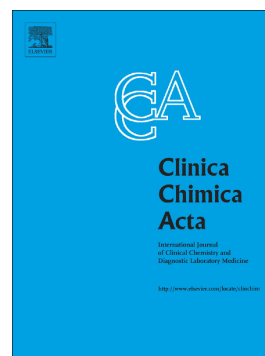


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In silico prediction of the pathogenic effect of a novel variant of BCKDHA leading to classical maple syrup urine disease identified using clinical exome sequencing

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***In silico* prediction of the pathogenic effect of a novel variant of *BCKDHA* leading to classical maple syrup urine disease identified using clinical exome sequencing**

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