Digitalislike Compounds Restore hNIS Expression and Iodide Uptake Capacity in Anaplastic Thyroid Cancer

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Anaplastic thyroid cancer (ATC) is a rare malignancy that accounts for 1%-2% of all thyroid cancers. ATC is one of the most aggressive human cancers, with rapid growth, tumor invasion, and development of distant metastases. The median survival is only 5 mo, and the 1-y survival is less than 20%. Moreover, as a result of severe dedifferentiation, including the loss of human sodium iodide symporter (hNIS) expression, radioactive iodide (RAI) therapy is ineffective. Recently, we have demonstrated beneficial effects of autophagy-activating digitalislike compounds (DLCs) on redifferentiation and concomitant restoration of iodide uptake in RAIrefractory papillary and follicular thyroid cancer cell lines. In the current study, the effects of DLCs on differentiation and proliferation of ATC cell lines were investigated. Methods: Autophagy activity was assessed in ATC patient tissues by immunofluorescent staining for the autophagy marker microtubule-associated protein 1A/1B-light chain 3 (LC3). In addition, the effect of autophagyactivating DLCs on the proliferation, gene expression profile, and iodide uptake capacity of ATC cell lines was studied. Results: Diminished autophagy activity was observed in ATC tissues, and in vitro treatment of ATC cell lines with DLCs robustly restored hNIS and thyroglobulin expression and iodide uptake capacity. In addition, proliferation was strongly reduced by induction of cell cycle arrest and, to some extent, cell death. Mechanistically, reactivation of functional hNIS expression could be attributed to activation of the transcription factors activating transcription factor 3 and protooncogene c-fos. Conclusion: DLCs could represent a promising adjunctive therapy for restoring iodide avidity within the full spectrum from RAI-refractory dedifferentiated to ATC.

Key Words: anaplastic thyroid cancer; autophagy; digitalislike compounds; hNIS; radioactive iodide

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naplastic thyroid cancer (ATC) is a rare type of nonmedullary thyroid cancer (TC) that accounts for 1%–2% of all thyroid malignancies. ATC is one of the most aggressive human cancers,

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with rapid growth, local tumor invasion, and distant metastases. Despite application of different treatment modalities, including surgery, chemotherapy, and radiotherapy, the median survival is only 5 mo and the 1-y survival is less than 20% (1-3). Furthermore, since functional expression of human sodium iodide symporter (hNIS), normally present on the thyrocyte membrane to facilitate iodide influx, is lost as a result of cellular dedifferentiation, the tumor becomes resistant to radioactive iodide (RAI) treatment (4).

For dedifferentiated TC, restoring functional hNIS expression on the cell membrane of TC tumor cells to reactivate iodide uptake, designated redifferentiation, represents a promising therapeutic strategy, mainly because of the high tumoricidal effects of RAI. Recently, several adjunctive therapies have been identified that are capable of improving iodide accumulation in RAIrefractory tumors, including inhibitors of mitogen-activated protein kinase (MAPK)/extracellular signal-regulated kinase (ERK) (5), v-RAF murine sarcoma viral oncogene homolog B1 (BRAF) (6) or mammalian target of rapamycin (7), and autophagy-activating compounds (8). However, no published studies have addressed the redifferentiation potential of any of these compounds in ATC.

Autophagy, an important mechanism for maintaining cellular homeostasis, is activated on cellular stresses, including nutrient deprivation and hypoxia. In addition, autophagy plays a role in carcinogenesis and the dedifferentiation process and is heavily intertwined with oncogenic pathways (9-12). In a previous study, autophagy activity was assessed in patients with poorly differentiated TC tumors, either follicular or papillary TC, and was correlated with clinical outcome parameters. This study demonstrated that TC tumors with high autophagy activity were responsive to RAI therapy and that the corresponding patients were more likely to reach remission, whereas low autophagy activity was associated with poor prognosis and RAI resistance (13). Moreover, autophagyactivating compounds, particularly digitalislike compounds (DLCs) such as digoxin, have been demonstrated to inhibit proliferation and to restore functional hNIS expression and iodide uptake capacity in poorly differentiated TC cell lines. Digoxin, directly inhibiting Na⁺/K⁺ adenosinetriphosphatase (ATPase), is currently prescribed to patients with cardiac failure and arrhythmia for its effects on cardiomyocyte contractility (14,15). Interestingly, several studies have addressed the potential of DLCs, digoxin in particular, as anticancer therapy and have revealed antiproliferative and proapoptotic effects on breast cancer, prostate cancer, melanoma, pancreatic cancer, lung cancer, and leukemia (16). Taken together, activation of autophagy by DLCs to restore hNIS expression and thereby reinduce

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sensitivity to RAI treatment might be a novel therapeutic strategy for poorly differentiated TC by multifaceted actions. However, this promising capacity of DLCs remains to be addressed for ATC.

In the present study, we assessed autophagy activity in ATC tissues and investigated the effect of DLC treatment concerning inhibition of proliferation or restoration of RAI uptake in 2 different ATC cell lines with distinctive genetic profiles. Based on the previous observations, we hypothesized that activation of autophagy by DLCs leads to inhibition of proliferation and to an increased capacity for RAI uptake in ATC cell lines.

MATERIALS AND METHODS

Cell Culture and Reagents

Two ATC cell lines, 8505C and Cal-62 (kindly provided by Prof. James Fagin, Memorial Sloan Kettering Cancer Center), harboring a BRAF^{V600E} or KRAS^{G12R} mutation, respectively, were cultured in Dulbecco modified Eagle medium supplemented with 10% fetal calf serum, 1 mM pyruvate, and gentamycin. Before use, the cell lines were authenticated by short tandem repeat profiling (17). The cells were incubated with dimethylsulfoxide (vehicle control); with the DLCs digoxin, strophanthin K, lanatoside C, digoxigenin (all from Sigma-Aldrich), or proscillaridin A (Santa Cruz Biotechnology); or with the MAPK/ERK-inhibitor selumetinib, the BRAF^{V600E} inhibitor dabrafenib, or the mammalian-target-of-rapamycin inhibitor rapamycin (all from Sigma-Aldrich) for indicated time points and concentrations. For all compounds, stock concentrations ranged between 50 and 100 mM in dimethylsulfoxide vehicle. Compound selection and applied concentrations were based on earlier studies (7,8,18,19).

Light Chain 3 (LC3)-II Immunofluorescence

To assess autophagy activity in ATC tissue specimens, fluorescent staining for the autophagy marker LC3 was performed on formalinfixed paraffin-embedded tissues derived from 10 ATC patients and compared with papillary TC cases with either low or high autophagy activity that were reported previously by our group (13). Ethical approval was obtained for the use of all patient tissues, which were obtained from the pathology archives of either Radboud University Medical Center, Nijmegen, The Netherlands, or collaborating pathology laboratories elsewhere in The Netherlands. LC3-II immunofluorescent 4',6-diamidino-2-phenylindole staining was performed as described previously (13). Staining was examined by a fluorescent microscope (DMI6000B; Leica). Autophagy activity was assessed by quantifying, using Fiji software, the number of LC3-II-positive puncta, representing LC3 incorporated in the autophagosomal membrane, per 100 cell nuclei. Scoring results were generated in quintuplet for each tissue section.

Real-Time Quantitative PCR

To investigate the potential capacity of autophagy activation by DLCs for reactivation of hNIS expression, the ATC cell lines Cal-62 and 8505C were treated for 24, 48, and 72 h with 5–50 μ M concentrations of the DLCs digoxin, strophanthin K, lanatoside C, digoxigenin, and proscillaridin A. In addition, both cell lines were treated with other previously identified TC redifferentiation agents: the MAPK/ ERK kinase inhibitor selumetinib (5), the mammalian-target-of-rapamycin inhibitor rapamycin (7), 8505C specifically with the selective *BRAF*^{V600E} inhibitor dabrafenib, or vehicle control (6). The treated ATC cell lines were lysed with TRIzol reagent (Invitrogen) followed by total RNA purification performed according to the manufacturer's instructions. iScript complementary DNA synthesis kits (Bio-Rad Laboratories) were used to transcribe isolated RNA into complementary DNA. Subsequently, quantitative PCR was performed using the SYBR Green method (Life Technologies). The primers used are listed in Supplemental Table 1 (supplemental materials are available at http://jnm. snmjournals.org).

Western Blotting

Western blotting of hNIS protein was performed as described previously (8). Cells (5×10^6) were lysed in 40 µL of lysis buffer (radioimmunoprecipitation assay buffer; Cell Signaling Technology). The membrane was blocked in Odyssey blocking buffer (Westburg) plus tris-buffered saline (TBS)/0.1% polysorbate 20 (1:1), followed by an overnight incubation at 4°C with a rabbit-antihuman hNIS antibody (1:500, 250552; Abbiotec) or a mouse-antihuman glyceraldehyde 3-phosphate dehydrogenase antibody (loading control, 1:10,000, ab8245; Abcam) in Odyssey blocking buffer plus TBS/0.1% polysorbate 20 (1:1). After overnight incubation, the membranes were washed with TBS/0.1% polysorbate 20 and incubated with goat-antirabbit 688 or goat-antimouse 688 for 1 h at room temperature. After being washed with TBS/0.1% polysorbate 20, membranes were analyzed by an Odyssey imager (Westburg).

Cell Proliferation and Viability Assays

For measurement of proliferation, 3-(4,5-dimethylthiazol-2-yl)-2,5diphenyltetrazolium bromide (MTT) assays were performed according to the manufacturer's instructions (Sigma-Aldrich). In brief, cells (1×10^3) were incubated with the DLCs for the indicated time points and concentrations. Subsequently, MTT substrate was added and the amount of converted substrate was measured by a plate reader at 570 nm. At first, the effects of DLCs on proliferation, cell survival, and cell cycle arrest were assessed. Separate MTT assays were performed at 24, 48, and 72 h after treatment with the selected DLCs at the indicated concentrations.

In addition, to determine whether inhibition of proliferation is attributable to induction of cell death, lactate dehydrogenase activity was measured in cell culture supernatants at the same time points for assessment of cell death in accordance with the suppliers' protocol (CytoTox 96 kit; Promega).

In Vitro Iodide Uptake

To determine whether increased expression of hNIS generates functional protein expression enabling iodide influx, in vitro ¹²⁵Iiodide uptake assays were performed as described previously (8,20). The ATC cell lines were treated for 48 h with one of the selected DLCs, at a 50 µM concentration. Subsequently, to determine the hNIS specific uptake of RAI, the cells were incubated for 30 min with 2.5 kBa of Na125I/mL (Perkin-Elmer) and 20 µM nonradioactive NaI as a carrier, with or without the competitive hNIS inhibitor sodium perchlorate, 80 µM, to control for hNIS-specific uptake. Radioactive medium was aspirated, and the cells were washed with ice-cold phosphate-buffered saline and lysed in 0.1 M NaOH buffer. Radioactivity was measured in the cell lysates in an automatic γ -counter (Wizard²; Perkin-Elmer). In parallel experiments, DNA was isolated from the cells (Puregene kit; Gentra Systems) and quantified by Nanodrop measurements (Thermo Fisher Scientific). Accumulated radioactivity was expressed as cpm per milligram of DNA.

Gene Expression Profiling of ATC Cell Lines after Treatment with DLCs

To elucidate the mechanism behind hNIS upregulation in ATC cell lines by DLCs, messenger RNA expression of thyroid-stimulating hormone receptor, thyroid transcription factor 1, thyroid transcription factor 2, paired box gene 8, activating transcription factor 3 (ATF3), and proto-oncogene c-fos (cFOS) was measured. In addition, expression of thyroglobulin and thyroid peroxidase was measured.

Statistical Analysis

Statistical significance was tested with the Student t test, Mann–Whitney U test, Spearman rank correlation coefficient, or Wilcoxon matched-pairs signed-rank test, when appropriate. P values below 0.05 were considered statistically significant. All statistical tests were performed using GraphPad Prism, version 5.0.

RESULTS

Autophagy Activity Is Diminished in ATC Tissues

All analyzed ATC tissue specimens exhibited low autophagy activity as demonstrated by the respective images of fluorescent LC3 staining (Fig. 1A), as well as by quantification of the number of LC3-II–positive puncta in ATC tumor cells (Fig. 1B).

DLCs Inhibit Proliferation and Induce Cell Death and Cell Cycle Arrest in ATC Cell Lines

All DLCs potently inhibited proliferation at and above concentrations of either 0.05, 0.5, or 5 μ M in both cell lines (Supplemental Fig. 1A). After 24 h of treatment, up to 37% of total lactate dehydrogenase was released by 8505C cells in a dose-dependent manner. The highest percentage of cell death was after treatment with 50 μ M proscillaridin A. For Cal-62, a dose-dependent increase in lactate dehydrogenase release was observed after incubation with digoxin and strophanthin K. Lanatoside C, digoxigenin, and proscillaridin A also induced lactate dehydrogenase release in the Cal-62 cell line, but largely independent of the administered dose



FIGURE 1. Autophagy activity in ATC tissues. (A) LC3-II immunofluorescent staining of formalin-fixed paraffin-embedded ATC specimens (1–6) and 2 papillary thyroid carcinomas with either low or high autophagy activity. LC3-II-positive puncta were visualized by fluorescent microscopy; cells were counterstained with 4',6-diamidino-2-phenylindole. (B) Autophagy activity in ATC tissues. Quantification of number of LC3-IIpositive puncta per 100 cells in papillary TC (PTC) control tissues and ATC tissues of all 10 patients. Data are based on 2 independent experiments, and images were analyzed in quintuplet. LC3-II = lipidated form of microtubule-associated protein 1A/1B LC3B.

(Supplemental Fig. 1B). The observation that 60%–90% of cells survived after DLC treatment for 48 h and that proliferation was completely inhibited in surviving cells indicated that these cells could have undergone cell cycle arrest. To provide additional arguments for this possibility, CDKN1A/p21 gene expression analysis was performed. Expression of CDKN1A/p21 was increased in both 8505C and Cal-62 after DLC treatment; the highest upregulation was observed in 8505C after 48 or 72 h of treatment, whereas in Cal-62 the highest upregulation was apparent after 72 h of treatment (Supplemental Fig. 1C).

DLCs Restore hNIS Expression in ATC Cell Lines

The diminished autophagy activity that was observed in ATC tissue sections uncovered a compelling opportunity to investigate the potential capacity of autophagy activation by DLCs for reactivation of hNIS expression. Whereas none of the targeted kinase inhibitors were able to induce hNIS expression, all tested DLCs were capable of reactivating hNIS messenger RNA expression in both cell lines, in most cases in a dose-dependent manner. In 8505C, the highest hNIS upregulation was observed at the 24- and 48-h time points; however, in Cal-62 the highest hNIS upregulation was maximally induced at 48 and 72 h (Fig. 2A). In addition, most DLCs induced hNIS protein expression after 48 h of treatment, except for digoxigenin-treated cells. Protein expression of nonglycosylated hNIS (56 kDa) and dimerized nonglycosylated hNIS (120 kDa) was especially elevated, whereas levels of the glycosylated monomer isoform (87 kDa) were increased to a lesser extent. Notably, in digoxin-treated 8505C solely increased expression of nonglycosylated hNIS was observed (Fig. 2B).

Iodide Uptake in ATC Cell Lines Is Restored by DLCs

In both 8505C and Cal-62, iodide uptake was strongly increased after treatment with digoxin, strophanthin K, and lanatoside C. In all these conditions, iodide uptake was efficiently inhibited by sodium perchlorate. In contrast, after treatment with proscillaridin A or digoxigenin, no increased RAI uptake was observed in the 2 ATC cell lines (Fig. 2C).

Expression of Some Genes in ATC Cell Lines Is Altered After Treatment with DLCs

No expression of thyroid-stimulating hormone receptor or thyroid peroxidase was observed in either cell line (data not shown). In 8505C, expression of thyroid transcription factor 1 and paired box gene 8 was not affected by DLC treatment. In contrast, expression of thyroid transcription factor 2 in 8505C and expression of paired box gene 8 and thyroid transcription factors 1 and 2 in Cal-62 were increased in some conditions, but this increased expression was not clearly associated with the degree of induced hNIS expression. In both 8505C and Cal62, increased thyroglobulin expression was observed after treatment with all DLCs (Fig. 3A). Interestingly, ATF3 and cFOS expression increased after DLC treatment and both were highly associated with the increased hNIS expression at all 3 time points in both cell lines (Fig. 3B).

DISCUSSION

ATC is a rare and devastating malignancy characterized by rapid proliferation, invasion, and dissemination. Patients diagnosed with ATC often present with inoperable locally advanced disease and have a poor prognosis with a median survival of only 5 mo and a 1-y survival rate of 20% (1,2). Genetic studies have



FIGURE 2. hNIS messenger RNA and protein expression and iodide uptake induced by digitalislike compounds. (A) hNIS messenger RNA expression in ATC cell lines 8505C and Cal-62 after 24, 48, and 72 h of treatment with selumetinib (10 μ M), dabrafenib (10 μ M), or rapamycin (50 μ g/mL) or with 5 μ M/50 μ M of the indicated digitalislike compounds. Data are mean ± SD. (B) hNIS protein expression in ATC cell lines 8505C and Cal-62 after treatment with 50 μ M of the indicated digitalislike compounds for 48 h. Data are representative of 3 independent experiments. (C) ¹²⁵I-iodide uptake by ATC cell lines 8505C and Cal-62 after treatment with 50 μ M of the indicated Cal-62 after treatment with 50 μ M of the indicated Cal-62 after treatment with 50 μ M of the indicated Cal-62 after treatment with 50 μ M of the indicated Cal-62 after treatment with 50 μ M of the indicated Cal-62 after treatment with 50 μ M of the indicated Cal-62 after treatment with 50 μ M of the indicated Cal-62 after treatment with 50 μ M of the indicated Cal-62 after treatment with 50 μ M of the indicated Cal-62 after treatment with 50 μ M of the indicated Cal-62 after treatment with 50 μ M of the indicated Cal-62 after treatment with 50 μ M of the indicated Cal-62 after treatment with 50 μ M of the indicated Cal-62 after treatment with 50 μ M of the indicated Cal-62 after treatment with 50 μ M of the indicated Cal-62 after treatment with 50 μ M of the indicated Cal-62 after treatment with 50 μ M of the indicated Cal-62 after treatment with 50 μ M of the indicated Cal-62 after treatment treatment with 50 μ M of the indicated Cal-62 after treatment with 50 μ M of the indicated Cal-62 after treatment with 50 μ M of the indicated Cal-62 after treatment with 50 μ M of the indicated Cal-62 after treatment with 50 μ M of the indicated Cal-62 after treatment with 50 μ M of the indicated Cal-62 after treatment with 50 μ M of the indicated Cal-62 after treatment with 50 μ M of the indicated Cal-62 after treatment with 50 μ



FIGURE 3. Gene expression profiling of ATC cell lines after treatment with digitalislike compounds. (A) TTF1, TTF2, and PAX8 and thyroglobulin (Tg) messenger RNA expression in ATC cell lines 8505C and Cal-62 after treatment with digitalislike compounds (5 μ M or 50 μ M) for 24, 48, and 72 h. Data are mean ± SD. (B) ATF3 and cFOS messenger RNA expression in ATC cell lines 8505C and Cal-62 after 24, 48, and 72 h of treatment with digitalislike compounds (5 μ M or 50 μ M). Data are mean ± SD.

revealed that ATC mostly originates from preexisting papillary TC with *BRAF* or *RAS* driver mutations, which have subsequently evolved toward ATC by acquiring additional genetic aberrations, particularly in *TP53*, *PIK3CA*, and *TERT* promoter (21,22). Despite recent insights into the etiology of ATC, currently no curative treatment options are available.

Conventional treatment of well-differentiated TC includes thyroidectomy followed by RAI ablation of tumor remnants (23). The response to RAI therapy highly depends on an intact iodide metabolism in which functional expression of hNIS, normally present at the basal membrane of thyroid follicular cells, is crucial. Currently, RAI therapy is not eligible for ATC patients, since tumor cells have lost their thyroid-specific gene expression related to iodide metabolism, leading to RAI refractoriness (4,24).

For poorly differentiated TC, novel therapeutic strategies have recently emerged that facilitate redifferentiation of tumor cells, thereby increasing their capacity for RAI uptake (5-8,25). Previously, loss of autophagy activity was shown to be associated with RAI resistance in dedifferentiated TC (13). Here, by assessment of autophagy activity in ATC tissue specimens, it was demonstrated that also in ATC tumor cells autophagy activity is clearly diminished. Considering the beneficial effects of autophagy activation observed in dedifferentiated TC, these similar findings in ATC specimens provide a compelling opportunity to study the effects of autophagy activation on ATC proliferation and differentiation.

Accordingly, the present study revealed that autophagy activation by DLCs such as digoxin, also designated as cardiac glycosides, enables reactivation of hNIS expression and iodide uptake in ATC cell lines. Although the targeted kinase inhibitors selumetinib, dabrafenib, and rapamycin, previously described as facilitating redifferentiation of RAI-refractory papillary and follicular TC, were not able to induce hNIS expression in ATC, several DLC analogs (at 5 µM and 50 µM concentrations) were demonstrated to reactivate hNIS expression and restore iodide uptake capacity in 2 ATC cell lines with different genetic backgrounds, that is, BRAF^{V600E}- and TP53-driven (8505C) and KRASG12R- and TP53-driven (Cal-62). As observed for dedifferentiated TC cell lines, similar intracellular mechanisms are activated by DLCs, including upregulation of early response genes ATF3 and cFOS, inhibition of proliferation and induction of cell cycle arrest and, to a lesser extent, cell death. Importantly, expression of the thyroid-specific transcription factors paired box gene 8 and thyroid transcription factors 1 and 2 was largely unaffected by DLC treatment, indicating their less pronounced role in restoring DLC-induced hNIS expression. In addition, expression of thyroglobulin increased after treatment with all DLCs in the 2 ATC cell lines, whereas expression of thyroid-stimulating hormone receptor and thyroid peroxidase remained undetectable. This finding indicates that the molecular machinery for iodide metabolism is partly restored by DLC treatment.

All DLC analogs used in this study, except digoxigenin, induced hNIS protein expression that was predictive for RAI uptake in both cell lines. Several hNIS protein isoforms were detected after DLC treatment: the 56-kDa nonglycosylated and 87-kDa glycosylated monomer isoforms and, with highest intensity, 120-kDa nonglycosylated hNIS dimers. Although the molecular characteristics and biologic relevance of hNIS dimerization are unclear, the occurrence and functionality of dimers and higher multimers has been described before in hNIS transfection studies and were shown to depend on the formation of disulfide bridges (26–28).

Of note, only the nonglycosylated monomeric form of the hNIS protein was induced after treatment of the 8505C cell line with digoxin, nevertheless facilitating increased iodide uptake comparable to the other DLC analogs. These findings suggest a relatively minor contribution of hNIS dimers in total iodide uptake capacity.

In contrast to dedifferentiated TC, redifferentiation agents inhibiting either MAPK/ERK, BRAF, or mammalian-target-ofrapamycin kinases were not capable of restoring functional hNIS expression in ATC cell lines. This observation could be explained by the multiple oncogenic pathways driving ATC in addition to constitutive BRAF/KRAS kinase activity, particularly caused by loss-of-function mutations in the tumor suppressor p53. Consequently, by inhibition of a single oncogenic pathway, others remain unaffected, and the intracellular processes driving proliferation and dedifferentiation are largely retained. In contrast, DLC treatment could modulate several intracellular signaling pathways in ATC cell lines simultaneously by activating intracellular Ca²⁺ signaling and downstream pathways, including autophagy, thereby robustly counteracting the ATC tumor phenotype.

Digoxin is a clinically used Na⁺/K⁺ ATPase inhibitor, inhibiting K⁺ influx and Na⁺ efflux for treatment of cardiac failure and arrhythmia. On Na⁺/K⁺ ATPase inhibition, intracellular Na⁺ concentrations rise, evoking passive exchange of intracellular Na⁺ for extracellular Ca²⁺ through the Na⁺/Ca²⁺ exchanger (NCX), ultimately leading to elevated intracellular Ca²⁺ concentrations. It is well established that intracellular Ca²⁺ acts as a second messenger and influences numerous cellular processes, including regulation of gene expression, proliferation, differentiation, migration, and apoptosis (29,30). Intracellular Ca²⁺ activates autophagy by triggering calcineurin, ERK, adenosine monophosphate-activated protein kinase, and the transcription factor EB (31,32). In addition, intracellular signaling is activated that partly resembles the thyroidstimulating hormone receptor pathway, involving cyclic adenosine monophosphate and its response-element-binding protein (33,34). Eventually, this process leads to activation of early-response genes ATF3 and cFOS, both transcription factors that are known to drive hNIS expression by triggering the CRE-like sequence of the hNIS upstream enhancer in cooperation with the thyroid-specific transcription factor paired box gene 8 (35,36).

Other previously reported efforts to restore functional hNIS expression in ATC comprise hNIS upregulation by gene transduction and, pharmacologically, by the use of histone deacetylase inhibitors or estrogen-related receptor γ -antagonists (*37,38*). It remains to be established whether combination therapies targeting Na⁺/K⁺ ATPases, histone deacetylases, and estrogen-related receptor γ induce synergistic effects on hNIS expression and iodide uptake in ATC.

CONCLUSION

Taken together, Na⁺/K⁺ ATPase inhibition by DLCs restores functional hNIS expression and the capacity for iodide uptake in ATC cell lines. Confirmation of these findings in mouse models and human studies is warranted. In addition, concerning the narrow therapeutic index of DLCs and the notion that the required DLC concentration for in vitro redifferentiation (5 μ M) exceeds the maximally tolerated dose in humans, future studies should address the dosage required for restoring functional hNIS in ATC in vivo in relation to its toxicity. DLCs could represent a promising adjunctive therapy for restoring iodide avidity within the full spectrum from RAI-refractory dedifferentiated to ATC.

DISCLOSURE

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