The CLN3 Gene is a Novel Molecular Target for Cancer Drug Discovery

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ABSTRACT

Juvenile Batten disease is a neurodegenerative disease caused by accelerated apoptotic death of photoreceptors and neurons attributable to defects in the CLN3 gene. CLN3 is antiapoptotic when overexpressed in NT2 neuronal precursor cells. CLN3 negatively modulates endogenous ceramide levels in NT2 cells and acts upstream of ceramide generation. Because defects in regulation of apoptosis are involved in the development of cancer, we evaluated the expression of CLN3 on both mRNA and protein levels in a variety of cancer cell lines and solid colon cancer tissue. We also observed the effect of the blocking of CLN3 protein expression on cancer cell growth, survival, ceramide production, and apoptosis by using an adenovirus-expressing antisense CLN3 construct. We show that CLN3 mRNA and protein are overexpressed in glioblastoma (U-373G and T98g), neuroblastoma (IMR-32 and SK-N-MC), prostate (Du145, PC-3, and LNCaP), ovarian (SK-OV-3, SW626, and PA-1), breast (BT-20, BT-549, and BT-474), and colon (SW1116, SW480, and HCT 116) cancer cell lines but not in pancreatic (CAPAN and As-PC-1) or lung (A-S49 and NCI-H520) cancer cell lines. CLN3 is also up-regulated in mouse melanoma and breast carcinoma colon cancer cell lines. We found CLN3 expression is 22–330% higher than in corresponding normal colon control tissue in 8 of 10 solid colon tumors. An adenovirus-expressing antisense CLN3 (Ad-AS-CLN3) blocks CLN3 protein expression in DU-145, BT-20, SW1116, and T98g cancer cell lines as seen by Western blot. Blocking of CLN3 expression using Ad-AS-CLN3 inhibits growth and viability of cancer cells. It also causes elevation in endogenous ceramide production through de novo ceramide synthesis and results in increased apoptosis as shown by propidium iodide and JC-1 staining. This suggests that Ad-AS-CLN3 may be an option for therapy in some cancers. More importantly these results suggest that CLN3 is a novel molecular target for cancer drug discovery.

INTRODUCTION

Mutations in the CLN3 gene are responsible for the JNCL (1). JNCL is a recessively inherited neurodegenerative disorder of childhood (1–3). The clinical hallmarks are progressive loss of vision, seizures, and mental deterioration. These symptoms are attributable to massive cortical neuronal death and gradual loss of photoreceptor cells (2, 3). Apoptosis has been shown to be the mechanism of neurodegeneration in the brain of patients with the juvenile form of Batten disease (4). Moreover, up-regulation of Bcl-2 and elevation of endogenous ceramide levels in the brain from affected individuals provides mechanistic evidence for apoptotic death of neurons in this disorder (5).

We have demonstrated previously that stable CLN3 overexpression protects NT2 neuronal precursor cells from serum starvation-induced growth inhibition and also rescues these cells from death caused by treatment with vincristine, etoposide, and staurosporine (6). We have also shown that CLN3 up-regulation decreases the level of the lipid second messenger, ceramide, in these cells and also attenuates vincristine-induced activation of ceramide (6). However, overexpression of CLN3 fails to protect NT2 cells from exogenous ceramide-induced killing. These facts place CLN3 upstream of ceramide in apoptosis signaling and suggest that CLN3 plays an important role in mechanisms of cell death and survival. The fact that CLN3 is highly conserved across species from human to yeast and also in Caenorhabditis elegans and Drosophila underscores its importance for cell function. CLN3 has also been found to be developmentally regulated in differentiating hNT neurons and in neonatal rat brain. Peaks of expression are noted just after hNT cells exit the cell cycle, and on day Po in neonatal rat brain, which corresponds to the period of maximum neuronal growth (7). This suggests that CLN3 is an oncofetal, anti-apoptotic gene. These facts led us to investigate whether CLN3 could be differentially expressed in some cancers.

Defects in proapoptotic events can contribute to cancer formation by allowing cells to survive and to proliferate beyond their normal life span (8–12). Regulation of apoptosis is involved in the development of tumors and plays an essential role in their treatment. A variety of chemotherapeutic agents and radiation kill tumor cells by inducing apoptosis (10, 13, 14). It has been established that many tumor types are resistant to chemotherapy-induced apoptosis. This can occur either because of inactivation of tumor suppressor genes, such as p53 or retinoblastoma, or because of overexpression of antiapoptotic onco-genes, such as Bcl-2 (B-cell lymphoma), Bcr-Abl (myelogenous leukemia), BUG-1 (breast cancer), and survivin in a number of cancers and lymphomas (15–18). Identification of novel antiapoptotic genes expressed in cancer cells provides a basis for a better understanding of the biology of these tumors and may lead to discovery of new targets for anticancer drug development (9, 19–21). Direct inactivation of antiapoptotic gene expression may promote cancer cell death. Suppression of antiapoptotic genes can be achieved by antisense strategies, which in some instances may improve the efficacy of conventional chemotherapy (9). Antisense-based therapies have already been developed to block Bcl-2 overexpression in non-Hodgkin’s lymphoma (22).

In this study we establish that CLN3 mRNA and protein are overexpressed in a number of cancer cell lines including breast, colon, malignant melanoma, prostate, ovarian, neuroblastoma, and glioblastoma multiforme but not lung or pancreatic cancer cell lines. We also show that CLN3 is overexpressed in 8 of 10 solid human colon cancer cases. Additionally, we demonstrate that blocking CLN3 protein expression in these cancer cell lines, using adenovirus-mediated antisense CLN3 methodology, inhibits cancer cell growth and viability. Treatment of cancer cells with Ad-AS CLN3 virus also affects the de novo ceramide synthetic pathway and results in ceramide elevation and cancer cell death by apoptosis.

MATERIALS AND METHODS

Cell Culture. Cancer cell lines were obtained from American Type Culture Collection (Manassas, VA) and maintained at 37°C and 5% CO2 except for...
SW1116 cells, which are cultured under CO₂-free conditions by sealing the flasks with parafilm. Human cancer cell lines were maintained in medium as described below. Normal fibroblasts and A 375 (melanoma) cells were cultured in DMEM with 10% FBS; SW 626 (ovarian adenocarcinoma) cells were cultured in DMEM with 1 mM sodium pyruvate, 10% FBS; PC3 (prostate adenocarcinoma) cells were propagated in Ham’s F12K medium with 7% FBS; A549 (lung carcinoma) cells were cultured in Ham’s F12K medium with 2 mM glutamine adjusted to contain 1.5 g/liter sodium bicarbonate, 10% FBS; SW1116 (colon adenocarcinoma) cells were cultured in Leibovitz’s L-15 medium with 10% FBS; SW-480 (colon adenocarcinoma) cells were cultured in Leibovitz’s L-15 medium with L-glutamine, 10% FBS; PC3 (prostate carcinoma) and PA 1 (ovary teratocarcinoma) cells were cultured in MEM with Earle’s BSS, 0.1 mM nonessential amino acids, and 10% FBS (heat inactivated for PA-1 cells); T98 G and U-373 MG (glioblastomas) cells and IMR-32 and SK-N-MC (neuroblastoma) cells were cultured in MEM with L-glutamine and Earle’s BSS adjusted to contain 1.5 g/liter sodium bicarbonate, 0.1 mM nonessential amino acids, 1 mM sodium pyruvate, 10% FBS; LNCaP (prostate carcinoma) and BT 549 (breast carcinoma) were cultured in RPMI 1640 with 10% FBS; A549 (lung carcinoma) cells were cultured in Ham’s F12K medium with L-glutamine, 10% FBS; SK-OV-3 (ovary adenocarcinoma) cells were propagated in Ham’s F12K medium with 7% FBS; CAPAN-1 and As-PC-1 (pancreas adenocarcinoma) cells were cultured in RPMI 1640 with 15% and 20% FBS, respectively; and NCI-H520 (lung carcinoma) and BT 474 (breast carcinoma) cells were cultured in RPMI 1640 with 2 mM glutamine adjusted to contain 1.5 g/liter sodium bicarbonate, 4.5 g/liter glucose, 10 mM HEPES, 1 mM sodium pyruvate, and 10% FBS. Penicillin-streptomycin (1%) were added in all of the mediums. Mouse B16 (melanoma) cells and 4T1 (breast carcinoma) cells were cultured in DMEM high glucose medium with 10% heat inactivated FBS, 1% penicillin-streptomycin-streptomycin-fungene; mouse embryo and contact sensitive C3H10 T1/2 cells were used as a control and were cultured in BME medium with Earle’s BSS, 10% heat inactivated FBS, and 1% penicillin-streptomycin-fungene.

**Human Colon Cancer and Normal Colon Tissues.** Samples of human colon cancer and normal colon tissue from the same patient were obtained using an Institutional Review Board-approved procedure for collecting leftover, “deidentified,” or “unlinked” specimens from surgical pathology. In brief, at the time of frozen section analysis of the patient tumor, samples of tumor and normal colon tissue not needed for diagnostic purposes were snap frozen in embedding medium and stored at −80°C. The samples were deidentified by removing all of the patient information from the samples except for a pathological diagnosis. Frozen sections of each specimen were stained with H&E and examined by a pathologist to insure that the frozen samples contained representative normal or neoplastic tissue. In all of the cases, the colon cancer specimens consisted mostly of neoplastic glands invading into the muscularis externa. The normal colon specimens contained mucosa, submucosa, and muscularis externa.

**Construction of Ad-AS-CLN3.** Ad-AS-CLN3 virus was constructed using Ad-Easy method (23). CLN3 cDNA was cloned in the antisense direction using cotransformation of the multiple cloning site of pShuttle-CMV vector. The Ad-AS-CLN3 vector was constructed by homologous recombination using cotransformation of the Ad-AS-CLN3 vector into BJ5183-competent cells. The Ad-AS-CLN3 vector was isolated, propagated, and purified exactly as described previously (24).

**Limiting Dilution Assay.** Serial dilutions of viral stock from 1 × 10⁵ to 1 × 10³ were made. B6 cells were plated to 80% confluency in two 24-well plates. Serial dilutions of virus were then added in each row of the 24-well plate and allowed to infect for 1 h. Additional medium was added, and the plates were incubated in a 37°C incubator at 5% CO₂ for 2 weeks. Limiting dilution was determined by the wells with the lowest viral titer that showed cytotoxic effect.

**Transduction of Cancer Cells with Control Vector or Ad-AS-CLN3 Virus.** Transduction of cancer cells was carried out in 60-mm dishes for Western Blot and in 12-well plates for growth curves and proliferation assays. At 70% confluency, the number of cells was determined. Cells were infected with the MOI (number of viral particles used per cell) of virus indicated in the figures by incubation with virus in a minimum amount of medium for 1 h and addition of medium to a final volume after that.

**RT-PCR for CLN3.** Total RNA was isolated from cancer cells using the RNeasy Mini kit (Qiagen Inc., Valencia, CA). Then mRNA was isolated from total RNA by the Oligotex mRNA spin-column Mini kit (Qiagen Inc.). First-strand cDNA was then synthesized from mRNA by reverse transcription using Omniscript Reverse Transcription kit (Qiagen Inc.). PCR reaction was set up with first-strand cDNA, 2.5 units of Ampli Taq Gold (PE Applied Biosystems, Foster City, CA), and 5 µCi of [α-³²P]dCTP in each reaction. The reaction was performed in PCR buffer (Applied Biosystems) containing 2.5 mM MgCl₂, 50 µM of dCTP, dATP, dTTP, and dGTP. The primers used for amplification of human CLN3 were 5’ primer, 5’-GGTGCAAGATCTCAAGGG-3’ (958–976), and 3’ primer, 5’-CTTGCGGAGAAAGCCAGACG-3’ (1229–1246). Cyclophilin was used as an internal control, and primers used for amplification of cyclophilin were the 5’ primer, 5’-AATGGTCTGGCCCAACAGTCTG-3’ (337–354), and 3’ primer, 5’-AACACACATCTGTTGCCC-3’ (384–401). The primers used for amplification of mouse CLN3 were 5’ primer, 5’-AGTTGTCAGGTGTCAGATCCTG-3’ (961–982), and 3’ primer, 5’-GAAAGTTATCAGCTAAGGCCG-3’ (1300–1321). The primers used for amplification of mouse cyclophilin were 5’ primer, 5’-ACAGCAAGTCTCCATCGTGTACATC-3’ (285–307), and 3’ primer, 5’-TGCTCTTTCTTCTTGCTGACATC-3’ (347–368). The reaction conditions used for human cDNA were 10 min at 95°C, and then 1 min at 94°C, 1 min at 50°C, and 2 min at 72°C, for 20 cycles. The reaction conditions used for mouse cDNA were 10 min at 95°C, and then 1 min at 94°C, 1 min at 50°C, and 2 min at 72°C, for 20 cycles. PCR amplified products were separated on 8% nondenaturing polyacrylamide gel. The bands of interest were quantitated using a Phospholmager (Molecular Dynamics Inc., Sunnyvale, CA). The results are expressed as the ratio of CLN3 signal to that of the internal control, cyclophilin.

**Immunocytochemistry.** Cells were grown on glass coverslips until confluent, fixed in 4% paraformaldehyde and 4% sucrose in PBS at 4°C, and permeabilized in 0.2% Triton X-100 in PBS at room temperature. After blocking in 3% BSA in PBS, cells were incubated with human CLN3 antisem. The CLN3 antibody used in this study is a polyclonal antibody raised against the peptide sequence AAHDILSHDRTSGNQSHVDP corresponding to amino acids 58–77 of the CLN3 protein (Research Genetics, Huntsville, AL). An additional glass coverslip of each cell type was exposed to rabbit IgG at the same concentration under the same conditions. Cells were washed and then incubated with 1:500 dilution of biotin-conjugated goat anti-rabbit IgG (Zymed, San Francisco, CA) for 1 h. This was followed with exposure to streptavidin-conjugated horseradish peroxidase (Zymed) 1:300 for 30 min. Cells were stained with 3,3’-diaminobenzidine in 0.2 M Tris-HCl, counterstained with hematoxylin blue, dehydrated in ethanol, washed in xylene, and mounted on glass slides with Permount.

**Western Blot.** Western Blot for CLN3 protein detection was performed as described previously (6). Cells were harvested on day 3 after infection and lysed in buffer containing 250 mM NaCl, 0.1% NP40, 50 mM HEPES (pH 7.0), 5 mM EDTA, 1 mM DTT, and 10% protease inhibitor mixture (Sigma Chemical Co.). After 40 min of incubation at 4°C, lysates were centrifuged at 12,000 × g, and the supernatants were quantitated for total proteins using the Bio-Rad protein assay. Equal amounts of protein (200 µg) were separated by SDS-PAGE using 4.5% and 12% acrylamide for stacking and resolving gels, respectively. Proteins were transferred to nitrocellulose membrane and probed with a polyclonal CLN3 antibody raised against the peptide sequence AAHDILSHDRTSNGSHTVD corresponding to amino acids 58–77 of the CLN3 protein (Research Genetics). CLN3 primary antibody complexes were detected using goat antirabbit IgG conjugated with horseradish peroxidase and visualized by SuperWest Pico Chemiluminescent Substrate (Pierce, Rockford, IL).

**[³H]Thymidine Incorporation Cell Proliferation Assay.** Cells were plated in 12-well plates at a density of 7 × 10⁴ cells/well and were incubated with 0.5 µCi/ml [³H]thymidine (DuPont) in medium for 2 h before harvesting. For viral transduction experiments, cells were infected with 5, 10, 15, 20, and 40 MOI of control or Ad-AS-CLN3 virus as described above. On day 2 after infection cells were incubated with 0.5 µCi/ml [³H]thymidine (DuPont) in medium for 2 h. The cells were washed twice with ice-cold PBS and the DNA was precipitated with 5% trichloracetic acid. The DNA precipitate was dissolved in 0.4 ml of 0.25 M NaOH, and incorporated [³H]thymidine was determined by liquid scintillation counting. Each time point was determined in triplicate.

**Measurement of Ceramide Levels.** DU145 prostate cancer cells were plated in 100-mm dishes and at 50% confluency were transduced with 40 MOI of Ad-AS CLN3 virus or control virus. Three days later cells were washed with...
PBS, and lipids were extracted using the Bligh and Dyer method. Fumonisin, an inhibitor of de novo ceramide biosynthesis (Biomol, Plymouth, PA) was added at a concentration of 25 μM to two dishes of DU 145 cells 12 h before transduction with virus. Ceramide levels in the cells were measured using diacylglycerol kinase assay, as described previously (6). The results are expressed as pmols of ceramide per nmol of total phospholipids and represent an average from three experiments.

Measurement of Sphingomyelin Content. Lipids were extracted from 10×10⁶ cells using the method of Bligh and Dyer. Lipid extracts in 1 ml of chloroform each were subjected to hydrolysis by adding 0.1 ml of 0.2 N NaOH in methanol. Aliquots were taken for quantitation of total phospholipids before hydrolysis. After neutralization with an equal amount of 2 N HCl, lipids were reextracted by adding 1 ml of chloroform and 1 ml of water and vortexing. The lower organic phase was then evaporated under nitrogen gas, resuspended in 75 μl of chloroform-methanol (2:1), and 50 μl out of it was spotted onto a TLC plate (Whatman Inc., Clifton, NJ). The plate was developed in CHCl₃–CH₃OH-acetic acid–H₂O (50:30:8:5). Lipids were visualized using iodine vapor, and bands corresponding to sphingomyelin standard were scraped. The amount of phosphor was determined using phosphomolybdate phosphate assay. The content of sphingomyelin was expressed in nmol per nmol of total phospholipids, and percentage change in sphingomyelin in patient cells in comparison to control fibroblasts was calculated.

PI Staining. DU145 prostate cancer cells were grown on glass coverslips and transduced with 40 MOI of Ad-AS CLN3 virus or control virus. After 72 h, cells were stained with PI in PBS at a concentration of 5 μg/ml for 5 min at 4°C. After that, cells were washed with PBS and mounted in PBS/glycerol (1:1). PI-positive apoptotic cells were observed under fluorescence (excitation wavelength: 525 nm, emission wavelength: 600 nm) at ×100 magnification.

JC-1 Staining. JC-1 is a cationic lipophilic dye and exhibits potential-dependent accumulation in mitochondria (25, 26). Cell were grown on glass coverslips, and transduced with 40 MOI of Ad-AS CLN3 virus or control virus. After 72 h, cells were stained with 1 μg/ml JC-1 in medium for 15 min at 37°C. After that, cells were washed twice with PBS and mounted on glass slides with PBS/glycerol (1:1). At the onset of apoptosis, a drop in potential was indicated by a fluorescence emission shift from green to red (525 to 590 nm) attributable to formation of J-aggregates. These appear as bright dots in Fig. 9. J-aggregates in Du145 prostate cancer cells are visualized using ×400 magnification. Percentage of apoptotic cells is calculated from two fields (60–150 cells/field), and the data are presented as the average of both counts. The statistical significance of the difference in the number of apoptotic cells after transduction with control or Ad-AS CLN3 virus is determined by the two-tailed Student t test.

RESULTS

CLN3 Is Overexpressed in Cancer Cell Lines Both at the RNA and Protein Levels. We had shown previously that CLN3 is involved in antiapoptotic pathways. Because defects in apoptosis play an important role in carcinogenesis, we compared CLN3 expression at the RNA level in neuroblastoma, glioblastoma, prostate, ovarian, colon, breast, melanoma, pancreas, and lung cancer cell lines to CLN3 expression in normal fibroblasts. Mouse CLN3 expression at the RNA level is also measured in cells derived from naturally occurring mouse melanoma and mouse breast carcinoma (Figs. 1 and 2). The results demonstrate that expression of CLN3 is increased 3.5-fold in glioblastoma, 1–4-fold in prostate cancer cell lines, 2–4-fold in ovarian cancer cell lines, and 2–4-fold in breast carcinomas in comparison to normal fibroblasts. CLN3 mRNA expression in C 32 (melanoma)

Fig. 1. CLN3 expression in human cancer cell lines by RT-PCR. mRNA was isolated from different cancer cell lines, cDNA synthesized, and PCR was performed in the presence of CLN3 5' and 3' primers as described in “Materials and Methods.” Cyclophilin was used as an internal control. Results are expressed as the ratio of the CLN3 signal to that of cyclophilin. a, neuroblastoma (SK-N-MC, IMR-32), glioblastoma (T98g, U373); b, prostate cancer cell lines; c, ovarian cancer cell lines; d, breast carcinomas; e, melanoma (C52, A375) and colon (SW1116) cancer cell lines; f, pancreas (Capan, As-PC-1) and lung (A-549, NCI-H520) cancer cell lines. Average of four separate experiments; bars, ± SD; *P < 0.05 versus the control; **P < 0.01 versus the control; au, arbitrary units.
demonstrate in 7 tumors that CLN3 expression is 50–330% higher than in the corresponding normal colon tissue. In an eighth case CLN3 expression was increased by 22% (Fig. 4).

**Ad-AS-CLN3 Virus Blocks CLN3 Protein Expression and Decreases Viability of Cancer Cells.** To demonstrate the impact of CLN3 overexpression on cancer cells we engineered an adenoviral vector with an antisense CLN3 cDNA construct capable of blocking CLN3 overexpression and studied its effect on cancer cell growth and apoptosis. Infection of prostate DU 145, neuroblastoma SK-N-MC, and breast BT-20 cancer cells with 40 MOI Ad-AS-CLN3 virus and colon SW1116 cells with 5 MOI appreciably block expression of the CLN3 protein in comparison to cells infected with the same amount of control virus (Fig. 5).

We then analyzed the effect of different MOIs of Ad-AS-CLN3 virus on cancer cell growth. Incorporation of [3H]thymidine was measured in DU 145, SK-N-MC, BT-20, and SW1116 cells infected with 5, 10, 15, 20, and 40 MOI of Ad-As CLN3 virus or control virus. Incorporation of [3H]thymidine into cancer cells was inversely related to an increase in MOI or the number of Ad-AS-CLN3 viral particles infecting each cell (Fig. 6, a, c, e, and g). Ad-AS-CLN3 at a dose of

![Image](http://cancerres.aacrjournals.org/)

**Fig. 3. CLN3 protein expression by immunocytochemistry.** Confluent cells were fixed, permeabilized, and incubated with human CLN3 antiserum for 12 h. After that cells were exposed to goat antirabbit biotin-conjugated antibody for 1.5 h and to horseradish peroxidase conjugated streptavidin for 30 min. Next, cells were stained with diaminobenzidine, dehydrated, and mounted in Permount. Brown staining correlates with CLN3 protein expression. Note increased amount of staining in Du145 prostate, SK-N-MC neuroblastoma, BT-549 breast, PA-1 ovarian, and T98g glioblastoma cancer cells compared with normal human fibroblasts.

and SW1116 colon cancer cell lines is 2-fold higher than in the normal control. Pancreas and lung cancer cells only show slight changes in CLN3 expression (Fig. 1). CLN3 expression in mouse melanoma and breast carcinoma cell lines is 25–38-fold higher in comparison to CLN3 expression in control mouse C3H10T1/2 cells (Fig. 2).

To compare the CLN3 protein levels in cancer cells to control fibroblasts we performed immunocytochemistry using polyclonal CLN3 antibody. Prostate DU145, ovarian PA1, glioblastoma T98G, neuroblastoma SK-N-MC and breast carcinoma BT-549 cancer cells show intense staining for CLN3 as opposed to light staining in normal fibroblasts (Fig. 3). Therefore, increase in CLN3 protein levels correlates with CLN3 mRNA overexpression in cancer cells.

**Expression of CLN3 Is Increased in 8 of 10 Solid Colon Tumors.** To corroborate the results obtained in cancer cells in solid human tumors, we analyzed CLN3 expression in 10 solid colon tumors. Normal colon tissue obtained from the same patient at the time of surgery and assessed pathologically is used as a control. We

![Image](http://cancerres.aacrjournals.org/)

**Fig. 2. CLN3 expression in mouse melanoma, breast carcinoma cell lines, and C3H10T1/2 control mouse fibroblasts.** mRNA was isolated from different cell lines, cDNA synthesized, and PCR was performed in the presence of CLN3 5' and 3' primers as described in “Materials and Methods.” Cyclophilin was used as an internal control. The results are expressed as the ratio of CLN3 signal to that of cyclophilin. Average of four separate experiments; bars, ± SD; *P < 0.05 versus the control; au, arbitrary units.
15 MOI was sufficient and effective in the killing of cancer cells as seen from the dose-response curves in Fig. 6, a, c, e, and g.

The growth profile of DU145, BT-20, SW1116, and SK-N-MC cancer cells infected with control virus practically matches that of untreated cells (Fig. 6, b, d, f, and h). In contrast, infection of the cells with Ad-AS-CLN3 halts cancer cell growth. This effect was noticeable on day 2 after transduction with virus. At the end of the treatment the amount of live cells drastically dropped compared with control, which indirectly indicates an increase in the number of dying cells.

Increase in Ceramide Levels in CLN3 Deficient Fibroblasts Is Not Sphingomyelin-derived. We also measured ceramide levels in fibroblasts obtained from patients with juvenile Batten disease that are deficient in CLN3. There was a 78.5% increase in endogenous ceramide in CLN3-deficient fibroblasts compared with normal fibroblasts. At the same time there was no simultaneous drop in sphingomyelin content. In fact, it was 26.9% higher than in normal cells (Fig. 7b). This suggests that the increase in ceramide levels seen with diminished expression of CLN3 is not attributable to sphingomyelin breakdown or the sphingomyelin cycle but more likely suggests an impact of CLN3 on the de novo ceramide pathway.

Blocking of CLN3 Expression in Prostate Cancer Cells Results in Increased Ceramide Levels. We have shown previously that CLN3 negatively modulates ceramide generation in NT2 cells. Ceramide levels in DU145 prostate cancer cells were increased by 91.4% after transduction with Ad-AS CLN3 virus compared with cells transduced with control virus (Fig. 7a). Moreover, pretreatment of these cells with fumonisin, an inhibitor of ceramide synthase, abrogated ceramide elevation. Fumonisin equally blocks ceramide elevation in DU145 cells after transduction with Ad-AS CLN3 virus and control virus by 93.3% and 91.4%, respectively. Therefore, blocking of de...
Ceramide synthesis suppresses ceramide level increases induced by Ad-AS CLN3 virus, and this suggests that CLN3 impacts the de novo ceramide pathway (Fig. 7a).

Transfection of CLN3-deficient Lymphoblasts with CLN3 Containing Plasmid Restores Growth but Has No Effect on Thymidine Incorporation. CLN3-deficient lymphoblasts from patients manifest slowed growth compared with normal lymphoblasts (Fig. 8a). Reintroduction of CLN3 into JNCL patient cells, as opposed to the empty vector, restores growth almost to normal levels but has no effect on thymidine incorporation (Fig. 8b). This indicates that the CLN3 gene impacts apoptotic pathways and not cell proliferation.

Blocking of CLN3 Expression in Prostate Cancer Cells Increases Apoptosis. Transduction of DU145 prostate cancer cells with Ad-AS CLN3 results in increased apoptosis. There is a higher number of apoptotic cells in DU145 prostate cancer cells after transduction with Ad-AS CLN3 virus compared with transduction with CLN3-deficient lymphoblasts. Thymidine uptake by CLN3-deficient lymphoblasts was measured 24 h after transfection with pGEM + 7f-CLN3 (gray triangles), as opposed to transfection with an empty vector, restores growth to normal levels. b, thymidine incorporation in CLN3-deficient lymphoblasts. Thymidine uptake by CLN3-deficient lymphoblasts was measured 24 h after transfection with pGEM + 7f-CLN3 (gray triangles) or empty plasmid (>). Note that CLN3 deficient lymphoblasts incorporate thymidine at a rate similar to lymphoblasts transfected with full CLN3 gene.

Fig. 7. a, ceramide levels in DU145 cells after transduction with Ad-AS CLN3 virus. DU145 cells were transduced with 30 MOI of Ad-AS CLN3 virus or control virus. Fumonisin was added 12 h before transduction with virus. Three days after transduction, lipids were harvested and ceramide levels were measured using the diacyl glycerol kinase assay. Results are an average of three experiments and are expressed in pmol of ceramide per nmol of total phospholipids. a, bar 1, DU145 cells + control virus; bar 2, DU145 cells + control virus + fumonisim; bar 3, DU145 cells + Ad-AS CLN3 virus; bar 4, DU145 cells + Ad-AS CLN3 virus + fumonisim. b, ceramide and sphingomyelin levels in CLN3 deficient fibroblasts. Ceramide and sphingomyelin were measured as described in “Materials and Methods” and expressed as percentage change compared with levels in normal fibroblasts. Note that the increase in ceramide levels is not accompanied by a decrease in sphingomyelin content suggesting that ceramide is not derived from activation of the sphingomyelin cycle; bars, ± SD.

Fig. 8. a, growth curves for CLN3-deficient lymphoblasts. Lymphoblasts (5 × 10⁴) are plated in 12-well plates and, after 24 h, transfected with pGEM + 7f-CLN3 or empty vector using LipofectAMINE 2000 kit (Stratagene, La Jolla, CA). The number of cells was counted using trypan blue dye exclusion assay starting 24 h after transfection. Note that JNCL patient cells (Œ) show diminished growth when compared with normal lymphoblasts (/H12135). Transfection of JNCL patient cells with the full CLN3 gene (gray triangles), as opposed to transfection with an empty vector, restores growth to normal levels. b, thymidine incorporation in CLN3-deficient lymphoblasts. Thymidine uptake by CLN3-deficient lymphoblasts was measured 24 h after transfection with pGEM + 7f-CLN3 (gray triangles) or empty plasmid (>). Note that CLN3 deficient lymphoblasts incorporate thymidine at a rate similar to lymphoblasts transfected with full CLN3 gene.

Fig. 9. PI (a, b) and JC-1 staining (c, d) of DU145 cells. On day 3 after transduction with control virus (a) or Ad-AS CLN3 virus (b), cells were washed, stained with PI (5 µg/ml) for 5 min, and observed under fluorescence (excitation wavelength: 525 nm, emission wavelength: 600 nm). Note an increase in the number of white apoptotic DU145 cells after transduction with Ad-AS CLN3 virus. On day 3 after transduction with control virus (c) or Ad-AS CLN3 virus (d), DU145 cells were washed, stained with JC-1 in medium (1 µg/ml) for 15 min at 37°C, and observed under fluorescence. At the onset of apoptosis there was a drop of potential indicated by a fluorescence emission shift from green to red (525 to 590 nm) because of formation of red fluorescent J-aggregates that appear as bright dots. The JC-1-stained cells are visualized using ×400 magnification. Apoptotic cells demonstrate formation of red J-aggregates. Note decrease in the total number of cells and an increase in those with red J-aggregate formation.
control virus shown by PI staining (Fig. 9, a and b). Also, the number of cells with J-aggregate formation is increased from 8.2 ± 3.7% to 34.5 ± 9% (P < 0.05) after transduction with Ad-AS CLN3 virus as shown by JC-1 staining. (Fig. 9, c and d).

**DISCUSSION**

Deregulation of antiapoptotic genes and/or mutated proapoptotic genes may contribute to carcinogenesis by allowing abnormal cells to survive and thereby become cancerous (21, 27, 28). Bcl-2 was one of the first antiapoptotic genes found to be up-regulated in cancer (29). Other genes have been discovered that also are overexpressed in cancer and that inhibit apoptosis. Examples include Bcr-Abl in chronic myeloid leukemia, **BUG-1** in breast tumors, and **Survivin** in a number of cancers and lymphomas (10, 15, 16, 30–34).

Previous work has shown that CLN3 is antiapoptotic (6). Ceramide levels are elevated in the brain of patients with the juvenile form of Batten disease compared with normal brain (5). NT2 cells stably or transiently overexpressing CLN3 protein have lower ceramide levels than control virus shown by PI staining (Fig. 9, a and b). Other genes have been discovered that also are overexpressed in cancer and that inhibit apoptosis. Examples include Bcr-Abl in chronic myeloid leukemia, **BUG-1** in breast tumors, and **Survivin** in a number of cancers and lymphomas (10, 15, 16, 30–34). CLN3-deficient lymphoblasts show no difference in thymidine incorporation when transfected with empty plasmid or plasmid carrying full-length CLN3 cDNA. This suggests that CLN3 impacts apoptotic pathways and not proliferative pathways.

We demonstrate that CLN3 is overexpressed at the mRNA and protein levels in a number of human cancer cell lines including prostate, glioblastoma, neuroblastoma, ovarian, breast, colon, and malignant melanoma. Pancreas and lung cancer cell lines do not show increased CLN3. CLN3 was also found to be up-regulated in mouse melanoma and breast carcinoma cell lines. On the basis of these observations, it may become possible to develop semiquantitative assays of CLN3 expression for screening or to monitor effectiveness of therapy in some cancers. Overexpression of CLN3 in 80% of solid colon cancers tested corroborates data obtained from human cancer cell lines. Discovery of novel antiapoptotic genes involved in cancer may lead to a better understanding of cancer biology. Also, these novel genes may prove to be effective targets for cancer drug development. Potential strategies for induction of apoptosis in cancer cells include inactivation of antiapoptotic genes by blocking their expression (9, 18). Reduction in Bcl-2 expression by antisense methods increases the susceptibility of cancer cells to induction of apoptosis by multiple chemotherapeutic agents (19, 22). Similarly, down-regulation of CLN3 expression with a resultant increase in ceramide generation may be explored as a potential cancer therapeutic strategy. We demonstrate that expression of the CLN3 protein drops significantly after transduction with fumonisin, an inhibitor of ceramide synthase, before transduction with the Ad-AS CLN3 virus completely blocks elevation in ceramide. This indicates that blocking of CLN3 protein expression induces an increase in ceramide levels not through sphingomyelin breakdown but via the de novo ceramide synthetic pathway. Additional evidence is the fact that increases in ceramide levels in CLN3-deficient brain and CLN3-deficient fibroblasts were not accompanied by a change in sphingomyelin levels.

Suppressing levels of CLN3 protein in cancer cells enhances ceramide production and results in death of cancer cells. There are a number of anticancer drugs of which the effects include boosting of ceramide levels in cancer cells (44).

In conclusion, CLN3 overexpression may be developed as yet another marker for some but not all cancers. Moreover, blocking CLN3 overexpression using an antisense adenoaviral CLN3 construct inhibits cancer growth, increases ceramide levels, and promotes death of cancer cells. The implications are 2-fold. The use of antisense CLN3 methodology may be valid for treatment of some cancers. Additionally, the CLN3 gene provides a novel molecular target to screen for new drugs effective in combating cancer.

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The CLN3 Gene is a Novel Molecular Target for Cancer Drug Discovery

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