

Combination of enzyme analysis, allele-specific PCR and sequencing to detect mutations in the *GALT* gene

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Summary Newborn screening can identify patients with classical galactosaemia, and their diagnosis needs to be confirmed with assay of the activity of galactose-1-phosphate uridylyltransferase (*GALT*). Unfortunately, in many cases the results can be ambiguous and further testing is required. Here we report a combination of biochemical analysis of *GALT* enzyme activity and mutation analysis of the most common mutations in the corresponding gene. Samples ($n=243$) submitted for confirmatory testing for classical galactosaemia were analysed simultaneously for *GALT* enzyme activity and allele-specific PCR/fragment analysis for seven mutations and two polymorphisms in the *GALT* gene (mutations IVS2-2A>G, p.S135L, p.T138M, p.L195P,

p.K285N, p.Q188R, p.Y209C; polymorphisms p.N314D, p.L218L). Mutation detection accorded with biochemical analysis in 93% of samples. Subsequently, a total of 34 samples with either discordant results between the above methods or low enzyme activity were fully sequenced, identifying previously reported pathogenic mutations and seven novel variations (p.P185H, p.R201C, p.E220K, p.R223S, p.I278N, p.L289F and p.L218X) in the *GALT* gene. This approach further increased concordance between genetic and biochemical analysis to 99% of all alleles tested. Our results indicate that DNA testing can help to verify biochemical enzymatic data and improve distinction of borderline enzyme activities where a patient may still benefit from treatment.

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Electronic Supplementary Material

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