

Metabolic Analysis Profile Support Guide



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Metabolic Analysis Profile

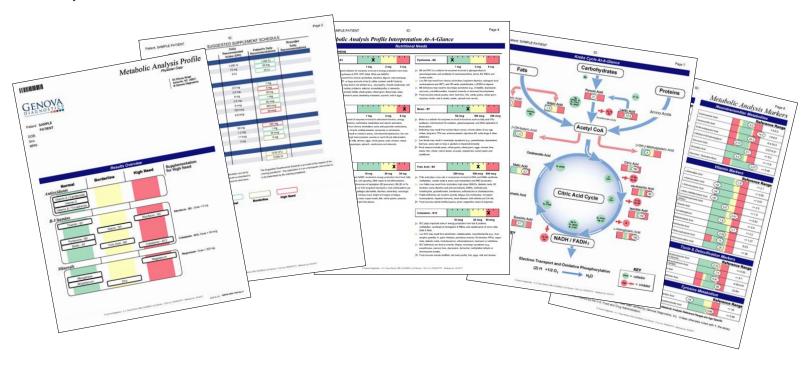
The **Metabolic Analysis Profile** is a functional nutritional assessment of urinary organic acids. Organic acids are a broad class of compounds formed during fundamental metabolic processes in the body. Metabolic reactions produce carboxylic acid compounds derived from the digestion of dietary protein, fat, and carbohydrates. The resulting organic acids are used by the body to generate cellular energy and provide many of the building blocks necessary for cell function.

What is a functional assessment?

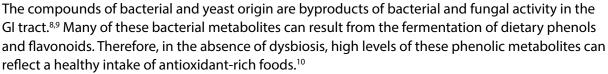
The quantitative measurement of specific organic acids in the urine offers a functional assessment of nutrient status. Enzymes that are responsible for metabolizing organic acids are vitamin and mineral dependent. With this, elevations in organic acids can speak to a functional need for these nutrients on a cellular and biochemical level, even despite normal serum levels.¹⁻⁷ Recommendations for nutrient supplementation based on elevated organic acid results are generated using a literature-based proprietary algorithm.

The Metabolic Analysis Profile report categorizes results into major metabolic areas:

- Malabsorption and Dysbiosis
- Cellular Energy and Mitochondrial Metabolites
- Neurotransmitter Metabolites
- Vitamin Markers
- · Toxin and Detoxification Markers
- Tyrosine Metabolites



Malabsorption and Dysbiosis Markers





Malabsorption and dysbiosis markers are usually evaluated as a group for overall trends rather than individually. When multiple markers are elevated, a stool test may provide further information regarding dysbiosis or other GI dysfunction.

MALABSORPTION MARKERS

Indolacetic Acid (IAA)

Indoleacetic acid (IAA), or indole-3-acetate, is produced by the bacterial fermentation of the amino acid tryptophan.¹¹ IAA can be formed from several common gut microbes such as *Clostridia* species, *Escherichia coli*, and species.¹²⁻¹⁴

High Levels:

Elevated IAA in the urine suggests incomplete digestion and absorption of tryptophan in the intestine, allowing colonic bacteria to convert tryptophan to IAA. Elevations may also reflect an overgrowth of bacteria acting on tryptophan.

Clinical Associations:

IAA elevations and altered tryptophan metabolism have been associated with systemic inflammation, psychologic and cognitive function, autism, and chronic diseases such as cardiovascular disease. Hartnup's disease, a genetically-linked dysfunction in the transport of freeform amino acids across the intestinal mucosa, can cause severe elevations of urinary IAA.

Phenylacetic Acid (PAA)

Phenylacetic acid (PAA) is produced by the bacterial metabolism of phenylalanine. Several bacterial strains are known to produce PAA, including *Bacteroidetes* and *Clostridium* species⁹. Dietary polyphenols may also contribute to PAA elevation.¹⁹

High Levels:

Elevated PAA in the urine suggests incomplete digestion and absorption of phenylalanine in the intestine, allowing colonic bacteria to convert phenylalanine to PAA. Elevations may also reflect an overgrowth of bacteria, which convert phenylalanine to PAA. Several bacterial strains are known to produce PAA, including Bacteroidetes and Clostridium species⁹. Dietary polyphenols may also contribute to PAA elevation.¹⁹

Clinical Associations:

There is a clinical correlation between decreased urinary PAA and depressive symptoms.²⁰⁻²²

BACTERIAL DYSBIOSIS MARKERS

Dihydroxyphenylpropionic Acid (DHPPA)

Dihydroxyphenylpropionic Acid (DHPPA), also known as **3,4 dihydroxyphenylpropionic acid**, is a byproduct of the fermentation of dietary phenols by several bacteria, including some *Clostridia* spp. and others. Although once thought to identify the presence of specific dysbiotic bacteria, ongoing research suggests there are several bacterial species potentially involved.

High Levels:

Elevated DHPPA levels may reflect dietary intake of polyphenols. They may also suggest dysbiosis or bacterial overgrowth, increasing dietary polyphenol conversion.

3-Hydroxyphenylacetic Acid and 4-Hydroxyphenylacetic Acid

3-Hydroxyphenylacetic acid and

4-hydroxyphenylacetic acid are produced by the bacterial fermentation of amino acids, much like IAA. ^{9,12}

High Levels:

Amino acids that are not digested and absorbed can be metabolized by bacteria in the gut to form these organic acids. Clinicians often use these markers to reflect protein malabsorption or dysbiosis. However, dietary intake of polyphenols such as wine, grapes, green tea, and grape seed extract can also contribute to increased levels.²³⁻²⁶

Clinical Associations:

These organic acid byproducts may exhibit free radical scavenging properties, which lends to further support for use of these organic acid markers as an indication of antioxidant consumption.²⁷⁻²⁹

Much like IAA and PAA, there is an inverse correlation between these markers and depressive symptoms.²⁰⁻²²

Benzoic Acid and Hippuric Acid

Benzoic acid and **hippuric acid** are formed from the bacterial metabolism of polyphenols. Urinary benzoic acid may also come from ingestion of food preservatives such as sodium benzoate. Hippuric acid is made when sodium benzoate is conjugated with glycine.³⁰

High Levels:

Increased metabolism by imbalanced gut flora may increase levels. Additionally, dietary intake of polyphenols or food preservatives can also increase levels of these organic acids.

Clinical Associations:

Elevated levels of urinary hippuric acid have been associated with several clinical conditions that may be linked to dysbiosis.^{31,32} For example, elevated urinary hippurate was associated with an increase in blood pressure, likely due to the direct effect of gut-microbial products on blood pressure. However, in other studies low hippuric acid excretion has also been attributed to dysbiosis, which supports its use as a biomarker for general microbial alterations³³

YEAST/FUNGAL DYSBIOSIS MARKERS

Arabinose, Citramalic Acid, and Tartaric Acid

Arabinose is a pentose sugar used by *Candida* to form D-arabinitol. Similarly, **Citramalic** and **Tartaric** acid are yeast metabolites that are also influenced by dietary intake of fruits, wine, and sugars³⁴⁻³⁸

High Levels:

Though often used by clinicians to gain insight into yeast overgrowth, it should be noted that fruit intake can influence levels. High levels may simply reflect a high dietary fruit intake. A high intake of sugars

feeds gastrointestinal yeast, which can promote yeast overgrowth. When these markers are elevated, and dietary influences have been ruled out, a stool test may be warranted to evaluate the presence of yeast in the GI tract.

As noted, the malabsorption and dysbiosis marker levels can also be influenced by common foods, supplements, or preservatives; correlation with the patient's dietary intake is encouraged. ^{25,26,34-37,39-58}

Urinary Metabolite	Common Dietary Sources	
Indoleacetic acid	High tryptophan intake, green/black tea	
Phenylacetic acid	Wine/grapes	
Dihydroxyphenopropionic acid	Whole-grains, chocolate, coffee, green/black tea, olives/olive oil, citrus fruits (animal stud-ies)	
3-Hydroxyphenylacetic acid & 4-hydroxyphenlyacetic acid	Wine/grapes, cranberries, green/black tea, berries, orange juice, grape seed extract	
Benzoic acid/Hippuric acid	Orange juice, elderberry, huckleberry, food preservative, berries, other flavonoids	
Arabinose	Widely distributed, grains, commercial sweetener	
Citramalic acid	Apples, cranberries, sugar beets	
Tartaric acid	Wine/grapes, chocolate, food additive/preservative	

Cellular Energy and Mitochondrial Metabolites

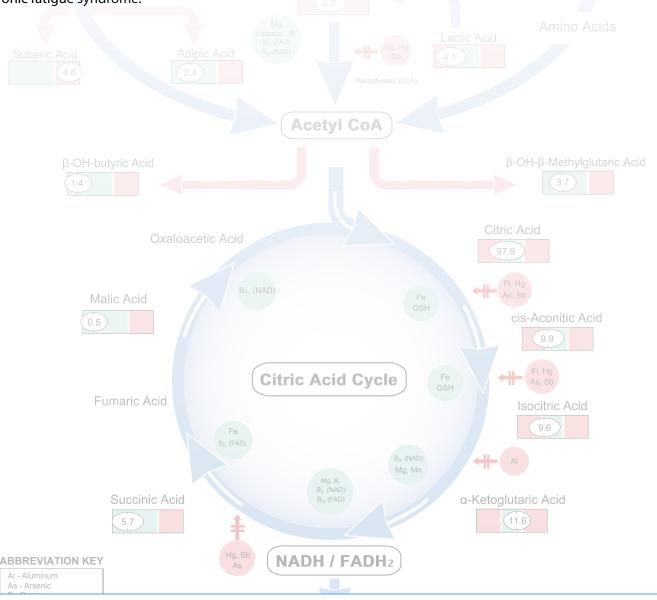
The cellular energy and mitochondrial metabolite markers reflect the body's ability to process dietary macronutrients to feed the **Krebs cycle (Citric Acid Cycle)** and subsequent energy production.

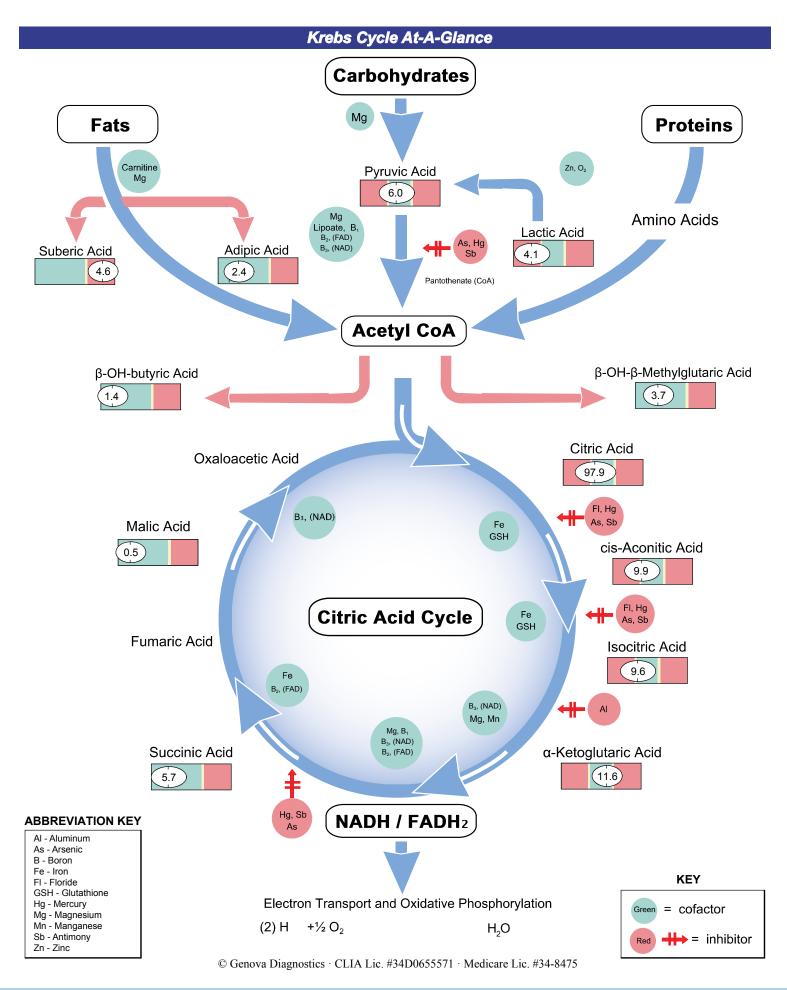
Abnormalities throughout the Krebs cycle, as well as in fatty acid oxidation, glycolysis, and protein metabolism may reflect enzymatic dysfunction and functional nutrient insufficiencies.

Various factors can alter mitochondrial enzymes such as nutrient and vitamin deficiency, toxins, genetic polymorphisms, and underlying disease. The enzymes catalyzing the transformation of these Krebs Cycle intermediates require a variety of nutrient cofactors, such as iron, niacin, magnesium, manganese, thiamin, riboflavin, pantothenic acid, and lipoic acid. ⁵⁹⁻⁶⁹ Toxic exposures and metals including, but not limited to, mercury, arsenic, and lead can interfere with mitochondrial function. ^{59,60,70}

Abnormal urinary excretion of these organic acids may provide a window into various clinical conditions, as well as potential therapeutic targets to correct mitochondrial dysfunction.⁷¹⁻⁷⁴

Mitochondrial dysfunction has been associated with several diseases. The presence of enzymatic antagonists within the Krebs cycle, or lack of specific nutrient cofactors for these enzymes, may contribute to mitochondrial dysfunction, and therefore conditions like neurocognitive disease, diabetes, cancer, mood disorders, cardiovascular disease, and chronic fatigue syndrome. 59,75,76





CARBOHYDRATE METABOLISM

Lactic Acid and Pyruvic Acid

Lactic Acid and Pyruvic Acid are byproducts of glycolysis. Carbohydrates, which contain glucose, are broken down through glycolysis to form pyruvate and two ATP molecules. Pyruvate can also be generated through the catabolism of various amino acids, including alanine, serine, cysteine, glycine, tryptophan and threonine⁷⁷ Magnesium is an important cofactor for a number of glycolytic enzymes necessary to produce pyruvate.⁷⁸ Optimally, pyruvic acid is oxidized to form Acetyl-Co-A to be used aerobically via the Krebs Cycle to produce energy. In an anaerobic state, lactic acid is formed instead.

High Levels:

An elevated pyruvic acid would reflect an inability to form Acetyl-Co-A to feed the Krebs Cycle. Pyruvate uses the pyruvate dehydrogenase complex to form Acetyl-Co-A. A different enzyme, pyruvate carboxylase, is responsible for the conversion of pyruvate into oxaloacetate. Nutrient cofactors, such as vitamin-B₁, B₂, B₃, B₇, magnesium, and lipoate are needed to support the pyruvate dehydrogenase and pyruvate carboxylase enzymes. ^{59,61,79-82} Insufficiency in any of these nutrients can raise levels of pyruvic acid. In vitro studies have shown there are some toxins that can also affect these enzymes, such as antimony, mercury, and cadmium. ^{83,84} Pyruvate elevations can also be seen with a high intake of carbohydrates, as well as rare genetic forms of pyruvate dehydrogenase deficiency. ⁷⁷

Any anerobic or low oxygen state, including pulmonary disease, anemia, sleep apnea, among others can lead to elevations of lactic acid. Elevations of urinary lactic acid can also be the result of strenuous exercise, insulin resistance, dysglycemia, and alcohol dependence. S5-89 Zinc is an essential component in the enzymes which regulate glycolysis, such as lactate dehydrogenase (LDH). LDH converts lactate back to pyruvate in the liver via the Cori cycle. T7,90,91 Elevations may be seen with a functional need for zinc.

Low Levels:

Low levels of pyruvic acid might imply low carbohydrate intake, lack of magnesium cofactors for glycolytic enzymes, or lack of insulin.^{78,92}

Clinical Associations:

Pyruvate metabolism abnormalities play important roles in cancer, heart failure, and neurodegeneration.⁷⁷

FATTY ACID METABOLISM

Adipic and Suberic Acid

Dietary fatty acids are metabolized into fuel sources using beta-oxidation. Fatty acid conversion into Acetyl-CoA requires transport across the mitochondrial membrane via the carnitine shuttle.⁹³ When beta-oxidation is impaired, fats are metabolized using an alternate pathway called omega-oxidation. Omega-oxidation results in elevated levels of dicarboxylic acids such as adipic and suberic acid.

Impaired beta-oxidation occurs in carnitine deficiency or enzymatic dysfunction due to lack of nutrient cofactors. 94,95 Vitamin B₂ and magnesium play a role in optimizing beta-oxidation. 96-101

High Levels:

Elevated levels of adipic and suberic acid may reflect insufficient carnitine or lack of nutrient cofactors for proper beta-oxidation.^{99,101,102}

Clinical Associations:

Increased omega-oxidation metabolites can be seen in ketosis, insulin resistance, diabetes, fasting, or low carbohydrate intake. Elevations of suberic and adipic acid can lead to further mitochondrial dysfunction by injuring the cell membrane and producing free-radical damage. 93,103,104

ENERGY METABOLISM

β-hydroxybutyric Acid

β-hydroxybutyrate is a ketone body. During periods of fasting, exercise, and metabolic disease, ketone bodies are generated in the liver and become an energy source instead of glucose.

High Levels:

Low carbohydrate intake and ketogenic diets may increase urinary levels of beta-hydroxybutyrate. The severity of ketosis is not accurately reflected by the degree of ketonuria. Only a small amount of the body burden of ketones is excreted in the urine; most must be oxidized in extrahepatic tissues using and depleting available oxygen.

Clinical Associations:

In the absence of dietary influence, elevations are sometimes used as an early indicator of diabetes, impaired glucose tolerance, and worsening glycemic control.¹⁰⁵⁻¹⁰⁷

β-hydroxy-β-methylglutaric Acid

β-hydroxy-β-methylglutaric acid (HMG) is a precursor to cholesterol and coenzyme Q10 synthesis. It is a product of hydroxymethylglutaryl-coenzyme A (HMGCoA). HMGCoA-reductase is a rate limiting enzyme in cholesterol production. Medications that interfere with this enzyme may result in elevated HMG and subsequent low levels of cholesterol and CoQ10.¹⁰⁸

High Levels:

Urinary β -hydroxy- β -methylglutaric acid is often elevated in patients taking statin medications and red yeast rice. CoQ10 supplementation has been shown to help ameliorate statin-associated myopathies. ¹⁰⁹

There are also inborn errors of metabolism which can elevate HMG. These affect the HMGCoA reductase enzyme with varying degrees of onset and clinical manifestations such as neurodevelopment disorders and cardiomyopathy.¹¹⁰

KREBS CYCLE (CITRIC ACID CYCLE)

Citric Acid, Isocitric Acid, and cis-Aconitic Acid

A two-carbon group from Acetyl-CoA is transferred to oxaloacetate to form citric acid. Citric acid is then converted to isocitric acid through a cis-aconitic intermediate using the enzyme aconitase. Aconitase is an iron-sulfate protein that controls iron homeostasis.¹¹¹

High Levels:

Iron deficiencies and overload at the systemic or cellular levels can negatively impact the aconitase enzyme and overall mitochondrial health and function. Due diligence with iron assessment is recommended when levels of these organic acids are abnormal. Glutathione may also be an important means of modulating aconitase activity during oxidative stress. Various toxins may influence mitochondrial enzymes and contribute to mitochondrial dysfunction, such as fluoride, aluminum, mercury, arsenic, and tin. 114-117

Low Levels:

Low levels of these analytes may reflect insufficient precursors, or suboptimal glycolysis or fatty acid oxidation.

α-Ketoglutaric Acid

Isocitric Acid is converted to **α-ketoglutaric acid** using the enzyme isocitrate dehydrogenase. Alpha-ketoglutarate is a rate-determining intermediate in the Krebs Cycle and provides an important source of glutamine and glutamate that stimulates protein synthesis and bone tissue formation, inhibits protein degradation in muscle, and constitutes an important metabolic fuel for cells of the gastrointestinal tract.¹¹⁸ Alpha-ketoglutaric acid is then converted to Succinyl CoA using the enzyme alpha-ketoglutarate dehydrogenase. This enzyme complex is very similar to the pyruvate dehydrogenase complex with similar nutrient cofactor needs.

High Levels:

Elevations can be seen with nutrient cofactor deficiencies needed for the enzymatic conversion of α -ketoglutarate such as vitamin B₃, zinc, magnesium, and manganese. Higher levels are seen in mitochondrial oxidative phosphorylation disorders and mitochondrial dysfunction. Genetic abnormalities with the enzyme itself can also limit conversion of alphaketoglutarate, causing elevations.

Low Levels:

Low levels of α -ketoglutarate may reflect lack of precursors higher up from enzymatic dysfunction due to lack of nutritional cofactors, genetic defects, or toxin exposures.

Succinic Acid

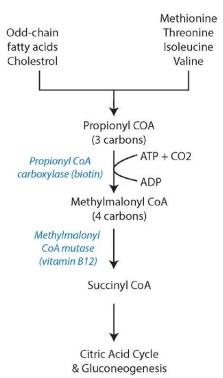
Succinyl CoA becomes **succinic acid** using succinyl CoA synthetase. This reaction produces NADH which directly provides electrons for the electron transport chain or respiratory chain.¹²⁰

Succinic acid requires the enzyme succinate dehydrogenase to become fumarate. This enzyme is iron-based and requires vitamin B₂ to support flavin adenine dinucleotide (FAD) as a redox coenzyme.¹²¹ Succinate dehydrogenase plays a critical role in mitochondrial metabolism. Impairment of this enzyme's activity has been linked to a variety of diseases such as cancer and neurodegenerative diseases.¹²²

High Levels:

Elevated levels of mitochondrial succinate are seen in nutritional cofactor insufficiencies of succinate dehydrogenase or primary enzymatic defects. Succinate can also be formed peripherally by microbes in the GI tract. The major producers of succinate in the gut are bacteria belonging to the *Bacteroidetes* phylum. However, it is typically detected at low rates in the gut lumen because it is rapidly converted to propionate, a major short chain fatty acid.¹²³

Several studies indicate that elevations in both succinate and fumarate play a role in oncogenesis by causing DNA damage and hypermethylation.¹²⁴



Low Levels:

Low levels of succinic acid can be seen with poor dietary intake or absorption of branched-chain amino acids. Branched-chain amino acids are catabolized to acetyl-CoA or succinyl-CoA to feed the Krebs cycle. Additionally, vitamin B_{12} deficiency can induce a defect in the conversion of methylmalonyl-CoA to succinyl-CoA at the distal end of the valine and isoleucine pathways which can then decrease succinyl-CoA. 125

Malic Acid

Fumaric acid uses the fumarase enzyme to become **malic acid**. Malate dehydrogenase catalyzes the conversion of malic acid into **oxaloacetate**. Two forms of this enzyme exist in eukaryotes. One operates within the mitochondria to contribute to the Krebs Cycle; the other is in the cytosol where it participates in the malate/ aspartate shuttle. ¹²⁶ Riboflavin is an important cofactor for this enzyme and overall mitochondrial energy production and cellular function. ¹²⁷

At the end of each Krebs cycle, the four-carbon oxaloacetate has been regenerated, and the cycle continues.

High Levels:

High levels of malic acid can be seen if its dehydrogenation to oxaloacetic acid is reduced from lack of vitamin B₃ as NAD. Malic acid also has many food sources, such as vegetables, as well as fruits like apples and pears. It is also an additive and preservative in beverages, throat lozenges, and syrups.¹²⁸

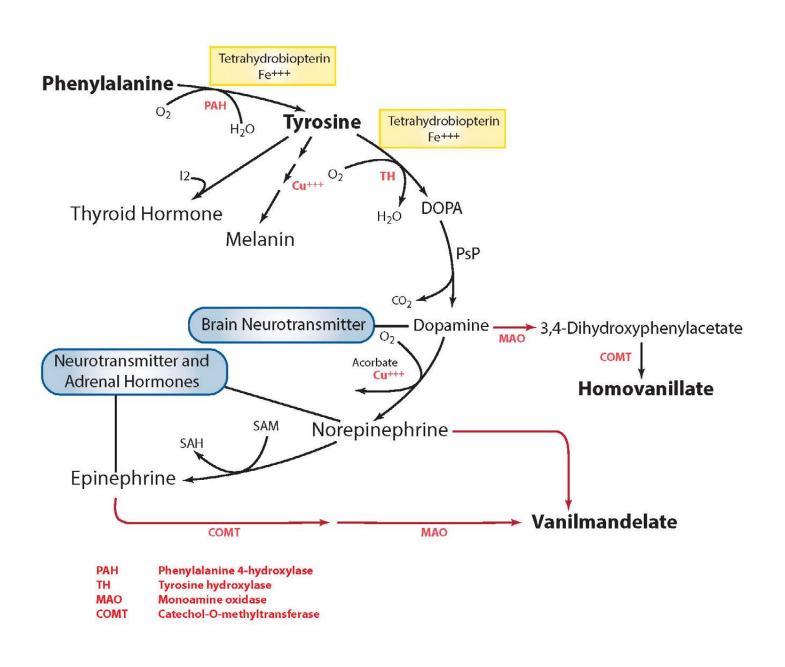
Urinary Creatinine

Urinary creatinine is commonly used as a laboratory standardization when evaluating urinary analytes. ¹²⁹⁻¹³¹ Creatinine excretion is influenced by muscle mass and body habitus since creatinine formation occurs in muscle. Dietary intake of proteins containing arginine and glycine (precursors of creatine) and creatine supplementation can elevate levels. ¹³² Hydration status may also play a role in urinary creatinine levels.

Neurotransmitter Metabolites

These organic acid compounds are down-stream metabolites of neurotransmitter synthesis and degradation. Many of the neurotransmitter metabolites in urine primarily reflect peripheral metabolism, as in the enteric nervous system. Elevations in these organic acids can represent altered neurotransmitter metabolism. This can be due to enzymatic nutrient cofactor needs, or genetic predispositions. Toxins, chronic illness, and stress can also influence results. 133,134

Neurotransmitters are synthesized from essential amino acids using various vitamins and minerals as cofactors. ¹³⁵ Dopamine is converted to norepinephrine using dopamine-beta-hydroxylase. This enzyme requires copper and vitamin C as cofactors. Norepinephrine is then converted to epinephrine through a methylation reaction. The enzymes MAO and COMT metabolize epinephrine and norepinephrine into vanilmandelic acid, which is excreted in the urine. Dopamine can also be metabolized using MAO to form 3,4-dihydroxyphenylacetate which is then metabolized to homovanillic acid using the COMT enzyme.



Vanilmandelic Acid

Vanilmandelic acid (VMA) is formed in the liver by the oxidation of 3-methoxy-4-hydroxyphenylglycol. 136 As a downstream metabolite of tyrosine-derived catecholamines, levels of VMA can reflect the overall synthesis and metabolism of catecholamines. 137 Whether norepinephrine or epinephrine are metabolized into VMA or 3-methoxy-4-OH-phenylglycol (MHPG) depends on the presence and specificity of various available aldehyde reductase and dehydrogenase enzymes. 138

High Levels:

Centrally-acting medications, such as antidepressants and stimulants used for ADHD can elevate overall catecholamines and therefore urinary metabolites. ^{139,140} Urinary levels have been shown to correlate with generalized anxiety disorder. ¹⁴¹ VMA is sometimes used in the work up of pheochromocytoma, neural crest tumors, renovascular hypertension, and neuroblastoma in the right clinical context. ¹⁴²⁻¹⁴⁵ Elevations in catecholamine urinary metabolites have been shown to correlate with the physiologic stress response, exercise, and PTSD. ¹⁴⁶⁻¹⁴⁹

Low Levels:

Low levels of catecholamine metabolites can reflect insufficient amino acid precursors for neurotransmitter production, nutrient cofactor insufficiencies for enzymatic conversion, and genetic abnormalities in enzyme function. Methylation is required for neurotransmitter creation and metabolism. Thus, methylation defects or lack of methylation cofactors may contribute to abnormal levels. Copper is an important cofactor for dopamine beta-hydroxylase, which forms norepinephrine from dopamine. In copper deficiency, norepinephrine formation can be impaired and potentially lower VMA levels.

Manganese released into the synaptic cleft may influence synaptic neurotransmission. Dietary manganese deficiency, which may enhance susceptibility to epileptic functions, appears to affect manganese homeostasis in the brain, probably followed by alteration of neural activity.¹⁵⁰

There are studies which evaluate the neurotoxicity of manganese. Elevated levels of VMA and HVA have been seen in manganese toxicity from occupational exposure which induces a CNS condition similar to Parkinson's disease. 151,152

Homovanillic Acid

Homovanillic acid (HVA), or 3-methoxy-4-hydroxyphenylacetic acid, is a metabolite of dopamine. Although dopamine is an important brain neurotransmitter, a substantial amount of dopamine is produced in the GI tract.¹⁵³

In neurotransmitter production, dopamine is formed from phenylalanine and tyrosine using several enzymes which require nutrient cofactors such as iron, tetrahydrobiopterin, and pyridoxal phosphate.¹⁵⁴ Dopamine then becomes norepinephrine using the enzyme dopamine beta-hydroxylase, which requires copper and ascorbic acid for optimal activity.¹⁵⁵

Dopamine can be metabolized to homovanillic acid using both monoamine oxidase (MAO) and catechol-Omethyltransferase (COMT). 153 MAO requires a vitamin B₂ (FAD) cofactor, while the COMT enzyme requires SAM, magnesium, and vitamin B₆. 156,157

High Levels:

Elevations of homovanillic acid can be seen with lack of vitamin cofactors for enzymes within the metabolism of dopamine or the production of norepinephrine. Quercetin supplementation can elevate plasma HVA and perhaps urinary excretion. Dietary flavanols, such as tomatoes, onions, and tea are also known to elevate urinary HVA.

Like VMA, urinary HVA is elevated in conditions such as neuroblastoma and neural crest tumors. And, since dopamine regulates emotional and motivational behavior, changes in dopamine levels, and subsequent HVA levels, have been studied in the overall stress response, PTSD, mood disorders, and autism. 162-167

Low Levels:

Low levels of urinary HVA imply deficient production of dopamine due to decreased amino acid precursors or lack of vitamin cofactors throughout the production cycle. It may also reflect impaired methylation of dopamine to HVA. Low dopamine turnover and low HVA levels are seen in some mood disorders and as an effect of various antidepressants. 168,169

5-Hydroxyindolacetic Acid (5-HIAA)

5-Hydroxyindolacetic acid (5-HIAA) is a downstream metabolite of serotonin, which is formed from the essential amino acid tryptophan. Most blood serotonin and urinary 5-HIAA comes from serotonin formation outside of the CNS, primarily the liver and enterochromaffin cells in the gastrointestinal tract. Serotonin is further metabolized by monoamine oxidase to become 5-HIAA.¹⁷⁰

High Levels:

Elevations, as well as low levels of urinary 5-HIAA, can reflect underlying intestinal microbial balance.¹⁷¹ Serotonin produced by intestinal enterochromaffin cells is necessary for GI motility.¹⁷² Because of this, antidepressants such as tricyclics and serotonin selective reuptake inhibitors have been used in treating IBS.¹⁷³ Enterochromaffin cells and their serotonin signaling are influenced by overall inflammatory responses to bacteria in the GI tract.

Diets rich in tryptophan and serotonin have been shown to increase urinary 5-HIAA. Bananas, plantains, kiwi, pineapple, nuts, and tomatoes, among other foods, can cause elevations of this urinary metabolite.¹⁷⁰

The excretion of 5-HIAA seems to vary among individuals who supplement with 5-hydroxytryptophan (5HTP).¹⁷⁰

Carcinoid tumors are well-differentiated neuroendocrine tumors derived from the enterochromaffin cells in the GI tract and lung. These tumors secrete vasoactive peptides, especially serotonin which causes flushing and diarrhea. Urinary 5-HIAA levels are elevated in patients with carcinoid syndromes.¹⁷⁴

It should be noted that certain medications may cause false abnormalities in urinary 5-HIAA, and/or interfere with electrochemical detection on chromatography. These include guaifenesin, aspirin, and acetaminophen. Many medications can alter serotonin levels and therefore impact urinary 5-HIAA levels. Due diligence is recommended to investigate medications as a possible etiology of abnormal levels. 170,178,179

Abnormalities, both high and low, in urinary 5-HIAA can be caused by methylation defects, as well as vitamin and mineral nutrient cofactor deficiencies.

Low Levels:

Decreased 5-HIAA levels can reflect low tryptophan intake, or malabsorption/maldigestion of tryptophan. Medications, like MAO inhibitors, decrease serotonin turnover and decrease 5-HIAA.¹⁸⁰ Low levels of urinary 5-HIAA have been observed in cardiovascular disease, metabolic syndrome, IBS patients, and those with mood disorders and migraines.¹⁸¹⁻¹⁸³

3-Methyl-4-Hydroxy-Phenylglycol (MHPG)

3-Methyl-4-OH-Phenylglycol (MHPG) is a byproduct of the central nervous system's norepinephrine metabolism. MHPG metabolizes to vanilmandelic acid (VMA) in the liver using the enzymes alcohol dehydrogenase and aldehyde dehydrogenase. Urinary MHPG was originally thought to represent CNS sympathetic output, but is now known to be principally derived from peripheral neuronal NE metabolism.¹⁸⁴

MHPG has been widely studied as a marker to predict response to medications used in mood disorders or as a biomarker to monitor pharmacotherapies.¹⁸⁵⁻¹⁸⁸

High Levels:

The role of hepatic alcohol and aldehyde dehydrogenase explains the clinical observations that ethanol consumption decreases the excretion of VMA, while increasing MHPG.^{189,190}

Because norepinephrine is involved in the pathophysiology of hot flashes in postmenopausal women, MHPG levels have been studied in this patient population. ^{191,192} Interestingly, folic acid was found to interact with receptors causing subjective improvement in symptoms. ¹⁹³

Sleep deprivation can act as a stimulus to the peripheral sympathetic nervous system, which can influence central nervous noradrenergic neurotransmitter levels and elevate MHPG.¹⁹⁴ As a central nervous system metabolite, levels can correlate with central catecholaminergic disturbances, as in anxiety and seizures.^{195,196} Elevated MHPG levels have also been associated with the stress response.¹⁹⁷

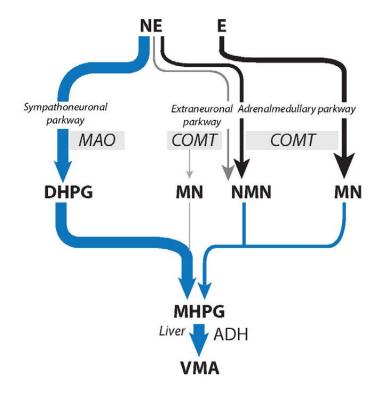
Pheochromocytomas are rare, mostly benign tumors of the adrenal medulla which can secrete catecholamines causing a wide array of sympathetic symptoms.

These tumors contain MAO and COMT. They can therefore produce MHPG. However, because peripheral sympathetic nerves can also contribute to high MHPG, using MHPG for diagnosis of pheochromocytoma is limited. VMA is also not very sensitive for diagnosis of pheochromocytoma because it can be made in the liver from MHPG. Although neither organic acid is diagnostic of pheochromocytoma, it is possible to see elevations of these analytes in the disease. 198

Low Levels:

Since catecholamines are made from dopamine, low levels of the MHPG metabolite can result from low levels of dopamine, dopamine amino acid precursors, nutrient enzymatic cofactor deficiencies in dopamine metabolism, and overall methylation defects.

Low levels of MHPG have been correlated to mood and behavioral disorders, anorexia, and ADHD. 199-201



*DHPG: 3,4-dihydroxyphenylglycol

*MN: metanephrine
*NMN: normetanephrine

*MHPG: 3-methyl-4-hydroxy-phenylglycol

*VMA: vanilmandelic acid *ADH: alcohol dehydrogenase *MAO: monoaminoxidase

*COMT: catechol-O-methyltransferase

Kynurenic Acid and Quinolinic Acid

Kynurenic acid and Quinolinic acid are tryptophan metabolites formed through the kynurenine pathway. Tryptophan is the amino acid precursor to serotonin; its major route for catabolism is the kynurenine pathway. Important products of the kynurenine pathway include xanthurenic acid and kynurenic acid, which can further metabolize into **quinolinic acid.**

The historical importance of this pathway has mainly been as a source of the coenzyme NAD+, which is important for all redox reactions in the mitochondria. However, it is now understood that kynurenic and quinolinic acid have physiologic implications. This alternate pathway is upregulated in response to inflammation and stress, which can lead to deficient serotonin production.²⁰²

Kynurenic acid has shown some neuroprotective properties in the brain, since it can stimulate NMDA receptors. However, its importance on the periphery is still not fully elucidated. Some studies outline anti-inflammatory, analgesic, antiatherogenic, antioxidative, and hepatoprotective properties to peripheral kynurenic acid.²⁰³⁻²⁰⁵ The correlation to levels of urinary excretion needs further study.

Quinolinic acid, in and of itself, can be inflammatory and neurotoxic.

High Levels:

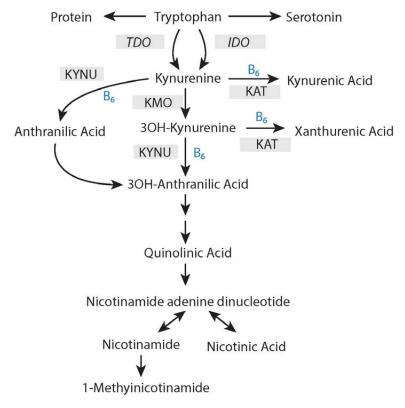
The kynurenine pathway is particularly sensitive to vitamin B₆ deficiency, which can elevate urinary kynurenic acid (and xanthurenic acid).²⁰⁶⁻²⁰⁸ Vitamin B₂ is also an important vitamin cofactor in the enzymatic conversion reactions within the pathway.²⁰⁹ Because a major-end product of this pathway is also NAD+, elevations in kynurenic and quinolinic acid may also reflect vitamin B₃ need.

Oral contraceptives and estrogen therapy have been implicated in increasing quinolinic acid excretion both from altered tryptophan metabolism directly, as well as vitamin B₆ insufficiency.²¹⁰

Many of the intermediates and products in the kynurenine pathway are implicated in numerous neurological and psychiatric diseases, such as depression. Alterations in this pathway also have some connection to the development of insulin resistance, diabetes, tumor growth and proliferation, and inflammatory myopathies.²¹¹⁻²¹⁵

Kynurenic/Quinolinic Acid Ratio

Because of the specific inflammatory component of quinolinic acid, as well as the potentially protective role of kynurenic acid peripherally (as outlined above), laboratories measure the ratio of kynurenic acid to quinolinic acid. This ratio can act as a measure of disturbed kynurenine pathway metabolism. It suggests that tryptophan is catabolized via the kynurenine pathway, rather than the serotonin pathway. There is literature regarding a low kynurenic/quinolinic ratio association with neurotoxicity and major depressive disorder.^{216,217}



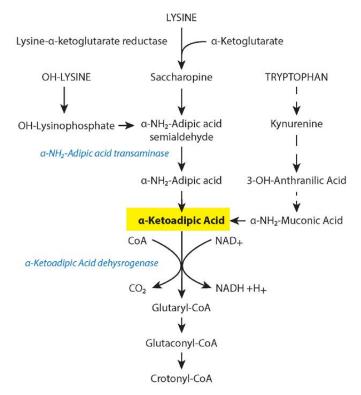
Vitamin Markers



There are groups of organic acids commonly used to assess the status of specific B-vitamins. By measuring organic acids that are known to rely on specific nutrients for enzymatic metabolism, clinicians can gain insight into functional vitamin needs.

α-Ketoadipic Acid (AKAA; 2-Oxoadipic Acid; 2-Ketoadipic Acid)

α-Ketoadipic Acid (AKAA; 2-Oxoadipic acid, 2-Ketoadipic acid) is an organic acid formed from α -aminoadipic acid (which originates with lysine) and also from α -aminomuconic acid (derived from tryptophan).²¹⁸ AKAA metabolizes to form glutaryl-CoA via oxidative decarboxylation. The cofactors needed in this step are Coenzyme A, NAD, thiamine pyrophosphate (vitamin B₁), lipoic acid, and vitamin B₂.¹⁸⁹

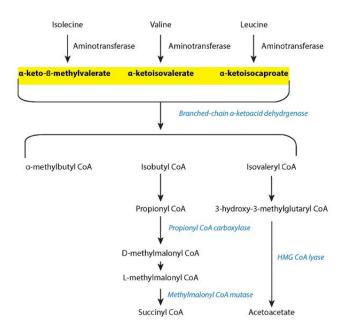


High Levels:

Elevations in urinary AKAA may reflect enzymatic dysfunction due to nutritional cofactor needs.⁶⁹ Mitochondrial oxidative phosphorylation disorders are also associated with higher levels of AKAA.¹¹⁹

α -Ketoisovaleric Acid, α -Ketoisocaproic Acid α -Keto- β -Methylvaleric Acid

Of the essential amino acids, there are three branched-chain amino acids (leucine, isoleucine, and valine). Unlike most amino acids, the initial step of branched-chain amino acid (BCAA) metabolism does not take place in the liver. They increase rapidly in systemic circulation after protein intake and are readily available for use. Skeletal muscle is where most of the initial catabolism of BCAA takes place using branched-chain aminotransferase enzymes to form α -ketoacids, which are then released from muscles back into the blood to be further metabolized, mainly in the liver. BCAA act as substrates for protein synthesis, energy production, neurotransmitter production, glucose metabolism, immune response, and many other beneficial metabolic processes. 219



- α-Ketoisovaleric Acid (AKIV) is produced from the essential amino acid valine. It then metabolizes to become succinyl Co-A. AKIV is glucogenic.
- α-Ketoisocaproic Acid (AKIC) is produced from leucine and further metabolizes to form acetyl-CoA and acetoacetate. AKIC is ketogenic.
- α-Keto-β-Methylvaleric Acid (AKBM) comes from isoleucine, and further metabolizes to form acetyl-CoA and succinyl-CoA. AKBM is therefore both glycogenic and ketogenic.

These α -ketoacids then require an enzyme complex called branched-chain α -keto acid dehydrogenase (BCKD) for further metabolism. ²²⁰ This enzyme complex requires multiple vitamin cofactors, such as vitamin B₁, B₂, B₃, B₅, and lipoic acid. ^{69,221-223}

High Levels:

Urinary elevations of these ketoacids can be the result of functional need for the vitamin cofactors to support BCKD.

A genetic defect of the α -keto acid dehydrogenase enzyme complex is responsible for maple syrup urine disease, which results in very elevated levels of AKIC, AKIV, AKMB.²⁰⁸

Elevated plasma levels of branched-chain amino acids have been associated with insulin resistance as a result of decreased catabolism for energy production. This metabolic disturbance may be compounded by further nutrient deficiencies limiting the activity of the BCKD enzyme^{212,213}

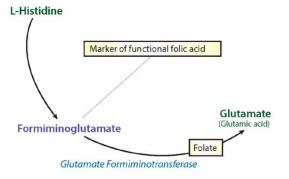
Formiminoglutamic Acid (FIGlu)

Formiminoglutamic Acid (FIGIu) is an intermediary organic acid in the conversion of the amino acid histadine to glutamic acid. This enzymatic conversion requires tetrahydrofolic acid.²²⁶

High Levels:

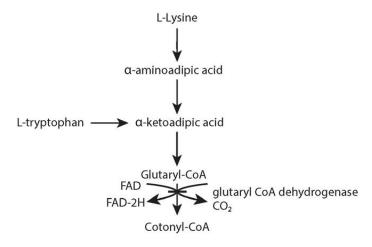
FIGIu elevations in urine have been used as a marker for folate deficiency dating back to the 1950's 2,227 In addition to folate deficiency, elevated urinary FIGIu may also reflect vitamin B_{12} status since folate recycling requires vitamin B_{12} as a cofactor and both are critical steps in the methylation cycle. 228

There are multiple clinical associations with elevated urinary FIGlu, including acute and chronic alcohol use, pregnancy, and oral contraceptive use.²²⁹⁻²³²



Glutaric Acid

Glutaric Acid is formed from the essential amino acids lysine and tryptophan through the intermediaries of alpha ketoadipic acid and glutaryl-CoA. Glutaryl-CoA is further metabolized to glutaconyl- and crotonyl-CoA by an enzyme called glutaryl-CoA dehydrogenase. This enzyme requires riboflavin (vitamin B₂) as a cofactor.



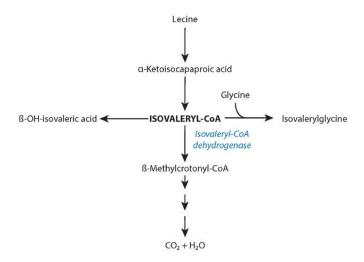
High Levels:

Elevations of urinary glutaric acid may reflect enzymatic insufficiency requiring vitamin B₂ or mitochondrial electron transport dysfunction.

Deficiencies of the enzyme glutaryl-CoA dehydrogenase, and multiple acyl-CoA dehydrogenase deficiency (MADD), are well-studied inborn errors of metabolism which result in significant glutaric aciduria. However, milder forms of this rare mitochondrial disorder exist and can result in adult-onset presentations. Late-onset forms can present as atypical beta-oxidation disorders with exercise intolerance, muscle weakness, and CNS dysfunction.^{221,222} In these cases, riboflavin, carnitine, and CoQ10 have been used therapeutically.²²²⁻²²⁴

Isovalerylglycine

Isovalerylglycine is produced from leucine catabolism. It is further metabolized via isovaleryl-CoA dehydrogenase. This enzyme requires vitamin B2 as a cofactor.^{237,238}



High Levels:

Acyl-CoA dehydrogenase enzymes are not only involved in branched-chain amino acid metabolism, but also beta-oxidation of fatty acids. ¹⁰² Enzymatic dysfunction and elevations in isovalerylglycine are seen when there is a functional nutrient cofactor need and in certain inborn errors of metabolism. However, elevations of isovalerylglycine are also seen in problematic mitochondrial fatty acid beta-oxidation. ^{239,240}

Carnitine, glycine, vitamin B₂, and antioxidants have been used therapeutically to treat abnormal levels of isovalerylglycine.²⁴¹⁻²⁴³

There is an association between elevated isovalerylglycine and anorexia nervosa. The mechanism is believed to be due to poor thyroid conversion of vitamin B₂ into active FAD, which normalized in some patients after a refeeding program.²⁴⁴

Methylmalonic Acid (MMA)

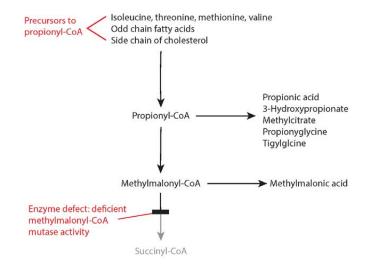
Methylmalonic Acid (MMA) is formed from propionyl-CoA via methylmalonyl-CoA. Major dietary sources of propionyl-CoA include valine, isoleucine, methionine, threonine, and odd chain fatty acids.²⁴⁵ Methylmalonyl-CoA is converted to succinyl-CoA to feed the Krebs cycle via the enzyme methylmalonyl-CoA mutase. This enzyme is very vitamin B₁₂ dependent. In B₁₂ deficiency, methylmalonyl-CoA is hydrolyzed to methylmalonic acid.²⁴⁶

High Levels:

The most common cause of MMA in the urine is vitamin B_{12} deficiency. However, a rare deficiency of the methylmalonyl-CoA mutase enzyme is another. Any underlying condition which results in vitamin B_{12} deficiency should be considered, such as reduced intestinal absorption, chronic alcoholism, or strict vegan diets.²⁴⁶

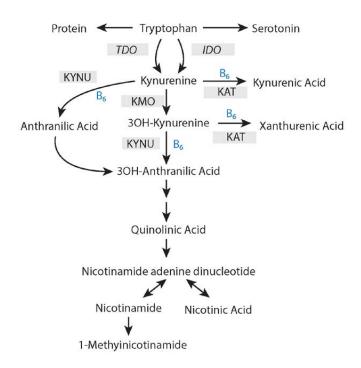
Methylmalonic acid, as a functional biomarker, is considered a more sensitive index of B_{12} status when compared to serum $B_{12}^{-247-252}$ Urinary MMA correlates with serum MMA, making the simple urine test a useful screening tool for B_{12} deficiency in at-risk populations, such as the elderly or patients with GI dysfunction. 249,253

Vitamin B_{12} therapy lowers MMA. Monitoring this metabolite may help prevent the consequences of B_{12} deficiency, such as cognitive decline and neuropathy.^{251,254,255}



Xanthurenic Acid

Xanthurenic acid is produced as part of the kynurenine pathway of tryptophan catabolism, along with kynurenic and quinolinic acid, as previously outlined.



High Levels:

Because this pathway is heavily dependent on vitamin B6, elevations of xanthurenic acid can reflect a functional need for vitamin B6.256 Kynurenine pathway metabolites may also become elevated when there are needs for vitamin B3.257,258

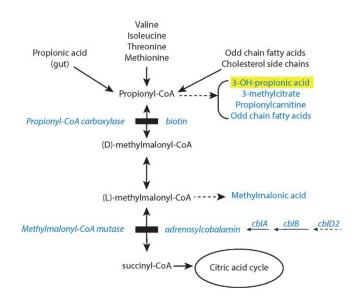
Elevations in urinary xanthurenic acid are seen with increased intake of tryptophan, and in high estrogen states. Pregnancy and oral contraceptive use is associated with elevated levels of urinary xanthurenic acid where a functional nutrient need for B-vitamins is pronounced.^{5,259}

Abnormalities in the kynurenine pathway have been associated with many clinical conditions including immune suppression, cancer, and inflammatory conditions.²⁵⁶

Administration of vitamin B₆ can decrease xanthurenic acid excretion. ^{260,261}

3-Hydroxypropionic Acid (3-HPA)

3-Hydroxypropionic Acid (3-HPA) is a major urinary metabolite of propionic acid. Propionic acid is derived from dietary branched-chain amino acids, odd-chain fatty acids, and can be produced in the gut by bacterial fermentation of fiber. The biotin-dependent enzyme propionyl CoA carboxylase is responsible for metabolizing propionic acid to methylmalonyl CoA, which is subsequently isomerized to succinyl CoA. Decreased activity of this enzyme shunts propionyl CoA into alternative pathways which form 3-HPA.



High Levels:

As noted, biotin is a cofactor in the propionyl-CoAcarboxylase enzyme. ²⁶² Reduced activity of this enzyme due to functional biotin deficiency can cause elevations of the urinary organic acid 3-hydroxypropionic acid. However, in isolation, it may not be as sensitive a marker as 3-hydroxyisovaleric acid to diagnose marginal biotin deficiency. ²⁶³

There are inborn errors of metabolism associated with this organic acid. When the propionyl-CoA-carboxylase enzyme is deficient, the result is propionic acidemia and elevated urinary 3-hydroxypropionic acid. Some isolated case reports reveal the possibility of a later onset in this enzyme deficiency.²⁶⁴

Because of the relationship between propionyl-CoA and methylmalonyl CoA, 3-HPA elevations have also been observed in inborn errors causing methylmalonic acidemia.²⁶⁵

Low Levels:

Low levels of urinary 3-hydroxypropionic acid may be seen with decreased amino acid and fatty acid precursors from maldigestion, malabsorption or impaired fatty acid oxidation. Because the propionic acid precursor is also made in the GI tract, decreased fiber intake or antibiotic use can result in lower urinary 3-hydroxypropionic acid as well.²⁶⁶ In fact, low protein diets and antibiotics are used acutely to treat inborn errors of metabolism which cause propionic acidemia.²⁶⁷

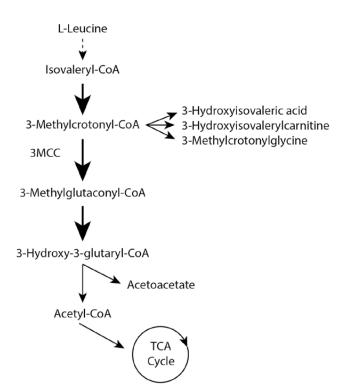
3-Hydroxyisovaleric Acid (3-HIA)

3-Hydroxyisovaleric Acid (3-HIA) is formed from the metabolism of the branched-chain amino acid leucine. Methylcrotonyl-CoA carboxylase catalyzes an essential step in this pathway and is biotin dependent. Reduced activity of this enzyme leads to an alternate pathway of metabolism resulting in 3-hydroxyisovaleric acid.

High Levels:

The urinary excretion of 3-HIA has been shown to be an early and sensitive indicator for marginal biotin deficiency.²⁶³

Elevated levels of 3-HIA in pregnant women reflect reduced or marginal biotin status.²⁶⁸ Smoking and anticonvulsant medication can also increase this metabolite as a reflection of accelerated biotin metabolism and therefore marginal deficiency.^{269,270}



Toxin and Detoxification Markers

These urinary markers can reflect exposure to environmental toxins, or up-regulation of detoxification pathways in response to exposures. When these markers are elevated, the recommendation is to identify, minimize, and remove exposures. Clinicians may consider the use of antioxidants and nutritional support of detoxification pathways. For further information on environmental toxins, the following websites may be helpful:

- ToxTown
- Environmental Working Group
- Agency for Toxic Substances and Disease Registry

α-Ketophenylacetic acid

α-Ketophenylacetic Acid, also known as phenylglyoxylic acid (PGA), is a urinary metabolite of styrene, toluene, xylenes, and ethylbenzene. It acts as a urinary marker of recent exposure via inhalation, contact, oral, and others.²⁷¹ The biologic half-life of styrene in humans is fairly short and corresponds with the disappearance of PGA from the urine.^{272,273}

Styrene is widely used for synthesis of polymers such as plastics, rubbers, and surface coating. It is also used in the pharmaceutical industry. Styrene is commonly applied in the manufacturing of paints, pigments, and glues. Coexposure to other solvents, like toluene and ethyl acetate is common in workplaces where styrene is a concern.²⁷⁴ Since toluene and xylene are components of unleaded gasoline, workers at gas stations are at potential risk of exposure, as well as the general population.²⁷⁵

Styrene exposure may interfere with peripheral metabolism of thyroid hormones by inhibiting conversion of T_4 to T_3 .²⁷⁶ It may also affect DNA repair capacity and damage.²⁷⁷ There are also clinical associations with insulin resistance, oxidative stress, and inflammation.²⁷⁸

α-Hydroxyisobutyric Acid

α-Hydroxyisobutyric Acid is a major urinary metabolite of the industrial solvent methyl tert-butyl ether (MTBE). MTBE was a gasoline additive discontinued in the early 2000's used to reduce automobile emissions. Due to significant ground water leakage from storage tanks, ongoing exposure to MTBE exists in ground water. There is also data available on levels of MTBE in ambient air. ²⁷⁹ Urinary α-hydroxyisobutryic acid is a marker of recent MTBE exposure. ^{280,281}

Although, MTBE was initially designated as "non-carcinogenic", recent studies suggest some interesting clinical associations. Exposure to MTBE has been linked to type 2 diabetes as a result of disrupted zinc homeostasis and glucose tolerance. There are also clinical associations with autism, DNA oxidative damage, and methylation defects. Studies on cancer, reproductive abnormalities, nonalcoholic fatty liver, and neurotoxicity have been either negative or inconclusive thus far.

Orotic Acid

Orotic Acid is an organic acid which serves as an intermediate in nucleotide synthesis and is linked to arginine metabolism as a urea cycle marker for nitrogen balance.²⁹⁰

It is formed from aspartic acid and carbamoyl phosphate.²⁹¹ Carbamoyl phosphate plays an important role in the body because it brings nitrogen into the urea cycle for detoxification and disposal. Carbamoyl phosphate enters the urea cycle to react with ornithine to form citrulline. When ammonia levels significantly increase or the liver's capacity for detoxifying ammonia into urea decreases, carbamoyl phosphate leaves the mitochondria and instead enters the pyrimidine pathway. This stimulates orotic acid biosynthesis and subsequent urinary excretion.²⁹²

Orotic acid can also be found in the diet. The richest dietary sources include cow's milk and dairy products. Most urinary orotic acid is synthesized in the body as an intermediate in the nucleotide synthesis. Although it is also linked with abnormalities in arginine metabolism as a urea cycle marker for nitrogen balance, orotic acid plays no direct role in the urea cycle, yet is increased in urea cycle disorders. Hyperammonemia is characteristic of all urea cycle disorders; orotic acid is only elevated in a few.

High Levels:

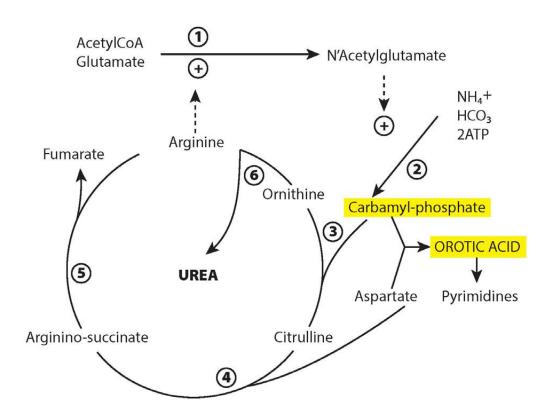
Elevations of orotic acid are seen in with hereditary deficiencies of urea cycle enzymes, ammonia overload as seen in high protein diets, and abnormalities in arginine metabolism.^{290,292}

Any hepatotoxin or underlying liver condition can affect ammonia metabolism and increase orotic acid. There are studies that show elevations in orotic acid after drinking alcohol, which then declined with abstinence.²⁹⁴ Orotic acid excretion is increased by allopurinol and 6-azauridine seemingly related to the action of these drugs on pyrimidine synthesis.²⁹⁵

There are animal studies which show a link between orotic aciduria and hypertension. Orotic acid can induce endothelial dysfunction by contributing to vascular and systemic insulin resistance which impacts nitric oxide production, leading to hypertension.²⁹⁶ Random case studies also show an association between megaloblastic anemia and orotic aciduria as a result of hereditary defects in pyrimidine synthesis.²⁹⁷

Low Levels:

There is no clinical significance to low levels of urinary orotic acid.



Pyroglutamic Acid (5-oxoproline)

Pyroglutamic acid (5-oxoproline) is produced and utilized in the gamma-glutamyl cycle. This cycle is needed to assist in the production and recycling of glutathione (GSH), a powerful antioxidant.

Glutathione is a tripeptide, consisting of glutamate, cysteine, and glycine. Using the gamma-glutamyl cycle, GSH is divided into cysteinyl glycine and a gamma-glutamyl molecule which attaches to another amino acid for transport across a membrane or into a cell. Gamma-glutamyl transferase then splits off that attached amino acid, and the glutamate becomes pyroglutamic acid (5-oxoproline). Cysteinyl glycine is also broken down and transported into the cell as cysteine and glycine. The entire GSH molecule needs to be reformed intracellularly from pyroglutamic acid by recombining cysteine, glycine, and glutamic acid using GSH synthetase.^{298,299} This enzymatic reformation requires cofactors such as ATP and magnesium.³⁰⁰

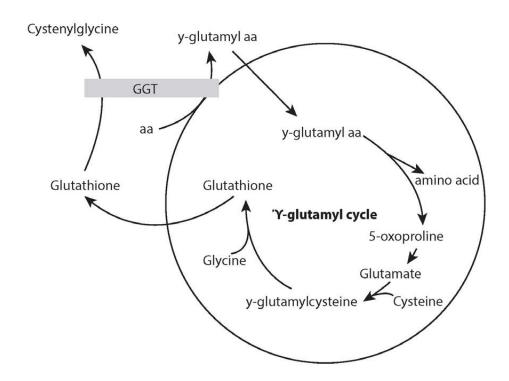
High Levels:

Elevations in pyroglutamic acid can reflect lack of precursors (glycine, cysteine, glutamine) or nutrient cofactors for GSH recycling (magnesium). Most specifically, pyroglutamic acid has been proposed as a measure of glycine availability. Oxidative stress, in general, can upregulate the detoxification pathways and result in elevated pyroglutamic aciduria. Significant toxic exposures, such as medication toxicities, can deplete ATP, interrupting GSH recycling and causing elevations in pyroglutamic acid. In rare cases, this can result in metabolic acidosis.

Deficiency in glutathione synthetase has also been described in literature as presenting with pyroglutamic aciduria.³⁰⁸

Low Levels:

Because pyroglutamic acid formation is dependent on glutathione entering the gamma-glutamyl cycle, an insufficient amount of GSH or its precursors and necessary cofactors can result in low pyroglutamic acid.



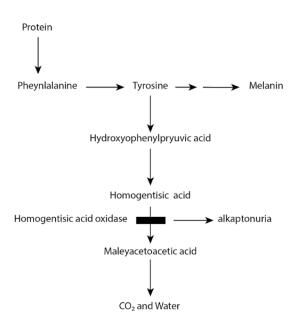
Tyrosine Metabolites

Tyrosine is a non-essential amino acid which can be produced in the body from phenylalanine when there is insufficient dietary intake. Tyrosine is an important neurotransmitter precursor and needed to produce thyroid hormone and melanin. The liver catabolizes any tyrosine that exceeds the amount needed in these metabolic processes.

No.

Homogentisic Acid (HGA)

Homogentisic Acid (HGA) is a downstream tyrosine metabolite also known as melanic acid. Tyrosine is converted to 4-hydroxyphenlylpyruvic acid (4-HPPA) which then becomes homogentisic acid. HGA requires homogentisate dioxygenase (homogentisic acid oxidase) for further breakdown. This is an iron dependent enzyme.

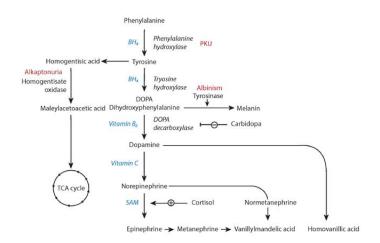


High Levels:

Elevations of HGA are most associated with the inborn error of metabolism, alkaptonuria. Alkaptonuria is a rare, recessive disorder lacking homogentisate dioxygenase (homogentisic acid oxidase). Patients accumulate HGA over time, and by approximately age 30, begin seeing ochronosis - pigmentation of connective tissue, arthritis, valvular heart disease, and kidney stones. ³⁰⁹⁻³¹¹ Urine darkens when exposed to air. Low protein diets and vitamin C are often used therapeutically, though results are mixed. ³¹²⁻³¹⁴

2-Hydroxyphenylacetic Acid (2-HPAA)

2-Hydroxyphenylacetic Acid (2-HPAA) is a metabolite of phenylalanine. Phenylalanine is converted to tyrosine using phenylalanine hydroxylase, which is a tetrahydrobiopterin (BH4)-dependent reaction. When phenylalanine hydroxylase is deficient, phenylalanine accumulates in circulation and is undergoes alternate conversion pathways to form the ketone phenylpyruvate, phenylacetylglutamine, or phenylacetic acid.



High Levels:

Phenylketonuria (PKU) is an autosomal recessive genetic mutation of the phenylalanine hydroxylase enzyme. Phenylpyruvate ketones accumulate causing acidosis, and phenylacetic acid accounts for the odor in PKU. Dietary restriction of phenylalanine and BH4 supplementation are the mainstays of treatment in PKU.³¹⁵

Nutritionally, elevated 2-HPAA can imply excessive dietary intake of phenylalanine, and possible BH4 insufficiency. Finally, 2-HPAA is a small phenolic molecule which may also be an organic acid formed by the gut microbiome, similar to 3,- and 4-HPAA (outlined previously). 316,317 Some use this marker as a dysbiosis marker to give insight into gut health.

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