Decreased Iron in Cancer Cells and Their Microenvironment Improves Cytolysis of Breast Cancer Cells by Natural Killer Cells

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Abstract. The association of iron with anticancer immunity is unclear. In order to determine the role of iron in anticancer immunity, we manipulated intracellular iron levels of the human MCF-7 and MDA-MB-231 breast cancer cell lines, and measured cytolysis of breast cancer cells by the natural killer cell line NK-92MI, nitric oxide (NO) production, tumor necrosis factor alpha (TNFa) production and gene expression of ferritin heavy chain (FTH1). We found that NK-92MI increased synthesis and release of NO and TNFa into the medium during co-culturing of NK-92MI cells with MCF-7 or MDA-MB-231 cells. Addition of iron inhibited the cytolysis of the breast cancer cell lines. The iron chelator deferoxamine (DFOM) increased NK-92MI cytolysis to MCF-7 or MDA-MB-231 cells. Iron reversed cytotoxicity to breast cancer cells induced by NO, released from S-nitroso-N-acetyl-penicillamine (NO donor). Real time quantitative polymerase chain reaction showed that iron upregulated the expression of FTH1 and iron chelator DFOM reduced FTH1 expression of MCF-7 and MDA-MB-231 cells. In conclusion, increased iron in cancer cells and their microenvironment protects cancer cells from natural killer cell cytolysis by antagonizing NO- and TNFα-associated cytotoxicity and by up-regulation of ferritin expression in breast cancer cells. Conversely, a decrease in iron concentration caused by DFOM improves natural killer cytolysis of tumor cells.

Iron is required for cell growth and proliferation, and changes in intracellular iron availability can have significant effects on cellular metabolism, cell-cycle regulation and cell division. Iron is a cofactor for several key enzymes in

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cellular respiration and metabolism, including enzymes of the citric acid cycle and ribonucleotide reductase. This latter enzyme catalyzes the reduction of ribonucleotides to deoxyribonucleotides, which is the rate-limiting step in DNA synthesis. Iron is also required for macromolecule biosynthesis, necessary for cell growth and division (1, 2). Neoplastic cells have an increased requirement for iron because of their rapid growth and proliferation. An increased requirement for DNA synthesis is accompanied by increased expression of the iron-dependent enzyme ribonucleotide reductase (RR) (3). This is achieved primarily through upregulation of transferrin receptors (TFRs). However, it has also been suggested that tumor cells may have alternate routes of iron uptake, and these routes may be critical in achieving increased intracellular iron levels under conditions in which TFRs are saturated (4). We previously reported that the MCF-7 human breast cancer cell line up-regulates the expression of iron importer genes including TFRs and solute carrier family 11, member 2 (SLC11A2) and down-regulates the expression of iron exporter SLC40A1 (ferroportin) (5). Free iron (Fe3+) is potentially hazardous to cells and has to be kept at very low concentrations. The protein ferritin has a high binding capacity for iron and protects cells from toxicity from free iron. Ferritin consists of two different subunits, known as heavy (H) and light (L) chains. H-Rich ferritin takes up iron faster than L-rich ferritin (6).

Iron deficiency has multiple effects on cell functions. Iron plays many critical roles in cancer progression through the cell cycle, and its deficiency can inhibit cell proliferation (7). Iron deficiency can significantly affect cellular metabolism by altering the expression and function of proteins that are central to the regulation of cellular metabolism and cellular respiration (8, 9). Therefore, the iron depletion induced by iron chelators is a potential method of cancer therapy (7, 10-13).

Both iron deficiency and iron excess can influence the functioning of the innate and adaptive arms of the immune system (14). Iron deficiency has been reported to be associated with increased susceptibility to infection in both experimental animals and humans, but the conflicting results

of these studies have made them hard to interpret. The conflicting results could be due to variations in baseline iron status, the severity of deficiency or possible co-existing nutritional problems (15). The situation is further complicated by the observation that iron supplementation in humans, particularly in the tropics, can increase the risk of infections such as malaria and tuberculosis (15-17). Iron overload caused by dietary excess, abnormal hemolysis or inherited disorders is also associated with greater susceptibility to infection (18-20). Studies of patients with hemochromatosis gene (HFE)-associated hemochromatosis demonstrated a decrease in circulating lymphocytes, with CD8+ cells being particularly affected (21, 22), but these abnormalities have not been consistently observed. Experimental iron overload in rodents is not associated with changes in lymphocyte numbers, but has been suggested to affect proliferative and cytokine responses (23,24). It has been shown that the iron-storing protein ferritin H-chain can inhibit tumor necrosis factor alpha (TNFα)-induced apoptosis by suppressing reactive oxygen species (25).

The association of iron in cancer cells and their microenvironment with host immunity is not clear. In the present study, we manipulated the level of iron in human breast cancer cell lines and their culture medium by iron supplementation or iron chelation. We also examined the cytolytic activity of the NK-92MI natural killer cell line against breast cancer cells.

Materials and Methods

Cell culture. The human breast cancer cell lines MCF-7 estrogen receptor-positive) and MDA-MB-231 (estrogen receptor-negative), and human NK-92MI natural killer cells were obtained from the American Type Culture Collection (Rockville, MD, USA). The MCF-7 and MDA-MB-231 cells were grown in alpha-minimum essential medium (α-MEM) supplemented with 10% fetal calf serum, 1 mM glutamine and 0.05 mg/ml gentamicin (Invitrogen, Carlsbad, CA, USA). NK-92MI is an interleukin-2 (IL2) independent natural killer cell line derived from the NK-92 cell line by transfection with human IL2 cDNA. The cell line is cytotoxic against a wide range of malignant cells (26). NK-92MI cells were grown in α -MEM without ribonucleosides and deoxyribonucleosides and with 2 mM L-glutamine, 1.5 g/l sodium bicarbonate, 0.2 mM inositol, 0.02 mM folic acid (Sigma Aldrich, St. Louis, MO, USA), 0.1 mM 2-mercaptoethanol, 12.5% horse serum, 12.5% fetal bovine serum and 0.05 mg/ml gentamicin (Invitrogen). All cell lines were incubated in 5% CO₂ at 37°C. Breast cancer cells were detached from tissue culture flasks by digestion with 0.05% trypsin and 0.53 mM EDTA.

Measurement of cell growth by colorimetric method. The MTT colorimetric method is based on the capacity of mitochondrial enzymes of viable cells to reduce the yellow soluble salt 3-[4,5-dimethylthiazol-2-yl]-2,5-diphenyltetrazolium bromide (MTT) to a purple-blue insoluble formasane precipitate which is quantified spectrophotometrically after dissolving in an organic solvent. The amount of MTT-formazan production is directly proportionate to

the number of viable cells. MCF-7 or MDA-MB-231 cells (20.000 cells/100µl per well) were plated in 96-well tissue culture plates. After 24 h of culture, the medium was replaced with 100 µl of media containing iron salt (FeCl₃) or iron chelator deferoxamine (DFOM). The plate was then incubated for 4 h. In order to measure the effect of iron on the cytotoxicity of the nitric oxide (NO) donor, S-nitroso-N-acetyl-penicillamine (SNAP) was added to breast cancer cells, 4000 cells/well were incubated with SNAP and iron for 72 h. The supernatant was then carefully removed and 100 µl of culture medium and 10 µl of 5 mg/ml of MTT solution was added to the wells. The plate was incubated for another 4 hours in 5% CO2 at 37°C. The medium was carefully removed then 150 µl of dimethyl sulfoxide (DMSO) was added to the wells. After 30 min the optical density (OD) was measured at a wavelength of 492 nm. Cell proliferation was expressed as a percentage that of the untreated control. All the experiments were carried out in triplicate wells and repeated three times independently.

Determination of cytolytic activity of human NK-92M natural killer cells. The MTT colorimetric method was adapted for evaluating the cytolytic activity of human NK-92MI cells towards breast cancer cells in vitro (27). We found that 100% of viable MCF-7 and MDA-MB-231 cells were able to attach to the wells of culture plates, but nonviable cells and cytolytic cells and NK-92MI cells floated in the medium after 4 h of incubation. NK cells, and dead and cytolytic breast cancer cells can therefore be removed by aspiration. Thus, we simplified the MTT colorimetric method for detection of cytolytic function of NK-92MI cells against breast cancer cells. First, 10,000 MCF-7 or MDA-MB-231 cells in 100 µl of medium were added to each well of a 96-well cell culture plate. NK-92MI (effector) cells were then added to the breast cancer target cells at effector/target cell ratios of 2.5:1, 10:1, and 40:1. Controls contained either breast cancer cells or NK-92MI cells alone. The final volume was 200 µl. Each plate was then incubated for 4 hours in 5% CO₂ at 37°C. The medium from each well was carefully collected. NO and $TNF\alpha$ concentration was measured in the supernatant. Then, 100 μl of complete αMEM and 10 µl of 5 mg/ml of MTT solution were added to each well. The plate was incubated for another 4 hours in 5% CO₂ at 37°C. The media were aspirated and 150 µl of DMSO was added to each well. The OD was measured at a wavelength of 492 nm. The percentage cytolysis by NK cells was calculated as follows: cytolysis (%) = [OD of the control of breast cancer cells - (OD of experimental wells with cancer cells and NK-92MI cells - OD of the control of NK-92MI cells)] ÷ (OD of the control of breast cancer cells) × 100. Each experiment was repeated three times and average cytolysis was calculated.

Effect of iron and DFOM on cytolysis of breast cancer cells by NK-92MI cells. Effector NK-92MI cells were co-cultured with breast cancer target cells at effector/target cell ratios of 2.5:1, 10:1 and 40:1. FeCl3 (400 μM) or DFOM (40 μM) was added to the mixture of NK-92MI and MCF-7 or MDA-MB-231 cells. Cells were incubated for 4 h in 5% CO₂ at 37°C. NK-92MI cytolysis of breast cancer cells was measured using the MTT colorimetric method. The media from wells of co-culturing NK-92MI and breast cancer cells were collected for measurement of NO and TNFα concentration. All experiments were repeated three times and the average cytolysis was calculated.

Measurement of NO in medium by the Griess Reagent System. Nitric oxide production was determined by measuring the nitrite (NO₂⁻) level, which is one of two primary, stable and nonvolatile breakdown products of NO, using the Griess Reagent System (Promega, Madison, WI, USA). Firstly, 50 µl of culture supernatant from controls and the co-cultured NK-92MI and breast cancer cells were added in duplicate to wells of a 96-well plate. Then 50 µl of sulfanilamide solution was dispensed to all experimental samples and wells containing the dilution series for the nitrite standard reference curve. The plate was incubated for 5-10 minutes at room temperature, protected from light. Finally, 50 µl of N-1napthylethylenediamine dihydrochloride solution was dispensed into all wells. The plate was incubated for 5-10 min at room temperature, protected from light. A purple/magenta color forms and its absorbance was measured within 30 minutes in a plate reader (Thermo Labsystems Multiskan MS, ThermoFisher, Waltham, MA, USA) with a 540 nm filter. The concentration of each sample was determined by comparison to the nitrite standard reference curve.

Human TNFα immunoassay. Human TNFα concentration was measured with a Quantikine HS TNFα ELISA kit (R&D Systems, Minneapolis, MN, USA). All reagents and standards were prepared as instructed. Fifty microliters of Assay Diluent RD1F was added to each well. Two hundred microliters of standards, supernatants of controls or the co-cultured NK-92MI and breast cancer cells were dispensed into duplicate wells. The plate was incubated for 3 hours at room temperature and washed six times. Two hundred microliters of Conjugate was added to each well. The plate was incubated for 2 hours at room temperature and washed 6 times. Fifty microliters of Substrate Solution was dispensed to each well and plates were incubated for 1 hour at room temperature, then 50 µl Amplifier Solution was added to each well and plates further incubated for 30 min at room temperature. Finally, 50 µl Stop Solution was added to each well and the OD of wells in plate was read at 492 nm within 30 min. The concentration of each sample was determined by comparison to the TNFα standard reference curve.

Real-time quantitative polymerase chain reaction (qPCR). We used real-time qPCR to determine the effect of iron (FeCl3) and iron chelator on ferritin heavy chain (FTH1) mRNA expression. The human MCF-7 and MDA-MB-231 breast cancer cell lines were treated with FeCl3 or the iron chelator DFOM for 4 hours at 37°C and 5% CO₂. The cells were removed from plates by trypsin-EDTA digestion. Total RNA of breast cancer cells was isolated with the PurLinkTM RNA Kit (Invitrogen). cDNA was then synthesized with High Capacity RNA-to-cDNA kit (Applied Biosystems, Grand Island, NY, USA). In brief, 2 µg of total RNA was mixed with 10 µl of 2x RT buffer and 1 µl of 20× Enzyme Mix and water to a total reaction volume of 20 µl. The reaction mixture was incubated for 60 minutes at 37°C and then for 5 minutes at 95°C to stop the reaction. The cDNA was ready for real-time PCR application or long-term storage in a freezer. FTH1 gene expression quantification was performed with TaqMan® Gene Expression Assays. The primers and probes (PrimeTime Mini qPCR assay) of FTH1 and. β-actin (ACTB) genes were synthesized by Integrated DNA Technologies (IDT, Coralville, IA, USA). The sequences were as follows: FTH1: TACCTGAA TGAGCAGGTGAAAG (forward), GATATTCCGCC AAGCCAGAT (reverse), and AGAATTGGGTGACCACGTGA CCAA (probe); ACTB: AGAAAGGGTGTAACGCAACTAA

(forward), GGATCA GCAAGCAGGAGTATG (reverse), and TCGTCCACCGCAAATG CTTCTAGG (probe). ACTB was used as internal gene control to normalize the PCRs for the amount of RNA added to the reverse transcription reactions. Then 40 µl of real-time qPCR reaction mixture containing 20 μl of TaqMan® universal PCR master mix (Applied Biosystems), 4 µl of 10× PrimeTime Mini qPCR assay (IDT) and 16 µl of cDNA (100 ng). The qPCR reaction was aliquoted in triplicate to wells of a 384-well PCR plate. The plate was sealed, briefly centrifuged, and reaction performed as using the 7900HT real-time PCR system (Applied Biosystems). The standard mode ran as 2 min at 50°C and 10 min at 95°C, and 40 cycles (15 sec at 95°C and 1 minute at 60°C). FTH1 gene expression was determined by relative quantification which related the signal of the target transcript in a treated group (iron- or DFOM-treated) to that of another sample such as an untreated control. We analyzed the relative quantification with RO Manager 1.2 software (Applied Biosystems). The relative expression of the studied samples was assessed using comparative delta-delta CT method and is presented as relative quantity (RQ) value.

Statistical analysis. The data were analyzed with a paired t-test and are presented as mean±standard deviation (SD). Findings were considered significant at p<0.05.

Results

NK-92*MI* cytolysis of breast cancer cells is associated with the synthesis and release of NO and $TNF\alpha$. NK-92MI and MCF-7 or MDA-MB-231 cells were co-cultured for 4 h in 5% CO_2 at 37°C with NK-92MI/cancer cell ratios of 2.5:1, 10:1 and 40:1. We found only low levels of NO and $TNF\alpha$ in the media of cultured NK-92MI cells. NO and $TNF\alpha$ were undetectable in the media from cultures of MCF-7 and MDA-MB-231 cells. When NK-92MI and cancer cells were co-cultured, levels of NO and $TNF\alpha$ were significantly increased (Table I). The addition of $FeCl_3$ (400 μM) or DFOM (40 μM) did not change the NO nor the $TNF\alpha$ level in the medium (Table II). These results suggest that cytolytic function of NK cells is associated with synthesis and release of nitric oxide and $TNF\alpha$.

Iron inhibits cytolysis of MCF-7 breast cancer cells by NK-92MI cells. In order to determine the effect of iron (FeCl₃) and iron chelator (DFOM) on cell cytolysis by NK killer, we first determined the appropriate concentration of iron or DFOM to be added to the culture media. FeCl₃ concentrations higher than 500 μ M significantly change the pH of the medium (turns media yellow). A FeCl₃ concentration of 400 μ M or less did not inhibit the growth of MCF-7 and MDA-MB-231 cells during 4 hours of incubation (all p>0.05). Thus we chose 400 μ M as the greatest concentration of iron to be used in culture. We cocultured 1×10^5 of NK-92MI and 1×10^4 MCF-7 or MDA-MB-231 cells (10:1) in media containing 50 μ M, 200 μ M and 400 μ M of FeCl₃. The control was a mixture of 1×10^5 of NK-92MI and 1×10^4 cancer cells without additional iron.

Table I. Concentrations of nitrite and tumor necrosis factor alpha (TNFa) in culture medium when NK-92MI cells were co-cultured with MCF-7 or MDA-MB-231 cells for 4 h.

| Cell line | Number of cells (ratio) ^a | Nitrite (μM) | TNFα (pg/ml) |
|--------------------|---|-----------------------|-----------------------|
| NK-92MI | 2.5×10 ⁴ | 0.50±0.09 | 0.24±0.12 |
| | 1×10 ⁵ | 1.10±0.16 | 0.52±0.22 |
| | 4×10 ⁵ | 1.50±0.18 | 1.36±0.45 |
| MCF-7 | 1×10 ⁴ | Undetectable | Undetectable |
| MDA-MB-231 | 1×10 ⁴ | Undetectable | Undetectable |
| NK-92MI+MCF-7 | $2.5 \times 10^4 + 1 \times 104 (2.5:1)$ | $1.20\pm0.26, p<0.05$ | $0.67\pm0.21, p<0.05$ |
| | $1 \times 10^5 + 1 \times 10^4 (10:1)$ | $3.29\pm0.68, p<0.01$ | $1.31\pm0.43, p<0.05$ |
| | $4 \times 10^5 + 1 \times 10^4 (40:1)$ | 4.30±0.75, p<0.01 | $2.50\pm0.48, p<0.05$ |
| NK-92MI+MDA-MB-231 | $2.5 \times 10^4 + 1 \times 1^{04} (2.5:1)$ | $0.99\pm0.25, p<0.05$ | $0.61\pm0.18, p<0.05$ |
| | $1 \times 10^5 + 1 \times 10^4 (10:1)$ | 2.89±0.96, p<0.05 | $1.40\pm0.48, p<0.05$ |
| | $4 \times 10^5 + 1 \times 10^4 (40:1)$ | 3.67±0.78, p<0.01 | 2.67±0.61, p<0.05 |

^aRatio of NK-92MI to MCF-7 or MDA-MB-231 cells. Data are the mean±standard deviation of three replicates. *p*-Values reported for comparison with NK-92MI cells by paired *t*-test.

Table II. Effect of iron (FeCl₃) and iron chelator deferoxamine (DFOM) on the level of nitric oxide and TNF α in culture medium when NK-92MI and breast cancer cells were co-cultured for 4 h.

| Cell line | | | Ratio of NK-92MI to MCF-7 or MDA-MB-231 cells | | | |
|------------|--------------|-------------|---|-------------------------|-------------------------|--|
| | Treatment | | 2.5:1 | 10:1 | 40:1 | |
| MCF-7 | Nitrite (μM) | 0 (control) | 1.20±0.26 | 3.29±0.68 | 4.30±0.75 | |
| | | 400 μM iron | 1.97±0.40 ^{NS} | 3.49 ± 0.57^{NS} | 3.89 ± 0.12^{NS} | |
| | | 40 μM DFOM | 1.05±0.31 ^{NS} | 2.73 ± 0.58^{NS} | 3.17±0.43 ^{NS} | |
| | TNFα (pg/ml) | 0 (control) | 0.67 ± 0.21 | 1.31±0.43 | 2.50±0.48 | |
| | | 400 μM iron | 0.83 ± 0.34^{NS} | 1.54 ± 0.63^{NS} | 2.63±0.63NS | |
| | | 40 μM DFOM | 0.51 ± 0.17^{NS} | 1.21±0.32 ^{NS} | 2.10±0.23NS | |
| MDA-MB-231 | Nitrite (µM) | 0 (control) | 0.99 ± 0.25 | 2.89±0.96 | 3.67±0.78 | |
| | , , | 400 μM iron | 1.23 ± 0.46^{NS} | 2.84±0.34 ^{NS} | 2.99±0.27NS | |
| | | 40 μM DFOM | 1.11 ± 0.37^{NS} | 3.12 ± 0.65^{NS} | 3.33±0.77NS | |
| | TNFα (pg/ml) | 0 (control) | 0.61±0.18 | 1.40±0.48 | 2.67±0.61 | |
| | 40 | 400 μM iron | 0.55±0.11 ^{NS} | 1.83±0.54 ^{NS} | 2.50±0.51 ^{NS} | |
| | | 40 μM DFOM | 0.81 ± 0.25^{NS} | 1.27±0.34 ^{NS} | 2.89±0.31 ^{NS} | |

NS: Not significant, compared to the medium control, p>0.05 by paired t-test. Data are the mean \pm standard deviation of three replicates.

After 4 hours of incubation, we used an MTT assay to measure cytolysis of breast cancer cells by NK-92MI cells. Both 200 μ M and 400 μ M of FeCl₃ significantly inhibited the cytolysis of cancer cells by NK-92MI cells (all p<0.05), but 50 μ M of FeCl₃ did not (p>0.05) (Table III).

The iron chelator DFOM increases NK-92MI cell cytolysis of breast cancer cells. DFOM concentration of 40 μ M or less did not suppresses the growth of MCF-7 or MDA-MB-231 cells (p<0.05). Therefore we used 40 μ M as the highest concentration of DFOM in the medium. MTT assays showed that both 10 μ M and 40 μ M of DFOM significantly increased the cytolysis of cancer cells by NK-92MI cells, but 2.5 μ M of DFOM did not (p>0.05) (Table III).

Iron neutralizes the cytotoxicity of NO to MCF-7 and MDA-MB-231 cells. To determine the effect of FeCl3 on the cytolytic function of NO during co-culturing of NK-92MI and breast cancer cells, we exposed breast cancer cells to NO released the NO donor SNAP with/without iron for 72 hours in 5% CO₂ at 37°C. MTT colorimetric assay showed SNAP alone inhibited the proliferation of breast cancer cells in a dosage-dependent manner. Addition of iron inhibited cytotoxicity of SNAP to MCF-7 cells (Figure 1). Iron similarly neutralized the cytotoxicity of SNAP to MDA-MB-231. Thus, NO from NK-92MI cells may deplete intracellular iron of MCF-7 and MDA-MB-231 cells. Iron inhibited the cytolysis of breast cancer cells by NK-92MI cells by reversing the iron-depleting effect of nitric oxide.

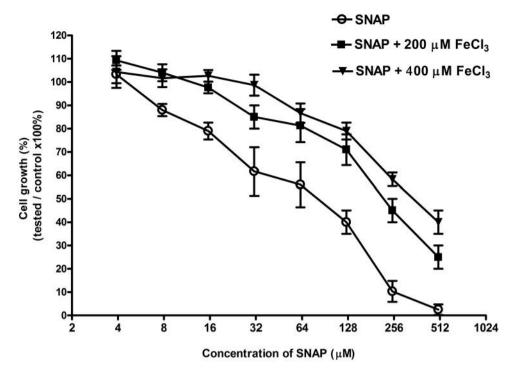


Figure 1. Iron inhibits the cytotoxicity of S-nitroso-N-acetyl-penicillamine (SNAP), a nitric oxide donor, to MCF-7 cells during 72 hours of incubation. Cell growth was determined by MTT assay.

Iron increases and DFOM reduces FHT1 mRNA expression in MCF-7 and MDA-MB-231. The relative expression of FHT1 was determined using the delta-delta CT method. Figure 2 shows that iron increases mRNA expression of FTH1 (Log₂RQ >0) in MCF-7 and MDA-MB-231 cells when treated with 50, 200 and 400 μ M of FeCl3 for 4 hours. On the other hand, the expression of FTH1 decreased (Log₂RQ<0) when breast cancer cells were incubated in media containing 2.5, 10 and 40 μ M of DFOM for 4 hours (Figures 2 and 3).

Discussion

Natural killer cells are large granular lymphocytes capable of destroying cells infected by virus or bacteria and susceptible tumor cells without prior sensitization and restriction by major histocompatibility complex antigens. Once the target is recognized by NK cells, their cytotoxic ability is mainly mediated *via* two main pathways. A membrane-disrupting protein, perforin, and a family of structurally related serine protease, granzyme, are secreted by exocytosis, which jointly induce apoptosis of the target cells. In the second pathway, caspase-dependent apoptosis takes place involving the association of death receptors (*e.g.* FAS/CD95) on target cells with their equivalent ligands such as FASL and tumor necrosis factor-related

Table III. Effects of iron (FeCl₃) and iron chelator deferoxamine (DFOM) on cytolysis of breast cancer cells by NK-92MI cells during 4 hours of incubation (details in the Materials and Methods).

| Cell line | Treatment | | Cytolysis (%) | <i>p</i> -Value |
|------------|-----------|-----|---------------|-----------------|
| MCF-7 | Control | | 26.1±1.4a | |
| | Iron (µM) | 50 | 22.2±1.5 | NS |
| | | 200 | 20.2±1.9 | < 0.05 |
| | | 400 | 18.9±1.3 | < 0.05 |
| | DFOM (µM) | 2.5 | 29.0±3.0 | NS |
| | · | 10 | 36.1±1.4 | < 0.01 |
| | | 40 | 33.9±3.7 | < 0.05 |
| MDA-MB-231 | Control | | 18.3±2.1 | |
| | Iron (µM) | 50 | 15.6±1.8 | NS |
| | • | 200 | 13.5±1.5 | < 0.05 |
| | | 400 | 12.1±2.2 | < 0.05 |
| | DFOM (µM) | 2.5 | 20.5±1.5 | NS |
| | | 10 | 24.6±3.1 | < 0.05 |
| | | 40 | 26.8±2.3 | < 0.01 |

^aData are the mean \pm standard deviation of three replicates. NS: Not significant, compared to the medium control, p>0.05 by paired t-test.

apoptosis-inducing ligand (TRAIL) on NK cells, resulting in caspase-dependent apoptosis. Cytotoxic factors, contained in the granules and released from NK cells after interaction with the target cells through mutual surface

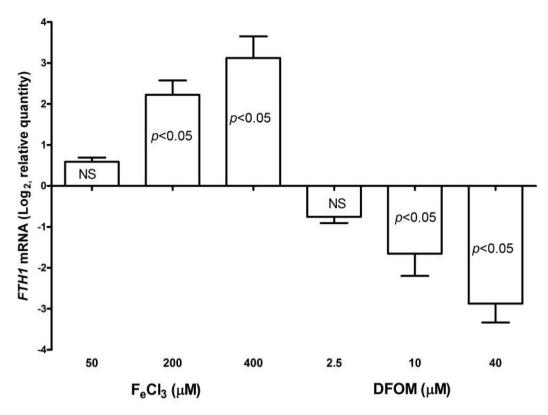


Figure 2. Iron up-regulates gene expression of ferritin heavy chain (FTH1); conversely, iron chelator deferoxamine (DFOM) inhibits FTH1 mRNA expression of MCF-7 cells during 4 h of incubation. NS: Not significant, compared to the control, p>0.05 by paired t-test (details are given in the Materials and Methods).

receptors, are able to induce both membrane and DNA damage in the target cells (28, 29). The function of NK cells, as well as their maturation and differentiation, are regulated by various stimuli, including IL2. In the present study, the cell line used, NK-92MI, is an NK cell line derived from the parent NK-92 cell line by transfection with human IL2 cDNA and therefore itself secretes IL2 (26). Great amount of research has shown that IL2stimulated NK cell tumoricidal activity correlates with increased NO production determined by measuring nitrite level in serum or culture media (30-33). NO can induce apoptosis and suppress tumor cell growth through depletion of intracellular iron in tumor cells. This effect of nitric oxide is similar to iron chelation by DFOM (34). In the current experiments, levels of NO are significantly increased when NK-92MI cells and MCF-7 or MDA-MB-231 cells are co-cultured for 4 hours (Table I). The increase in NO can rapidly deplete intracellular iron and induce apoptosis of breast cancer cells. Thus iron depletion by NO is one mechanism of NK cell cytolysis. NO not only suppresses target cells but also protects NK cells themselves from activation-induced cells death and maintains their lytic capacity (35). Figure 1 shows that iron supplementation can significantly mitigate the cytotoxicity of NO, when released from the NO donor SNAP, in breast cancer cells. These results agree with a previous report by Feger *et al.* (34). We also found that addition of iron to cocultured NK-92MI cells and MCF-7 or MDA-MB-231 cells significantly inhibited the cytolysis of MCF-7 or MDA-MB-231 cells by NK cells (Table III). However, iron and the iron chelator DFOM did not change the level of NO in the medium (Table II). Thus, iron partially suppressed cytolysis by NK-92MI cells by abrogating NO-induced iron depletion of MCF-7 and MDA-MB-231 cells.

Moreover, the availability of a high iron pool increases ferritin synthesis and, conversely, a low intracellular iron pool reduces ferritin synthesis by a post-transcriptional mechanism. This process is mediated by an interaction between RNA-binding proteins and a region in the 5' untranslated region of ferritin H and L chain mRNA termed the iron-responsive element (36,37). Real time qPCR shows the addition of iron increases mRNA of *FTH1* in MCF-7 and MDA-MB-231 cells (Figures 2 and 3). Ferritin plays a role in immune suppression of NK cells.

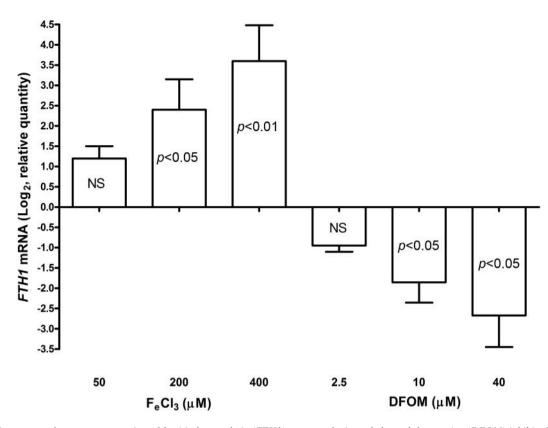


Figure 3. Iron up-regulates gene expression of ferritin heavy chain (FTH1); conversely, iron chelator deferoxamine (DFOM) inhibits FTH1 mRNA expression of MDA-MB-231 cells during 4 h of incubation. NS: Not significant, compared to the control, p>0.05 by paired t-test (details are given in the Materials and Methods section).

NK cells also release pro-inflammatory cytokines such as TNF α to control tumor and virus-infected cells. In this study, we found that TNFα significantly increased during NK cells lysizing MCF-7 and MDA-MB-231 cells. TNFa was produced by NK-92MI cells and was undetectable in MCF-7 and MDA-MB-231 cells (Table I). TNFα binds to cell surface TNFα receptor and then activates both cell-survival and celldeath mechanisms simultaneously. Activation of NF-kBdependent genes regulates the survival and proliferative effects of TNF, whereas activation of caspases and induction of reactive oxygen intermediates, ceramide, phospholipases and serine proteases, regulates the apoptotic effects (38, 39). Reactive oxygen species (ROS) produced by TNFα have an important function in cell death by activating c-Jun Nterminal kinase (JNK). However, the exact mechanism of mitochondrial ROS production, after TNFα stimulation, is not clearly understood. ROS modulator 1 is a molecular bridge between TNFa signaling and the mitochondria for ROS production that triggers TNFα-mediated apoptosis (38). Nuclear factor-kappa B (NF-kB) transcription factor antagonizes apoptosis induced by TNFα. NF-κB inhibits JNK activation and also up-regulates FTH1 which suppresses ROS accumulation through the mechanism of iron sequestration. The suppression of ROS accumulation consequently inhibits JNK signaling and apoptosis (40). These data support that upregulation of ferritin protects cells from ROS and TNF α -induced apoptosis. Down-regulation of ferritin may increase TNF α -induced apoptosis. We previously found that antisense oligonucleotides inhibiting the expression of FTH1 mRNA synergistically increased the cytotoxicity of recombinant TNF α to MCF-7 cells (41). It has been shown that DFOM depletes iron including free iron and bound-iron in ferritin and hemeprotein, and reduces the level of ferritin including H and L chains (42-44). Real-time qPCR showed that iron chelator DFOM reduced *FTH1* mRNA levels in MCF-7 and MDA-MB-231 cells (Figures 2 and 3).

In the present study, addition of iron into the cell culture medium increased the intracellular iron and ferritin of breast cancer cells. Increased iron in cells and the medium neutralized NO-induced cancer cytolysis by NK cells. Elevated ferritin can protect breast cancer cells from $TNF\alpha$ -induced cytotoxicity by NK cells. On the other hand, the iron chelator DFOM can tightly bind to iron in cells and culture medium, and down-regulate expression of FTH1.

Thus, DFOM reduces the antiapoptotic activity of iron and ferritin during the interaction of NK cells with targets such as MCF-7 and MDA-MB-231 cells. These results are very important for clinical practice as patients with chronic infectious diseases, such as chronic hepatitis and cancer, often have increased levels of iron and ferritin, that may impair the patient's immune surveillance (45-48). Manipulation of iron and ferritin by nutrition and iron chelation may improve the anti-infectious and antitumor function of the immune system.

Conflicts of Interest

No conflicting financial interest exists.

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