

Poster Presentation

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Crystallographic evidence for oxidative Met→Asp conversion in human hemoglobin

J. Soman¹, M. Strader², W. Hicks², T. Kassa², E. Singleton¹, J. Olson¹, M. Weiss³, T. Mollan², M. Wilson⁴, A. Alayash²

¹Rice University, Department of Biochemistry and Cell Biology, Houston, USA, ²Food and Drug Administration, Laboratory of Biochemistry and Vascular Biology, Bethesda, USA, ³Childrens Hospital of Philadelphia, Philadelphia, USA, ⁴University of Essex, Department of Biological Sciences, UK

The mutants HbA Bristol-Alesha (β V(E11)67M) and HbF Toms River (γ V(E11)67M) [1,2] are examples of a 'silent' posttranslational modification in which the side chain of the substituted amino acid is chemically modified (Met→Asp) resulting in a disparity between the DNA and protein sequences. In both cases the patients' hemolysate contained both V67M and V67D isoforms. But in the analogous α subunit mutant, Hb Evans α V(E11)62M, the conversion to Asp was not identified and DNA sequencing confirmed the Met replacement [3]. Our crystal structures of the three (ferrous) CO-bound recombinant V(E11)M mutants show the MetE11 side chain in similar conformations. But the air-oxidized β mutant crystals clearly showed a 'bifurcated' and smaller electron density pattern for the E11 side chain, indicating the appearance of Asp. Also, the ligand electron-density at the iron atom in the oxidized β subunit appears to be an oxoferryl Fe⁴⁺=O rather than a Fe³⁺+OH₂ ferric complex. In contrast, there was little change in the electron density for α MetE11 in oxidized α V62M crystals. The ligand in the ferric α subunit is clearly a coordinated water molecule. But again, a ferryl Fe⁴⁺=O complex appears to occur in the wild-type β subunit. This strongly suggest that β subunits have a greater propensity to form highly reactive ferryl species, and that the ferryl species play a role in the Met→Asp conversion. Our autoxidation and proteomics studies showed that although all three recombinant VE11M mutants had similar, high rates of autooxidation and a strong H₂O₂ dose dependence on sulfoxide and sulfone formation, no Asp formation was detected in α subunits whereas MetE11 is converted to Asp to levels as high as 15% in vitro in β and γ subunits. We propose that the Met→Asp conversion specifically involves H₂O₂ mediated oxidation of the ferrous heme to an oxoferryl state, and because the transient ferryl intermediates are much less stable in the α subunits, there is no oxidative conversion.

[1] D.C. Rees, J. Rochette, C. Schofield et al., *Blood*, 1996, 88, 341-348, [2] M.A. Crowley, T.L. Mollan, O.Y. Abdulmalik et al., *N Engl J Med*, 2011, 364, 1837-1843, [3] M.I. Zanotto, K. Calvo, G. Schwartzman et al., *Archivos argentinos de pediatria*, 2010, 108(6), 130-133

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