The Acute Inhibitory Effect of Iodide Excess on Sodium/Iodide Symporter Expression and Activity Involves the PI3K/Akt Signaling Pathway

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lodide (I⁻) is an irreplaceable constituent of thyroid hormones and an important regulator of thyroid function, because high concentrations of I⁻ down-regulate sodium/iodide symporter (NIS) expression and function. In thyrocytes, activation of phosphatidylinositol 3-kinase (PI3K)/protein kinase B (Akt) cascade also inhibits NIS expression and function. Because I⁻ excess and PI3K/Akt signaling pathway induce similar inhibitory effects on NIS expression, we aimed to study whether the PI3K/Akt cascade mediates the acute and rapid inhibitory effect of I⁻ excess on NIS expression/activity. Here, we reported that the treatment of PCCI3 cells with I⁻ excess increased Akt phosphorylation under normal or TSH/ insulin-starving conditions. I⁻ stimulated Akt phosphorylation in a PI3K-dependent manner, because the use of PI3K inhibitors (wortmannin or 2-(4-Morpholinyl)-8-phenyl-4H-1-benzopyran-4-one) abrogated the induction of I⁻ effect. Moreover, I⁻ inhibitory effect on NIS expression and function were abolished when the cells were previously treated with specific inhibitors of PI3K or Akt (Akt1/2 kinase inhibitor). Importantly, we also found that the effect of I⁻ on NIS expression involved the generation of reactive oxygen species (ROS). Using the fluorogenic probes dihydroethidium and mitochondrial superoxide indicator (MitoSOX Red), we observed that I⁻ excess increased ROS production in thyrocytes and determined that mitochondria were the source of anion superoxide. Furthermore, the ROS scavengers N-acetyl cysteine and 2-phenyl-1,2-benzisoselenazol-3-(2H)-one blocked the effect of I⁻ on Akt phosphorylation. Overall, our data demonstrated the involvement of the PI3K/Akt signaling pathway as a novel mediator of the I⁻-induced thyroid autoregulation, linking the role of thyroid oxidative state to the Wolff-Chaikoff effect. (Endocrinology 155: 1145-1156, 2014)

odide (I⁻) is essential for thyroid hormone biosynthesis. I⁻ is actively transported across the basolateral membrane of thyroid follicular cells by the sodium/iodide symporter (NIS), a plasma membrane glycoprotein that plays a critical role in thyroid physiology and pathophysiology (1, 2). NIS-mediated I⁻ uptake constitutes the first step in the biosynthesis of the only iodine-containing hormones in vertebrates, T_3 and T_4 .

NIS expression in thyroid cells is regulated by several signal pathways, including the phosphatidylinositol 3-kinase (PI3K)/protein kinase B (Akt) pathway, which has an important role in the proliferation and differentiation of

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Abbreviations: Akt, protein kinase B; Akti_{1/2}, Akt1/2 kinase inhibitor; DHE, dihydroethidium; DMSO, dimethylsulfoxide; Ebselen, 2-phenyl-1,2-benzisoselenazol-3-(2H)-one; FBS, fetal bovine serum; GAPDH, glyceraldehyde 3-phosphate dehydrogenase; HA, hemagglutinin; HBSS, Hanks' balanced salt solution; H₂O₂, hydrogen peroxide; I⁻, iodide; LY294002, 2-(4-morpholinyl)-8-phenyl-4H-1-benzopyran-4-one; MitoSOX Red, mitochondrial superoxide indicator; NAC, N-acetyl cysteine; Nal, sodium iodide; NHS-SS-biotin, succinimidyl 2-(biotinamido)-ethyl-1,3-dithiopropionate-biotin; NIS, sodium/iodide symporter; O₂⁻, superoxide; PEG-SOD, superoxide dismutase-polyethylene glycol; PI3K, phosphatidylinositol 3-kinase; PTEN, phosphatase and tensin homolog; PTU, propylthiouracil; ROS, reactive oxygen species.

thyrocytes (3–5). Several studies have demonstrated that PI3K/Akt activation negatively regulates NIS function and expression in thyroid cells (6–9). Growth factors, such as insulin or IGF-I, activate PI3K and inhibit the TSH/cAMP-regulated I[–] transport and NIS gene transcription (9, 10). Additionally, the use of selective PI3K inhibitors significantly increase NIS expression and function in thyrocytes (7–9, 11–13).

In addition to its essential role as a thyroid hormone constituent, I⁻ is one of the most important regulators of thyroid function. Since 1944 it is well known that I⁻ excess inhibits the biosynthesis and secretion of thyroid hormones (14). This phenomenon, known as the Wolff-Chaikoff effect (15), is transitory, and in rats, thyroid hormone biosynthesis and secretion are restored 2 days after I⁻ administration (16). The critical determinant for the Wolff-Chaikoff effect is the intracellular rather than the blood concentration of I^{-} (17). The "escape" of the Wolff-Chaikoff effect has been ascribed to an inhibition of NIS function (18-20). In fact, I⁻ excess inhibits NIS expression and activity, both at the transcriptional and posttranscriptional levels (21-26). Moreover, we recently reported that I⁻ excess promptly reduces NIS mRNA expression, stability, and translation efficiency both in vivo and in vitro (27, 28).

Although basal level of reactive oxygen species (ROS) production is important for maintaining thyroid hormone biosynthesis (29, 30), I⁻ excess increased production of ROS in thyrocytes (26, 31, 32). Several authors suggested that the increased generation of ROS triggered by I⁻ excess is responsible for its antiproliferative and cytotoxic effect on thyrocytes (29, 33, 34). Furthermore, the in-

creased production of ROS has been associated to the activation of PI3K/Akt signaling pathway in different cellular lineages (35–39).

Because I⁻ excess and PI3K/Akt signaling pathway trigger similar inhibitory effects on NIS expression and activity in thyrocytes, this study aimed to investigate whether the PI3K/Akt cascade mediates the acute I⁻-induced NIS inhibition.

Materials and Methods

Reagents and antibodies

Bovine TSH, insulin, transferrin, hydrocortisone, protease inhibitor cocktail, Akt1/2 kinase inhibitor (Akti1/2), wortmannin, 2-(4-morpholinyl)-8-phenyl-4H-1-benzopyran-4-one (LY294002), N-acetyl cysteine (NAC), 2-phenyl-1,2-benzisoselenazol-3-(2H)-one (Ebselen), dihydroethidium (DHE), superoxide dismutase-polyethylene glycol (PEG-SOD), antirabbit IgG-fluorescein isothiocyanate (FITC) antibody, and sodium iodide (NaI) were purchased from Sigma Chemical Co. Mitochondrial superoxide indicator (MitoSOX Red) was obtained from Molecular Probes. Affinity-purified polyclonal antirat NIS antibody was kindly provided by Dr Nancy Carrasco, as reported (40). Rabbit polyclonal antihemagglutinin (HA) antibody was purchased from Abcam. TRIzol, fetal bovine serum (FBS), OPTI-MEM, Platinum SYBR Green qPCR SuperMix, and LipofectAMINE 2000 reagent were from Invitrogen. Sulfo-succinimidyl 2-(biotinamido)-ethyl-1,3-dithiopropionate-biotin (NHS-SS-biotin) and streptoavidin agarose beads were purchased from Pierce Chemical Co. Polyclonal anti-Akt and antiphospho-Akt (Ser473) antibodies were from Cell Signaling Technology. Monoclonal anti-E-cadherin and anti-α-tubulin antibodies were from BD Transduction and Sigma Chemical Co, respectively. Enhanced

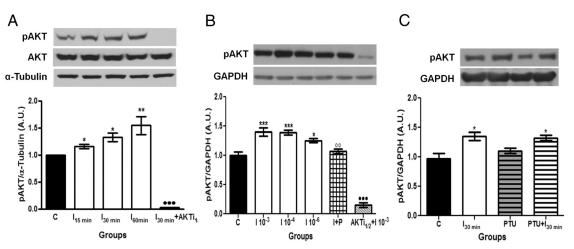


Figure 1. I^- excess induces Akt phosphorylation in PCCI3 cells. A, Western blot analysis of total or phosphorylated Akt in PCCI3 cells treated with 10^{-3} M Nal for different periods of time (15–60 min) or preincubated with the Akt inhibitor Akti_{1/2} (10µM) for 1 hour before Nal treatment for 30 minutes. α -Tubulin was used for assessing of equal loading. B, Western blot analysis of Akt phosphorylation in PCCI3 cells treated with different concentrations of Nal (10^{-6} M to 10^{-3} M) or concomitantly treated with equal concentrations of Nal and NaClO₄ (I+P) for 30 minutes. GAPDH was used as loading control. C, Western blot analysis of Akt phosphorylation in PCCI3 cells preincubated with 1mM PTU for 1 hour before 10^{-3} M Nal treatment for 30 minutes. In all cases, immunoblots shown are representative of at least 3 independent experiments. Results are expressed as means ± SEM in arbitrary units (A.U.). *, P < .05; **, P < .01; ***, P < .001 vs control (C); °°, P < .01 vs 110^{-3} , 110^{-4} ; •••, P < .001 vs 1_{15} min, 1_{30} min, 1_{60} min, 110^{-3} , 110^{-4} , 110^{-6} (ANOVA, Student-Newman-Keuls).

chemiluminescence kit was purchased from Amersham Biosciences. Monoclonal antiglyceraldehyde 3-phosphate dehydrogenase (GAPDH) and secondary horseradish peroxidase-conjugated antibodies were from Santa Cruz Biotechnology, Inc. I⁻ carrier-free Na¹²⁵I was purchased from PerkinElmer.

Cell culture and treatments

PCCl3 thyroid cells were cultured in Ham's F12 medium supplemented with 5% FBS, 1-mU/mL bovine TSH, $10-\mu g/mL$ insulin, $5-\mu g/mL$ transferrin, and 10nM hydrocortisone at 37°C in a 95% air/5% CO₂ atmosphere. For experiments under starving conditions, PCCl3 cells at 70% confluence were shifted to the same medium without TSH and insulin, supplemented with 0.2% of FBS and cultured for 5 days before treatment (41, 42).

Cultured cells were treated with 10^{-3} M NaI (23, 25, 26) for 15, 30, and 60 minutes. Cells were also treated with different doses of NaI (10^{-3} M to 10^{-6} M) for 30 minutes. To evaluate the role of intracellular I⁻ excess, cells were concomitantly treated with equal concentrations of sodium iodide (10^{-3} M) and so-dium perchlorate (10^{-3} M), a competitive NIS inhibitor (43, 44). To investigate the participation of the organified I⁻ in the induction of Akt phosphorylation, cells were incubated with 1mM propylthiouracil (PTU), which blocks I⁻ organification, for 1

hour before NaI treatment $(10^{-3}M)$ for 30 minutes. When signaling pathways were studied, cells were preincubated with vehicle (dimethylsulfoxide [DMSO]) or specific inhibitors ($10\mu M$ Akti_{1/2}, $0.5\mu M$ wortmannin, and $10\mu M$ LY294002) 1 hour before NaI treatment.

Transient transfection assays and efficiency analysis

PCCl3 cells cultured in starving medium for 3 days were transiently transfected with LipofectAMINE 2000 reagent following the manufacturer's recommendation. Briefly, cells seeded into 12-well dishes were transfected with HA-tagged rat NIS cDNA cloned into pcDNA3.1 vector (45). After transfection, cells were maintained in starving medium for 48 hours before NaI treatment.

The transfection efficiency was accessed by flow cytometry, as previously reported (45). Cells were fixed in 2% paraformaldehyde and stained with anti-HA tag antibody under nonpermeabilized conditions. Thereafter, the cells were incubated with an antirabbit IgG-fluorescein isothiocyanate (FITC) antibody. The fluorescence of 10^4 cells was assayed on a FACSCalibur flow cytometer. The results were evaluated using the Flowing Soft-

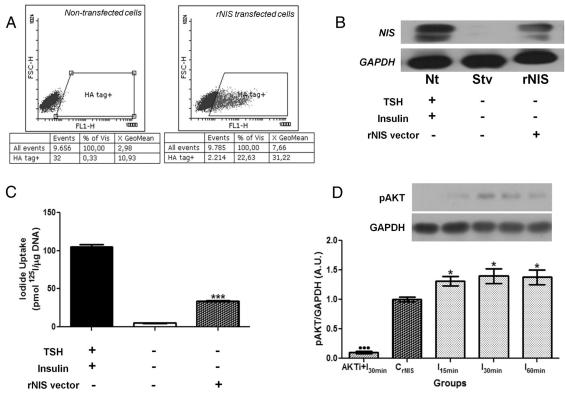


Figure 2. High concentration of I⁻ induces Akt phosphorylation in TSH/insulin-starved NIS-transfected PCCI3 cells. A, Transfection efficiency analysis performed by flow cytometry analysis under nonpermeabilized condition using an anti-HA antibody. B, Western blot analysis of total NIS expression. Immunoblots are representative of at least 3 independent experiments. GAPDH was used as loading control. Nt, nontransfected cells; Stv, cells starved for 5 days; rNIS, rat NIS-transfected cells starved for 5 days. C, Steady-state I⁻ uptake in starved PCCI3 either transfected or not with rat NIS cDNA. I⁻ uptake was expressed as pmol I⁻/µg DNA. Each value represents the mean ± SEM of 3 independent experiments performed in triplicate. ***, P < .005 vs Stv (ANOVA, Student-Newman-Keuls). D, Western blot analysis of Akt phosphorylation in starved PCCI3 transiently expressing NIS cDNA treated with 10^{-3} M Nal for the indicated time points. One group of cells was preincubated with 10μ M Akti_{1/2} for 1 hour before Nal treatment (Akti+I₃₀ min). GAPDH was analyzed as loading control. Immunoblots are representative of at least 3 independent experiments. Results are expressed as means ± SEM in arbitrary units (A.U.). *, P < .05 vs C rNIS; •••, P < .001 vs I_{15 min}, I_{30 min}, I_{60 min}, and C rNIS (ANOVA, Student-Newman-Keuls). C rNIS, control PCCI3 expressing NIS cDNA.

ware 2.4.1 (Turku Centre for Biotechnology, University of Turku, Finland).

Measurement of intracellular superoxide (O_2^-) production

Detection of O_2^- generation in PCCl3 cells was performed using the O_2^- indicator DHE. Briefly, cells were incubated with 5μ M DHE for 10 minutes before NaI treatment. Fluorescence was evaluated under a rhodamine filter in a ZEISS Axiovert 100M fluorescence microscope and Image-Pro Plus software. PEG-SOD enzyme (100 U/mL) was used to confirm that the red signal observed was related to O_2^- production (46).

To evaluate the role of ROS production on Akt phosphorylation, PCCl3 cells were pretreated with the nonspecific antioxidant NAC (1mM) (47), or the glutathione peroxidase mimetic Ebselen (30μ M) (48), for 1 hour before NaI treatment.

Measurement of mitochondrial O₂⁻ production

 O_2^- production was evaluated by flow cytometry and confocal fluorescence microscopy analysis using MitoSOX Red probe (49). Briefly, PCCl3 cells were treated with NaI and then incubated with 5µM MitoSOX Red probe for 20 minutes. The fluorescence of 10⁴ events per sample was acquired using Guava easyCyte flow cytometer and analyzed with GuavaExpress Pro Software. Confocal microscopy analysis was performed on a Zeiss LSM 510 laser-scanning confocal microscope (Carl Zeiss).

RNA extraction and real-time PCR assay

Total RNA was extracted using TRIzol following the manufacturer protocol. NIS mRNA level was evaluated by real-time PCR, as previously described (27, 28). Real-time PCR amplifications were performed in duplicate using Platinum SYBR Green qPCR SuperMix-UDG. Gene-specific primers sequences are described in Supplemental Table 1, published on The Endocrine Society's Journals Online web site at http://endo.endojournals. org. Relative changes in NIS expression were calculated using the $2^{-\Delta\Delta Ct}$ method using ribosomal protein L19 as internal loading control. Analysis of melting curves resulted in a single peak for each pair of primers, indicating that a single PCR product was amplified.

Protein extraction and Western blot analysis

PCCl3 cells were lysed in an ice-cold homogenization solution containing 1% Triton X-100 in PBS supplemented with 1mM NaOV₃, 1mM NaF, 0.1mM phenylmethanesulfonyl fluoride, and protease inhibitor cocktail. Thirty micrograms of protein were diluted in loading buffer and boiled for 5 minutes. Proteins were resolved by 10% sodium dodecyl sulfate-PAGE and then transferred onto nitrocellulose membranes. Membranes were blocked and incubated with 1- μ g/mL anti-phospho-Akt (Ser473) or 1- μ g/mL anti-Akt antibodies. Equal loading was evaluated by stripping and reprobing the same membrane with 1- μ g/mL anti-GAPDH or 0.2- μ g/mL anti- α -tubulin antibodies. Blots were developed using the enhanced chemiluminescence kit. Blots densitometry was analyzed using the ImageJ software (National Institutes of Health).

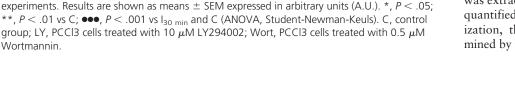
Cell surface biotinylation

PCCl3 cells were washed with ice-cold PBS supplemented with 1mM MgCl₂ and 0.1mM CaCl₂ (PBS-CM) and incubated in 2mM CaCl₂, 150mM NaCl, and 20mM HEPES (pH 8.5) containing 1.0-mg/mL Sulfo-NHS-SS-biotin for 30 minutes at 4°C. Sulfo-NHS-SS-biotin excess was quenched in PBS-CM containing 100mM glycine. Cells were lysed in 50mM Tris-HCl (pH 7.5), 150mM NaCl, 5mM EDTA, 1% Triton X-100, 0.1% sodium dodecyl sulfate, and protease inhibitors; 150 μ g of total protein were incubated overnight at 4°C with streptavidin agarose beads. Beads were washed, and adsorbed proteins were eluted with sample buffer at 75°C for 5 minutes. NIS content at the plasma membrane was analyzed by Western blotting, using 0.4- μ g/mL anti-NIS antibody. Equal loading was evaluated

by stripping and reprobing the same membrane with $0.2-\mu$ g/mL anti-E-cadherin antibody.

I⁻ uptake assay

I⁻ transport assays were performed as previously described (50). PCCl3 cells were treated with DMSO (vehicle), Akti_{1/2} (10 μ M), LY294002 (10 μ M), or wortmannin $(0.5 \mu M)$ for 1 hour. Thereafter, the cells were incubated with 10^{-3} M sodium iodide for 30 minutes. Cells were washed 10 times with Hanks' balanced salt solution (HBSS) to remove any intracellular remaining free I⁻ and incubated with 20µM NaI supplemented with 50 μ Ci/ μ mol I⁻ carrier-free Na¹²⁵I in HBSS for 30 minutes. Cells were washed twice with cold HBSS. The amount of ¹²⁵I⁻ accumulated in the cells was extracted with ice-cold ethanol and quantified in a γ -counter. For standardization, the DNA amount was determined by the diphenylamine method af-



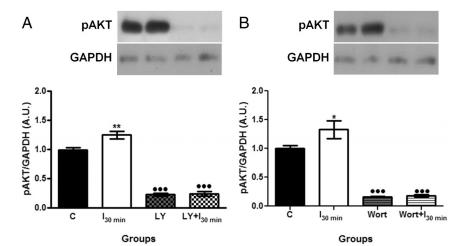


Figure 3. I⁻-induced Akt phosphorylation is PI3K dependent. Western blot analysis of Akt

GAPDH expression. Immunoblots shown are representative of at least 3 independent

phosphorylation in PCCI3 cells preincubated with 10μ M LY294002 (A) or 0.5μ M wortmannin (B)

for 1 hour before 10⁻³M NaI treatment for 30 minutes. Equal loading was evaluated assessing

ter trichloroacetic acid precipitation. Results were expressed as picomoles of I⁻ per μ g DNA (pmol/ μ g DNA).

Statistical analysis

All data were reported as means \pm SEM. Statistical analysis was performed by GraphPad Prism (GraphPad Software) from at least 3 independent experiments. Data were subjected to unpaired one-way ANOVA followed by Student-Newman-Keuls post hoc test. Differences were considered statistically significant at P < .05.

Results

I⁻ excess increases Akt phosphorylation

PI3K/Akt signaling pathway has been reported as a significant down-regulator of NIS gene expression in thyroid cells under different functional conditions (8, 9, 11). Therefore, we sought to study whether this signaling pathway is the mediator of decreased NIS expression induced by I⁻ excess in thyroid cells. We examined Akt phosphorylation, a well-know target of PI3K, in I⁻ excess-treated thyroid cells. Incubation of PCCl3 cells with 10^{-3} M NaI significantly increased Ser473 Akt phosphorylation at all time points analyzed (15–60 min) (Figure 1A). I⁻ excess-induced Akt phosphorylation was completely abolished by the specific Akt inhibitor Akti_{1/2} (Figure 1A). Total Akt levels were not changed by NaI treatment (Figure 1A).

We also observed that different concentrations of NaI $(10^{-6}M \text{ to } 10^{-3}M)$ induced Akt phosphorylation, although $10^{-3}M$ presented the most prominent induction (Figure 1B). This effect was abolished in the presence of the competitive NIS inhibitor perchlorate (Figure 1B).

Preincubation of PCCl3 cells with PTU has not altered the induction of Akt phosphorylation by NaI treatment (Figure 1C), suggesting that the effect of I⁻ does not require an organification step. Moreover, the effect of I⁻ on Akt phosphorylation was not related to a change in the osmolarity of the medium, because the treatment of PCCl3

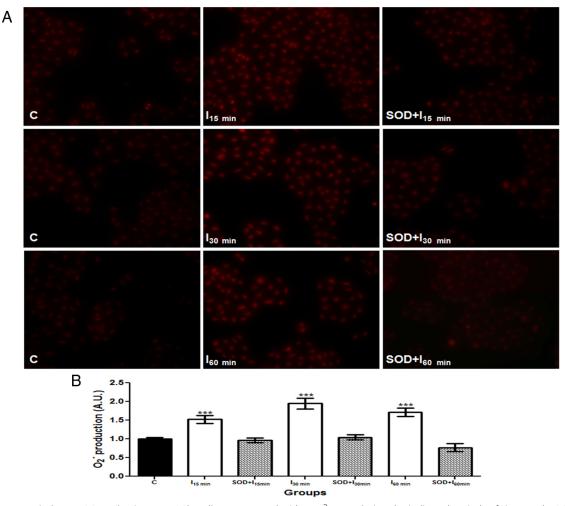


Figure 4. I⁻ excess induces ROS production. A, PCCI3 cells were treated with 10^{-3} M Nal during the indicated periods of time, and ROS production was evaluated using the fluorescent dye DHE (5 μ M). Specificity of ROS production was evaluated in the presence of 100-U/mL PEG-SOD (SOD). B, Relative quantification of O₂⁻ production in PCCI3 cells treated with 10^{-3} M Nal. Results are expressed as means ± SEM in arbitrary units (A.U.). Four independent experiments were performed, in triplicate. ***, *P* < .001 vs C (ANOVA, Student-Newman-Keuls). C, control group.

cells with 10^{-3} M NaCl has not induced Akt phosphorylation (Supplemental Figure 1).

In thyroid cells, Akt phosphorylation is stimulated by TSH or insulin/IGF-I treatment (9, 11). Therefore, we evaluated I⁻-induced Akt phosphorylation in PCCl3 cells cultured in the absence of TSH and insulin for 5 days (starved condition). Because NIS expression is severely reduced in the absence of TSH (51), PCCl3 cells were transiently transfected with NIS cDNA to ensure I⁻ accumulation. After transfection, we observed that 20%–25% of the transfected cell successfully expressed NIS at the plasma membrane (Figure 2A). Moreover, the partial recovery of NIS expression and activity was confirmed by Western blotting (Figure 2B) and ¹²⁵I⁻ uptake assay (Figure 2C). Interestingly, NaI treatment induced Akt phosphorylation, at all analyzed time points, even in the absence of TSH and insulin (Figure 2D). It is worth noting that under starved condition in non-NIS-transfected PCCl3 cells, NaI treatment was able to induce a minimal, although significant, Akt phosphorylation (Supplemental Figure 2).

PI3K-dependent I⁻-induced Akt phosphorylation

The canonical pathway for Akt phosphorylation involves the upstream activation of PI3K (52, 53). Therefore, we evaluated whether PI3K signaling is involved in the phosphorylation of Akt induced by I⁻ excess. PCCl3 cells were preincubated with the well-known PI3K inhibitors, wortmannin and LY294002 before NaI treatment. As shown in Figure 3, PI3K inhibition markedly reduced Akt phosphorylation in response to I⁻ excess, indicating that I⁻ induction of Akt phosphorylation is PI3K dependent.

I⁻-induced Akt phosphorylation relies on ROS production

Recently, Yao et al (32) demonstrated that high concentrations of I⁻ significantly increased ROS production in FRTL-5 thyroid cells. Therefore, we evaluated ROS production in PCCl3 cells in response to I⁻ excess and in shorter periods of time than the ones assessed before. As shown in Figure 4, the treatment of PCCl3 cells with NaI for 15–60 minutes increased the ROS production as evaluated by oxidation of the fluorescent indicator DHE. Preincubation of cells with PEG-SOD supported that NaI

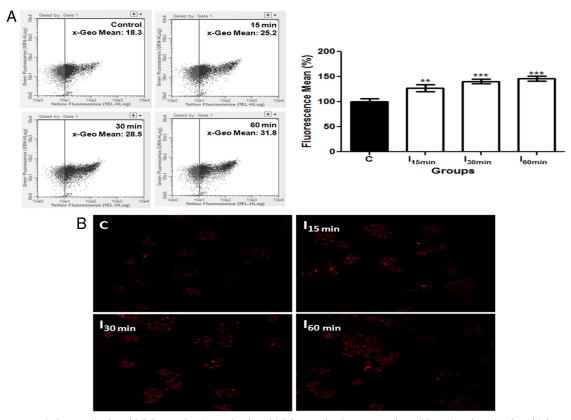


Figure 5. I^- excess induces mitochondrial O_2^- production. Mitochondrial O_2^- production was evaluated by using the mitochondrial superoxide indicator MitoSOX Red. A, left panel, Representative dot plots analysis of flow cytometry data obtained from cells incubated with 10^{-3} M Nal for 15, 30, and 60 minutes. Right panel, Relative quantification of fluorescence intensity means from flow cytometry data. Ten thousand events were evaluated per sample. Three independent experiments were performed in triplicate. **, P < .01; ***, P < .001 vs C (ANOVA, Student-Newman-Keuls). B, Evaluation of mitochondrial O_2^- production by confocal fluorescence microscope. Pictures are representative of 2 independent experiments performed in triplicate. C, control group.

treatment specifically increased O_2^- production, a precursor of hydrogen peroxide (H₂O₂) (Figure 4). Furthermore, we have observed that the treatment with lower doses of NaI (10⁻⁶M and 10⁻⁴M) also increased ROS production (Supplemental Figure 3). In addition, mitochondria seem to be the source of O_2^- production in PCCl3 cells exposed to I⁻ excess, as demonstrated by using the fluorogenic probe MitoSOX Red in flow cytometry (Figure 5A) and confocal fluorescence microscopy (Figure 5B) assays. As apoptosis interfere with mitochondrial function, we evaluated whether I⁻ excess affects PCCl3 cells viability or induces apoptosis. As shown in Supplemental Figure 4, NaI (10⁻³M) treatment did not produce any significant effect on the analyzed parameters.

Thereafter, we evaluated the role of I⁻-induced ROS production on Akt phosphorylation. PCCl3 cells were incubated with NaI in the presence of ROS scavengers, NAC or Ebselen. As shown in Figure 6A, ROS scavengers impaired the induction of Akt phosphorylation by NaI treatment (Figure 6A). Ebselen treatment also reduced the I⁻-induced Akt phosphorylation in NIS-transfected PCCl3 cells cultured in starving medium for 5 days (Figure 6B).

PI3K/Akt signaling pathway mediates I⁻-induced NIS down-regulation

Inhibition of NIS expression is one of the main effects of I⁻ excess in thyrocytes. Thus, we evaluated whether the PI3K/Akt signaling pathway was involved in the reduction of NIS mRNA levels induced by I⁻ excess. As expected, a rapid reduction of NIS mRNA content was observed in PCCl3 cells treated with 10^{-3} M NaI for 30 minutes, but interestingly, the effect of I⁻ excess was abolished in the presence of PI3K or Akt inhibitors (Figure 7).

Previous studies have reported that high concentrations of I⁻ reduce NIS content at the thyrocytes plasma membrane (26, 54). As such, we observed a significant and rapid decrease of NIS protein expression at the plasma membrane of thyroid cells in response to NaI treatment for 30 minutes (Figure 8, A and B). This effect was prevented when the PCCl3 cells were pretreated with the Akt inhibitor Akti_{1/2} (Figure 8A) or the PI3K inhibitor wortmannin (Figure 8B).

Inhibition of I⁻ uptake by high concentrations of I⁻ seems to be an adaptive response to reduce the intracellular I⁻ content (18, 19). In agreement with a reduced expression of NIS at the plasma membrane, NaI treatment rapidly (30 min) decreased I⁻ uptake in thyrocytes, and this effect was abolished in the presence of PI3K/Akt inhibitors (Figure 8, C–E). Interestingly, our data also suggest that, at short periods of time, inhibition of PI3K/Akt signaling did not affect I⁻ uptake in control cells.

Discussion

 I^- is one of the most important NIS regulators in thyrocytes, and several studies have investigated the mechanisms that underlie the reduction of NIS expression and activity in response to high I^- concentrations (19–26). Transcriptional repression induced by I^- has been postulated as responsible

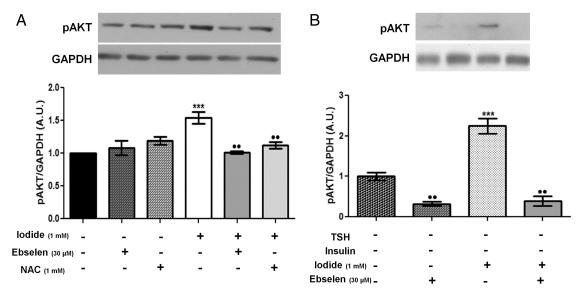


Figure 6. I⁻-induced ROS production increases Akt phosphorylation. A, Western blot analysis showing Akt phosphorylation in PCCI3 cells preincubated with 1mM NAC or 30μ M Ebselen for 1 hour before 10^{-3} M Nal treatment for 30 minutes. GAPDH was used as loading control. B, Western blot analysis of Akt phosphorylation in rat NIS-transfected PCCI3 cells cultured under starving medium. Cells were preincubated with 30μ M Ebselen for 1 hour and then treated with 10^{-3} M Nal for 30 minutes. Immunoblots shown are representative of at least 2 independent experiments. Results are expressed as means ± SEM in arbitrary units (A.U.). ***, P < .001 vs C; ••, P < .01 vs I_{30 min} (ANOVA, Student-Newman-Keuls). C, control group.

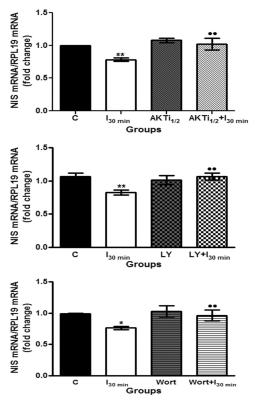


Figure 7. PI3K/Akt signaling pathway mediates I[–]-reduced NIS mRNA abundance. PCCI3 cells were preincubated with the Akt inhibitor Akti_{1/2} (10 μ M), or the PI3K inhibitors LY294002 (10 μ M) and wortmannin (0.5 μ M) for 1 hour before NaI treatment (10^{–3}M) for 30 minutes. NIS mRNA levels relative to those of RPL19 were evaluated by real-time PCR. Values are indicated as fold change relative to the mRNA levels of cells treated with vehicle (DMSO). *, *P* < .05; **, *P* < .01 vs C; ••, *P* < .01 vs I_{30 min} (ANOVA, Student-Newman-Keuls). C, control group; LY, PCCI3 cells treated with 10 μ M LY294002; Wort, PCCI3 cells treated with 0.5 μ M Wortmannin; RPL19, ribosomal protein L19.

for decreasing NIS mRNA levels in thyroid cells (22, 55). However, recent studies demonstrated that prompt post-transcriptional events, such as reduction of NIS mRNA stability and translation efficiency, are triggered in response to I^- administration (26–28, 56).

In this study, we demonstrated that the rapid inhibitory effect of high concentrations of I⁻ on NIS expression and function involve the participation of PI3K/Akt signaling pathway. In thyrocytes, PI3K/Akt signaling cascade has been associated with cell-cycle progression, cell survival, and differentiation (3–5, 57). However, the knowledge concerning PI3K/Akt pathway in regulating thyroid cell differentiation is still limited. Several studies have shown that PI3K/Akt pathway reduces TSH-stimulated NIS expression through transcriptional mechanisms (9, 11, 12). However, TSH, the main positive regulator of NIS expression in thyroid cells, activates PI3K/Akt signaling pathway through the binding on TSH receptor (6, 11). Stimulation of TSH receptor leads to the dissociation of trimeric G proteins into G α and G $\beta\gamma$ subunits. The release of G $\beta\gamma$ dimmers stimulates PI3K/Akt signaling pathway in a cAMP-independent manner and down-regulates NIS gene expression by decreasing paired box gene 8 binding to the NIS promoter (11).

Our results demonstrated that I⁻ excess rapidly increased Akt phosphorylation in PCCl3 cells. This effect was observed even in the presence of low concentrations of I⁻ and in short periods of time. Although the important role of iodolipids on I⁻-induced effects in thyrocytes is well described (58), our results indicated that the iodolipids production through I⁻ organification may not be relevant for the induction of Akt phosphorylation by I⁻ excess. Moreover, the effect of I⁻ was dependent on intracellular I⁻ excess, because it was prevented by blocking I⁻ transport with perchlorate treatment (43, 44). We also evaluated whether I⁻ effects on Akt phosphorylation were dependent on TSH or insulin action. Our data demonstrated that NaI treatment has induced Akt phosphorylation even in the absence of both hormones, indicating a direct role of I⁻ on this cascade. In addition, we confirmed that I⁻-induced Akt phosphorylation was dependent on PI3K activity, because I⁻ effect was abolished when the cells were preincubated with PI3K inhibitors, wortmannin or LY294002.

Hyperactivity of PI3K/Akt signaling pathway has been implicated in the progression of thyroid cancer and correlates with reduced NIS expression at the plasma membrane (59-61). In addition, previous studies have demonstrated that PI3K/Akt cascade activation is involved in the internalization of proteins (62, 63). Furthermore, in breast carcinoma cells, PI3K signaling impairs NIS glycosylation and cell surface trafficking (64). In concordance, the blockage of PI3K signaling increased NIS targeting to the plasma membrane in normal and neoplastic thyrocytes (7). In agreement, here, we demonstrated that specific PI3K/Akt inhibitors abrogated the effects of I⁻ excess on NIS mRNA and plasma membrane expression. These data strongly suggest that the PI3K/Akt signaling pathway constitutes a novel mediator of the posttranscriptional effects induced by I[–] excess.

The rapid reduction of NIS activity induced by I⁻ excess seems to be coherent, because I⁻ uptake is mainly mediated by NIS molecules placed at the plasma membrane (65). Moreover, the reduction of NIS activity observed when the intracellular concentration of I⁻ raises seems to be an efficient adaptation to prevent the deleterious effect of I⁻. Interestingly, we observed that blockage of PI3K/ Akt signaling pathway prevent the down-regulation of NIS function induced by I⁻. Therefore, this signaling pathway participates in the adaptation mechanism of thyroid cells to I⁻ excess. Although we demonstrated that a reduction of NIS expression at the plasma membrane closely correlate with the reduction of NIS activity observed in

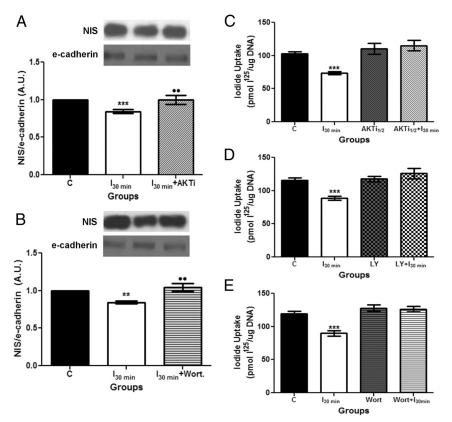


Figure 8. PI3K/Akt signaling pathway mediates I⁻-reduced NIS plasma membrane content and activity. PCCI3 cells were treated with 10 μ M Akti_{1/2}, 0.5 μ M wortmannin, or 10 μ M LY294002 for 1 hour before 10⁻³M Nal treatment for 30 minutes. A and B, NIS expression at the plasma membrane was evaluated by cell surface biotinylation, followed by Western blot analysis. The plasma membrane marker E-cadherin was used as loading control. Immunoblots shown are representative of at least 3 independent experiments. Results are expressed as means ± SEM in arbitrary units (A.U.). **, P < .01; ***, P < .001 vs C; ••, P < .01 vs I_{30 minutes} (ANOVA, Student-Newman-Keuls). C–E, Steady-state I⁻ uptake assay was performed in PCCI3 cells previously treated with Akti_{1/2} (C), LY294002 (D), or wortmannin (E). I⁻ uptake was expressed as pmol I⁻/ μ g DNA. Results are expressed as mean ± SEM of 3 independent experiments performed in triplicate. ***, P < .001 vs C; •••, P < .001 vs I_{30 min} (ANOVA, Student-Newman-Keuls). C, control group; LY, PCCI3 cells treated with 10 μ M LY294002; Wort, PCCI3 cells treated with 0.5 μ M Wortmannin.

response to NaI treatment, we do not discard that additional mechanisms might be involved in the regulation of NIS activity, such as inactivation of NIS molecules at the plasma membrane. Particularly, Leoni et al (26) predicted 2 intracellularly oriented and potentially redox-sensitive cysteine residues whose oxidation may result in a rapid inactivation of NIS protein.

NIS up-regulation induced by PI3K/Akt inhibition involves a transcriptional effect and, therefore, requires long periods of time to be observed (7–9). These results are consistent with our data, because the treatment of control PCCl3 cells with PI3K/Akt inhibitors for 1 hour did not alter NIS expression or activity. However, we have observed that the inhibitory effect of I⁻ excess on NIS was reversed by the preincubation of thyroid cells with PI3K/Akt inhibitors. Altogether, these data suggest that besides the well-described transcriptional events, the activation of

PI3K/Akt signaling pathway is also involved in the I⁻-induced posttranscriptional events.

The mechanism by which I⁻ induced Akt phosphorylation in our experimental model was an intriguing issue. Previous studies have demonstrated that angiotensin I, cadmium, and other factors were able to induce PI3K/Akt signaling pathway in different cell lineages through the ROS production (35–39). In agreement, our data support the hypothesis that I⁻-induced ROS generation is crucial for stimulating Akt phosphorylation in thyrocytes, as pretreatment of PCCl3 cells with ROS scavengers, such as NAC or Ebselen, abrogated the effect of I⁻ on Akt phosphorylation.

ROS generation is extremely important for thyroid hormone biosynthesis, particularly H_2O_2 , which plays an essential role in normal thyroid physiology, as a cofactor required for thyroid peroxidase-mediated I⁻ organification (66). The exposure of thyrocytes to I⁻ excess increases ROS production, intracellular oxidation levels, and therefore cell toxicity (29, 33, 34). However, the timing of I⁻-induced ROS generation is still controversy. Leoni et al (26) did not observe significant alteration on ROS production until 6

hours after I⁻ treatment, a modest effect after 24 hours (1.2-fold), reaching the maximum of ROS production at 48 hours. Nevertheless, the authors evaluated ROS production using the general oxidative stress indicator carboxy-2',7'-dichlorodihydrofluorescein diacetate, which is not very sensitive to O_2^- anion (67). In sharp contrast, Yao et al (32) demonstrated increased mitochondrial $O_2^$ generation in FRTL-5 after 2 hours of I⁻ treatment, supporting our hypothesis that I⁻ could rapidly induce ROS production. In fact, posttranscriptional events are rapid and transiently elicited to keep cellular homeostasis, but they can be followed by different events that require longer periods of time to be triggered. Moreover, in agreement with Yao et al (32), our results demonstrated an impairment of ROS production in thyrocytes previously treated with PEG-SOD, suggesting that the source of ROS could be the mitochondria. In addition, the oxidation of MitoSOX Red probe confirmed increased mitochondrial O_2^- production in response to NaI treatment.

It has been described that redox status interferes with the activity of several members of protein tyrosine phosphatases through the interaction with reactive cysteine residues (68-71). Phosphatase and tensin homolog (PTEN) is a well-known phosphoinositide-3-phosphatase that inhibits Akt activation (72). Several studies demonstrated that PTEN activity is regulated by different posttranslational mechanisms, as its inhibition through the reversible oxidation of catalytic cysteine residues in the presence of H_2O_2 (73, 74). Thus, we hypothesize that the increased ROS production induced by I⁻ might interfere with PTEN or other phosphatase activity leading to an increased Akt phosphorylation in thyrocytes. Therefore, these data reinforce the important role of thyroid redox state in the inhibition of NIS expression/activity induced by I^- excess (26).

In summary, our study points to a novel role of PI3K/ Akt signaling pathway as a mediator of thyroid autoregulation induced by I⁻ and links the role of thyroid oxidative state to the Wolff-Chaikoff effect.

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Supplemental Material Manuscript EN-13-1665R1

"The acute inhibitory effect of iodide excess on sodium/iodide symporter expression and activity involves the PI3K/Akt signaling pathway" by Caroline Serrano-Nascimento, Silvania da Silva Teixeira, Juan Pablo Nicola, Renato Tadeu Nachbar, Ana Maria Masini-Repiso, and Maria Tereza Nunes.

Supplemental Material and Methods

Cell viability and DNA fragmentation assay. Cell viability was analyzed as previous described (28). Briefly, PCC13 cells were treated with 10^{-3} M sodium iodide for 1 h, the longest period of treatment used in this study. To assess plasma membrane integrity cells were trypsinized and re-suspended in 500 µl PBS containing 50 µl of 20 µg/ml propidium iodide. To evaluated DNA fragmentation, PCC13 cells were treated with a lysis buffer intended to disrupt the plasma membrane and allow propidium iodide to bind to intact DNA. Ten thousand events were analyzed per condition were evaluated using a FACSCalibur Flow Cytometer (Becton–Dickinson, San Juan, CA, USA). Data was analyzed using Cell Quest Software (Becton–Dickinson, San Juan, CA, USA).

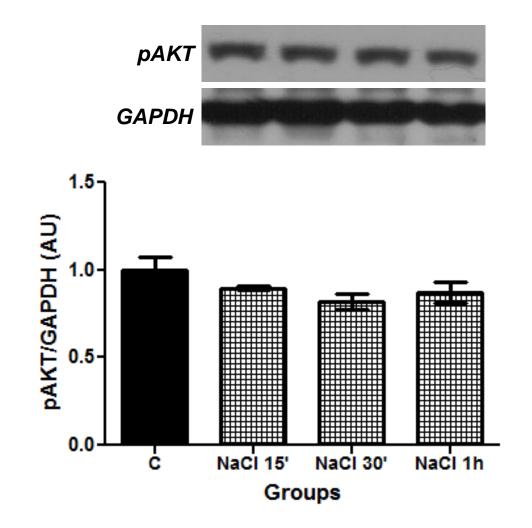
Supplemental Figures Legends

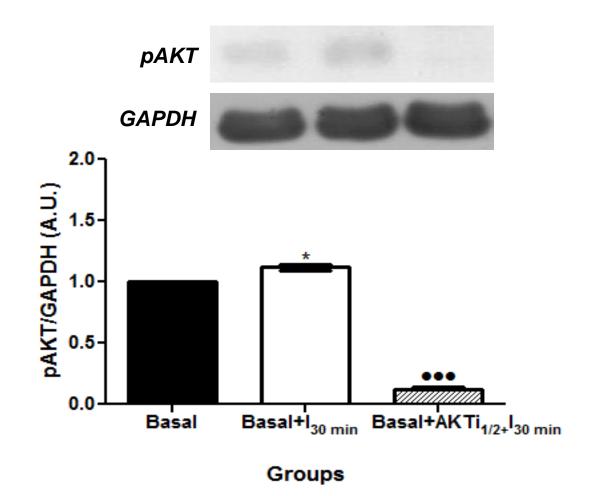
Figure 1 – *Change in osmolarity does not induce Akt phosphorylation in PCCl3 cells.* Western blotting analysis of Akt phosphorylation (p-Akt) in PCCl3 cells treated with 10^{-3} M NaCl for different periods of time (15 to 60 min). GAPDH was used for assessing equal loading. Immunoblots shown are representative of two independent experiments. Results are expressed as means ± SEM in arbitrary units (AU). P> 0.05 (ANOVA, Student–Newman–Keuls).

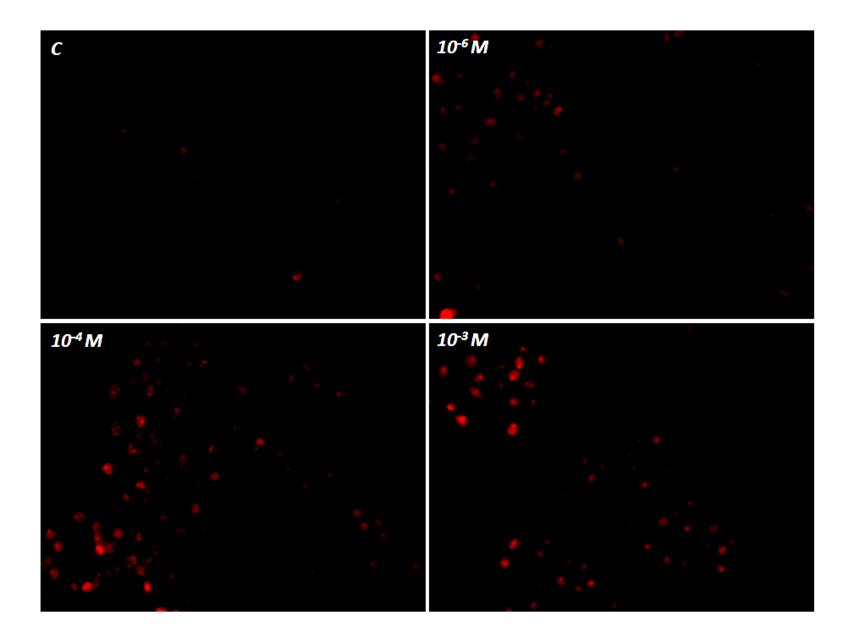
Figure 2 – *I induces Akt phosphorylation in starved PCCl3 cells.* Western blotting analysis of Akt phosphorylation (p-Akt) in PCCl3 cells cultured under starving condition for 5 days and treated with 10^{-3} M NaI for 30 min. GAPDH was used as loading control. Immunoblots shown are representative of at least two independent experiments. Results are expressed as mean \pm SEM in arbitrary units (AU).* P< 0.05 vs. Basal;••• P< 0.001 vs. Basal+I_{30min}(*ANOVA, Student–Newman–Keuls*).

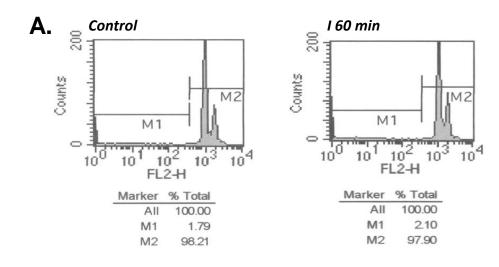
Figure 3 – *ROS production is stimulated by low concentrations of* Γ . PCC13 cells previously incubated with the fluorescent dye DHE (5 μ M), were treated with 10⁻³ to 10⁻⁶ M NaI for 30 min. ROS production was evaluated under a rhodamine filter in a common fluorescence ZEISS Axiovert 100 M Microscope and Image-Pro Plus software. Pictures are representative of two independent experiments performed in triplicate.

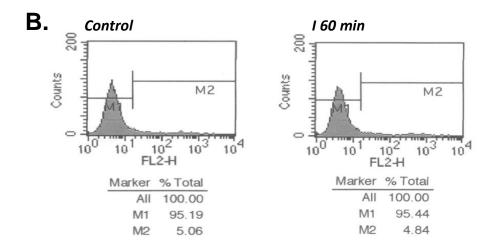
Figure 4 – Γ excess does not impair plasma membrane integrity and DNA fragmentation in PCCl3 cells. Membrane integrity and DNA fragmentation were evaluated in PCCl3 cells treated with 10⁻³ M NaI for 1 h by flow cytometry, using propidium iodide. Results are representative of two independent experiments performed in triplicate. **A.** M1 and the M2 axis represent cells containing fragmented or intact DNA, respectively. DNA integrity was preserved after iodide excess treatment. P> 0.05 (unpaired two-tailed Student's t-test). **B.** M1 axis represents the number of cells with intact plasma membrane, whereas M2 axis shows cells with altered plasma membrane integrity. No alteration of PCCl3 plasma membrane integrity was induced by iodide excess treatment. P> 0.05 (unpaired two-tailed Student's t-test).











SUPPLEMENTAL TABLE: Primers information

2

1

3 Supplemental Table 1 – Primer sets used for NIS mRNA expression analysis in PCCl3 cells by Real-

4 Time PCR.

Primer	GenBank
	Reference
FWD - 5'-GGCCCGCAAGAAGCTACTG-3'	NM_031103
REV - 5'-TTTCGTGCTTCCTTGGTCTTAGA-3'	
FWD – 5'-AGCCTCGCTCAGAACCATTC-3'	NM_052983
REV – 5'-GTGTACCGGCTCCGAGGAT-3'	
	FWD - 5'-GGCCCGCAAGAAGCTACTG-3' REV - 5'-TTTCGTGCTTCCTTGGTCTTAGA-3' FWD – 5'-AGCCTCGCTCAGAACCATTC-3'