



Title	Understanding the molecular pathogenesis of SOX9 Y440X campomelic dysplasia
Author(s)	Cheah, KSE; Au, T; Szeto, IYY; Wynn, S; Chan, YS; Cheung, K; Chan, WY; Lovell-Badge, R; Chan, D; Fritzschn, B
Citation	The 2010 Academic Symposium on Developmental Studies in Health and Diseases, The Chinese University of Hong Kong, Hong Kong, 18-21 October 2010.
Issued Date	2010
URL	http://hdl.handle.net/10722/140080
Rights	Creative Commons: Attribution 3.0 Hong Kong License

Understanding the molecular pathogenesis of *SOX9* Y440X campomelic dysplasia

Kathryn SE Cheah^{1,2}, Tiffany Au¹, Irene Y.Y. Szeto¹, Sarah Wynn¹, Y.S. Chan³, Kenneth Cheung^{4,2}, Wood Yee Chan⁵, Robin Lovell-Badge⁶, Danny Chan^{1,2}, Bernd Fritzscht⁷

¹Department of Biochemistry, ²Centre for Reproduction, Development & Growth, ³Department of Physiology, ⁴Department of Orthopaedic Surgery, The University of Hong Kong, Hong Kong, China, ⁵ Department of Anatomy, Chinese University of Hong Kong, Hong Kong China, ⁶Division of Developmental Genetics, MRC National Institute for Medical Research, London, UK, ⁷Department of Biological Sciences, University of Iowa, Iowa City, USA.

Human *SOX9* mutations cause the skeletal malformation syndrome campomelic dysplasia (CD). Complete inactivation of the *Sox9* gene in mice results in failure of cartilage formation. Studies in zebrafish and *Xenopus* suggest that *Sox9* may be crucial for specification of the otic placode. In mice, loss of *Sox9* results in failure of otic placode invagination. Heterozygous mutations in human *SOX9* result in conductive and sensorineural deafness in some CD patients, implying a later morphogenetic role but phenotypic details are limited. *Sox9*^{-/-} null mice die before morphogenesis of the inner ear is complete, precluding investigation of the role of *Sox9* later in ear development. Because all the *SOX9* mutations are heterozygous and appear to cause loss of function, the CD phenotype has been attributed to haploinsufficiency of *SOX9*. However *SOX9* proteins containing an intact HMG box and a truncated activation domain may act dominant negatively by competition with the wild-type for binding to target genes and interfere with interaction with partner factors via the transactivation domain. To assess whether such mutations in *SOX9* may act via a dominant interference mechanism we generated transgenic and conditional knock'in mice expressing a mouse equivalent of a CD mutation, a Y440X nonsense mutation causing premature termination within the trans-activation domain of *SOX9* (*Sox9*^{Y440X}). We compared the phenotypic impact of the *Sox9*^{Y440X} mutation with a *Sox9* null mutation. These studies point to an essential role for *Sox9* in inner ear and intervertebral disc development and context dependent mechanisms for the Y440X nonsense mutation.