

MOLECULAR ENDOCRINOLOGY

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COVER: Molecular model of human 21-hydroxylase, CYP21, with heme and steroid-substrate. All disease-causing mutations are color-coded corresponding to the different degrees of clinical severity (SW, SV, and NC) seen in congenital adrenal hyperplasia, CAH. A structural explanation for corresponding phenotype was found for all but two mutants, illustrating the possibility to predict degree of enzyme impairment caused by missense mutations using bioinformatic means. From the article by Robins *et al.*, in this issue, pp. 2946–2964.

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The editors would also like to acknowledge the design for our page ender which appears at the end of each article. The symbol represents interaction between ligand, receptor, and DNA. It evolved from an original concept designed by Lois B. Thompson for MOLECULAR ENDOCRINOLOGY.