stress, inflammatory processes, or enhanced DNA repair activity, and may also be influenced by environmental exposures, including certain xenobiotics.

While 5-MC provides insight into methylation turnover, it does not directly assess methylation capacity. Interpretation should be made in conjunction with clinical findings and, where appropriate, other markers of one-carbon metabolism and oxidative stress.

Suggested Treatment Considerations:

Review potential contributors to increased oxidative stress and environmental exposures. Consider assessment of one-carbon metabolism and methyl donor status (e.g. folate, vitamin B12, choline) where clinically indicated. Supportive strategies may include optimisation of nutritional status and antioxidant capacity. Further evaluation should be guided by the overall clinical context and associated laboratory findings.

Methodology

Automated Chemistry/Immunochemistry, Gel Electrophoresis, Ion-Selective Electrodes (ISE), Inductively Coupled Plasma Mass Spectrometry (ICP-MS), Liquid Chromatography-Mass Spectrometry (LC-MS/MS/MS), Gas Chromatography-MS (GC/MS)