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Abstract

Small cell lung cancer (SCLC) is an aggressive disease, representing 15% of all cases of lung cancer, has high metastatic potential and low prognosis that urgently demands the development of novel therapeutic approaches. One of the proposed approaches has been the down-regulation of BCL2, with poorly clarified and controversial therapeutic value regarding SCLC. The use of anti-BCL2 small interfering RNA (siRNA) in SCLC has never been reported. The aim of the present study was to select and test the in vitro efficacy of anti-BCL2 siRNA sequences against the protein and mRNA levels of SCLC cells, and their effects on cytotoxicity and chemosensitization. Two anti-BCL2 siRNAs and the anti-BCL2 G3139 oligodeoxynucleotide (ODN) were evaluated in SCLC cells by the simultaneous determination of Bcl-2 and viability using a flow cytometry method recently developed by us in addition to Western blot, real-time reverse-transcription PCR, and cell growth after single and combined treatment with cisplatin. In contrast to previous reports about the use of ODN, a heterogeneous and up to 80% sequence-specific Bcl-2 protein knockdown was observed in the SW2, H2171 and H69 SCLC cell lines, although without significant sequence-specific reduction of cell viability, cell growth, or sensitization to cisplatin. Our results question previous data generated with antisense ODN and supporting the present concept of the therapeutic interest in BCL2 silencing per se in SCLC, and support the growing notion of the necessity of a multitargeting molecular approach for the treatment of cancer.

Key words: Small cell lung cancer; Gene silencing; Bcl-2, siRNA; Antisense ODN; Flow cytometry; Chemosensitization

Introduction

Small cell lung cancer (SCLC) is an aggressive and highly metastatic neuroendocrine histological subtype of lung cancer. The 5-year survival rate for patients diagnosed with invasive SCLC and bronchus cancer in the USA, from 1996 to 2004, was 5.9% (1). This clearly demonstrates the need for the development of novel therapeutic approaches. The BCL2 gene is overexpressed in most SCLC specimens and cell lines, as opposed to other histological lung cancer types, a fact that has led to the hypothesis that it could be implicated in the pathogenesis of this tumor (2,3). After the establishment of the oncogenic potential of Bcl-2 (4), its anti-apoptotic function further attracted the attention of researchers (5) because apoptosis deregulation has been implicated in cancer, therefore highlighting BCL2 as a potential therapeutic target in a wide variety of tumors.

In the meantime, high expectations have been created regarding cancer therapy, with the possibility of targeting any single protein within a cell using gene silencing approaches. However, concerning SCLC, combined treatment of SCLC patients with G3139 oligodeoxynucleotide (ODN; oblimersen sodium) in phase II clinical trials did not improve any of the clinical parameters assessed, with insufficient BCL2 silencing being the reason suggested to explain the results obtained (6,7). Although ODNs are still undergoing preclinical and clinical evaluation, other gene silencing molecules such as small interfering RNAs (siRNAs) are being used extensively in research both as powerful tools to decipher gene function and as promising therapeutic agents. The assessment of the therapeutic potential of anti-BCL2 gene silencing in SCLC has been restricted to the use of ODN (7-9), whereas no reports with siRNA are available. It is widely accepted that gene silencing therapies depend on the development of adequate nanocarriers to be used to achieve adequate intracellular delivery. This strategy is likely to profit from potent gene
silencing molecules such as siRNA.

The objective of the present study was to select and validate the activity of an anti-BCL2 siRNA sequence for further use in the development of ligand-mediated targeted nanocarriers, in order to silence BCL2 in SCLC. In contrast to previous reports (8,9) in which gene silencing was assessed in bulk cells by Western blot, a great advantage was obtained from a recently established flow cytometry procedure that simultaneously evaluates Bcl-2 knockdown and viability in SCLC cells (10).

Material and Methods

Material

All salts and paraformaldehyde were purchased from Merck (Germany). 4-(2-Hydroxyethyl)-1-piperazine-ethanesulfonic acid (HEPES), sodium azide, N,N',N'-tetramethylethlylenediamine, ammonium persulfate, digitonin, bovine serum albumin, RPMI 1640 with L-glutamine and without sodium bicarbonate (RPMI 1640), and penicillin-streptomycin solution (10,000 U/mL and 10 mg/mL, respectively) were from Sigma (Sigma-Aldrich Chemie GmbH, Germany). 7-Aminoactinomycin D (7-AAD) and actinomycin D were from Fluka (Sigma-Aldrich). Fetal bovine serum (FBS) was from Gibco (Invitrogen S.A., Spain).

Cell lines

The human variant SCLC cell line SW2 was kindly provided by Drs. U. Zangemeister-Wittke and R. Stahel (University Hospital of Zurich, Switzerland). The classic NCI-H69 and the variant NCI-NCI-H2171 human SCLC cell lines were from the American Type Culture Collection. Cells were cultured in RPMI 1640 supplemented with 10% (v/v) heat-inactivated FBS, 1% (v/v) penicillin-streptomycin solution, 10 mM HEPES and 0.2% NaHCO3 and maintained at 37°C in a humidified atmosphere (90%) containing 5% CO2. Cells were maintained in the exponential growth phase and periodically tested for mycoplasma contamination using the MycoAlert® mycoplasma detection kit from Cambrex Bio Science Verviers, Inc. (Belgium).

ODN and siRNA sequences

All ODN sequences exhibit full length phosphorothioate modifications. The desalted forms of ODN G3139 (5'-TCTCCCCAGCGTGCAGCATGCAATG-3') and G4126 (5'-TCTCC CAGCATGTGCCAT-3') were purchased from Microsynth (Balgach, Switzerland). The human variant SCLC cell line SW2 was kindly provided by Drs. U. Zangemeister-Wittke and R. Stahel (University Hospital of Zurich, Switzerland). The classic NCI-H69 and the variant NCI-NCI-H2171 human SCLC cell lines were from the American Type Culture Collection. Cells were cultured in RPMI 1640 supplemented with 10% (v/v) heat-inactivated FBS, 1% (v/v) penicillin-streptomycin solution, 10 mM HEPES and 0.2% NaHCO3 and maintained at 37°C in a humidified atmosphere (90%) containing 5% CO2. Cells were maintained in the exponential growth phase and periodically tested for mycoplasma contamination using the MycoAlert® mycoplasma detection kit from Cambrex Bio Science Verviers, Inc. (Belgium).

Lipoplex preparation and cell transfection

Lipoplexes were prepared with Oligofectamine™ Reagent (Invitrogen). Oligofectamine (at 15 µL/nmol ODN or at 140 µL/nmol siRNA) was diluted 5 times in HEPES buffer saline (HBS) for 5 min, mixed with ODN or siRNA solutions diluted in HBS (25 mM HEPES, 140 mM NaCl, pH 7.4) and incubated for 20 min. Complexes were added to cells in 1/5 of the final volume followed, 4 h later, by the addition of RPMI 1640 supplemented with 30% FBS to achieve a final concentration per well of 10% (v/v).

Effect on cell growth of single and combined treatments with Bcl-2 silencing agents and cisplatin

Sixteen-thousand viable cells/well in RPMI without phenol red, buffered with 25 mM HEPES and 2 g/L NaHCO3, were plated onto 96-flat well plates in quadruplicate. Cells were treated for 4 h with a fixed concentration of the silencing agent or the corresponding control complexed with Oligofectamine. After 4 h, cells were washed 3 times with RPMI without serum and further maintained in culture medium with 10% FBS. In the combined treatment schedules, cisplatin (Faulplatin® solution for iv injection, Mayne Pharma Ltd., Portugal), kindly donated by the University Hospital of Coimbra, was added 28 h after the first treatment. Ninety-six hours after the beginning of the experiment, relative cell growth inhibition was evaluated by the resazurin reduction method adapted from the AlamarBlue™ instructions (BioSource, Belgium). Briefly, plates were centrifuged, and 100 µL of the supernatant was replaced with 100 µL of a 30% (v/v) solution of resazurin (Sigma) in RPMI 1640 without FBS from a recently prepared stock solution at 0.1 mg/mL of resazurin in Dulbecco's phosphate buffer saline (PBS; 1.4 mM KH2PO4, 2.7 mM KCl, 4.3 mM Na2HPO4, 137 mM NaCl, pH 7.4). After incubation at 37°C for 2 h, absorbance was measured at 540 and 630 nm in a plate spectrophotometer (Multiscan Ex, Thermo Labsystem, Finland). Percent resazurin reduction was calculated taking into consideration the extinction coefficients of resazurin at the aforementioned wavelengths. The results were normalized with respect to the untreated control wells (% cell growth).

BCL2 mRNA evaluation by real-time reverse-transcription PCR

Cells were plated onto 6-well (0.25 or 0.5 million cells/well) or 12-well plates (0.125 or 0.25 million cells/well) and transfected as described above with the indicated concentration of ODN or siRNA. At 24 h post-transfection, cells were harvested, washed with PBS and kept at -80°C. Total RNA was isolated with the RNeasy® mini kit (QIAGEN GmbH, Germany), with in-column DNase digestion (QIAGEN GmbH) according to manufacturer instructions. The Ribogreen kit was used to quantify the isolated RNA solutions and 0.4 or 1 µg RNA was reverse transcribed using the SuperScript® III First-strand Synthesis SuperMix (Invitrogen) according to the manufacturer protocol. Quan-
titative real-time reverse-transcription PCR (qRT-PCR) was performed using the iQ SYB Green Supermix (Bio-Rad Laboratories, USA), and the iQ5 PCR detection system/iCycler® Thermal Cycler (Bio-Rad Laboratories). The optimized condition included an annealing and extension temperature of 55°C and primers at 300 nM. The following PCR primers were used: BCL2, sense 5'-CATGTGTGTTGG AGAGGCTCAA-3', and antisense, 5'-GCCGGTCCAGG TACTCAGTCA-3', and β-2-microglobulin (B2M), sense 5'-GAGTATGCGTGGCTGTG-3', and antisense, 5'-AAT CCAATGCGGACATCT-3' (Microsynth, Switzerland). The melting curve protocol was started immediately after amplification to confirm the specificity of the reaction, which was further confirmed by agarose gel analysis. The percentage of BCL2 mRNA was calculated by the Pfaffl method (11), \[ \frac{E(\text{target})}{E(\text{reference})} \times 100 \], using B2M as reference gene.

**Evaluation of Bcl-2 protein level and cell viability**

Cells were plated and treated as described in the previous section. The Bcl-2 protein level was measured by flow cytometry simultaneously with cell viability in a BD FACSCalibur™ system (BD Biosciences). Bcl-2 protein levels were determined with a phycoerythrin (PE)-conjugated mouse anti-human Bcl-2 mAb (Bcl-2 mAb-PE, IgG1k, clone Bcl-2/100) and the corresponding PE-conjugated isotype control (IgG1k, clone MOPC-21) were BD Pharmingen™ (BD Biosciences) (10).

Cell viability (%) was calculated from the ratio between the percentage of viable treated cells and viable untreated cells. In each cell region (total cell population or viable cells), the specific geometric mean of the Bcl-2 signal (%) was normalized to the corresponding region of untreated cells. The specificity of the Bcl-2 signal in treated cells was also evaluated by considering the nonspecific signal, obtained upon staining with the mAb-PE isotype. The percentage of Bcl-2-silenced cells was calculated by subtracting the frequency histogram of viable untreated cells from that of viable treated cells and normalized to the total viable population. All subtractions were performed in histograms with the same number of events. The Bcl-2 signal in silenced cells (%) was calculated from the ratio between the specific geometric mean of the Bcl-2 signal of silenced and untreated cells.

**Statistical analysis**

Data are reported as means ± SEM. Differences between sequences were evaluated by matched or unmatched two-way analysis of variance (ANOVA), with Bonferroni’s post-test analysis, or by the two-tailed unpaired t-test, depending on the study design, and significance levels were indicated with asterisks (\( P < 0.05 \), \( **P < 0.01 \), and \( ***P < 0.001 \)). Differences between treated and untreated cells were evaluated by two-tailed one sample t-test, and significance levels were indicated with the number sign (#).

**Results**

Bcl-2 down-regulation in SW2 SCLC cells is heterogeneous

Under comparable conditions, the Bcl-2 signal of ODN-treated cells determined by flow cytometry was similar to the Bcl-2 levels measured by Western blot when evaluated in the total cell population (data not shown). However, flow cytometry offers the additional possibility to evaluate Bcl-2 protein levels and heterogeneity of expression in direct correlation with cell viability by creating regions in the FSC/7-AAD plot (10).

Instead of a small protein down-regulation (about 30% at 200 nM G3139 at 72 h; data not shown) in the “bulk” population, the single cell approach provided by flow cytometry showed a strong reduction in the Bcl-2 signal in a reduced percentage of viable cells. The subtraction of the histogram corresponding to untreated viable cells from the histogram of G3139-treated viable cells revealed that about 20% of...
the cells presented an 85% Bcl-2 knockdown (Figure 1A). This potent but cell percentage-limited Bcl-2 down-regulation did not lead to a specific decrease of viability (Figure 1B), and nonspecific toxicity was observed with treatments at 400 nM. These findings at the protein level demonstrate the limited information provided by Western blot regarding the pattern of protein expression. Such heterogeneous Bcl-2 silencing might be correlated with different extents of complex internalization by SW2 cells, as observed with FITC-labeled nucleic acids at 400 nM, where only 36% of tumor cells had internalized complexes after 4 h of incubation (data not shown). The representation of the mean Bcl-2 signal demonstrated a significant sequence-specific effect of G3139 ODN on Bcl-2 knockdown in a viable cell subpopulation (Figure 1C, \( P < 0.001 \), matched two-way ANOVA). Consecutive transfections (at 0 and 24 h) with 100, 200 or 400 nM G3139 ODN did not result in significant improvements of sequence-specific Bcl-2 knockdown (data not shown).

These surprising results of incomplete and heterogeneous silencing of Bcl-2 were further corroborated by transfections carried out with anti-BCL2 siRNA. At the optimized oligofectamine/siRNA ratio (4 times higher than the optimal proportion for ODN complexation in terms of charge ratio), one of the tested siRNA sequences (sequence \( b \)) produced a potent and sequence-specific Bcl-2 down-regulation in a dose-response manner (\( P < 0.0001 \)), reaching a maximum of 63% in viable cells compared to untreated cells (Figure 2A). The Bcl-2-silencing potency of another siRNA sequence, siRNA \( a \), was significantly lower than that of siRNA \( b \) (data not shown). Moreover, no further improvements in Bcl-2 silencing were observed when the double-treatment schedule with siRNA \( b \) was tested (data not shown).

The observation of the frequency histograms of the Bcl-2 signal clearly shows that knockdown of Bcl-2 was heterogeneous also with siRNA (Figure 2B), and the overall improvement in Bcl-2 silencing, either by increasing the oligofectamine/siRNA ratio (data not shown), or siRNA concentration, always correlated with an increased percentage of silenced viable cells (up to 60%, Figure 2C). In the silenced cells, the Bcl-2 signal reduction was between 76 and 82%, regardless of siRNA concentration and percentage of silenced cells (Figure 2C).

The same optimized oligofectamine/siRNA \( b \) ratio was further tested in H2171 and H69 SCLC cells, which exhibit lower and higher constitutive Bcl-2 levels than SW2 cells, respectively. A partial Bcl-2 signal reduction was observed in both cell lines, although to a lesser extent than in SW2 cells (Figure 3A and B). The difference in sequence effect was statistically significant under two-way ANOVA (\( P < 0.0001 \)). *\( P < 0.05 \) and ***\( P < 0.001 \) represent the significance level of the difference between the effects of sequence \( b \) and the respective concentration of non-targeting (nt) sequence (two-way ANOVA, Bonferroni’s post-test). **\( P < 0.05 \), ###\( P < 0.01 \), and ####\( P < 0.001 \) represent the significance level of the tested sequences relative to untreated cells (one sample \( t \)-test).

**Effect of strong BCL2 silencing on cell viability and growth**

Interestingly, siRNA \( b \) and siRNA nt exhibited a different effect on the viability of SW2 cells (\( P < 0.001 \), two-way ANOVA). At 75 and 100 nM, corresponding to the best BCL2-silencing conditions, and with which nonspecific toxicity was
siRNA silencing of Bcl-2 in SCLC

associated (evidenced from the incubation with siRNA nt), siRNA b induced less reduction of cell viability than siRNA nt (Figure 4A). The absence of the expected cytotoxicity of the anti-BCL2 sequence was also confirmed in the cell growth assay using resazurin reduction. At this level, however, the effect of the BCL2-targeting and non-targeting sequences was not significantly different (Figure 4B).

In agreement with that observed with SW2 cells, Bcl-2 down-regulation did not lead to a reduction in viability of H2171 and H69 SCLC cells at 72 h post-transfection (Figure 4C and D).

Bcl-2 mRNA down-regulation in SW2 SCLC cells

To gain further insights into the cellular effects of ODN and siRNA transfection the BCL2 mRNA levels were measured by qRT-PCR in SW2 cells 24 h after the beginning of the experiment. A sequence-specific reduction of the BCL2 mRNA level was observed, reaching approximately 70% at

Figure 4. Effect of siRNA concentration on the viability and growth of small cell lung cancer cells. Tumor cells were treated with different concentrations of siRNA complexed with oligofectamine at 140 µL/nmol siRNA, as described in Material and Methods. The viability of SW2 cells (A) and cell growth (B) were assessed 72 and 96 h after treatment. siRNA = small interfering RNA. Each point represents the mean ± SEM for N = 3 or 4. The difference in sequence effect was statistically significant in viable H2171 and H69 cells (P < 0.0001, two-way ANOVA). **P < 0.01 and ***P < 0.001 represent the significance level of the difference between the effects of sequence b and the respective concentration of non-targeting (nt) sequence (two-way ANOVA, Bonferroni’s post-test). #P < 0.05 and ##P < 0.01 represent the significance level of the difference of the tested sequences relative to untreated cells (one sample t-test).
50 nM siRNA and 400 nM G3139 ODN (Figure 5).

Increases of Bcl-2 as a result of oligofectamine-mediated transfection of SW2 SCLC cells

From the previous results, nonspecific sequence effects at the protein level were evident in viable cells when incubated with control sequences, as demonstrated by the increase of the Bcl-2 signal with concentration. This increase was significant at 200 and 400 nM G4126 ODN (Figure 1C) and 25 and 50 nM siRNA nt (Figure 2A). Conversely, a reduction of the Bcl-2 signal was evident in the dead cell subpopulation compared with the same subpopulation of untreated cells, when treatments were carried out at the highest concentration of control ODN (data not shown). This nonspecific increase of Bcl-2 in viable cells might contribute to masking the average reductions in Bcl-2 signal in these cells compared to untreated cells. When assessment is performed with methods like Western blot this nonspecific effect might not be noticed, conditioning the interpretation of the results.

Assessment of inhibition of SW2 SCLC cell growth by a combination of BCL2 silencing and cisplatin treatment

Although the BCL2-silencing approach was not deleterious to SCLC cells per se, it still could enhance the sensitivity of these tumor cells to cytotoxic drugs like cisplatin widely used in the treatment of SCLC patients. In this regard, the effect of the combined treatments with siRNA and cisplatin on cell growth was evaluated by the resazurin reduction assay. For this purpose, cells were treated with siRNA or nt under optimized silencing conditions (50 nM siRNA and 140 µL oligofectamine/nmol siRNA), followed by cisplatin 28 h after the first treatment. The combined treatment with siRNA and cisplatin was as toxic as cisplatin alone, whereas the combination of siRNA nt with cisplatin resulted in an even lower cytotoxicity than the one caused by the latter treatment (Figure 6, P = 0.0003, two-way ANOVA), suggesting the occurrence of an antagonist effect.

Discussion

BCL2 is an important molecular target whose silencing has been attempted in tumors of different histological origin,
aiming at cancer treatment. The G3139 ODN has been one of the most relevant BCL2-silencing molecules tested, as illustrated by the promising results achieved in several clinical trials, notably in phase III trials against melanoma and chronic lymphocytic leukemia (12,13). Another phase III trial (AGENDA trial) was recently initiated in patients with advanced melanoma. Mechanistically, however, the G3139 ODN has revealed significant effects unrelated to Bcl-2. Its in vitro cytotoxicity to PC3 prostate cancer cells is independent of Bcl-2 down-regulation and correlates with the expression of stress inducible genes (14,15). In melanoma cells, it triggers cytochrome c release from mitochondria and apoptosis, presumably by interacting directly with a mitochondrial channel (voltage-dependent anion selective channel, VDAC) (16).

The application of anti-BCL2 siRNA for therapeutic purposes has never been tested in SCLC. In the present study, it was demonstrated that the best siRNA sequence tested (sequence b) provided a high absolute value of mean BCL2 silencing under optimized conditions (Figure 2C). However, despite multiple treatments and protein evaluation at different times post-treatment, BCL2 silencing analyzed in bulk SW2 SCLC cells did not go beyond 50 and 70% for the Bcl-2 protein knockdown and mRNA level reduction, respectively (Figures 2A and 5). Silencing was in fact potent (80% of protein knockdown, approximately) and heterogeneous (Figure 2B and C). This result correlates with the reduced fraction of successfully targeted cells, as assessed in the present study with FITC-labeled nucleic acid, and in agreement with the low transfection efficiencies previously reported for SCLC cell lines (17,18). It is important to note that, in a previous report, transfection of SW2 cells with anti-survivin siRNA associated with oligofectamine led to the same level of protein down-regulation (50%) (19). Besides heterogeneous transfection, an additional reason that could justify the absence of a full protein (Bcl-2) knockdown is its high stability (reviewed in Ref. 20). In the present study, the use of flow cytometry analysis provided an opportunity to gain insights for the first time into Bcl-2 down-regulation at the single-SCLC cell level upon treatment with siRNA simultaneously with cell viability.

Surprisingly, it was possible to observe an increase in the Bcl-2 protein signal in viable cells after treatment with ODN (Figure 1C) and siRNA (Figures 2A and 3B) control sequences, an effect likely caused by oligofectamine. Its manufacturers do recommend the use of ODN concentrations of 200 nM and alert to the occurrence of nonspecific effects at higher concentrations. This indication is consistent with the observed nonspecific cytoxic effects within the range of 400-800 mM for ODN (Figure 1B). It is important to emphasize that at the mRNA level, and in contrast to that observed with the protein, there was a sequence-specific increase in down-regulation when the dose of G3139 ODN was increased from 200 to 400 nM (Figure 5). These results provide evidence that the observed nonspecific increase of Bcl-2 protein is probably due to a post-translational effect, and could account for the lower Bcl-2 down-regulation compared to the corresponding mRNA.

The most important aspect emerging from the present study was that the down-regulation of Bcl-2 with ODN or siRNA did not result in a sequence-specific decrease in viability or in inhibition of cell growth (Figure 4), or in sensitization of tumor cells to cisplatin treatment (Figure 6). Intriguingly, the most efficient conditions for promoting Bcl-2 down-regulation resulted in a higher cell viability than the corresponding treatment with siRNA nt (Figure 4A). In contrast, there was a similar activity in terms of cell growth under all tested conditions (Figure 4B). Since proliferation and viability are parameters that directly contribute to the overall cell growth of any cell population, the previously mentioned observations imply that proliferation should be reduced in siRNA b-treated cells compared to siRNA nt-treated cells. These observations may be due either to the strong Bcl-2 silencing, revealing an alternative function of Bcl-2 to the currently described anti-apoptotic one, or to the off-target effects derived from the siRNA sequence itself. Although the latter hypothesis cannot be excluded, there is evidence in the literature supporting the former. The notion that Bcl-2 is a multifunctional protein, rather than being merely anti-apoptotic, is gaining strength. In distinct cell contexts, Bcl-2 has been reported to interfere with a variety of biological processes. In SCLC, there is evidence of Bcl-2 interaction with c-Myc in the cell nucleus and in the outer mitochondrial membrane, enhancing c-Myc half-life (21). c-Myc is a protein that has a positive effect on tumor cell proliferation, whereas upon a toxic stress to the cell, such as transfection in the present study, it can decrease tumor cell viability (22,23). Thus, it can be hypothesized that, upon transfection, Bcl-2 silencing by siRNA b leads to a decrease in the levels of c-Myc, causing a reduction in cell proliferation and an increase in cell viability (as compared to siRNA nt).

The present in vitro observations seem to contradict existing expectations regarding the impact of BCL2 on the viability of SCLC cells, namely raised from its role in hematological malignancies and from reports involving the use of anti-BCL2 ODN in SW2, H69 and H82 SCLC cells (8). However, the contribution of the in vivo environment should also be considered, since it may determine a different role for Bcl-2. This was the case in melanoma, where in vitro drug-induced cytotoxicity and resistance were independent of Bcl-2, but stable Bcl-2 silencing reduced the in vivo tumorigenic potential of melanoma cells (24,25). Nevertheless, despite controversial data, existing evidence about the specific role of Bcl-2 in SCLC is consistent with the results obtained in the present study. It has been demonstrated that, regardless of the frequently observed Bcl-2 overexpression in SCLC cell lines (2) and tumors (3), there is no correlation between Bcl-2 expression and resistance to therapy and/or patient survival (26-30). In fact, Bcl-2
overexpression, mainly with nuclear localization, together with a high Bcl-2/Bax ratio, was correlated directly with an elevated spontaneous apoptosis index (31-33), and with the absence of metastasis in SCLC tumors (34). The relationship between Bcl-2 overexpression and tumor stage is less clear, but an inverse correlation between Bcl-2 levels and cancer progression has already been pointed out (29,34-36). In addition, there are cases of reduced Bcl-2 levels upon in vitro acquisition of resistance to cisplatin (37). Overall, it is not surprising that an immunosilent derivative of G3139 ODN did not reveal any therapeutic activity when tested in a murine model of human SCLC (38).

With the development of small molecules capable of inhibiting anti-apoptotic Bcl-2 family proteins, BCL2 regained a new protagonism (39). Results from studies with these inhibitors support the idea that in order to achieve a deleterious effect on SCLC cells it will be necessary to interfere with more than one of the Bcl-2 family proteins. In fact, the Bcl-2 family inhibitor ABT-737 is effective against some SCLC lines by simultaneously inhibiting three anti-apoptotic proteins (Bcl-2, Bcl-xL, and Bcl-w). In addition, in the case of the expression of a fourth anti-apoptotic protein (Mcl-1), its knockdown is further required to achieve increased tumor cell death, thus reinforcing the importance of a multitargeted approach (40).

Gene silencing represents an opportunity to change the treatment paradigm of many diseases. With such strategy it is theoretically possible to inhibit any molecular target within a cell, as opposed to what happens with small molecular weight drugs or protein therapeutics. The present study, which reports for the first time the effect of anti-BCL2 siRNAs on SCLC, benefited from single-cell analysis. Among the sequences tested, siRNA b significantly down-regulated Bcl-2 in a sequence-specific manner both at the protein and mRNA levels. Surprisingly, Bcl-2 knockdown was heterogeneous at the individual cell level and did not translate into a specific decrease of cell viability, specific inhibition of cell growth or chemosensitization to cisplatin treatments. These results raise questions about the in vitro (anti-apoptotic) role of Bcl-2 in SCLC claimed in previous reports, and support the growing notion of the necessity of a molecular multtargeting approach, especially when considering anti-apoptotic strategies, in the fight against cancer.

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