

## **Studying a rare disease from a biophysical point of view: the example of cblC**

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The cblC disease is a rare inborn disorder of the vitamin B12 (cobalamin, Cbl) metabolism characterized by combined methylmalonic aciduria and homocystinuria. The clinical consequences are devastating and, even when early treated with current therapies, the affected children manifest symptoms involving vision, growth, and learning. The molecular genetic cause of the disease was found in the mutations of the gene coding for MMACHC, a 282 amino acid protein that transports and processes the various forms of Cbl.

Here, by using biophysical methods including spectroscopy, microcalorimetry and molecular dynamics we investigated the differences in stability, binding properties and functionality between MMACHC wild type and the pathological variant p.R161Q, resulting from the most common missense mutation found in cblC patients.

Overall, our results reveal how a biophysical approach based on the complementarity of computational and experimental methods can offer new insights in the study of the specific effects of the pathological cblC mutation and help prospecting new routes for the cblC treatment.

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