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# QAP for qualitative urinary organic acid analysis

# Annual Report (Amended) 2016(Sheffield)

#### Original Date of issue: 07th February 2017

# Amended report issued: 2<sup>nd</sup> March 2017

Changes: Data in table 3 was incorrect for 2015 & 2014 in the original report. This would have affected the total scores and the average scores. Please discard the original report dated 7<sup>th</sup> February 2017.

# Participation

Active participants (reporting on at least one set of samples in the year) are shown in Table 1. The numbers of participants continues to increase. New applicants are distributed between the Sheffield and Heidelberg qualitative urinary organic acid schemes which are run separately. The two organising laboratories each participate in the other's scheme.

	2016	2015	2014	2013	2012	2011	2010
Argentina	4	3	3	2	2	2	2
Australia	6	6	6	6	6	6	6
Belgium	6	6	6	6	5	5	6
Brazil	2	2	2	2	2	2	-
Canada	1	1	1	1	1	1	1
Columbia	1	1	1	1	1	1	1
Chile	1	1	1	-	-	-	-
China	7	11	11	9	8	10	7
Finland	2	2	2	2	1	1	1
France	17	15	15	15	15	15	13
Germany†	1	1	1	1	1	1	1
Israel	3	3	3	3	3	3	4
Japan	2	2	2	1	1	1	1
Lebanon	1	1	1	1	1	1	1
Malaysia	3	3	3	3	3	3	4
New Zealand	1	1	1	1	1	1	1
Portugal	3	2	2	2	2	2	2

#### Table 1: Geographical distribution of registered participants

*Qualitative urinary organic acid scheme Annual Report 2016 (amended report issued: 02 March 2017)* 

Republic of Korea	-	1	1	1	1	1	1
Republic of Ireland	1	1	1	1	1	1	1
Republic of Singapore	1	1	1	1	1	1	1
South Africa	2	2	2	2	2	2	2
Spain	7	6	6	6	6	6	6
Turkey	10	7	3	3	3	3	3
United Kingdom	16	18	18	18	18	18	19
USA	6	5	5	5	3	3	3
Uruguay	1	1	1	-	-	-	-
Pakistan	1	-	-	-	1	1	1
Kingdom of Saudi Arabia	1	-	-	-	-	-	-
Vietnam	1	1	1	1	1	1	-
TOTAL	108	104	100	94	91	95	90

ERNDIM

† Heidelberg laboratory

#### Samples and results

Three sets of three samples (numbered 232-240) were dispatched together in April 2016. Laboratories were asked to analyse the sets at intervals during the year as if they were separate circulations. Ninety-nine laboratories returned results for all three sets; seven returned two, one laboratories made only a single return, and one made no return. Two participants are educational participants.

### Scoring of results

To enable data reduction the results were scored as shown below:

Satisfactory	4	Helpful but incomplete	3
Not helpful	2	Slightly misleading	1
Misleading	0		

One point was deducted for each transposed sample number

# **Table 2: Distribution of scores for individual samples** (laboratories making returns)

	Scores							
Sample	0	1	2	3	4			
Sample 232       48 year old male with muscle weakness         Nothing specifically diagnostic         .	3	0	0	0	99			
Sample 2331 year old boy frequent vomiting and epilepsySignificantly increased excretion of malonic acid with increased methylmalonic acid. These results would be consistent with malonic	4	1	0	5	93			



aciduria and the most likely defect would be Malonyl CoA decarboxylase deficiency.					
Sample 2344 year old boy, unwell, drowsy and vomitingA large peak of 2-hydroxyisovalerate is present together with lesser amounts of 2-hydroxy-3-methylvalerate and 2-hydroxyisocaproate, consistent with a defect in branched-chain 2-oxoacid dehydrogenase activity. The clinical details suggest a mild or intermittent variant of Maple Syrup Urine Disease. Increased excretion of <i>p</i> -hydroxyphenyl- lactate indicates some degree of liver disorder.	5	3	1	1	93
Sample 235       3 year old microcephaly         Nothing specifically diagnostic	23	0	1	2	76
Sample 23639 year old woman. Mother of S235Nothing specifically diagnostic	19	0	1	1	81
<ul> <li>Sample 237 5 year old girl with recurrent kidney stones</li> <li>Significantly elevated peaks of glycolate and oxalate. Absence of glycerate.</li> <li>Results consistent with a diagnosis of Primary Hyperoxaluria Type1</li> </ul>	9	1	3	0	89
<i>Sample 238</i> 9 month old girl, failure to thrive Increased orotic acid which may indicate an inborn error of metabolism, check plasma ammonia and urine and plasma amino acids as a matter of urgency. Discuss with a metabolic consultant.	17	0	0	0	87
Sample 2396 year old female – mild dysmorphic featuresIncreased excretion of 3-methylglutaconic acid and 3-methlyglutaricacid. Results indicative of 3-methlyglutaconic aciduria. Suggest furtherfollow up tests to confirm and suggest discussion with a metabolicconsultant.	13	0	4	8	79
Sample 24035 year old female: Muscle weaknessIncreased excretion of 3-hydroxyisovaleric acid and 3-methylcrotonyl glycine. Results consistent with 3-Methylcrotonyl-CoA Carboxylase deficiency.	0	0	3	0	101



# Table 3: Cumulative scores for 2014 - 2016 (current Sheffield participants only)

Lab	Total	Number	Total	Number	Total	Numb	Total	Average
Number	Score 2016 <sup>*</sup>	of returns 2016	Score 2015 <sup>*</sup>	of returns 2015	Score 2014 <sup>*</sup>	er of returns	Score over 3	score per year (max
						2014	years (max	36)
							(max 108)	
1	36	3	-	-	-	-	36	36
2	32	3	36	3	36	3	104	35
3	32	3	36	3	31	3	99	33
4	32	3	34	3	35	3	101	34
5	32	3	36	3	36	3	104	35
6	32	3	34	3	36	3	102	34
7	36	3	34	3	36	3	106	35
8	32	3	35	3	36	3	103	34
9	32	3	36	3	32	3	100	33
10	32	3	36	3	36	3	104	35
11	36	3	36	3	36	3	108	36
12	32	3	32	3	35	3	99	33
13	36	3	34	3	36	3	106	35
14	26	3	30	3	23	3	79	26
15	36	3	36	3	30	3	102	34
16	32 29	3	28 36	3	4 35	1	64 100	21 33
17	29 36	3	36	3	35	3	100	33
18 19	28	3	30	3	27	3	91	30
19 20	35	3	36	3	34	3	105	30
20	33	3	36	3	35	3	103	33
21	36	3	36	3	36	3	108	36
22	35	3	36	3	35	3	106	35
23	36	3	32	3	36	3	100	35
25	36	3	36	3	35	3	107	36
26	36	3	35	3	36	3	107	36
27	28	3	36	3	36	3	100	33
28	36	3	34	3	24	2	94	31
29	36	3	35	3	36	3	107	36
30	36	3	36	3	24	2	96	32
31	36	3	36	3	35	3	107	36
32	36	3	35	3	36	3	107	36



33	22	3	32	3	34	3	88	29
34	32	3	36	3	33	3	101	34
Lab Number	Total Score 2016	Number of returns 2016	Total Score 2015	Number of returns 2015	Total Score 2014	Numb er of returns 2014	Total Score over 3 years (max 108)	Average score per year (max 36)
35	34	3	12	1	36	3	82	27
36	28	3	36	3	36	3	100	33
37	32	3	35	3	34	3	101	34
38	32	3	36	3	35	3	103	34
39	32	3	36	3	36	3	104	35
40	36	3	36	3	36	3	108	36
41	28	3	36	3	36	3	100	33
42	36	3	36	3	35	3	107	36
43	36	3	36	3	36	3	108	36
44	36	3	36	3	36	3	108	36
45	32	3	36	3	32	3	100	33
46	32	3	36	3	34	3	102	34
47	32	3	36	3	35	3	103	34
48	35	3	36	3	35	3	106	35
49	36 27	3	36 32	3	36 36	3	108 95	36 32
50 51	35	3	32	3	30	3	103	32
51	35	3	35	3	36	3	103	34
52	31	3	24	2	26	3	81	27
55	28	3	35	3	36	3	99	33
55	34	2	36	3	36	3	106	35
56	29	3	36	3	33	3	98	33
57	28	3	30	3	28	3	86	29
58	(edu) 12	2	-	-	-	-	12	12
50	32	3	33	3	31	3	96	32
60	36	3	26	3	31	3	93	31
61	32	3	36	3	35	3	103	34
62	32	3	22	2	35	3	89	30
63	36	3	36	3	32	3	104	35
64	36	3	36	3	34	3	106	35
65	34	3	32	3	33	3	99	33
66	35	3	-	-	31	3	66	33
67	36	3	32	3	34	3	102	34
68	32	3	24	2	36	3	92	31
69	36	3	35	3	31	3	102	34
70	32	3	36	3	28	3	96	32



71	32	3	32	3	36	3	100	33
72	32	3	32	3	12	1	76	25
73	32	3	20	2	35	3	87	29
Lab Number	Total Score 2016 <sup>*</sup>	Number of returns 2016	Total Score 2015 <sup>*</sup>	Number of returns 2015	Total Score 2014 <sup>*</sup>	Numb er of returns 2014	Total Score over 3 years (max 108)	Average score per year (max 36)
74	24	2	20	2	24	2	68	23
75	19	3	10	1	34	3	63	21
76	28	3	36	3	36	3	100	33
77	24	3	30	3	30	3	84	28
78	28	3	34	3	34	3	96	32
79	29	3	36	3	- 24	-	65	33
80	36 35	3	34 32	3	34 30	3	104 97	35 32
81	28	3	32	3	30	3	97	32
82 83	32	3	20	2	21	3	73	24
84	28	3	36	3	36	3	100	33
85	36	3	10	1	16	2	62	21
86	31	3	26	3	-	-	57	29
87	30	3	20	3	-	_	50	25
88	20	3	36	3	22	3	78	26
89	10	3	36	3			46	23
90	36	3	32	3	36	3	104	35
91	24	3	-	-	-	-	24	24
92	-	-	31	3	31	3	62	31
93	18	3	22	2	-	-	40	20
94	8	1	32	3	34	3	74	25
95	34	3	36	3	35	3	105	35
96	20	2	33	3	32	3	85	28
97	33	3	35	3	35	3	103	34
98	22	3	-	-	-	-	22	22
99	35	3	36	3	35	3	106	35
100	36	3	30	3	34	3	100	33
101	24	2	28	3	31	3	83	28
102	32	3	24	3	-	-	56	28
103	26	3	-	-	-	-	26	26
104	36	3	36	3	-	-	72 28	36
105	28 17	3	-	-	-	-	28 17	28 17
106	(edu)	-	-	-	-	-	- 17	- 17
107	(edu) 36	- 3	-	-	-	-	- 36	- 36
108	30	3	-	-	-	-	30	30

\* Maximum total score is 36 for three returns and 24 for two returns. Maximum scores over 3



years = 108 and is only applicable if all three returns have been received over the previous three years.

Your Laboratory OA Number in the above Table is 0

# <u>Please use your ERNDIM number in all communication with the</u> <u>scientific advisors, laboratory numbers are only used for the purpose</u> <u>of this table.</u>

# Commentary

As in previous years, samples were sent out to cover the spectrum of what is typically observed in the metabolic laboratory. A mix of clearly diagnostic profiles and some more challenging profiles were provided. As in previous years normal profiles were also sent out. The requirement to interpret a normal profile, as such, is as important as correctly identifying abnormal profiles. Correctly identifying a profile as normal can avoid unnecessary further investigation and distress to the patient and family. For the samples sent out in 2016 the majority of participants correctly assigned the normal profile however as in previous years there are still a significant proportion of participants who over interpret based on clinical details. This was clearly the case with samples 236 & 237 which were donated by an unaffected mother and child. Participants, in view of the clinical picture, appropriately looked for metabolites associated with phenylketonuria however even in the absence of any metabolites a number still incorrectly assigned the disorder without any biochemical evidence. It is useful to be guided by clinical details but the likely diagnosis should only be suggested in view of the biochemical evidence. In contrast, highlighting the complexity of the role of metabolic interpretation, the hyperoxaluria type 1 case (237) had strong clinical details; 5 year old girl with recurrent kidney stones. The majority of participants correctly identified increase in both glycolate and oxalate. Some participant's identified increased glycolate only. Which is not uncommon, as oxalate extraction for organic acids is problematic and highly variable, given this knowledge and clinical details further testing such as quantitative method for oxalate measurement would be expected.

Of the abnormal samples, two proved particularly challenging; the orotic acid (sample 238) and the sample with 3-methylglutaconic aciduria (sample 239). Over the past few years the detection and identification of orotic acid has continued to be challenging for a number of laboratories. The detection of orotic acid is crucial in the diagnosis of inherited defects of metabolism and laboratories need to have internal procedures in place to ensure this peak is not missed.

The detection of a slightly increased 3-methylglutaconic acid also proved to be challenging. The significance of a persistently elevated excretion of this metabolite in the absence of a known protein defect is open to debate, but generally it is agreed this finding requires further follow up and investigation. The majority of laboratories agreed and did identify this as abnormally increased and suggested appropriate follow up. Traditionally in Sheffield we have used the numerical classification for the classification of type IV methlyglutaconic acid. However, we draw the attention of



our participants to manuscript of Saskia Wortmann on an updated classification. For reporting purposes one should distinguish between primary 3-methylglutaconic aciduria (previously known at type I) due to deficiency of 3-methylglutaconyl-CoA Hydratase deficiency and secondary 3-methlyglutaconic aciduria. The secondary group should be identified by the defective protein or historical name (e.g. TAZ or Barth syndrome, SERAC1 defect or MEGDEL syndrome, OPA3 defect or Costeff syndrome, DNAJC19 defect or DCMA syndrome and TMEM70 defect). The remaining patients that do not fit the above criteria should be referred to as not otherwise specified (NOS) 3-methylgutaconic aciduria. The full classification can be found in J Inherit Metab Dis (2013) 35:923-928. This sample (239) was donated from a patient with NOS 3-methylgutaconic aciduria.

It is appreciated that in all the cases the urine organic acid profile is only part of the diagnostic investigation and where the organic acid profile does not give a clear diagnosis the further investigations box is key when it comes to scoring. The 'Further investigations' box should indicate any additional investigations you consider necessary to interpret or confirm conclusions based on the analytical results. The 'Additional comments' box may also be used for caveats or to suggest other lines of investigation based on the clinical presentation rather than the analytical findings. Suggestions should follow a logical hierarchy with simple group investigations such as amino acid chromatography or blood-spot acylcarnitine profiling (if indicated) taking precedence over much more specific investigations such as gene sequencing.

We are somewhat limited in the samples we send out by the volume of sample required and the number of participants within our scheme. We are also limited by the number of samples with appropriate amounts of abnormal metabolites in. We appeal to any of our participants to donate samples for inclusion in our scheme or any of the other ERNDIM schemes.

#### *Certificates of Participation and Performance*

We are required to define "Participation" and "Satisfactory Performance" for the purpose of the ERNDIM Annual Certificate which covers all ERNDIM schemes. For this urinary organic acid scheme we have defined "Participation" as requiring at least two returns during the year. Defining "Satisfactory Performance" is more problematical as in some years there are more difficult samples than in others.

The criteria for satisfactory performance: for three returns a score of 22/36 and for two returns a score of 15 or more. On this basis four laboratories were deemed to fall below the satisfactory criteria. Satisfactory Performance" criteria are always somewhat arbitrary and in practice even a single missed or wrong diagnosis can be highly damaging. Thus the reason(s) for failure to correctly report on <u>any</u> of the samples in the scheme should be investigated locally and appropriate remedial action taken.

A critical error is an error that would be unacceptable to the majority of labs and would have a serious adverse effect on patient management. The introduction of critical error is on the advice of the Genetic Services Quality Committee (GSQC) of the European Society of Human Genetics (ESHG), which wants to see harmonisation



across all European genetic EQA providers. A confirmed critical error will mean automatic classification as a poor performer. The final scoring of all qualitative schemes will be discussed at the Spring meeting of the Scientific Advisory Board (SAB) and all proposed critical errors will need to be ratified by the SAB before being confirmed.

We thank Lynne Wolstenholme for administering our participant database and dealing with the returns, and Jennifer Watkinson for preparing the samples. We hope that you continue to find this scheme useful.

Yours sincerely

Mrs Camilla Scott & Dr Jane Dalley Scheme organisers