

ERNDiM



**Amino Acids Interpretation Scheme -
What can we learn, how can we improve?**

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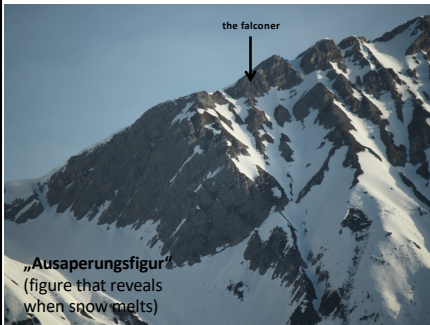
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Outline

1. Key points of the amino acid interpretation scheme
2. International classification of inherited metabolic disorders
3. Cases of 2024 scheme
4. Take to work message

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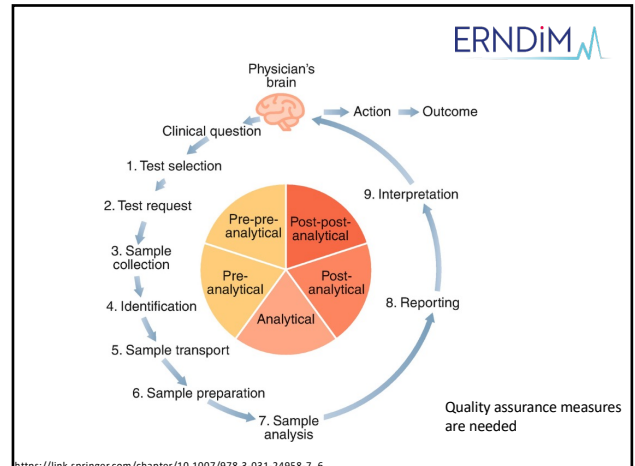


the falconer

**Key points of the
amino acid
interpretation
scheme**

„Ausaperungsfigur“
(figure that reveals
when snow melts)

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Various types of EQA are offered

- Quantitative EQA schemes (QA)
- Qualitative EQA schemes (QL)
- Hybrid proficiency tests (HP)

Quality assurance measures aim at

★ ★ ★ ★ ★

O Ordering

P Preanalytical conditions

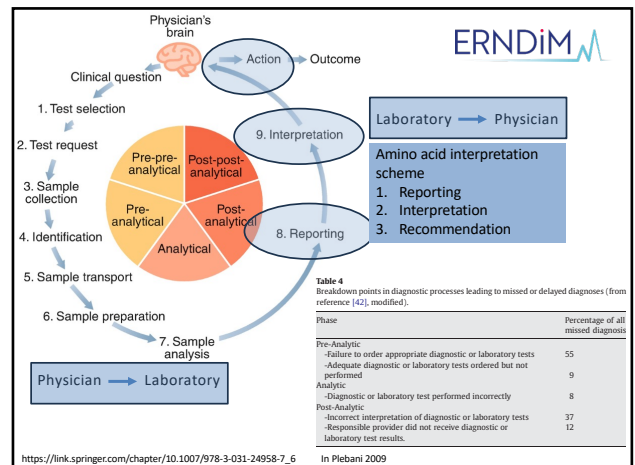
A Analysis

I Interpretation of results

An interpretative EQA scheme, which does not require physical samples, represents a new approach.

EQA = external quality assessment

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Interface 1: Optimal situation

Physician → Laboratory

- What symptoms does the patient have?
- Acute or chronic?
- Which ward is the patient in?
- Is there a clinical suspicion of a diagnosis?
-

Physician Laboratory
staffNo question –
no answer!

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- Different languages
- Symptoms vs. pathways

The diagnosis of inborn disorders of metabolism is complex.

<https://www.alamy.com>

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Interface 2: Optimal situation

Laboratory → Physician

- KISS (keep it short and simple)
- Sometimes an explanation is needed
-

Physician Laboratory
staff

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Amino acid interpretation scheme

1. Reporting

2. Interpretation

3. Recommendation

Case 2025-4	
clinical information	noticeably tired and pale for several weeks, hypochromic
age at time of sample collection	1.5 years
sex	female
pregnancy	normal
birth	normal
family history	
initial symptoms	
treatment	none
miscellaneous	

amino acids in plasma, done with ion exchange chromatography with ninhydrin detection			
	μmol/L		reference range
Taurine	27	19	- 130
Threonine	186	33	- 130
Serine	273	24	- 178
Asparagine	142	25	- 150
Glutamic Acid	23	5	- 80
Glutamine	666	200	- 600
Proline	517	51	- 185
Glycine	286	56	- 308
Alanine	698	99	- 350
Citrulline	28	5	- 24
Valine	92	57	- 270
Cystine	22	0	- 40
Methionine	611	3	- 29
Cystathionin/Alloisoleucin	3	0	- 0
Isoleucine	31	26	- 94
Leucine	36	45	- 155
Tyrosine	211	11	- 120
Phenylalanine	125	23	- 70
Homocystine	n.d.	0	- 0
Ornithine	57	10	- 107
Lysine	207	45	- 144
Histidine	69	24	- 112
Arginine	56	11	- 75
n.d. not detected			

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Amino acid interpretation scheme

1. Reporting

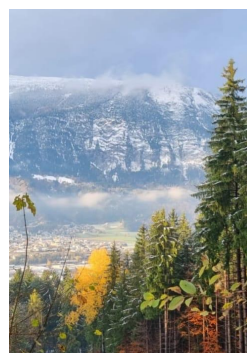
2. Interpretation

3. Recommendation

An example for the answer (anonymously for scorers, no ERN number):

Part. No.	Case 4 abnormalities	Case 4 diagnosis	Case 4 further testing recommendations
	Each 5 points: elevated Tyr and Met, maximum 2 points	Tyrosinemia type 1 (2 points), liver failure (due to other TMS like galactosemia, mitochondrial FA, etc) 2 points, tyrosinemia type 1 (0 points), maximum 2 points	each 1 point (maximum 2 points): alpha-ketoglutarate, organic acids (urine, inclusive succinylacetone), 5-aminolevulinic acid porphobilinogen synthase activity, molecular genetic analysis of FAH gene
x	High levels of Tyrosine, Methionine and slightly increased levels of Phenylalanine High levels of Alanine, Proline and Glycine (Lactic Acidosis) and slight increase on Glutamine, Threonine and Serine. Leucine levels are low or borderline (Ile and Val) related with liver disease.	Tyrosinemia Type 1 (High levels of Tyr, Met, Phenyl). Tyrosinemia is a progression between altered urea cycle and congenital liver failure. Lactic acidosis (liver failure and renal) could be due to an hepatic failure and Glutamine by the secondary urea cycle failure. Other causes: mitochondrial diseases or liver failure	urine organic acids with succinylacetone, MS succinylacetone, urine alpha-ketoglutarate, transaminases, coagulation test, urine ammonia, Ammonia FAH gene or exome sequencing searching for another disease like mitochondrial diseases

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Amino Acid Interpretation scheme

- since 2023 full scheme
- 6 (2x3) cases per year
- 6 (3x2) points per case (findings, interpretation, recommendation)
- 56% cut off for poor performance
- concept of critical error

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The diagram features the ERNDiM logo at the top right, consisting of the text 'ERNDiM' in blue and a stylized 'M' with a blue and red gradient. Below the logo is a horizontal bar with a rainbow gradient. The main title, 'Amino acid interpretation scheme', is centered in a large, bold, black font. Below the title is a horizontal flowchart with five chevron-shaped boxes connected by arrows, representing the process steps: 'application', 'case details', 'submission', 'scoring', and 'reports'. Each box is light blue with a darker blue border. Below each box is a corresponding section header in bold, followed by a list of details or criteria.

```
graph LR; A[application] --> B[case details]; B --> C[submission]; C --> D[scoring]; D --> E[reports]
```

application

- there is a limit to the number of participants

case details

- a short clinical description and
- amino acid concentrations inclusive reference ranges
- there are two rounds of 3 cases per year

submission

- description of results
- diagnosis
- recommendation for further investigations
- each part scores a total of two points

scoring by two independent and blinded experts

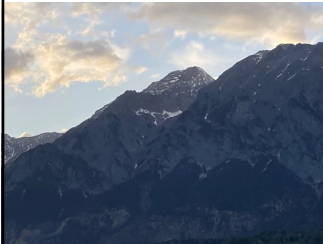
- the scoring criteria and the criteria for critical error are predetermined
- in addition to sufficient participation, there are poor performers, partial submitters and non submitters


reports

- circulation reports
- meeting of the scientific advisors to validate the results and critical errors
- annual report together with the report on critical errors

Sabine Scholl-Bürg
Apolline Imbard
Oliver Braissant
Rachel Carling
Alistair Horman
Daniela Karall
Anke Schumann
(Mary Anne Preece
Brian Fowler)

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Nearly 2000 inherited metabolic disorders are known!

(a classification to get an overview is therefore necessary)

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International classification of inherited metabolic disorders

- **hierarchical, group-based collation** of (all) currently known inherited metabolic disorders
- includes any primary genetic condition in which **alteration of a biochemical pathway** is intrinsic to specific biochemical, clinical and/or pathophysiological features
- aims to **facilitate an improved understanding** of the interconnections between conditions that share functional, clinical and diagnostic features

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ICIMD

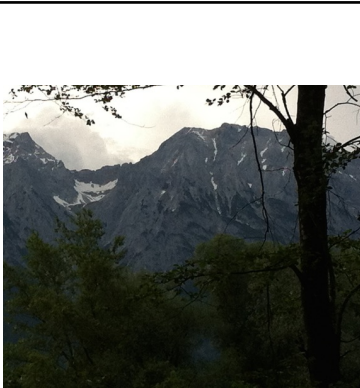
- > 1450 IMD known
- Classification in 6/23 groups
 - **Intermediary metabolism**
 - Lipid metabolism and transport
 - Metabolism of heterocyclic components
 - Complex molecule and organelle metabolism
 - Cofactor and mineral metabolism
 - Metabolic cell signaling
- Of these, approx. 17% have a treatment option.

<http://www.icimd.org>

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[illegible]

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A photograph of a mountain landscape. In the foreground, a large, dark tree trunk and some green foliage are visible. In the background, there are steep, rocky mountains with patches of snow or ice. The sky is overcast with grey clouds. The overall scene is a natural, mountainous environment.

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Case 2024-1

Case 2024-1	
clinical information	deterioration in general condition on the second day of life, suspected sepsis, antibiotic
age at time of sample collection	three days old
sex	female
pregnancy	normal
birth	normal
family history	unremarkable
initial symptoms	
treatment	
miscellaneous	

amino acids in plasma, done with biochrome 30		
	μmol/L	reference range
Taurine	130	10 - 95
Threonine	83	33 - 128
Serine	108	24 - 178
Glutamic acid	341	6 - 93
Glutamine	1711	200 - 550
Proline	506	52 - 277
Glycine	263	70 - 430
Alanine	555	99 - 380
Citrulline	589	0 - 29
Valine	68	57 - 262
Methionine	77	3 - 41
Isoleucine	3	26 - 53
Leucine	peak overlaid	46 - 109
Tyrosine	142	11 - 112
Phenylalanine	71	23 - 110
Ornithine	56	10 - 151
Histidine	155	25 - 114
Lysine	246	45 - 240
Arginine	11	10 - 85
Argininosuccinic acid	detectable	
n.d. not detected		

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Case 2024-1: further details

Ammonia + Bicarbonate and Phosphate → Carbamylphosphate synthetase (defective in CPS deficiency) → Carbamylphosphate

Carbamylphosphate + Ornithine → Ornithine transcarbamylase (defective in OTC deficiency) → Citrulline

Citrulline + Aspartate → Argininosuccinate synthetase (defective in ASS deficiency) → Argininosuccinate

Argininosuccinate → Argininosuccinate lyase (defective in ASL deficiency) → Arginine + Fumarate

Arginine → Arginase → Ornithine + Urea

Ornithine → Carbamylphosphate synthetase (defective in CPS deficiency) → Carbamylphosphate

findings, abnormalities, maximum 2 points (has to be mentioned for 2 points)	interpretation	scores (points)
elevated gln, cit, met	argininosuccinic acid	1
low urea	argininosuccinic acid	1
diagnosis, maximum 2 points	argininosuccinic aciduria (ASL deficiency)	2
further tests (if molecular genetic recommended specify the gene), maximum 2 points	genetic analysis of ASL gene	1
comments	As the detection of argininosuccinic acid is typical for the diagnosis of ASL (argininosuccinate lyase) deficiency, other urea cycle defects are not recognised as differential diagnoses and are therefore not awarded points. In addition, the mention of molecular genetic analysis (ASL gene) without additional metabolite diagnosis (such as determination of orotic acid) was only awarded one point.	1
critical error	Overlooking an inborn disorder of metabolism is considered a critical error.	

order	author	CE	level of difficulty (1- easy, 3- difficult)	estimated overall proficiency	
ASL deficiency	2024_1	SSB, 93%	n=0	1	>90%

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Case 2024-2

Case 2024-2	
clinical information	Speech delay, mental retardation, eating difficulties, foot edema with lameness
age at time of sample collection	5 years
sex	♂
pregnancy	No particularities. Born by caesarean due to medical antecedent of the mother (several co-sections)
birth	Nothing
family history	Nothing
initial symptoms	
treatment	No treatment
miscellaneous	2 older siblings without particularities

amino acids in plasma, done with LC-MS/MS		
	μmol/L	reference range
Glutamine	1032	334 - 666
Alanine	296	122 - 456
Glycine	354	135 - 274
Proline	196	57 - 266
Valine	129	158 - 310
Threonine	179	66 - 170
Lysine	59	94 - 224
Serine	238	75 - 179
Glutamic Acid	87	14 - 106
Leucine	54	68 - 168
Taurine	29	42 - 134
Histidine	62	48 - 112
Ornithine	12	27 - 93
Arginine	22	47 - 123
Tyrosine	31	38 - 94
Phenylalanine	35	33 - 74
Isoleucine	33	37 - 89
Cystine	41	48 - 120
Methionine	18	11 - 43
Citrulline	41	16 - 40

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Case 2024-2: further details

- during hospitalization: detection of anaemia due to a homozygous sickle cell disease, biological work-up also revealed a hyperferritinaemia and increased LDH
- after diagnosis treatment with protein restricted diet, ammonia scavenger and citrulline
- persistant feeding difficulties - gastrostomy
- really hard diagnosis (but as often for LPI) - probable malnutrition and unspecific decrease of some amino acids, but

→ importance of asking for amino acids in urines if there is any doubt

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Lysinuric protein intolerance (LPI)

y ⁺ LAT1 4F2hc	SLC7A7 SLC3A2	y ⁺ L	LPI = déficit y ⁺ LAT1 (SLC7A7)
b ⁰ ·AT rBAT	SLC7A9 SLC3A1	b ⁰ ·+	Cystinurie type III = déficit b ⁰ ·AT (SLC7A9) Cystinurie type I = déficit rBAT (SLC3A1)

Fig. 1 Membranous transport of cationic amino acid(CAA): y⁺LAT and b⁰·AT transport both CAA and neutral amino acids. In addition, y⁺LAT is Na⁺ dependent. a CAA transport at an polarized cell. b CAA transport at a non-polarized cell. CAT1/2 (cationic transporter-1/2) are Na⁺ independent

A) Polarized (Epithelial) cell: y⁺LAT (Na⁺ dependent) on apical side, b⁰·AT on basolateral side. CAA and neutral amino acids are transported.

B) Non-polarized cell: CAT1/2 (Na⁺ independent) on apical side, y⁺LAT on basolateral side. CAA and neutral amino acids are transported.

Noguchi et al., J of Hum Genetics 2019

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Lysinuric protein intolerance (LPI)

- presentation at all ages (neonatal →adult)
- classic hepatobiliary presentation with spontaneous avoidance of protein in diet and secondary malnutrition
 - nausea, vomiting, and diarrhea
 - hepatosplenomegalia – hypotonia
 - altered consciousness (NH₃)
- Complications (may lead to diagnosis)
 - growth retardation, osteopenia
 - hematological and immunological abnormalities: cytopenia, macrophage activation syndrome, susceptibility to infections
 - renal involvement: tubulopathy, renal insufficiency, glomerulonephritis, nephrocalcinosis, proteinuria, hematuria
 - pulmonary involvement: interstitial alveolar pneumonia

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Lysinuric protein intolerance (LPI)

- plasma amino acids
 - ↓ of Lys, Orn, Arg
 - +/- non-specific signs of hyperNH₃: ↑Gln + Glu +/- Ala
- urine amino acids
 - ↑ Lys, Orn, Arg, without associated cystinuria
- increased urinary orotic acid
- cytopenia: anemia, thrombocytopenia
- increased ferritin, LDH, TG
- hyperammonemia (variable)
- genetics (AR): mutations only in the *SLC7A7* gene, not in *SLC3A2*

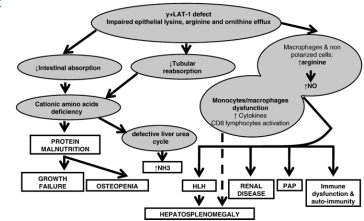
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Lysinuric protein intolerance (LPI)

- complex pathophysiology that is still poorly understood
- intracellular accumulation of arginine in macrophages/monocytes disrupts their activity → immune damage
- Arg = substrate of NOS
→ neurotransmission, vasodilation, inflammation
- role of malnutrition



Ogier de Baulny et al., MGM, 2012

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Case 2024-2: Results

Case 2 abnormalities

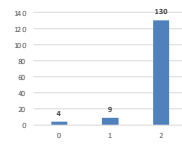
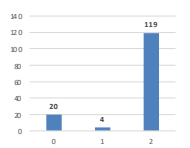
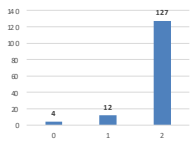
elevated Gln or Cit 1 point
Reduced Lys, Arg, Orn 1 point
Maximum 2 points

Case 2 diagnosis

Lysinuric protein intolerance (LPI) 2 points,
Malnutrition 1 point
Maximum 2 points

Case 2 further testing recommendations

each 1 point, maximum 2 points: orotic acid
(organic acids) in urine; amino acids in urine;
ammonia, ferritin, triglycerides; molecular
genetic analysis *SLC7A7* gene



critical error If elevated Gln concentration doesn't lead to further action or if the findings are quoted as unspecific.

N=1

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Case 2024-2: Results

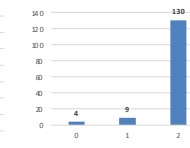
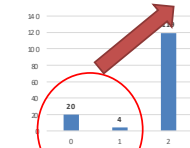
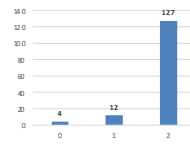
Case 2 abnormalities

elevated Gln or Cit 1 point
Reduced Lys, Arg, Orn 1 point
Maximum 2 points

- Kwashiorkor or low protein intake
- Ca glutamine synthetase deficiency
- glutaminase deficiency
- Lys ornithine transcarbamylase deficiency
- Me carbamoyl phosphate synthetase I
- BCKD kinase deficiency

recommendations

ints: orotic acid
in acids in urine;
ides; molecular
genetic analysis



critical error If elevated Gln concentration doesn't lead to further action or if the findings are quoted as unspecific.

N=1

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Case 2024-3

Case 2024-3		amino acids in plasma, done with LC-MS/MS			
		μmol/L	reference range		
clinical information	Cutis laxa, axial hypotonia, IUGR (height and weight < 3SD), microcephaly, joint hyperlaxity, inguinal hernia	Glutamine	873	515	- 651
age at time of sample collection	4 months	Alanine	670	258	- 400
sex	male	Glycine	337	190	- 268
pregnancy		Proline	78	144	- 220
birth	IUGR (intrauterine growth retardation)	Valine	140	174	- 246
family history		Threonine	147	91	- 167
initial symptoms		Lysine	116	129	- 191
treatment	/	Serine	189	110	- 168
miscellaneous		Glutamic Acid	127	44	- 80
		Leucine	92	117	- 142
		Taurine	37	36	- 82
		Histidine	73	63	- 91
		Ornithine	19	43	- 83
		Arginine	33	64	- 92
		Tyrosine	69	51	- 85
		Phenylalanine	53	46	- 82
		Isoleucine	36	48	- 80
		Cysteine	19	72	- 100
		Methionine	25	21	- 33
		Aspartic Acid	13	5	- 13
		Citrulline	5	18	- 42

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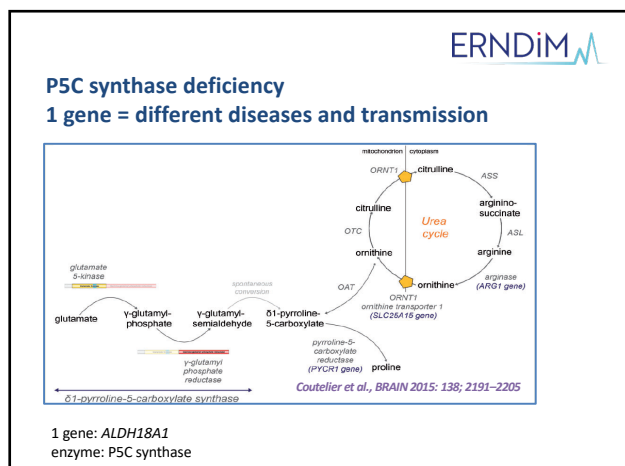
Case 2024-3

- The results are from a 4 months-old boy from a non-consanguineous family.
- He presented an intrauterine growth retardation and joint hyperlaxity, inguinal hernia and cutis laxa at birth.
- At 2 months he presented a neurological deterioration with axial hypotonia, pyramidal syndrome and a cataract on the left eye was diagnosed.
- His growth was also severely impaired leading to naso-gastric feeding. He died at 7 month of age.
- Decreased Cit, Arg, Orn, Pro → De Barys syndrome due to D1-Pyrroline-5-carboxylate (P5C) synthase deficiency (*ALDH18A1* gene).
- This is an ultrarare metabolic disorder, but with the clinical symptoms (and pubmed search) and the description of the amino acid changes, the diagnosis could be made.

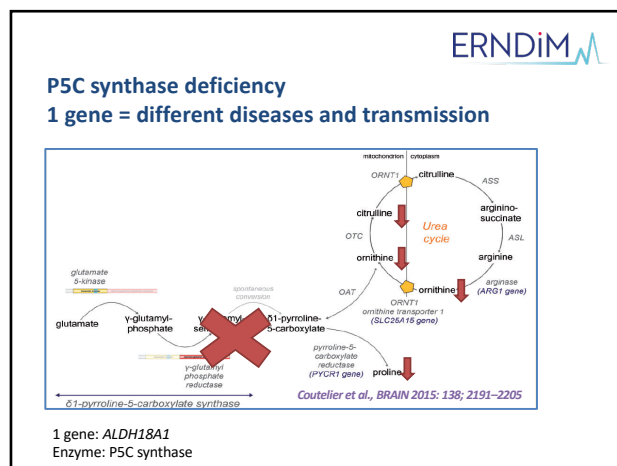
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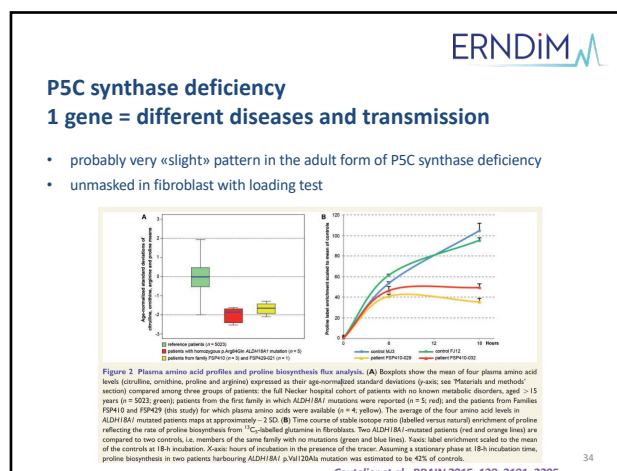
P5C synthase deficiency

1 gene = different diseases and transmission

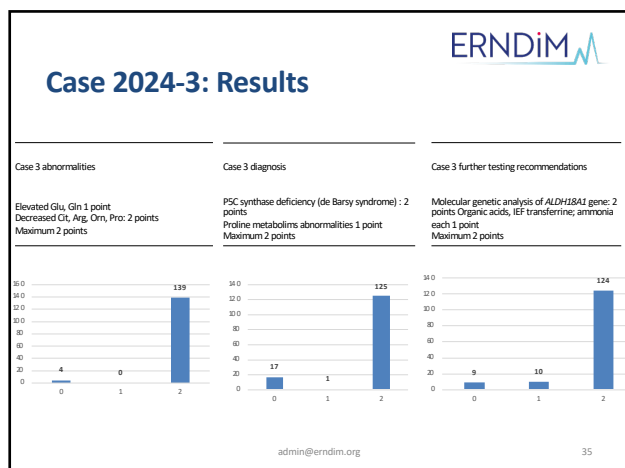
- severe form: cutis laxa & hyperammonemia
- = de Barsy syndrome
 - AD or AR (mutation dependent)
 - symptoms
 - severe psychomotor retardation
 - cutis laxa
 - joint abnormalities
 - small size
 - cataract
 - microcephaly
 - rare (< 20 cases)
 - very severe (early death ++)
- adult form: spastic paraparesis
- autosomal dominant or autosomal recessive (mutation dependent)
- symptoms
 - Pyramidal syndrome with paraparesis and spasticity
- found by exome sequencing in large cohorts of families with spastic paraparesis

Fisher et al., Molecular Genetics and Metabolism 112 (2014)

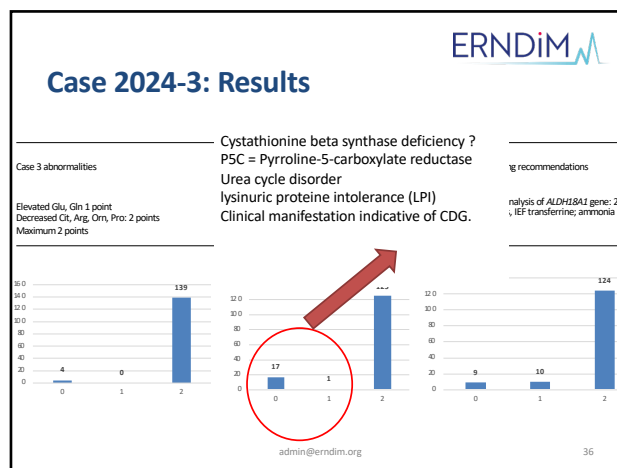
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Case 2024-4

clinical information	seizures, liver failure with disturbed clotting in the age of eight months
age at time of sample collection	8 months old
sex	male
pregnancy	normal
birth	normal
family history	unremarkable, two healthy brothers
initial symptoms	ictus
treatment	
miscellaneous	
laboratory results	
amino acids	19 µmol/L (25-70 µmol/L)

amino acids in plasma, done with biochrome 30	µmol/L	reference range
Glutamine	576	200 - 350
Alanine	279	99 - 320
Glycine	188	56 - 308
Proline	174	51 - 185
Valine	97	57 - 262
Threonine	195	33 - 128
Lysine	196	45 - 150
Serine	109	24 - 178
Glutamic Acid	24	6 - 70
Leucine	58	45 - 155
Taurine	17	10 - 130
Histidine	61	24 - 112
Ornithine	82	10 - 107
Arginine	75	11 - 75
Tyrosine	222	11 - 112
Phenylalanine	55	23 - 70
Isoleucine	29	26 - 94
Cystine	49	10 - 40
Methionine	174	3 - 29
Aspartic acid	n.d.	-
Citrulline	48	5 - 24
	n.d. not detected	

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Case 2024-4: further details

interpretation	scores (points)
findings, abnormalities, maximum 2 points	1
diagnosis, maximum 2 points	1
further tests (if molecular genetic recommended specify the gene), maximum 2 points	1
comments	
critical error	

order	author	CE	level of difficulty (1-5)	estimated overall proficiency
2024-4	SBH	87%	nd	3
				<80%

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Case 2024-5

clinical information	microcephaly, psychomotor retardation, failure to thrive
age at time of sample collection	5 years
sex	male
pregnancy	
birth	born preterm (gestational week 35+6)
family history	consanguineous parents
initial symptoms	
treatment	
miscellaneous	MRI: enlargement of the peri-cerebral spaces of the lateral ventricles and delayed myelinisation

amino acids in plasma, done with LC MS/MS	µmol/L	reference range
Glutamine	541	334 - 666
Alanine	313	122 - 456
Glycine	190	135 - 274
Proline	126	57 - 266
Valine	49	158 - 310
Threonine	127	66 - 170
Lysine	174	94 - 224
Serine	131	75 - 179
Glutamic Acid	66	14 - 106
Leucine	16	68 - 168
Taurine	69	42 - 134
Histidine	59	48 - 112
Ornithine	61	27 - 93
Arginine	54	47 - 123
Tyrosine	79	38 - 94
Phenylalanine	57	33 - 74
Isoleucine	7	37 - 89
Cystine	44	48 - 120
Methionine	19	11 - 43
Citrulline	18	16 - 40

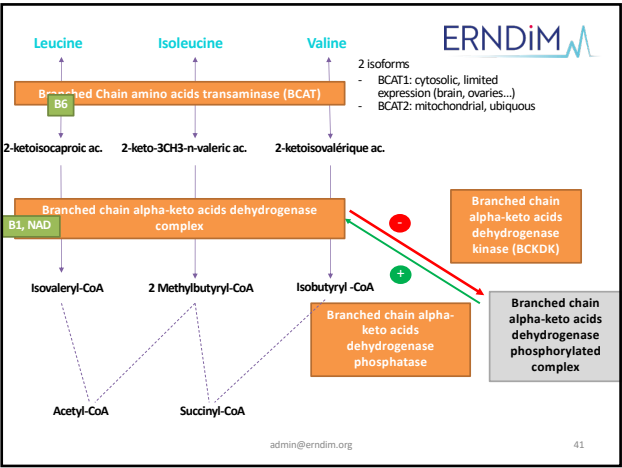
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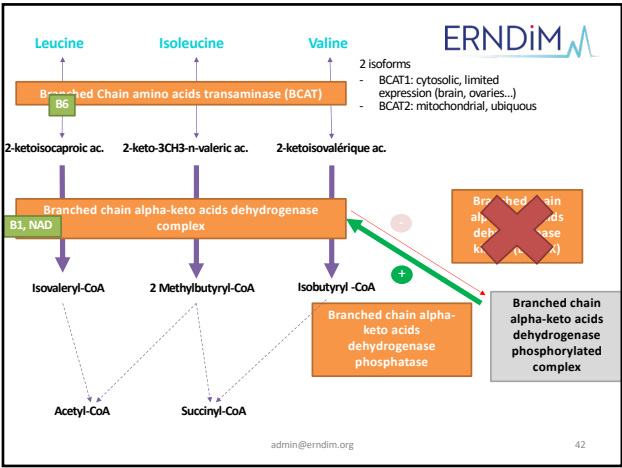
Case 2024-5

- The sample was taken from a five-year-old boy who had microcephaly, mental retardation and a failure to thrive.
- In the extended work-up, cerebral MRI showed enlargement of the peri-cerebral spaces of the lateral ventricles and delayed myelinisation.
- The analysis of amino acids showed isolated reduced concentrations of branched-chain amino acids.
- BCKDK deficiency

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Case 2024-5



- First description of BCKDK deficiency in 2012

Mutations in BCKDK-kinase Lead to a Potentially Treatable Form of Autism with Epilepsy

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- Large cohort (17 patients) described in 2023

BCKDK deficiency: a treatable neurodevelopmental disease amenable to newborn screening

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Table 1 BCKDK phenotype at diagnosis

Phenotype	HPO code	Frequency n (%)
Progressive microcephaly	HP:0000253	1/20 (5%)
Cerebral developmental delay	HP:0011263	21 (100%)
Motor delay	HP:0011270	21 (100%)
Language impairment	HP:0004463	17 (100%)
Intellectual disability	HP:0012489	16 (100%)
Intellectual disability, severe	HP:0010854	15/16 (93.8%)
Developmental regression	HP:0002235	1/16 (6.2%)
Behavioural abnormality	HP:0000708	20/21 (95.2%)
Autistic behaviour	HP:0000129	14/21 (66.7%)
Autism	HP:000017	14/21 (66.7%)
Self injurious behaviour	HP:000716	5/17 (29%)
Aggressive behaviour	HP:0000718	8/18 (44.4%)
Hyperactivity	HP:0000752	5/21 (23.8%)
Stereotyped	HP:0000711	4/21 (19.0%)
Attention deficit hyperactivity disorder	HP:000018	4/18 (22.2%)
Seizures	HP:0001250	9/21 (42.9%)
Bilateral tonic-clonic seizure	HP:000069	3/18 (16.7%)
Generalised tonic-clonic seizure	HP:000123	1/18 (5.6%)
Typical absence seizure	HP:001147	3/18 (11.1%)
Generalised-onset seizure	HP:000137	9/21 (42.9%)
Focal-onset seizure	HP:000759	1/21 (4.8%)
Bilateral epileptiform activity	HP:001182	12/18 (66.7%)
Hypotonia	HP:000124	5/17 (29.4%)
Abnormality of movement	HP:000022	3/20 (15%)
Dystonia	HP:000132	2/10 (20%)
Ataxia	HP:0001251	1/20 (5%)
Hyperkinetic movements	HP:000487	1/20 (5%)
Chorea	HP:000312	1/17 (5.9%)
Feeding difficulties	HP:001168	5/20 (25%)
Sensorineural hearing impairment	HP:000067	3/21 (14.3%)
Hyperreflexia	HP:000147	5/17 (29.4%)
Polysomnopathy	HP:000171	1/17 (5.9%)
Abnormal facial shape	HP:000199	10/21 (47.6%)
Full cheeks	HP:000028	4/21 (19%)
Thin upper lip vermillion	HP:0000219	4/21 (19%)
Abnormal nasal bridge morphology	HP:000422	2/21 (9.5%)
Hypoplastic philtrum	HP:000326	2/21 (9.5%)
Small forehead	HP:000050	3/21 (14.3%)
Crooked hair	HP:000036	1/17 (5.9%)
Dry skin	HP:000058	2/17 (11.8%)
Inflammatory abnormality of the skin	HP:001123	2/17 (11.8%)
Hypotrichiasis	HP:000026	1/17 (5.9%)

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Case 2024-5

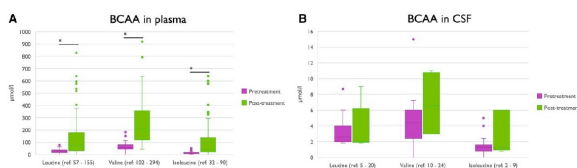


Figure 3 Branched chain amino acids (BCAA) before BCAA supplementation and a follow-up. (A) Pretreatment: n=5/21, 1/5 patients did not have CSF leucine levels pretreatment. No significant differences between the samples from before and after treatment. Mean of 1.2 samples per patient pretreatment and 1 sample post-treatment. (B) Pretreatment: n=19/21; post-treatment: n=14/21; P<0.001 for leucine, valine and isoleucine independently. Mean of 1.8 samples per patient pretreatment and 6.4 samples post-treatment.

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- !!! Sample must be taken while fasting

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Case 2024-5



- Possibly treatable disease if treatment is started early

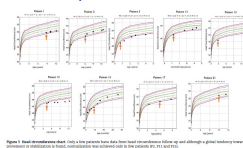


Figure 1 Individual patient data (IPD) for 17 patients with BCKDK deficiency. The figure shows 17 line graphs, each representing a patient's clinical course over time. The x-axis represents age in years, and the y-axis represents various clinical parameters. The graphs show a general trend of improvement in clinical parameters over time, particularly in the first 10 years of life.

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- Neonatal screening ?

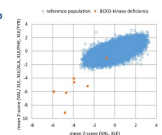


Figure 2 Neonatal screening results for BCKDK deficiency. The figure shows a scatter plot of the ratio of branched chain amino acids (BCAA) to total amino acids (TAA) in newborns. The x-axis represents the ratio of BCAA to TAA, and the y-axis represents the ratio of BCAA to TAA. The plot shows a clear separation between the BCKDK deficiency group (red dots) and the control group (blue dots).

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Case 2024-5: Results



Case 5 abnormalities

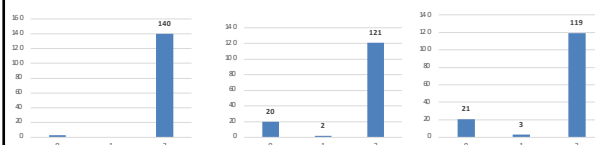
Decreased BCAA (Ile, Val, Leu): 2 points

Case 5 diagnosis

BCKDK (branched-chain 2-ketoacid dehydrogenase kinase) deficiency 2 points,
Glucose infusion/anabolism (hyperinsulinism) 1 point
Maximum 2 points

Case 5 further testing recommendations

Repetition of the determination of amino acids in plasma: 1 point
Molecular genetic analysis of BCKDK: 2 points
Maximum 2 points



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Case 2024-5: Results



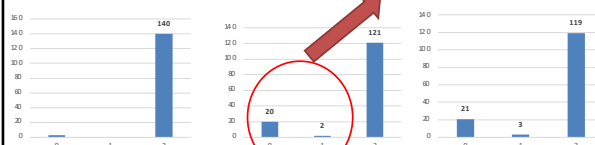
Case 5 abnormalities

Decreased BCAA (Ile, Val, Leu): 2 points

MSUD in treatment
Congenital hyperinsulinism
Denutrition
lysosomal or peroxysomal diseases
treated organic aciduria
Autosomal recessive microcephaly
Hartnup disease
Leigh syndrome

Recommendations

Repetition of the determination of amino acids in plasma: 1 point
Molecular genetic analysis of BCKDK: 2 points



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Case 2024-6

Case 2014-6		amino acids in urine, done with Biochrome 30	
		mmol/mol Creat	reference range
clinical information	seizures (begin 6 days old), responsive to pyridoxine		
age at time of sample collection	7 days	Glutamine 157	52 - 205
sex	female	Alanine 205	75 - 244
		Glycine 825	283 - 1097
pregnancy	uneventful	Proline 194	21 - 213
birth	normal	Valine 20	3 - 26
		Threonine 52	20 - 138
family history	unremarkable	Lysine 44	22 - 171
		Serine 132	80 - 282
initial symptoms		Glutamic Acid 15	0 - 30
		Leucine 20	3 - 25
		Taurine 442	8 - 226
treatment		Histidine 199	80 - 295
		Ornithine 6	0 - 19
		Arginine 5	0 - 14
miscellaneous		Tyrosine 27	6 - 55
		Phenylalanine 16	4 - 32
		Isoleucine 4	0 - 6
		Cystine 15	24 - 78
		Methionine 4	7 - 27
		Aspartic acid 7	2 - 12
		Citrulline 5	0 - 13
		Phosphoethanolamine ★	705 n.s. - n.s.
Laboratory results in plasma			
ASAT	24 U/L (R: 10-50)		
ALAT	7 U/L (R: 3-38)		
α-GT	96 U/L (R: 0-265)		
alkaline phosphatase ★	<20 U/L (R: 48-406)		
Ammonia	14 μmol/L (R: 31-104)		
Lactate	18 mg/dL (R: 1-26)		

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Case 2024-6: further details

	interpretation	scores (points)
findings, abnormalities, maximum 2 points	elevated PEA decreased AP	2 2
diagnosis, maximum 2 points	Hyphosphosphatasia Pyridoxine-responsive seizures	1 2
further tests (if molecular genetic recommended specify the gene), maximum 2 points	phosphoethanolamine (P, CSF, U) alkaline phosphatase (AP) in plasma determination of vitamin B6 metabolites molecular genetic analysis of ALPL gene molecular genetic analysis of ALDH7A1 gene	1 1 1 1 1
comments	The participant should recognize that it is a case of vitamin B6-dependent epilepsy and begin further clarification of the cause.	
critical error	If the participant does not recognise that an inborn metabolic disorder is present.	

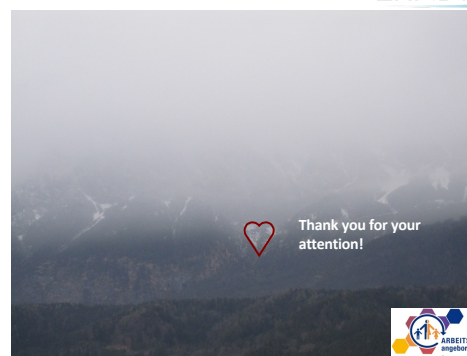
	order	author	CE	level of difficulty (1-easy, 3-difficult)	estimated overall proficiency
hypophosphatasia	2024_6	SSB, 98%	no CE	1	>90%

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Take to work message

- The aai aims to improve the interpretation of complex laboratory results.
- Real-life cases are used.
- The performance of the participating laboratories is generally good.
- Errors arise from overlooking a relevant laboratory result, misinterpretation, or a lack of recommendations for further investigation (the term 'molecular genetic analysis' as the only recommendation is insufficient).
- We hope that the cases have been instructive for you so far.

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