

# Learning from the worm: the effectiveness of protein-bound Moco to treat Moco deficiency

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**Molybdenum cofactor (Moco) is synthesized endogenously in humans and is essential for human development. Supplementation of Moco or its precursors has been explored as a therapy to treat Moco-deficient patients but with significant limitations. By using the nematode *C. elegans* as a model, Warnhoff and colleagues (pp. 212–217) describe the beneficial impact of protein-bound Moco supplementation to treat Moco deficiency. If such an effect is conserved, this advance from basic research in worms may have significant clinical implications as a novel therapy for molybdenum cofactor deficiency.**

In recent years, the genetic model organism *C. elegans* has increasingly been used to address fundamental biological problems in the fields of cell metabolism and nutrient homeostasis, taking advantage of the conservation of most of the basic pathways and mechanisms between that in the nematode and mammals. While the conceptual advances of the studies are commonly found in the connections of metabolic events to animal development, behavior, and aging, some research results have more direct medical relevance. The study on molybdenum cofactor deficiency in this issue of *Genes & Development* Warnhoff et al. (2021) presents an excellent example of using *C. elegans* as a unique and effective system to study the problem of nutrient deficiency.

Molybdenum (Mo) is a transition metal that has long been known as an essential micronutrient in animals and plants (Leimkühler et al. 2011). Mo has a cofactor (Moco) and this cofactor-bound form is the catalytically active form that is used by the majority of Mo-dependent enzymes across phyla (Schwarz 2005; Mendel 2013). Moco deficiency (MoCD) is a rare genetic disorder caused by loss of Moco biosynthesis, a conserved multistep process, and results in the subsequent loss of all Mo-dependent enzyme activities; patients suffer from severe neurological damage and die prematurely (Fig. 1; Mendel

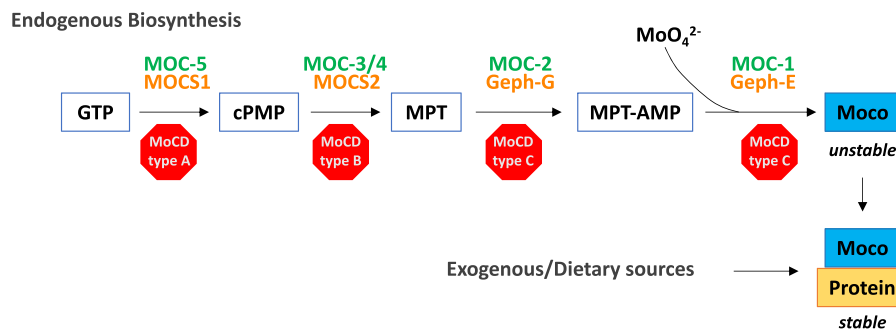
2013). While it is clear that the patients need additional Moco, the deficiency cannot be directly treated with Moco supplementation due to the instability of the Moco complex, lasting only a few minutes in neutral, aqueous environments when not bound to a protein (Schwarz 2005). Free Moco is readily oxidized and is therefore expected to be bound to protein, and thereby stabilized, immediately after synthesis.

Recent efforts using Moco precursor supplementation or gene therapy have yielded promising results for MoCD patients. Most human MoCD patients have a defect in the first step of Moco biosynthesis caused by mutations in the *MOCS1* gene, referred to as MoCD type A. Researchers supplemented MoCD type A mutants with the *MOCS1* product cPMP (isolated from *E. coli*), a stable Moco precursor, and observed significant improvements in both mouse models and human patient studies (Schwarz et al. 2004; Veldman et al. 2010; Schwahn et al. 2015). Gene therapy has also been shown to be effective to rescue Moco biosynthesis in mouse models for both type A and type B MoCD (Kügler et al. 2007; Reiss 2019). If a stable, downstream Moco supplement could be identified, it would be an ideal alternative treatment for all types of MoCD. The work by Warnhoff et al. (2021) presents a promising treatment for a MoCD deficiency model in *C. elegans*.

A previous study from the same laboratory demonstrated that *C. elegans* can use exogenous, dietary Moco from *E. coli* in addition to synthesizing endogenous Moco (Warnhoff and Ruvkun 2019), which provided the opportunity to search for a particular form of exogenous Moco that is effectively used by worms. They found that protein-bound Moco, recombinantly expressed and isolated from *E. coli*, red bread mold *Neurospora crassa*, or green algae *Volvox carterii*, or purified from cow's milk, was sufficient to support the growth of *moc-1(-)/Geph-E* mutant worms grown on Moco-deficient *E. coli*. They showed that the benefit of protein-bound Moco to worms is likely executed through

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**Figure 1.** Diagram of Moco biosynthesis and stabilization with protein binding. The Moco biosynthesis pathway is conserved in eukaryotes, including *C. elegans* and humans. Loss of Moco biosynthetic enzymes results in the indicated types of Moco deficiency (MoCD). Free, unbound Moco is unstable but is stabilized when bound to proteins (e.g. sulfite oxidase). Warnhoff et al. (2021) showed that dietary supplementation with protein-bound Moco was an effective treatment for Moco deficiency in *C. elegans*. (MOCS1/MOCS2) Molybdenum cofactor synthesis enzymes, (Geph-G) gephyrin G-domain, (Geph-E) gephyrin E-domain (human; orange text), (MOC-1,2,3,4,5) molybdenum cofactor biosynthesis enzymes (*C. elegans*; green text), (GTP) guanosine triphosphate, (cPMP) cyclic pyranopterin monophosphate, (MPT) molybdopterin, (MPT-AMP) adenylated molybdopterin, (Moco) molybdenum cofactor.

an *E. coli*-independent mechanism that also did not rely on the endogenous Moco biosynthesis pathway in worms. The investigators further showed that the supplemented protein-bound Moco was also sufficient to rescue the defects caused by a partial loss-of-function mutation in the *suox-1* gene, encoding sulfite oxidase. Sulfite oxidase is a conserved Mo-containing enzyme that is essential for both *C. elegans* and human development (Leimkühler et al. 2011; Warnhoff and Ruvkun 2019).

This work demonstrated that protein-bound Moco is both stable and bioavailable to *C. elegans* and provides a therapeutically relevant demonstration of previous observations that Moco from plant, animal, or bacterial sources can be used interchangeably and that Moco is stabilized when bound to protein (Schwarz 2005). Just as *E. coli*-derived cPMP was successfully used by both mice and humans in supplementation tests (Schwarz et al. 2004; Veldman et al. 2010; Schwahn et al. 2015), Warnhoff et al. (2021) demonstrated that *C. elegans* can use protein-bound Moco from a spectrum of exogenous origins. These findings raise a delightful possibility that protein-bound Moco from diverse sources may serve as an effective supplement for all types of MoCD patients.

More mechanistic questions remain to be addressed in future studies. It is still not understood how Moco or protein-bound Moco is transported into intestinal cells and what becomes of it in the cellular environment. Does the complex remain intact or is it broken down for utilization? Is it transported to other tissues, and by what mechanism? The investigators' previous work (Warnhoff and Ruvkun 2019) showed that MOC-1 and SUOX-1 expression in the worm hypodermis was sufficient to rescue the *moc-1(-)* mutant, suggesting that the hypodermis (analogous to the epidermis) is the tissue of action for Moco/SUOX-1 in *C. elegans*. Since dietary protein-bound Moco also rescues the *moc-1(-)* mutant, it may be necessary for that complex to be transported from the intestinal lumen to the hypodermis by a yet unknown mechanism. Alternatively, it is also possible that the protein-bound

Moco complex is metabolized after entering the intestinal cell for utilization by the worm. Further efforts in multicellular models and human patients will reveal the benefit and impact of this finding in *C. elegans*.

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