

## INVITED REVIEW

## Sodium/iodide symporter: a key transport system in thyroid cancer cell metabolism

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### Abstract

The recent cloning of the gene encoding the sodium/iodide symporter (NIS) has enabled better characterization of the molecular mechanisms underlying iodide transport, thus opening the way to clarifying its role in thyroid diseases. Several studies, at both the mRNA and the protein expression levels, have demonstrated that TSH, the primary regulator of iodide uptake, upregulates NIS gene expression and NIS protein abundance, both *in vitro* and *in vivo*. However, other factors, including iodide, retinoic acid, transforming growth factor- $\beta$ , interleukin-1 $\alpha$  and tumour necrosis factor  $\alpha$ , may participate in the regulation of NIS expression. Investigation of NIS mRNA expression in different thyroid tissues has revealed increased levels of expression in Graves' disease and toxic adenomas, whereas a reduction or loss of NIS transcript was detected in differentiated thyroid carcinomas, despite the expression of other specific thyroid markers. NIS mRNA was also detected in non-thyroid tissues able to concentrate radioiodine, including salivary glands, stomach, thymus and breast.

The production of specific antibodies against the NIS has facilitated study of the expression of the symporter protein. Despite of the presence of high levels of human (h)NIS mRNA, normal thyroid glands exhibit a heterogeneous expression of NIS protein, limited to the basolateral membrane of the thyrocytes. By immunohistochemistry, staining of hNIS protein was stronger in Graves' and toxic adenomas and reduced in thyroid carcinomas.

Measurement of iodide uptake by thyroid cancer cells is the cornerstone of the follow-up and treatment of patients with thyroid cancer. However, radioiodide uptake is found only in about 67% of patients with persistent or recurrent disease. Several studies have demonstrated a decrease in or a loss of NIS expression in primary human thyroid carcinomas, and immunohistochemical studies have confirmed this considerably decreased expression of the NIS protein in thyroid cancer tissues, suggesting that the low expression of NIS may represent an early abnormality in the pathway of thyroid cell transformation, rather than being a consequence of cancer progression.

The relationship between radioiodine uptake and NIS expression by thyroid cancer cells require further study. New strategies, based on manipulation of NIS expression, to obtain NIS gene reactivation or for use as NIS gene therapy in the treatment of radiosensitive cancer, are also being investigated.

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## Role of iodine in thyroid physiology and pathophysiology

### *Iodine in thyroid physiology*

Iodine represents an essential element in thyroid physiology, being a critical component of thyroxine and tri-iodothyronine molecules, and a key regulator of thyroid gland function. Thus dietary iodide supply influences the functional activity of the thyroid gland, iodine deficiency being the main cause of endemic goiter. Furthermore, iodine directly modulates thyroid sensitivity to thyroid-stimulating hormone (TSH), and

intrathyroidal iodine has a generally inhibitory effect on thyroid function, independently of the TSH concentration. This autoregulatory effect of iodine encompasses many aspects of thyroid function, including iodide transport mechanism, iodide organification, thyroid hormone synthesis and secretion, and thyroid intermediary metabolism.

The first step of iodine thyroid metabolism is represented by the thyroid trapping and concentration of iodide from the blood, which are achieved by an active, energy-dependent transport process across the basolateral plasma membrane of the thyrocytes (Fig. 1). The recent cloning (1, 2) of the gene encoding the protein responsible

for this process, the sodium/iodide symporter (NIS), enabled better characterization of the molecular mechanisms underlying the iodide transport, thus opening the way to clarify its role in thyroid diseases (Table 1).

Radioactive isotopes of iodine have been used first as tracers of thyroid function and, subsequently, for treatment of hyperthyroidism and benign thyroid diseases. The evidence that iodine transport is present in thyroid neoplastic tissues has been the basis for the use of radioiodine in the diagnosis and treatment of patients with thyroid cancer.

In this review, we will discuss the more recent knowledge concerning the NIS system and its regulation in neoplastic thyroid cells, attempting to correlate the molecular mechanisms underlying the defective iodide transport in cancer cells with the clinical management of thyroid cancer patients. Prospectively, the availability of an efficient iodide transport system is a prerequisite

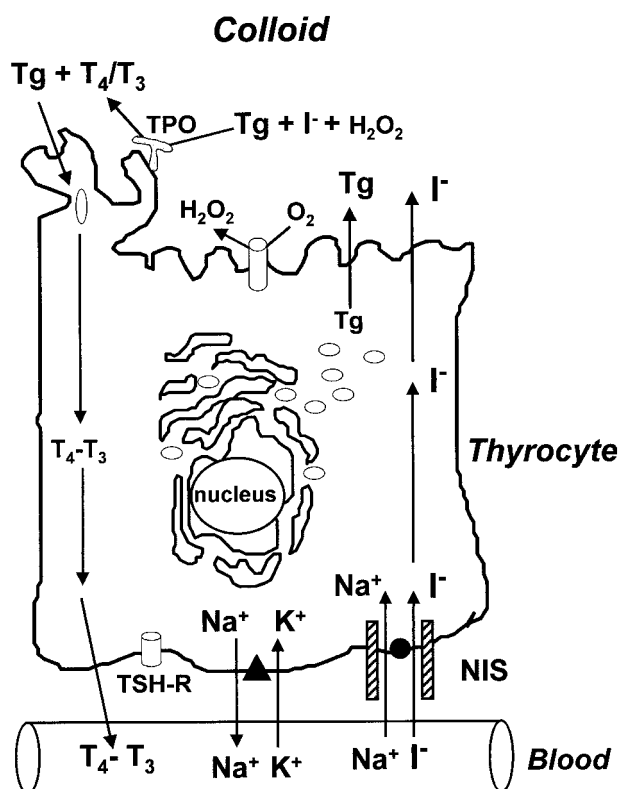
to obtaining optimal benefit from radioiodine treatment of thyroid diseases, including primary and metastatic papillary and follicular carcinomas and, by the use of *NIS* gene therapy, to extending this 'targeted radiotherapy' to radiosensitive non-thyroid carcinomas.

### NIS as carrier of iodide transport in thyroid cells

Iodide concentration occurs in the thyroid gland by 20- to 40-fold with respect to the anion concentration in plasma. Then, iodide ( $I^-$ ) is transported transcellularly from the basolateral to the apical membrane of the thyrocyte, where it is organified. NIS-catalyzed active accumulation of  $I^-$  from the interstitium into the cell is achieved against an electrochemical gradient, whereas apical  $I^-$  efflux (translocation of  $I^-$  to the follicular lumen) is passively achieved through a putative  $I^-$  channel. NIS couples the energy released by the inward 'downhill' translocation of  $Na^+$  down its electrochemical gradient, generated by  $Na^+/K^+$ -ATPase, to driving the simultaneous inward 'uphill' translocation of  $I^-$  against its electrochemical gradient (3). This process is competitively inhibited by thiocyanate, perchlorate and other anions (4). Differently from other tissues able to concentrate iodide, the thyroid gland alone is able to achieve prolonged  $I^-$  accumulation, as a result of the prompt organification of  $I^-$  in the tyrosyl residues of thyroglobulin, through a reaction catalyzed by thyroid peroxidase (TPO).

### Iodide concentration and utilization by thyroid cells, and its regulation

Many studies, using different experimental models, have elucidated the role of TSH and the activation of the cAMP pathway as the principal regulator of iodide uptake. Many other factors, including insulin, insulin-like growth factor (IGF)-I, epidermal growth factor and iodide itself also influence iodide uptake in the thyroid gland (3). The molecular characterization of the regulation of iodide transport has become possible since the cloning of the *NIS* gene. Indeed, in recent years, the regulation of *NIS* expression has been extensively studied, at both mRNA and protein levels, by using either *in vivo* or *in vitro* experimental models (Table 2) (5-14). In primary cultured human thyrocytes, TSH and forskolin upregulate *NIS* gene expression and *NIS* protein abundance, with 2.5- to 2.7-fold and 2.4- to 2.7-fold increases after 24 h and 48 h respectively (15). These findings are concordant with results obtained in FRTL-5 and PC Cl3 rat thyroid cells (9, 10). Re-addition of TSH to TSH-negative FRTL-5 cells significantly increased *NIS* mRNA (approximately sixfold maximal stimulation after 24 h), followed by a slower increase in *NIS* protein synthesis (9). Forskolin and dibutyryl-cAMP mimicked the stimulatory effect of TSH on both the  $I^-$  transport activity and mRNA levels.



**Figure 1** Schematic representation of the iodide uptake and biosynthetic pathway of thyroid hormones in thyrocytes.  $I^-$  is actively accumulated across the basolateral plasma membrane of the thyrocyte in a process catalyzed by the NIS. This transport is driven by the  $Na^+$  gradient generated under ATP hydrolysis by  $Na^+/K^+$ -ATPase. The iodide is passively translocated across the apical membrane of the thyrocyte into the colloid, where it is used to iodinate the thyroglobulin (Tg). This reaction, called ' $I^-$  organification', is catalyzed by TPO and requires  $H_2O_2$ . The iodinated Tg, containing thyroid hormones, is stored in the colloid. Thyroid hormones, thyroxine ( $T_4$ ) and tri-iodothyronine ( $T_3$ ) are released from Tg and secreted in the blood. All steps in the thyroid hormone biosynthetic pathway are stimulated by TSH.

**Table 1** Characteristics of iodide transport and human *NIS* gene and NIS protein.

<b>Iodide transport</b>
<i>K<sub>m</sub></i> ~ 36 μmol/l
Active, against electrochemical gradient: 30–40:1
Cotransport with sodium: 2 Na <sup>+</sup> /1 I <sup>-</sup>
Other substrates: ClO <sub>3</sub> <sup>-</sup> , SCN <sup>-</sup> , NO <sub>3</sub> <sup>-</sup> , Br <sup>-</sup> , BrO <sub>3</sub> <sup>-</sup> , IO <sub>4</sub> <sup>-</sup>
Specific inhibitors: ClO <sub>4</sub> <sup>-</sup>
<b><i>NIS</i> gene</b>
OMIM 601843
Gene map locus 19p13.2-p12
Gene structure: >20 Kb, 15 exons, 14 introns
mRNA: 3.9Kb, two forms, including a smaller one, lacking exon 5 and stopping close to 5' end of exon 6
Promoter-responsive element: AP-1, Sp1, CRE, TTF-1 like
<b>NIS protein</b>
643 aminoacid residues (84% homology with the <i>NIS</i> rat gene)
Three sites of glycosylation
~70–80 kDa
12 or 13 transmembrane domains

OMIM, Online Mendelian inheritance in man; AP-1, activating protein-1; Sp1, stimulating protein-1; CRE, cAMP response element; TTF-1, thyroid-specific transcription factor-1.

However, iodide uptake was enhanced 27-fold, with an increase evident after a 24-h stimulation, whereas the concentration of NIS protein increased only after 36 h, by ~2.6-fold. Taken together, these observations suggest that other mechanisms, including cAMP-dependent phosphorylation and coregulation by other activating or inhibiting proteins, may participate in the post-translational regulation of NIS activity, as proposed previously (16).

NIS upregulation by TSH has also been described *in vivo* (5). In hypophysectomized rats with markedly low TSH

concentrations, NIS protein expression is low, but increases promptly after a single injection of TSH. Accordingly, in rats maintained on an I<sup>-</sup>-deficient diet or treated with propylthiouracil, increased TSH circulating concentrations paralleled increased NIS protein expression. Also, in dog thyroid glands (6), the expression of NIS mRNA is upregulated by goitrogenic treatment. Furthermore, administration of iodide, even in low doses, inhibited the expression of TPO and NIS (Table 2) mRNAs without affecting either the TSH concentrations or the expression of thyroglobulin or TSH receptor

**Table 2** Regulation of NIS mRNA and protein expression.

Agent	Experimental model	Effect on NIS		Reference
		mRNA	Protein	
TSH	<i>In vivo</i> , rats		+	5
	<i>In vivo</i> , dogs	+		6
	Human thyrocytes	+	+	7, 8
	FRTL-5 cells	+	+	9
	PC C13 cells	+		10
Forskolin/db-cAMP	Human thyrocytes	+	+	7, 8
	FRTL-5 cells	+	+	9
	PC C13 cells	+		10
Iodide	<i>In vivo</i> , rats		-	5
	<i>In vivo</i> , dogs	-		6
Retinoic acid	FRTL-5 cells	-		11
	Human thyroid carcinoma cells	+		11
TGF-β	FRTL-5 cells	-*		12
IL-1/IFN <sub>γ</sub> /TNF <sub>α</sub>	FRTL-5 cells	-*		13
	Human thyrocytes	-*		7
Oncostatin M	FRTL-5	-*		14

db-cAMP, dibutyryl-cAMP; TGF-β, transforming growth factor β; IL-1, interleukin-1; IFN<sub>γ</sub>, interferon-γ; TNF<sub>α</sub>, tumor necrosis factor α.

-\*, inhibition of TSH stimulation.

(TSH-R) mRNAs. This suggests that an increase in iodide, probably through its intracellular concentration, may downregulate expression of the *NIS* gene.

Factors other than TSH and iodide may participate in the regulation of *NIS* expression. Retinoic acid was found to upregulate iodide transport in human follicular thyroid carcinoma cell lines, whereas a down-regulation was observed in non-transfected FRTL-5 cells (11). In the same cells, transforming growth factor- $\beta$  inhibited TSH stimulation of *NIS* mRNA and protein expression (12). In both FRTL-5 and primary thyroid follicular cell cultures, interleukin-1 $\alpha$ , tumour necrosis factor  $\alpha$  and interferon- $\gamma$  inhibited TSH-induced *NIS* gene expression and iodide uptake (7, 13) (Table 2).

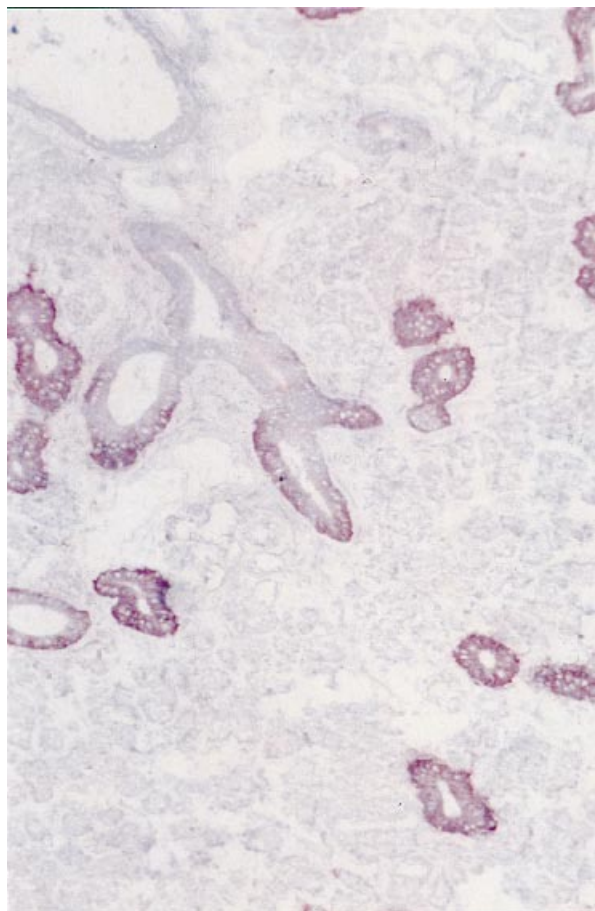
## ***NIS* tissue-expression studies**

### ***NIS* gene promoter characterization and *NIS* gene expression**

To investigate the transcriptional level of *NIS* mRNA regulation, the 5'-flanking region of the *NIS* gene has been characterized. The tentative human *NIS* gene transcriptional start site has been located 375 nucleotides relative to the ATG site (17–19), and potential binding sites for activating protein (AP)-1, AP-2, stimulating protein-1 and cAMP response element binding protein have been identified (17), in addition to two sites similar to the consensus sequence for thyroid-specific transcription factor (TTF)1 (18). However, the most relevant finding concerns the absence of the regulatory elements necessary to obtain cell-type specific transcription in a region located proximal to the transcription start site. Differently from that observed in promoters of other genes encoding thyroid-specific proteins, such as thyroglobulin, TPO and TSH receptor, elements within 2 kb in the 5'-flanking region of recombinant *NIS* are not sufficient to confer thyroid-selective transcription. However, a less stringent thyroid-specific control of *NIS* gene expression may be expected, considering that iodide transport also occurs in other tissues (see below). Recently, Ohno *et al.* (20) reported the presence of an enhancer that is located between nucleotides 2264 and 2495 in the 5'-flanking region of the rat *NIS* gene. It stimulates transcription in a thyroid-specific and cAMP-dependent manner, in the presence of the transcription factor, PAX-8.

### ***NIS* protein expression in thyroid tissue**

The production of specific antibodies against the *NIS* has facilitated the study of expression of the symporter protein. Despite of the presence of high levels of human (h) *NIS* mRNA, normal thyroid glands exhibit heterogeneous expression of *NIS* protein. Immunohistochemical studies showed that only a minority (approximately 30%) of follicular cells express detectable



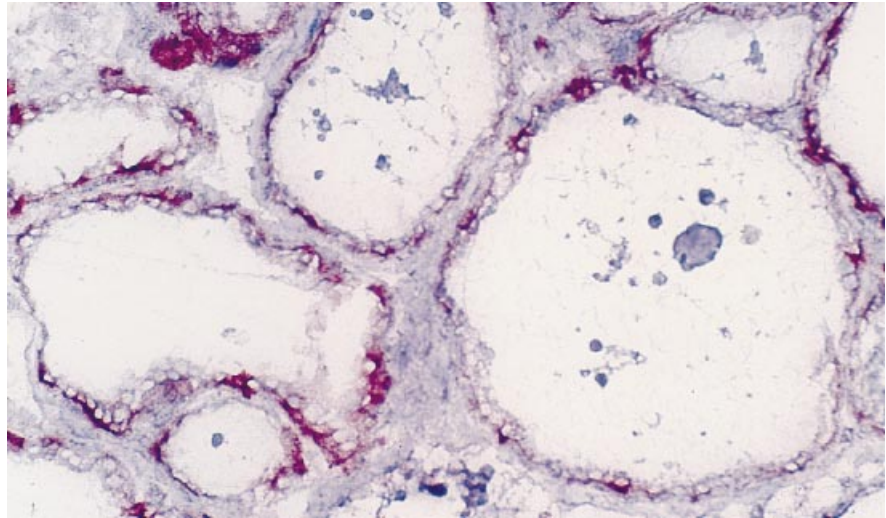
**Figure 2** Expression and localization of *NIS* protein in normal human thyroid tissue (original magnification,  $\times 200$ ). Only some follicular cells are stained.

amounts of hNIS protein (21, 22) (Fig. 2). Differently, in Graves' disease thyroid glands and toxic adenomas staining of hNIS protein was stronger and was detected in the majority of thyrocytes, corresponding to the high levels of *NIS* mRNA detected in these tissues (7, 15). As expected, cell polarization has been observed, with immunostaining of the *NIS* protein limited to the basolateral membrane of the thyrocytes (Fig. 3).

In immunoblot experiments, the human *NIS* protein is detected as a single band of approximately 70–80 kDa (8, 23, and our unpublished observations), indicating that glycosylation accounts for  $\sim 15$  kDa, along with the  $\sim 65$  kDa polypeptide backbone, as deduced from its primary structure.

### ***NIS* expression in extrathyroidal tissues**

*NIS* mRNA was also detected in non-thyroid tissues able to concentrate radioiodine (Table 3), including salivary glands, stomach, thymus and breast. Lower levels of expression were detected in prostate, ovary,



**Figure 3** Basolateral localization of the NIS protein in follicular cells from Graves thyroid tissue (original magnification,  $\times 1000$ ). Note that all follicular cells are stained, and that staining is more intense in the basal part.

adrenal gland, lung and heart. By contrast, no expression of *NIS* gene was detected in colon, normal orbital fibroblasts and nasopharyngeal mucosa (7, 24, 25).

Salivary glands do express hNIS protein, as shown by immunohistochemical studies using a specific anti-hNIS antibody (21, 26). Immunostaining of the NIS protein occurs only in the ductal cells, where it is diffusely expressed within cells, and not in the acinar cells (Fig. 4). In the human stomach, expression of NIS was restricted to parietal cells. Recently, characterization of rat gastric NIS by Western blot analysis showed the presence of a small amount of immature gastric NIS, with different molecular weights compared with the thyroid protein (27). This suggests the existence of tissue-specific post-translational mechanisms in the functional regulation of NIS expression.

## Biochemical properties of thyroid cancer tissues

### Alteration of iodine metabolism in cancer tissue

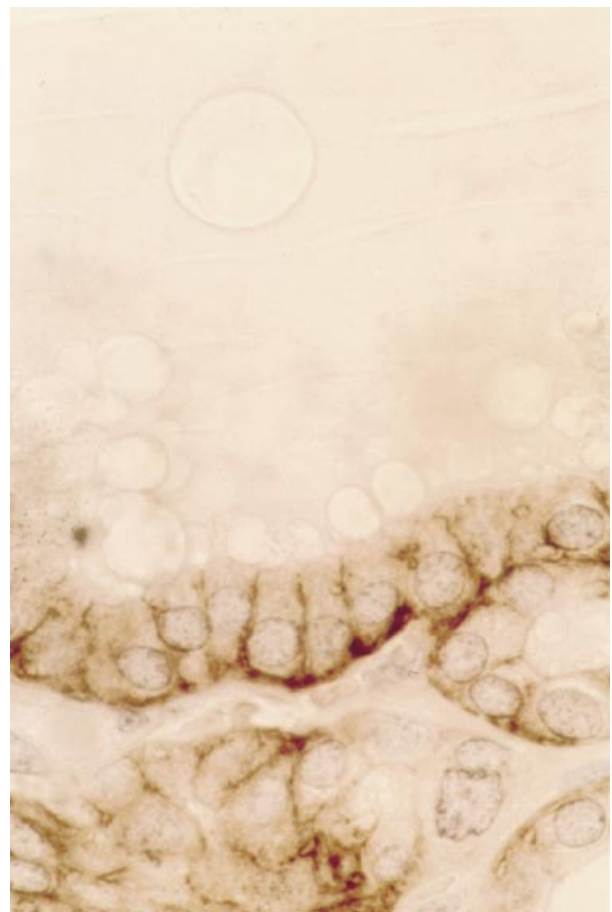
Although papillary and follicular thyroid carcinomas retain the majority of biological properties of normal

**Table 3** NIS transcript and protein expression in human tissues.

NIS mRNA*	+++	Thyroid, salivary gland, parotid gland, submandibular gland
	+	Pituitary gland, pancreas, mammary gland, gastric mucosa, prostate, testis, ovary, adrenal gland, thymus, lung
	-	Orbital fibroblasts, colon, nasopharyngeal mucosa
NIS protein**	++	Thyroid, salivary gland

\* Detected by RT-PCR and Southern hybridization (24).

\*\* Detected by immunohistochemistry (21, 22).



**Figure 4** Expression and localisation of NIS in the submaxillary gland (original magnification,  $\times 200$ ). Only ductal cells are stained, with diffuse staining within cells.

thyroid cells, a variety of biochemical defects has been demonstrated (28–30). Indeed, impaired intrathyroidal iodine metabolism represents one of the most peculiar abnormalities present in thyroid neoplastic tissue. The biological activity of peroxidase, although normal in benign cold adenomas, was decreased or absent in thyroid carcinomas, resulting in a low iodine organification. As a consequence, in thyroid cancer tissues, a low intrathyroidal iodine concentration, a low degree of iodination of thyroglobulin and a low rate of thyroid hormones synthesis were observed. The altered iodine intrathyroidal metabolism may confer on tumor cells a proliferative advantage as a result of the loss of the iodine autoregulating process (31).

How and to what extent do the abnormalities of iodine affect the management of patients with thyroid cancer? The derangement of thyroid iodine metabolism accounts for the appearance of adenomas and of the majority of malignant thyroid tumors as hypofunctioning ('cold' nodules) on scintiscan, indicating an impaired radioiodine trapping ability *in vivo*. Moreover, *in vivo* studies in patients with metastases have shown that uptake of radioiodine is always less in neoplastic tissue than in its normal counterpart. In fact, tissue uptake of iodine is about 1%/g of the administered activity in normal thyroid tissues, whereas it ranges from 0.1% to 0.001%, or even less in neoplastic tissues. Furthermore, the effective half-life of iodine in neoplastic tissues is, on average, equal to 3–5 days, but sometimes much shorter, whereas it ranges from 6 to 8 days in normal thyroid tissues. This short half-life may be due to abnormalities in the organification process that are, at least in part, related to a defect in the peroxidase system. Increasing the effective half-life of iodine-131 in the thyroid cells by the use of lithium salts significantly increases the dose of radiation to the neoplastic tissue (32). Uptake of iodine-131 can be heterogeneous among foci in which uptake occurs, but also in a given focus; this is related to the speckled distribution of radioiodine, which is different from one neoplastic thyroid cell to another, as shown by autoradiographic studies, and is in accordance with the heterogeneous expression of the NIS protein, as shown by immunohistochemistry (see above).

## Iodine and thyroid cancer

### ***Iodide transport and NIS expression in oncogene-transformed thyroid cells***

Proto-oncogene activation, over-expression, or both, have a role in thyroid tumorigenesis and determine a loss of differentiation of thyroid function (33, 34).

PC Cl 3 and FRTL-5 are differentiated rat thyroid cells that possess specific markers of thyroid function – namely, thyroglobulin synthesis and secretion, the ability to trap iodide and TSH-dependent growth (35, 36). Moreover, the oncogene-transformed cell lines display different degrees of malignancy and differentiation (10). They

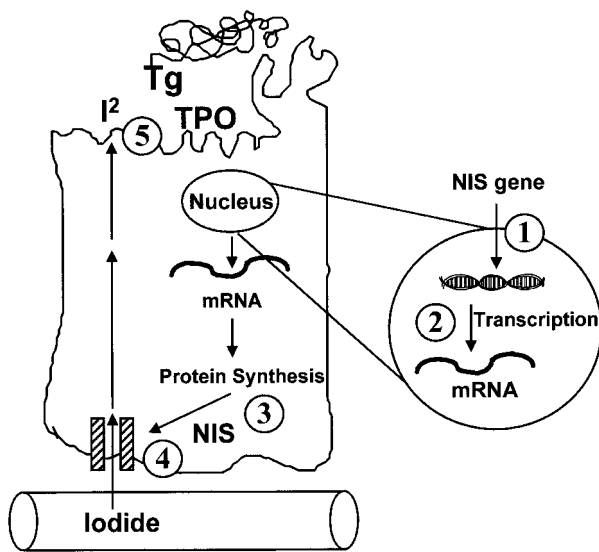
exhibit a TSH-independent growth, even if some clones require thyrotropin hormone for optimal proliferation when cultured in conditions of low serum concentration. However, all transformed thyroid cells, exhibiting either a high or a low degree of malignancy (10; D Russo and S Filetti, unpublished data), display a loss of iodide uptake function. Moreover, whereas all thyroid-specific differentiation markers are lost in highly malignant rat thyroid cell lines (that is, they are transformed by KiMSV, Kirsten murine sarcoma virus), the de-differentiation process appears to be restricted to the loss of iodide uptake in transformed thyroid cells of low malignancy; this suggests that, *in vitro*, the loss of ability to concentrate iodide is an early effect of oncogenic rat thyroid cell transformation.

Likewise, the phenotype of transformed thyroid cells may make it possible to define, at the molecular level, the mechanisms whereby oncogene-transformed rat thyroid cells lose the ability to trap iodide. The possibility exists that activation of a specific oncogenic pathway may alter the iodide-trapping ability of transformed cells, through an alteration in NIS gene expression. In this regard, we found that some oncogene-transformed cell lines showed very low or absent amounts of NIS transcript compared with normal, non-transformed thyroid cells (10). Oncogenic transformation determines also the loss of other thyroid-specific transcripts – thyroglobulin and thyroperoxidase in Ras transformed cells – suggesting that the transcription factors controlling their expression are either not present or inactive (37). Indeed, in K-Ras-transformed cells, both PAX-8 and TTF-1 mRNA are undetectable (38), and in RET/PTC1-transformed cells, PAX-8 is expressed at reduced levels and TTF-1 is inactive (39). PAX-8 expression is lost also in oncogenic transformed cells mediated by p53 and Polyomavirus middle T antigen (37, 39); TTF-2 DNA binding activity is lost in transformed cells by the polyomavirus middle T antigen (38, 41). Other oncogene-transformed cell lines (PC v-erbA, PC HaMSV, PC v-raf, PC E1A) showed reduced NIS mRNA levels when compared with non-transformed cells. These studies suggest the presence of a transcriptional or post-transcriptional event, or both, responsible for the complete loss of iodide uptake function (10).

Figure 5 summarizes the possible targets of alterations in the tumorigenic process that are responsible for the loss of iodide uptake in thyrocytes.

### ***NIS expression and impaired iodine metabolism in thyroid cancer cells***

The recent cloning of the hNIS has provided the possibility to examine, at the molecular level, the mechanism through which thyroid iodide trapping is hampered in human thyroid cancer cells. Using RT-PCR, Smanik PA *et al.* (25) found NIS expression to variable extents in papillary thyroid carcinomas. However, NIS mRNA was not detected in several thyroid cancer cell lines

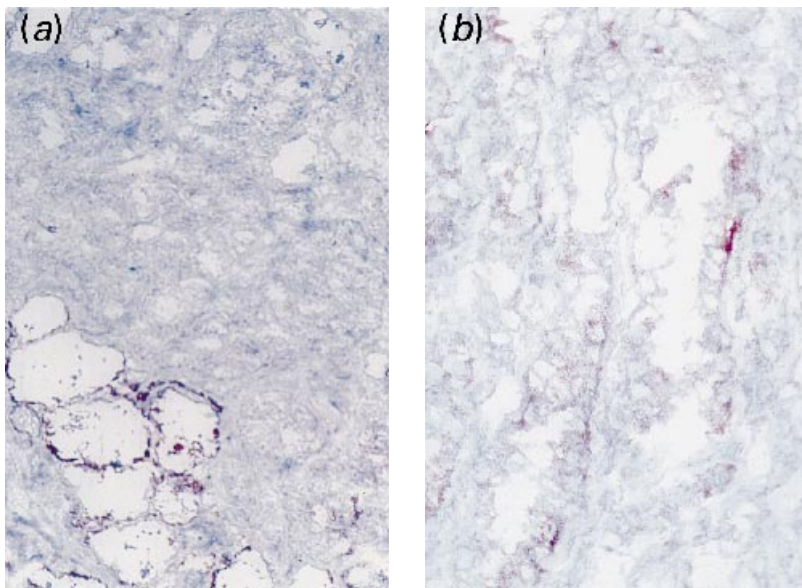


**Figure 5** The possible targets of the alterations occurring in the tumorigenic process responsible for the loss of iodide uptake in thyrocytes. 1, Alterations in *NIS* structural gene; 2, alterations in regulation of *NIS* transcript expression; 3, alterations in *NIS* protein synthesis; 4, alterations in *NIS* transport to the plasma membrane; 5, alterations in iodide organification.

that have lost the ability to concentrate iodide (2). By using non-quantitative RT-PCR in a series of 24 differentiated carcinomas, we found a loss of expression of the iodide symporter gene in six primary thyroid tumors (42); in contrast, all neoplastic tissues studied expressed the *Tg* (thyroglobulin) gene. The absence of *NIS* mRNA transcript, however, was not restricted to the malignant phenotype, being found also in one of 11 'cold' benign follicular adenomas. To identify more precisely the whole array of defects in thyroid-specific gene expression in thyroid tumors, quantitative methods of analysis were applied, based on the fluorescent TaqMan methodology and real-time measurement of fluorescence. This method, by providing accurate and reproducible quantitation of gene copies, enabled examination of the expression of the four functional parameter-encoding genes, including *NIS*, thyroid peroxidase, thyroglobulin and thyrotropin receptor, in a large number of well-characterized microdissected tissues. Whereas in cold benign adenomas the expression of *Tg*, *TSH-R* and *TPO* genes was similar to that observed in normal thyroid tissue, *NIS* mRNA levels were decreased (2- to 700-fold lower) in the majority of samples studied. In neoplastic thyroid tissues, more extended alterations were detected: *NIS* mRNA expression, although normal in three samples, was 10- to 1200-fold lower than in normal tissues (median 100-fold) in another 38 samples; *TPO* mRNA expression was reduced by 5- to 500-fold and thyroglobulin mRNA by 2- to 300-fold. Interestingly, the expression of *TSH-R* transcripts was normal in most

tumors (V Lazar *et al.*, unpublished observations). Furthermore, later tumor stages (stage >1 as opposed to stage 1) were associated with lower levels of expression of *NIS* ( $P=0.03$ ) and *TPO* ( $P<0.01$ ), but not with the expression of thyroglobulin or *TSH-R*. Taken together, these data indicate an extreme, peculiar and specific alteration of *NIS* gene expression in thyroid tumors. Having already been detected in cold follicular adenomas, the low expression of *NIS* may represent an early abnormality in the thyroid cell transformation pathway, and not be a consequence of cancer progression. Conversely, *Tg* gene expression appears to be well conserved in cancer cells, being absent only in undifferentiated cancer tissue (unpublished observations). More interestingly, in human thyroid tumorigenesis, thyroid-specific genes generally display a specific pattern in their alteration ( $NIS > TPO > Tg > TSH-R$ ), and this behavior is similar to that observed *in vitro* in oncogene-transformed rat thyroid cells (see above). Immunohistochemistry using anti-*NIS* antibodies confirmed the much lower expression of the *NIS* protein in thyroid cancer tissues, and also demonstrated that its expression was heterogeneous, being detected in only a few malignant papillary or follicular thyroid cells (Fig. 6). In these studies, the serum TSH concentration was in the normal range in all patients at surgery and the amounts of *NIS* mRNAs obtained, therefore, represent the basal and unstimulated concentrations (22).

These data suggest, in accordance with clinical data, that an intensive TSH stimulation should be performed in patients with thyroid cancer, before any administration of iodine-131, in order to increase *NIS* expression and thus the ability of thyroid cancer tissue to take up the iodine-131 (43, 44). Obviously, the complete/partial reversibility of the uptake function requires the structural integrity of the *NIS* gene and *NIS* protein – an issue confirmed by the absence of point mutations or other genetic alterations in a series of thyroid carcinomas screened for genetic abnormalities (M Derwahl, personal communication). All the published studies involved primary cancers only, and thus the possibility exists that their metastases may display a different pattern of expression of thyroid-specific proteins. Although systematic studies on this issue are not yet available, lymph-node metastases arising from differentiated thyroid carcinoma retain, as primary carcinomas, some features of epithelial thyroid cells and transcribe both *Tg* and *TSH-R* genes (45). In contrast, about one-third of thyroid cancer metastatic tissues, although showing *Tg* and *TSH-R* transcripts, did not express the *NIS* gene (F Arturi, D Russo, S Filetti, unpublished observations). Another issue is whether a discrepancy in *NIS* expression may exist between metastases and primary cancer. In this regard, dot blot analysis demonstrated that, although *NIS* mRNA expression was maintained in thyroid tumor lymph-node metastases, a lower *NIS* expression was detected in metastatic tissue, in comparison with the respective



**Figure 6** NIS expression in papillary thyroid carcinoma. (a) NIS expression is lower than in normal thyroid tissue (magnification,  $\times 200$ ); (b) NIS expression is observed in only some papillary cells (original magnification,  $\times 400$ ).

primary cancer tissue (Fig. 7). The possibility exists, therefore, that a de-differentiation process may occur in metastases; however, additional studies on this issue are necessary with respect to the clinical application of such a finding. Moreover, in the absence of specific NIS mRNA in the primary tumor, there was no detectable expression of the NIS gene in corresponding lymph-node metastases (Table 4). However, the only data available at present are from tissues collected in the absence of TSH stimulation.

At the present time, the low expression of NIS and of peroxidase genes in thyroid cancer tissues is largely unexplained, although promoter-specific protein:protein or protein:DNA interactions are likely to be involved. Studies of transcription factors and of abnormal methylation of genes are, indeed, being performed.

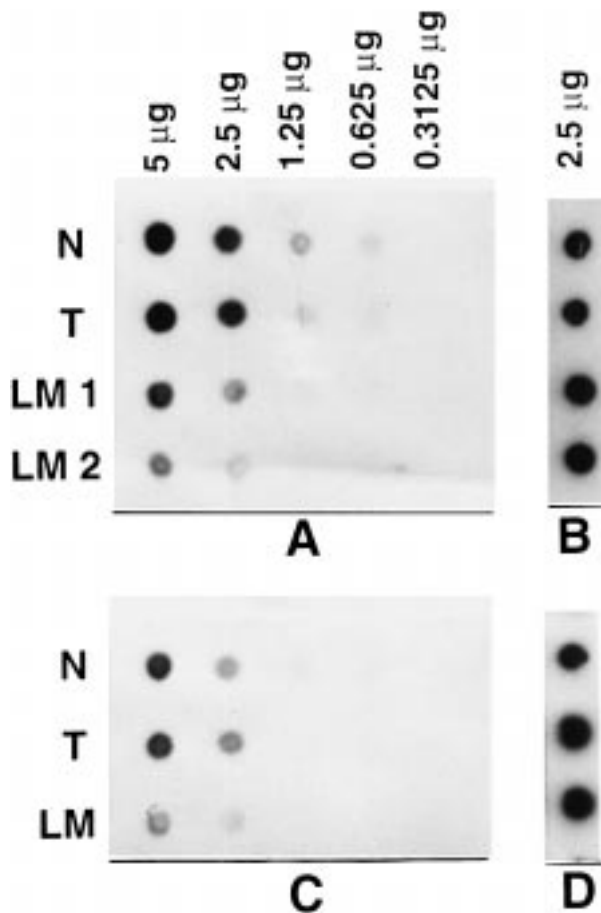
## Radioiodine in the follow-up and treatment of differentiated thyroid carcinoma

### Radioiodine in the follow-up

Iodine uptake by thyroid cancer cells and blood levels of thyroglobulin are the two cornerstones in the follow-up of patients with thyroid cancer. Indeed, whereas serum thyroglobulin is detectable in most patients with persistent or recurrent disease, radioiodine uptake is found only in about two-thirds of patients with metastatic disease. Detectable production of thyroid hormones is, however, infrequent, unless genetic alterations, such as activating mutations in the TSH receptor, may determine the rare association of thyroid carcinoma and thyrotoxicosis (46–49). These clinical findings are in agreement with the molecular analysis (see above). However, differentiation of normal

and of neoplastic thyroid cells depends on TSH stimulation. In fact, iodine uptake is absent during thyroid hormone-suppressive treatments; after withdrawal of thyroid hormone treatment, a close relationship has been observed between the degree of TSH stimulation and the increase of radioiodine uptake in metastases (50). Similarly, serum concentrations of thyroglobulin increase after withdrawal of thyroid hormone treatment in almost all patients with persistent or recurrent disease, even in the absence of radioiodine uptake in the metastases (51). This clearly shows that all these tumor tissues can respond to TSH stimulation, demonstrating the presence of functional TSH receptors in their plasma cell membrane. Indeed, the presence of the TSH-R gene has been exploited also as marker for the early identification of lymph-node metastases from differentiated thyroid carcinoma (45). As thyroglobulin is produced only by thyroid follicular cells, a detectable serum concentration of thyroglobulin after total thyroid ablation indicates persistent or recurrent disease. In fact, during thyroid hormone-suppressive treatment, the majority (about 98%) of patients with differentiated thyroid carcinoma considered to be in complete remission exhibit an undetectable serum thyroglobulin concentration, whereas high levels were found in the majority of patients with large metastases. However, a serum thyroglobulin concentration was undetectable during L-thyroxine-suppressive treatment in about 20% of patients with isolated neck lymph-node metastases and in about 5% of patients with small lung metastases, not observable on standard chest radiographs. After withdrawal of thyroid hormone treatment, the serum concentration of thyroglobulin remained undetectable in most patients in complete remission and increased to a high level in most patients with disease. Therefore, increased TSH concentrations obtained after





**Figure 7** Dot blot hybridization of normal (N), tumoral (T), and metastatic (LM) tissue RNA of two patients with thyroid papillary carcinoma. In panels A and C, total RNA was hybridized with labeled iodide carrier cDNA (EcoRI-HindIII, 1-2046 fragment of rat NIS gene). In panels B and D, total RNA was hybridized with labeled murine actin probe to ascertain equal RNA loading (from Arturi F, Russo D, Schlumberger M, DuVillard JA, Caillou B, Vigneri P, Wicker R, Chieffari E, Suarez HG & Filetti S. Iodide symporter gene expression in human thyroid tumors. *Journal of Clinical Endocrinology and Metabolism* 1998 **83** 2493–2496. © The Endocrine Society, with permission).

withdrawal of thyroid hormone treatment increase the sensitivity of serum thyroglobulin determination for the detection of neoplastic disease (44, 51).

The localization of neoplastic tissue relies firstly on iodine-131 total body scanning ( $^{131}\text{I}$ -TBS) (Fig. 8). The results of  $^{131}\text{I}$ -TBS depend on the ability of thyroid cancer tissue to take up iodine-131 in the absence of iodine contamination and in the presence of high serum TSH concentrations (e.g. >25–30 mU/l), which are achieved by withdrawing thyroxine for 4–6 weeks. Intramuscular injection of recombinant human (rh) TSH is a promising alternative, because thyroxine does not need to be discontinued and the side effects are minimal. It provides a short but intense stimulation. The results of  $^{131}\text{I}$ -TBS and of serum thyroglobulin

determination performed after the administration of rhTSH and after withdrawal of thyroid hormone treatment are similar in most patients (52).

Iodine uptake in neoplastic tissues is greater and more frequent in younger patients, in those with papillary or well-differentiated thyroid carcinoma and in those with small metastases (53). This suggests an accumulation of metabolic defects with both age and tumor progression. Total-body scanning may also show physiological concentration of iodine-131 in salivary glands, in the stomach, in the liver (attributable to iodoproteins) and accumulation of iodine-131 in the colon (Fig. 9).

As already mentioned, fractional uptake is often low in thyroid cancer tissue and scanning with a diagnostic dose may be negative. This problem can be overcome by administering a high dose of iodine-131. In fact, assuming equivalent fractional uptake after the administration of a diagnostic or a therapeutic dose of iodine-131, an uptake too low to be detected with 2–5 mCi may be detectable after the administration of 100 mCi. This is the rationale for administering 100 mCi iodine-131 to patients with increased serum thyroglobulin concentrations, even in the absence of any other evidence of disease, including a negative diagnostic  $^{131}\text{I}$ -TBS. When this is done, total-body scanning should be performed 4–7 days later. This strategy enabled the discovery of yet unknown foci of uptake in 60–80% of those patients treated in this way (44).

In the absence of uptake of iodine-131, computed tomography or magnetic resonance imaging of the neck and lungs, and bone scintigraphy may be useful. Scintigraphy with a less specific tracer has been advocated for the follow-up of patients with thyroid cancer. Such tracers include thallium, meta-iodobenzyl iodothyrosine (MIBI), tetrofosmin and somatostatin analogs (54, 55). They can be used while the patient is receiving thyroxine treatment. In some reports, the sensitivity of these techniques appeared to be high, but they cannot replace  $^{131}\text{I}$ -TBS in patients with suspected disease, as they do not avoid the use of  $^{131}\text{I}$ -TBS; therefore, they should be performed only in patients with a negative  $^{131}\text{I}$ -TBS. Positron emission tomography (PET) using (F18) fluorodeoxyglucose (FDG) is more promising. Enhanced glucose metabolism is a non-specific feature of tumor cells. It can be performed while the patient is receiving thyroxine treatment. However, FDG uptake was found to be greater when thyroxine was withdrawn (56). FDG uptake was detected more frequently in patients with poorly differentiated thyroid carcinoma, in whom no detectable iodine-131 uptake could be demonstrated. It was detected in neck lymph-node metastases, and even in nodes of less than 1 cm in diameter (57, 58). Although highly useful in specific contexts, FDG PET scan cannot supersede  $^{131}\text{I}$ -TBS, but can and should be performed in patients with a high likelihood of persistent or

**Table 4** Clinical features and *Tg*, *TPO* and *NIS* gene expression in a series of differentiated thyroid carcinomas with lymph-node metastases.

Histotype	Sex	Age (yr)	Metastases	Uptake of iodine-131	Thyroglobulin (ng/ml)		<i>Tg/TPO/NIS</i>
					On T <sub>4</sub>	Off T <sub>4</sub>	
Papillary	F	23	+	+	2.5	24	+ / + / +
Papillary	F	35	+	+	9	54	+ / + / +
Follicular	F	55	+	+	10	38	+ / + / +
Follicular	M	28	+	+	17	135	+ / + / +
Papillary	M	75	+	-	41	360	+ / + / -
Papillary	M	22	+	-	32	205	+ / + / -
Papillary	F	25	+	-	6	37	+ / + / -
Follicular	F	53	+	-	31	116	+ / + / -
Papillary	F	28	+	-	25	330	+ / + / +
Papillary	F	40	+	-	1	30	+ / + / +
Papillary	F	28	+	-	4	18	+ / + / +
Follicular	M	57	+	-	23	100	+ / + / +

The *NIS* and mRNA expression data are referred to primary tumors. When absent from the primary tumor, the *NIS* transcript was not detectable in corresponding lymph-node metastases. T<sub>4</sub>, thyroxine.

recurrent disease and a negative high-dose <sup>131</sup>I-TBS. In some patients showing a negative <sup>131</sup>I-TBS, we demonstrated absent or very low expression of *NIS* gene in the neoplastic tissue. This defect appeared to be intrinsic, being present already in the primary thyroid tumor, and not acquired in the metastatic tissues through a further de-differentiation during the process of tumor progression. In some of these tumors with no demonstrable uptake of iodine-131, we found an increased expression of the type 1 glucose transporter (V Lazar *et al.*, unpublished observations). These findings may have a clinical impact. In these cases, alternative methods with which to detect metastases, such as PET scan, may be performed.

### Radioiodine in the treatment of differentiated thyroid cancer

The outcome of iodine-131 treatment is related to the effective dose of radiation delivered to the thyroid cancer tissues, which in turn depends on the effective half-life and the radioactive concentration, namely, the ratio of total uptake to the mass of thyroid tissue. A radiation dose greater than 80 Gy should be delivered to obtain cure; if radiation doses are less than 35 Gy, there will be little chance for success. This shows how TSH stimulation and the absence of iodine contamination are important (59, 60). The treatment dose of radiation is usually 3.7–5.5 GBq (100–150 mCi) in adults, and approximately 37 MBq (1 mCi)/kg body weight in young children. Successive treatments with iodine-131 are given to achieve sufficient doses of radiation to thyroid cancer tissues, and until total ablation of residual uptake is attained on post-treatment total-body scanning (44).

Complete responses to iodine-131 treatment have been observed in 33–50% of patients with distant metastases that take up iodine-131. The overall survival rate at 10 years from the time of detection of

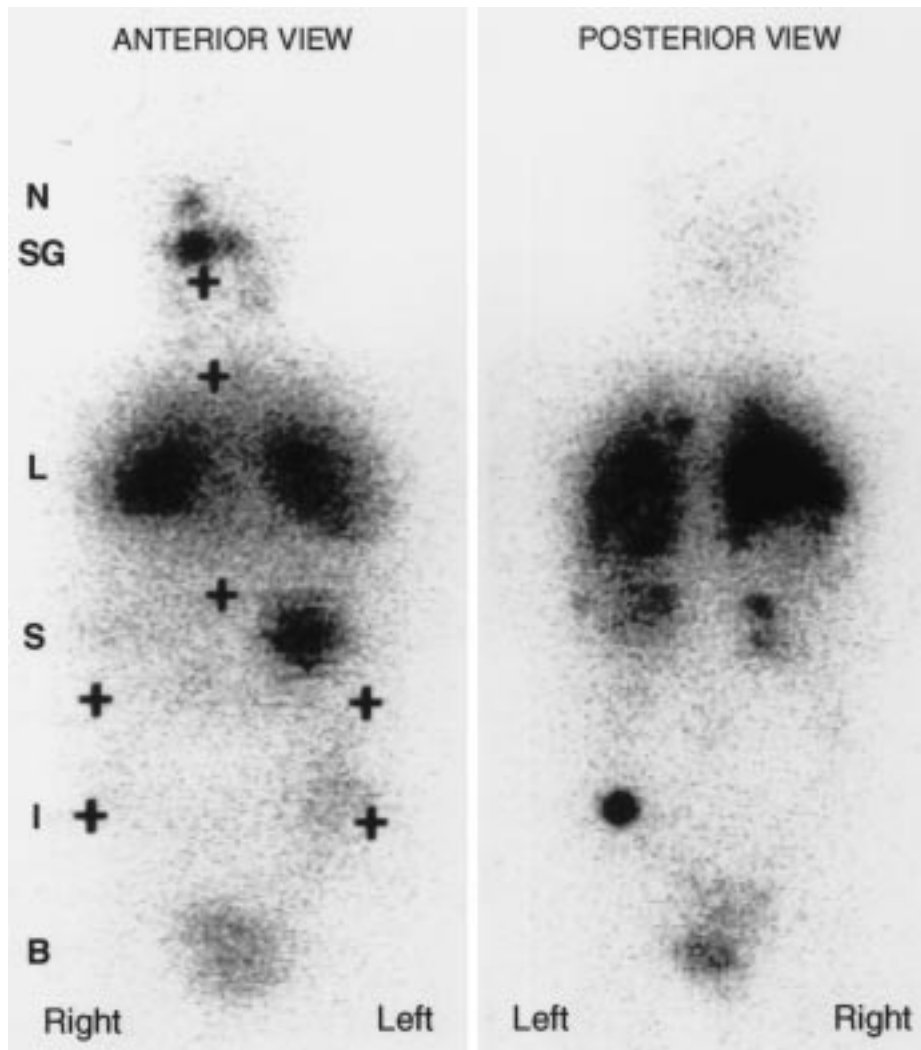
metastases ranges from 25 to 40%. A younger age, a well-differentiated histotype, a positive uptake of iodine-131 and a small extent of disease have a favorable impact on outcome (44, 53). The disappearance of iodine-131 uptake, despite the persistence of tumor masses, may be related to the speckled distribution of iodine-131 in tumor tissues, which may be due to a heterogeneous expression of the *NIS* gene; in fact, the maximal path of beta-rays of iodine-131 in biological tissues is 2–3 mm.

In conclusion, iodine-131 treatment is beneficial in some patients with iodine-131 uptake, and mainly in those with small tumor foci. This underlines the need to treat patients with metastases at an early stage, and this is made possible by the combined use of measurement of serum thyroglobulin, and of <sup>131</sup>I-TBS. This also shows that iodine-131 treatment should be considered as a therapeutic adjunct to surgery, in particular in patients with isolated neck lymph-node metastases. In patients with bone metastases, surgery should be performed, when feasible; in case of radiological abnormalities, external irradiation should be combined with iodine-131 treatment. External radiotherapy may, in fact, deliver homogeneous and high doses of radiation to limited fields and therefore efficiently complement iodine-131 treatment. In patients not amenable to these therapeutic modalities, no other forms of treatment proved to be beneficial. This, again underscores the paramount importance of iodine-131 in the treatment of patients with differentiated thyroid carcinoma (53).

### Manipulating *NIS* expression: gene reactivation and gene treatment

#### Gene reactivation

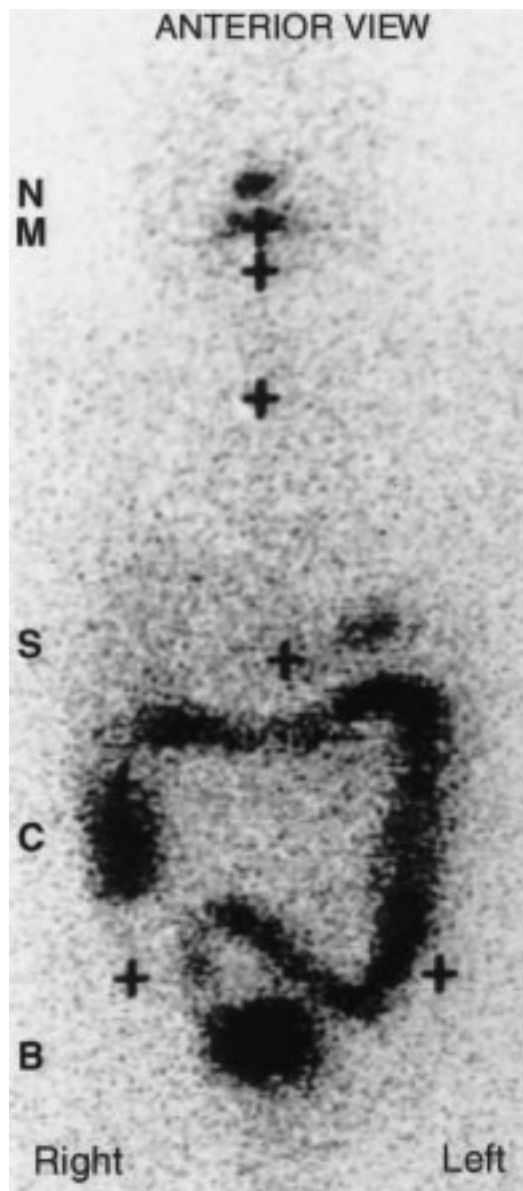
The possibility to achieve expression and functional stimulation of the *NIS* gene in thyroid tumor cells has



**Figure 8** <sup>131</sup>I-TBS in a patient with functioning lung metastases (L) and an iliac bone metastasis (I). N, nose; SG, salivary glands; S, stomach; B, bladder.

been exploited for a long time, by using physiological (TSH) or pharmacological agents (retinoic acid, protein kinase activators) (50, 61–63). Moreover, they suggest that transcriptional or post-translational events, or both, and not a defective structural *NIS* gene, occur in the transformation process responsible for the low iodide uptake of the thyrocytes. The goals of several studies, therefore, were to find a means to increase *NIS* gene expression in order to reverse the loss of iodide uptake observed in some transformed thyroid cells. In fact, expression of *NIS* mRNA is maintained in some transformed thyroid cell lines, including follicular (64) and anaplastic (19) carcinoma cell lines and oncogene-expressing rat thyroid cell clones (10). However, although an increase in *NIS* mRNA has been described *in vitro* in some follicular carcinoma cell

lines (FTC-133 and FTC-238) stimulated with retinoic acid, neither an increase in *NIS* protein nor a reactivation of iodide transport was detected (64). Studies are in progress in our laboratory and by other groups, taking advantage of the outstanding findings on the transcriptional regulation of the *NIS* gene (20) in order to modulate *NIS* gene transcription and reverse the loss of iodide uptake in transformed thyroid cells. Elucidation of other post-transcriptional and post-translational events that occur to achieve the complete function of the thyroid iodide transport system, including the involvement of cofactors or other transporter molecules, such as the recently identified pendrin (65), will be necessary to find the optimal conditions in which to target the thyroid tumoral tissues with radioiodine.



**Figure 9** <sup>131</sup>I-TBS in a patient with no evidence of disease. Only physiological uptake is seen, in the nose (N), mouth (M) and stomach (S), with accumulation in the colon (C) and in the bladder (B).

### Gene therapy

Combining targeting and expression of the *NIS* gene together with radioiodine treatment constitutes a powerful new strategy for the treatment of malignant thyroid diseases, in addition to the non-thyroid malignant diseases. The promise of *NIS* gene therapy lies in the low risk of adverse reactions of gene therapy procedures, in the specific iodide-concentrating effect induced by *NIS* expression, and in the long-standing experience of radioiodine treatment of patients with thyroid cancer.

Recent papers provided evidence that this approach may prove beneficial in the treatment of radiosensitive cancers. Transfection of rat *NIS* cDNA into malignantly transformed rat thyroid cells enabled the transfected cells to concentrate iodide, resulting in a 60-fold accumulation of iodine-125 (66). Furthermore, injection of transfected malignant cells into subcutaneous tissues of rats induced solid tumors that were capable of concentrating iodine-125 by 11- to 27-fold. Mandell *et al.* (67) demonstrated that *NIS* targeting may be used to deliver  $\beta$ -emitting radioisotopes to non-thyroid tumors. Retroviral transfer of the *NIS* gene into human and murine tumor cells, from melanoma, colon, liver or ovarian origin, resulted in an iodide-concentrating ability of infected cells. This approach allowed *in vivo* radioimaging of human tumor xenografted in mice. Moreover, *in vitro* experiments indicated that *NIS*-infected tumoral cells can be selectively killed by the induced accumulation of iodine-131.

Although the future of this approach appears promising for *in vivo* clinical applications, several questions remain unanswered and must be addressed. First, functional *NIS* targeting requires both the expression of the *NIS* gene and the adequate post-translational modification and trafficking of the *NIS* protein to plasma membrane in target tumor cells, which are known often to lack differentiated functions. Gene expression can be efficiently monitored by engineering appropriate tissue-specific promoters or controlling regulation as described above. Functional expression of the *NIS* gene in several extrathyroidal tissues appears to be different from that in thyroid tissue. Indeed, characterization of gastric *NIS* in rat shows that, whereas large amounts of *NIS* transcripts are expressed, little protein is detected, indicating a rapid degradation of the immature gastric *NIS* protein (27). In salivary glands, *NIS* protein is diffusely expressed within cells, suggesting a loss of the polarisation of the symporter. The significance of the variants of the *NIS* transcripts observed in different tissues, resulting from alternative splicing, remains unknown (25). Assuming that the *NIS* gene is functionally expressed at a sufficient level, the efficacy of the iodide-concentrating activity has to be demonstrated. Uptake, efflux and saturation kinetics of iodide in tumor cells, particularly those of extrathyroidal origin, are likely to be different from those in normal thyroid cells. Moreover, extra-thyroidal tissues do not efficiently accumulate and organify iodide as does the thyroid tissue, resulting in a short half-life of the iodide. This would suggest that the accumulation of radioactive iodide in tumor cells may not be adequate to achieve sufficient doses of radiation for therapeutic efficacy. Finally, problems also lie in the development of safe and efficient gene-delivery systems (68). Recombinant retroviruses are known to be efficient in *in vitro* experiments, but result in low infection efficiency *in vivo*. *NIS* gene transfer based on adenoviral vectors (which have

greater transfer efficiencies), liposomes or other vehicle remains to be investigated.

## Conclusion

Available data demonstrate that a decrease, loss, or both, of NIS gene expression may have a central role in the defective concentration of iodine by neoplastic thyroid carcinomas, in both primary or metastatic tissues (11, 25, 42, 69). Although the molecular mechanisms of this loss are unclear, the possibility exists that alterations in the iodide symporter may represent a primary defect; thus, in an originally heterogeneous population, a small number of cancer cells carrying this defect may overgrow over time and determine the appearance of an NIS-negative cell population. Conversely, the clinical utility of administration of iodine-131 in the diagnostic and therapeutic management of patients with differentiated thyroid carcinoma has been well established. In this regard, efforts to elucidate the molecular mechanisms underlying the iodide transport system in thyroid cancer cells will lead to the development of new diagnostic and therapeutic strategies, by allowing a more precisely tailored management of differentiated thyroid cancer.

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