

Quantum mechanical considerations on the mechanism of the P450 conversion of 11-deoxycorticosterone to Aldosterone

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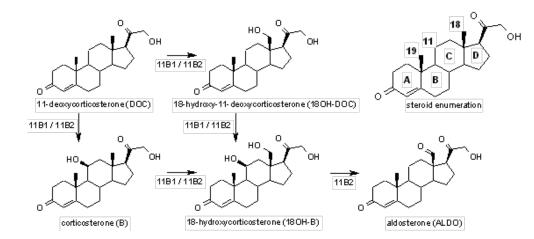
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10th European ISSX Meeting (May 18-21, 2008): Quantum Mechanical Considerations On The Mechanism Of The P450 Conversion Of 11-deoxycorticosterone To Aldosterone



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The final steps of the biosynthesis of the steroid aldosterone involve C_{11b}- and C₁₈-hydroxylations by cytochrome P450 11B1 and 11B2 (figure), and C18-aldehyde production by 11B2 [1]. Generally regioselective substrate and intermediate conversions are driven by enzyme-ligand sterical and energetic factors. To evaluate in how far energetic considerations play a role in determining the sequential conversions of DOC, we have analysed multiple steroid conformations using Quantum Mechanics calculations. Therefore, we have conducted: (1) conformational analyses for the prediction of the optimal steroid configurations and (2) Fukui index calculations to determine the inherent reactivity of the individual carbon atoms for prediction of carbon oxidation preference. Molecule geometries were optimised using the QM package Gaussian03. Mulliken population analysis was used to determine the electron density distribution. Accurate computation of the ionic state required for the Fukui index of a nucleophilic attack, was conducted using the B3LYP functional with the 6-31+g basis set. For the lowest energy conformations of all four steroids, the Fukui index indicated that C19 and in some cases C12 were more reactive than C11 and C18. We conclude that the 11B1 and 11B2 active site conformations must prohibit C12 and C19 interactions with the active site heme. For DOC, the Fukui index of C₁₁ (0.088) is higher than that of C₁₈ (0.043), indicating that B is more likely to be produced than 180H-DOC. The resulting conformation of B portrays an unfavourable C₁₈ Fukui index (-0.016), but the resulting 180H-DOC conformation possesses a favourable C₁₁ Fukui index (0.020). This can indicate two things: (1) either B is the most abundant product from DOC hydroxylation whereas 180H-B and aldosterone are formed via the less abundant 180H-DOC or (2) active site-ligand interactions result in a conformational change in the hydroxyacetyl group of B to allow further conversion to 180H-B and aldosterone. In a next step the complexity of the steroids was reduced by neglecting the A and B rings. Conformational space analyses of the simplified steroids and their calculated Fukui indices suggest that the hydroxyacetyl group of 180H-B requires an anti-periplanar conformation for conversion into aldosterone as well as an internal hydrogen bond with the C₁₈ hydroxyl group.



[1] A Fisher, EC Friel, R Bernhardt, C Gomez-Sanchez, JM Connell, R Frazier, E Davies, Effects of 18-hydroxylated steroids on corticosteroid production by human aldosterone synthase and 11beta-hydroxylase, J Clin Endocrinol Metab (2001), 86 (9), p4326-p4329

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