Analysis of adenomatous polyposis coli gene in thyroid tumours

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Summary Familial adenomatous polyposis (FAP) is known to be associated with neoplasia of various tissues, including thyroid carcinoma. Germline mutations of the tumour-suppressor gene APC, responsible for the predisposition to FAP, may therefore be involved in the pathogenesis of these tumours. In this report the structure of the APC gene has been investigated in 26 thyroid tumours, at different stages of dedifferentiation, that were surgically excised from patients with a negative history of FAP. Approximately 35% of the APC gene coding region, where most of the mutations are clustered, has been analysed by a combination of single-strand conformation polymorphism and direct sequencing. No significant alterations could be demonstrated in any sample examined. It is concluded that, at least in patients not affected by FAP, APC gene abnormalities do not seem to play a relevant role in the pathogenesis of thyroid carcinoma.

In thyroid tumorigenesis the only evidence in favour of the alteration of tumour-suppressor genes concerns the occurrence of p53 mutations, which appear to be restricted to poorly differentiated and undifferentiated carcinomas of the thyroid gland (Ito et al., 1992; Nakamura et al., 1992; Don-ghi et al., 1993; Fagin et al., 1993). Germline mutations of the tumour-suppressor gene APC, located on the long arm of chromosome 5, are responsible for the predisposition to familial adenomatous polyposis (FAP). FAP is an autosomal dominant disorder characterised by the development, in young adults, of hundreds to thousands of adenomatous colonic polyps. FAP may be associated with osteomas, epidermoid cysts, fibromas and desmoid tumours, as well as with tumours of other tissues, including the thyroid, outlining the clinical picture of Gardner's syndrome (Gardner & Richards, 1953; Järvinen & Sipponen, 1986; Jagelman et al., 1988). Somatic mutations of the APC gene are also considered to be an early event in the development of sporadic gastrointestinal tumours (Powell et al., 1992).

The association between FAP and thyroid carcinoma was first observed in 1949 (Crail, 1949), and since then several cases have been reported in the literature (Delamarre *et al.*, 1988; Ono *et al.*, 1991; Bell & Mazzaferri, 1993). The importance of this association has not been well established, but the development of a thyroid carcinoma in two sisters affected by FAP (Camiel *et al.*, 1968), and the high incidence of thyroid carcinoma observed in two different large series of patients with FAP (Plail *et al.*, 1987; Bülow *et al.*, 1988), has suggested that the concurrence of the two diseases may not have arisen by chance.

In addition, APC has been found to be expressed in normal as well as in neoplastic human thyroid tissue, in which multiple forms of specific RNA transcripts have been detected (Horii *et al.*, 1993; Zeki *et al.*, 1993). Considering all the above observations, it may be hypothesised that alterations in APC are likely candidates for a pathogenetic role in thyroid tumorigenesis.

To test this hypothesis, a series of 26 thyroid tumours of different histological grades were analysed, by using a combination of single-strand conformation polymorphism (SSCP) and direct sequencing, in the search for structural alterations in APC.

Patients and methods

Twenty-six female patients, with a mean age of 44.8 years and a negative clinical history for colorectal as well as for other gastrointestinal neoplasms, were treated by surgery for the presence of thyroid neoplasia. Histological diagnosis revealed the presence of a thyroid carcinoma in 18 cases (13 papillary, three follicular, two anaplastic) and a follicular adenoma in the remaining eight patients. After surgical removal, the tumoral and the corresponding extratumoral tissues were quickly frozen in liquid nitrogen and stored at $- 80^{\circ}C$.

Genomic DNA was extracted from all thyroid samples by the standard SDS-proteinase K digestion followed by phenol-chloroform extraction. Exons 7-10 and portions of exon 15 (codons 653-751, 998-1,141 and 1,260-1,547), which represent approximately 35% of the *APC* coding sequence and in which about 90% of all *APC* gene mutations are clustered (Miyoshi *et al.*, 1992; Nakatsuru *et al.*, 1991), were amplified by the polymerase chain reaction (PCR) using primer pairs and incubation conditions previously described (Miyoshi *et al.*, 1992).

One microlitre of a 1:1,000 dilution of each PCR product was further amplified in the presence of $1 \mu Ci$ of $[^{32}P]dCTP$ for a total of 28 cycles. Labelled PCR products were diluted (1:10) in a solution containing 95% deionised formamide, 0.1% bromophenol blue and 0.1% xylene cyanol, and denatured at 95°C for 5 min. SSCP analysis was performed by electrophoresing denatured samples through a nondenaturing 6% polyacrylamide gel at 10 W constant power at either 4°C or 24°C in the presence of 5% glycerol (Cama *et al.*, 1993).

DNA samples showing an abnormal SSCP electrophoretic profile were further analysed by direct sequencing of PCR products using the dideoxy chain-termination method (Kadowaki *et al.*, 1990).

Results and discussion

SSCP analysis of APC sequences showed an altered band pattern only in one case of multifocal papillary carcinoma (Figure 1, lane 7). This alteration was observed in the exon 15 region corresponding to codons 1,389-1,547 and was also present in a different DNA sample obtained from a distinct focus of the same neoplastic lesion (Figure 1, lane 27). No alterations in the APC sequence were detected in the thyroid extratumoral tissue obtained from the same patient (Figure 1, lane 28) or in non-tumoral specimens from any of the other

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patients (data not shown). Direct sequencing of the two samples demonstrated, in both cases, the presence of a guanine to adenine transition at nucleotide position 4,497 of the *APC* coding sequence, corresponding to a CG dinucleotide. This nucleotide change, however, represented a silent mutation since it did not cause any amino acid change in the primary structure of the APC protein (data not shown). The

Figure 1 SSCP analysis of APC gene exon 15. A portion of exon 15 of the APC gene (codons 1,389–1,547) was amplified by PCR and scanned by SSCP for the presence of mutations in 13 papillary (lanes 1–13), three follicular (lanes 14–16) and two anaplastic (lanes 17 and 18) carcinomas and in eight follicular adenomas (lanes 19–26). In lane 27 is shown the SSCP profile of DNA extracted from a different focus of the same papillary tumour shown in lane 7. The altered SSCP pattern (lanes 7 and 27) reflected a conservative nucleotide substitution that was not present in the extratumoral tissue from the same thyroid gland (lane 28) or in two other non-neoplastic samples (lanes 29–30).

remaining samples did not show alterations of the APC gene in any of the mutation cluster regions analysed (Figure 1).

Since 1949 the concurrence of FAP and thyroid carcinoma has been observed in 48 cases (Table I). Most of the studies published have pointed to the fact that thyroid carcinoma occurs with an unexpectedly high frequency in patients affected by FAP. Recently, statistical analysis of data obtained from an English Polyposis Register has indicated that the risk for a young female affected by FAP of developing thyroid carcinoma, particularly of the papillary type, is about 160-fold higher than expected (Plail et al., 1987). Similar conclusions have been obtained in a Danish population of FAP patients in which the risk of developing thyroid carcinoma has also been estimated to be 100-fold greater than in the general population (Bülow et al., 1988). Thyroid carcinoma associated with FAP has been more frequently found in young female patients (F/M = 6.3:1) than sporadic thyroid carcinoma (F/M = 2-3:1). The thyroid neoplasia has usually been discovered within 1-7 years after FAP was diagnosed. Papillary carcinoma represented the predominant histotype (8.5%) with a 2-fold higher than expected frequency of multifocal lesions (Table I).

The patients with thyroid carcinoma examined in the present study were all females, did not show evidence of an altered bowel function and always had a negative family

 Table I
 Association between familial adenomatous polyposis (FAP) and thyroid carcinoma (TC)

	Reference		Age		Histological type of
Case		Sex	TC	FAP	thyroid carcinoma
1	Crail (1949)	М	24	24	Papillary
2	Ogata et al. (1964)	М	31	_	Adenocarcinoma
3	Ravnham & Louw (1966)	F	20	20	Unknown (multifocal)
4	Smith (1968)	M	29	39	Papillary (multifocal)
5	5	_	_	-	Unknown
6		_	_	_	Unknown
7	Camiel et al. (1968)	F	10	28	Papillary
8	Camer er al. (1966)	F	20	20	Papillary (multifocal)
0	Smith & Kern (1073)	F	10	29	Papillary (multifocal)
10	Alm & Licznersci (1973)	F	19	20	Linknown
10	Ann & Elezherser (1973)	L. L.	_	_	Unknown
11		Г	-	_	
12		-	-	-	Unknown
13	M (1) 0 0 (1077)	-	~~~~	-	Unknown
14	Mathias & Smith (1977)	F	< 30	-	Papillary
15	Keshgegian & Enterline (1978)	F	21	14	Papillary (multifocal)
16	Takahashi <i>et al.</i> (1976)	M	58	-	Papillary
17	lida <i>et al.</i> (1977)	Μ	26	-	Unknown
18		F	26	-	Unknown
19	Ushio <i>et al.</i> (1977)	Μ	27	-	Unknown
20	Harada <i>et al.</i> (1977)	F	22	-	Papillary
21	Okamura <i>et al.</i> (1979)	F	29	-	Papillary
22	Hamilton <i>et al.</i> (1979)	F	18	17	Papillary
23	Miura et al. (1980)	F	27	-	Papillary
24	Lee & Mackinnon (1981)	F	23	32	Papillary (multifocal)
25	Delamarre et al. (1982)	F	21	27	Follicular
26	Thompson et al. (1983)	F	24	22	Papillary (multifocal)
27	Schneider et al. (1983)	F	37	33	Papillary
28	Masuvama et al. (1986)	F	26	_	Papillary
29	Plail et al. (1987)	F	22	21	Papillary (multifocal)
30		Ē	26	19	Papillary (multifocal)
31		F	34	31	Papillary
32		F	23	27	Papillary (multifocal)
32		F	20	20	Linknown (multifocal)
34		F	16	20	Unknown (multilocal)
25		F	24	17	Unknown
36	Differ (1088)	Г	24	17	Ealliantan (multifa cal)
27	Filler (1988)	Г	20	24	Poincular (multilocal)
29		г Г	20	-	Papillary
20	Delements at $al (1099)$	Г	33	1	
37 40	Defamarie et al. (1988)	Г	29	10	Papillary (multifocal)
40	B #1	F	26	21	Papillary (multifocal)
41	Dulow et al. (1988)	F	19	17	Papillary
42 42		F	_	_	Papillary
43		F	40	26	Papillary
44	van Erpecum <i>et al.</i> (1988)	F	34	31	Papillary
45	Herrera <i>et al.</i> (1989)	F	27	23	Follicular
46	•	F	-	-	Papillary
47	Ono et al. (1991)	F	50	50	Follicular
48	Bell & Mazzaferri (1993)	F	24	24	Papillary

history for the presence of gastrointestinal diseases. Only one among the 13 cases of papillary thyroid carcinoma was multifocal. In this case, the occurrence of the same mutation in DNA extracted from two distinct tumoral foci, localized in opposite lobes of the same thyroid gland, but not in the corresponding extratumoral tissue (Figure 1, lane 28), suggested a common clonal origin of the thyroid tumour.

The frequent association between FAP and thyroid carcinoma suggests the involvement of common mechanisms in the pathogenesis of these two diseases. However, the absence of germline and somatic APC gene defects in our series of 26 thyroid neoplasms, at different stages of dedifferentiation, suggests that alterations of this tumour-suppressor gene do not represent a frequent event in thyroid tumorigenesis.

The small number of cases analysed does not allow any statistically significant conclusion to be drawn, since a low incidence of APC mutations cannot be ruled out. However, our results are in agreement with recent unpublished observations (Varesco *et al.*, 1993; Zeki *et al.*, 1993). SSCP analysis of a population of 73 benign and malignant thyroid tumours, performed on a 1,200 bp stretch of exon 15, also failed to

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detect any mutation in the APC gene. Only a nucleotide insertion, leading to a premature stop codon, was identified in one out of four thyroid carcinoma cell lines examined, namely the highly undifferentiated ARO carcinoma cells (Zeki *et al.*, 1993). APC gene mutations were also absent in a small group of thyroid tumours in which most of the APC gene coding region was investigated (Varesco *et al.*, 1993).

In conclusion, our own results and those of others, gathered from a total of more than 100 thyroid tumours, suggest that APC mutations do not play a pathogenetic role in thyroid tumorigenesis, at least in patients not affected by FAP. However, true estimates of the incidence of APC alterations in thyroid tumours will require the collection of molecular information from a larger number of cases.

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