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Genetics of Hyperparathyroidism, Including Parathyroid Cancer

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KEYWORDS

- Tumor suppressor
 Oncogene
 Multiple endocrine neoplasia
 MEN1
 MEN2A
- CDC73 CCND1 RET

KEY POINTS

- Primary hyperparathyroidism, caused by parathyroid tumors, is mostly sporadic.
- The molecular genetic investigation of rare syndromic forms of hyperparathyroidism has nevertheless led to significant advances in the understanding of both familial and sporadic parathyroid neoplasia.
- Both oncogenes and tumor suppressors have been implicated in the cause of parathyroid tumors.
- The discovery of novel parathyroid tumor susceptibility genes is likely to result from the application of next-generation sequencing methods to the analysis of sporadic parathyroid tumors and nonsyndromic familial cases of hyperparathyroidism.

INTRODUCTION

Primary hyperparathyroidism (HPT) is a disorder of mineral metabolism, typically manifesting in hypercalcemia, that results from the excessive secretion of parathyroid hormone from 1 or more neoplastic parathyroid glands. Although HPT is mostly sporadic, familial forms of HPT represent some 2% to 5% of total cases, most of which are caused by germline mutation of known HPT-susceptibility genes (Table 1). Investigation of the molecular genetics underlying these rare familial syndromes has yielded significant insight into the pathophysiology of both sporadic and familial parathyroid neoplasms. Signaling involving the G

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Gene	Protein Encoded	Associated Hyperparathyroid Syndrome: Main Syndromic Manifestations	Features of Syndromic Parathyroid Tumors	Defect in Sporadic Parathyroid Tumors
MEN1	Menin	MEN1: anterior pituitary, parathyroid, enteropancreatic, foregut carcinoid tumors	Multiple, asymmetric tumors typical (>99% benign)	Inactivation in ~25%–35% of benign tumors; mutation exceedingly rare in cancer
CDC73/HRPT2	Parafibromin	HPT-jaw tumor syndrome: fibro- osseous jaw, parathyroid, uterine tumors; renal cysts	Single tumor common (~15% malignant)	Inactivation in ~70% of cancers; mutation rare in sporadic adenomas
CDKN1B	P27(Kip1)	MEN4: anterior pituitary, other involvement varies	Single to multiple glands (benign in reports to date); can be recurrent	Loss-of-function mutation in ~5% of sporadic adenomas; including germline mutation in sporadic presentation
CASR	Calcium-sensing receptor	FHH1 with heterozygous inactivation; NSHPT with homozygous inactivation	FHH1: near-normal size and surgical pathology; altered serum calcium set-point for PTH release NSHPT: marked enlargement of multiple glands by polyclonal (nonneoplastic) mechanism	Decreased expression common; mutation exceedingly rare
GNA11	G protein α11 subunit	FHH2	ND	ND
AP2S1	Adaptor protein-2 sigma subunit	FHH3: hypercalcemia more severe than in FHH1	ND	ND
RET	c-Ret	MEN2A: medullary thyroid cancer, pheochromocytoma, parathyroid tumors	Single tumor common (>99% benign)	Mutation exceedingly rare
CCND1/PRAD1	Cyclin D1	NA	NA	Overexpression results from DNA rearrangement involving PTH gene

Abbreviations: MEN1, multiple endocrine neoplasia type 1; MEN4, multiple endocrine neoplasia type 4; NA, not applicable; ND, not determined (lack of relevant published studies); NSHPT, neonatal severe hyperparathyroidism; PTH, parathyroid hormone.

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