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Genes underlying delayed puberty

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20 21	Abstract
22	The genetic control of pubertal timing has been a field of active investigation for
23	the last decade, but remains a fascinating and mysterious conundrum. Self-
23 24	limited delayed puberty (DP), also known as constitutional delay of growth and
25	puberty, represents the extreme end of normal pubertal timing, and is the
26	commonest cause of DP in both boys and girls. Familial self-limited DP has a
27	clear genetic basis. It is a highly heritable condition, which often segregates in an
28	autosomal dominant pattern (with or without complete penetrance) in the
29	majority of families. However, the underlying neuroendocrine pathophysiology
30	and genetic regulation has been largely unknown. Very recently novel gene
31	discoveries from next generation sequencing studies have provided insights into
32	the genetic mutations that lead to familial DP. Further understanding has come
33	from sequencing genes known to cause GnRH deficiency, next generation
34	sequencing studies in patients with early puberty, and from large-scale genome
35	wide association studies in the general population. Results of these studies
36	suggest that the genetic basis of DP is likely to be highly heterogeneous.
37	Abnormalities of GnRH neuronal development, function, and its downstream
38	pathways, metabolic and energy homeostatic derangements, and transcriptional
39	regulation of the hypothalamic-pituitary-gonadal axis may all lead to DP. This
40	variety of different pathogenic mechanisms affecting the release of the puberty
41	'brake' may take place in several age windows between fetal life and puberty.
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