## **Chitayat Syndrome (ERF Variant)**

https://pubmed.ncbi.nlm.nih.gov/27738187/

Methods:
Through ongoing collaboration, we had collected patients with strikingly-similar phenotype.
Trio-based exome sequencing was first performed in Patient 2 through Deciphering Developmental
Disorders study. Proband-only exome sequencing had previously been independently performed in
Patient 4. Following identification of a candidate gene variant in Patient 2, the same variant was
subsequently confirmed from exome data in Patient 4. Sanger sequencing was used to validate this
variant in Patients 1, 3; confirm paternal inheritance in Patient 5.
Results:
A recurrent, novel variant NM_006494.2:c.266A>G p.(Tyr89Cys) in
ERF
was identified in five affected individuals: de novo (patient 1, 2 and 3) and inherited from an affected
father (patient 4 and 5). p.Tyr89Cys is an aromatic polar neutral to polar neutral amino acid

substitution, at a highly conserved position and lies within the functionally important ETS-domain of
the protein. The recurrent
ERF
c.266A>C p.(Tyr89Cys) variant causes Chitayat syndrome.
Discussion:
ERF
variants have previously been associated with complex craniosynostosis. In contrast, none of the
patients with the c.266A>G p.(Tyr89Cys) variant have craniosynostosis.
Conclusions:

We report the molecular aetiology of Chitayat syndrome and discuss potential mechanisms for this distinctive phenotype associated with the p.Tyr89Cys substitution in

ERF

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