A Synergistic Antitumor Effect of Rituximab and Gamma Interferon Combined Therapy on Human CD20+ B-Cell Lymphoma Cells

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Abstract: Rituximab (RTX) is an anti-CD20 human-mouse chimeric monoclonal antibody that exhibits antibody-dependent, cell-mediated cytotoxicity and complement-dependent cytotoxicity, resulting in an antitumor effect with immune cells or complement. RTX is approved for the treatment of many diseases including B cell lymphoma and rheumatoid arthritis. We examined whether combined RTX and gamma interferon (IFNγ) therapy provides a higher antitumor effect than RTX single therapy using B-cell lymphoma cells. In addition, we investigated the mechanisms underlying the antitumor effect. We treated tumor-derived cell lines with RTX alone, IFNγ alone, or a combination of RTX and IFNγ (RTX-IFNγ). Untreated cells served as controls. We experimentally examined in vitro cell proliferation, conducted apoptosis and cell cycle assays, performed Western blotting to identify changes in the levels of proteins related to the cell cycle, and investigated tumor growth in a mouse xenograft experiment. Cell proliferation experiments indicated that RTX or IFNγ alone did not significantly suppress cell growth compared with the control, whereas treatment with RTX-IFNγ significantly suppressed cell proliferation. In vivo mouse experiments also showed that the administration of RTX-IFNγ significantly suppressed tumor growth compared to the single therapies. Some tumors in mice treated with RTX-IFNγ were completely resolved. The cell cycle assays revealed a significantly increased rate of cells in the G0/G1 phase following treatment with RTX-IFNγ compared with the other groups, and the levels of p27kip1 increased and the levels of cyclin E and Cdk 2 decreased in cells treated with RTX-IFNγ. Our findings suggested that RTX-IFNγ combined therapy directly affects cells by arresting the cell cycle at the G1/S checkpoint and had a synergistic antitumor effect compared to RTX single treatment of B-cell lymphomas. This combined therapy may change the mortality rate for B-cell lymphomas.

Key words: rituximab, gamma interferon, combined therapy, cell cycle arrest

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Introduction

Rituximab (RTX) is an anti-CD20 human-mouse chimeric monoclonal antibody that shows antibody-dependent cell-mediated cytotoxicity (ADCC) and complement-dependent cytotoxicity (CDC). It is also the first monoclonal antibody to be used for clinical cancer therapy. RTX is combined with several chemotherapies for treating B cell lymphoma and is used alone to treat autoimmune diseases such as rheumatism. Clinical trials have already provided evidence on the safety of RTX1-4.

Historically, it was anticipated that gamma interferon (IFNγ) would exhibit antitumor effects; however, clinical trials in the 1980s on its efficacy as a cancer therapy showed no antitumor effects, thus IFNγ was not pursued as a cancer therapy5. IFNγ is a type II interferon secreted as a cytokine by natural killer cells, dendritic cells, and CD8 T cells6-8. Several reports indicate that selected molecular target drugs require type I and II interferon for efficacy9.

We previously reported that combined anti-HER2 antibody and IFNγ therapy showed a higher antitumor effect than anti-HER2 antibody single therapy10. Her2 is regarded as an oncogene because overexpression of HER2 causes cells to become malignant. In contrast, CD20 is not an oncogene. In addition, cancers that overexpress HER2 form solid tumors, whereas lymphomas are humoral tumors.

In this study, we examined whether combined RTX and IFNγ therapy (RTX-IFNγ) provides a higher antitumor effect than RTX single therapy using B-cell lymphoma cells. We also investigated the mechanisms related to any synergistic antitumor effects observed.

Materials and methods

Cell lines and culture conditions

The Raji and RAMOS cell lines are Burkitt’s lymphoma cells that originated from B lymphocytes. Both cell lines were obtained from the American Type Culture Collection (ATCC, Manassas, VA) and were cultivated in RPMI-1640 growth medium containing L-glutamine and 10% fetal bovine serum (Biosera, Kansas City, MO).

Drugs

RTX was obtained from Chugai (Chugai Pharmaceutical Co., Ltd., Tokyo, Japan) and human IFNγ was obtained from PROSPEC (PROSPEC Protein Specialists, Rohovot, Israel). We diluted the RTX with phosphate-buffered saline (PBS) to a final concentration of 1 mg/ml and IFNγ to a final concentration of 1×10^5 IU/ml. For all RTX and IFNγ combined treatment experiments, IFNγ was added 30 minutes after the addition of RTX.

Cell growth assay

We used four groups of Raji and RAMOS cells: control group, RTX group, IFNγ group, and RTX-IFNγ group. Cells (1×10^5 of Raji or 5×10^5 of RAMOS) were seeded onto 6-well plates (2 ml of medium for Raji cells and 3 ml of medium for RAMOS cells), one for each of the
four groups, and cultured in RPMI-1640 containing L-glutamine and 10% fetal bovine serum. Drug treatment was initiated 8 hours after the cell passage, and then the cells were cultured for a further 24, 48, or 72 hours. The cells were counted at each time using an automated cell counter (Bio-Rad Laboratories, Inc., Hercules, CA) and a cell growth curve was constructed. No drugs were added to the control group, the RTX group was treated with 10 µg/ml of RTX, the IFNγ group was treated with 100 IU/ml of IFNγ, and the RTX-IFNγ group was treated with 10 µg/ml of RTX and 100 IU/ml of IFNγ.

*Cell apoptosis assay*

Cells (5×10^5) were seeded onto 6-well plates, and then treated for 48 hours with RTX, IFNγ, or RTX-IFNγ; the control group was not treated with drugs. Following treatment, the cell number was adjusted to between 1×10^6 and 1×10^7/100 µl in culture medium, and then stained using a Mouse Annexin V and Dead Cell Assay Kit® (Merck Group, Darmstadt, Germany). The cells were analyzed using a flow cytometer (Merck Group).

*Cell cycle analysis*

Cells (1×10^5) were seeded onto 6-well plates, then 2 mM hydroxyurea was added to each well to synchronize the cell cycle phase for 12 hours. The cells were then treated for 8 hours with RTX, IFNγ, or RTX-IFNγ, or left untreated as control. The cells were then washed with PBS, collected by pipetting, and fixed with 75% cold ethanol for 4 hours at −20°C. The cells were incubated and the cell cycle phase was determined using a flow cytometer and a Cell Cycle Assay Kit® (Merck Group).

*Lysate preparation and Western blotting*

Cells (1×10^6) were seeded onto 6-well plates and cultured with each drug or the drug combination for 48 hours, and then lysed in RIPA buffer (Wako Pure Chemical Industries, Ltd., Osaka, Japan) containing a protease inhibitor cocktail (Sigma-Aldrich, St. Louis, MO), before analysis by Western blotting. In brief, 40 µg of protein per lane was separated by 12% SDS-polyacrylamide gel electrophoresis, and then electroblotted onto PVDF membranes (Bio-Rad Laboratories) for 1 hour in PBS containing 5% nonfat milk (Wako Pure Chemical Industries, Ltd.) and 0.1% Tween 20 (Wako Pure Chemical Industries, Ltd.). The membranes were incubated overnight at 4°C with a rabbit antibody against p27<sup>kip1</sup> (Cell Signaling Technology, Danvers, MA), a mouse antibody against cyclin E (Santa Cruz Biotechnology, Dallas, TX), or a mouse antibody against Cdk 2 (Santa Cruz Biotechnology). The membranes were then washed three times in PBS containing 0.1% Tween 20 and incubated for 1 hour with either a goat anti-rabbit IgG antibody conjugated with horseradish peroxidase (SeraCare Life Sciences, Milford, MA) as a secondary antibody for the p27<sup>kip1</sup> primary antibody or a goat anti-mouse IgG antibody conjugated with horseradish peroxidase (Santa Cruz Biotechnology, Dallas, TX) as a secondary antibody for the cyclin E and Cdk 2 primary antibodies. Protein bands were measured using a LI-COR 3600®
computerized digital imaging system (LI-COR, Inc., Lincoln, NE).

Mouse in vivo experiment

Eight-week-old female C.B-17 SCID mice, which have a severe combined immunodeficiency affecting both B and T lymphocytes, were purchased (CHARLES RIVER LABORATORIES JAPAN, Inc., Yokohama, Japan) and bred for one week to allow adjustment prior to the experiment. Raji cells (1×10^7) were injected subcutaneously into both sides of the back of each mouse.

We divided the mice into four groups so that the average tumor size in each group was similar and initiated drug treatment 14 days after cell inoculation. The control group was treated with 100 µl PBS, the RTX group with 100 mg/100 µl RTX in PBS, the IFNγ group with 10,000 IU/100 µl IFNγ in PBS, and the RTX-IFNγ combined therapy group with both drugs. The drugs were delivered by intraperitoneal injection. For the combined therapy group, IFNγ was injected 30 minutes after the RTX injection. The experiment was performed for four weeks. Tumors were measured three times a week using a digital caliper and tumor size was calculated as length×width×height. The drugs were administered three times a week.

Statistical analysis

Statistical analysis was carried out using YSAT 2013 (Igakutosho-shuppan Ltd., Toda, Japan). Differences in mean values were statistically analyzed using non-repeated measures ANOVA, followed by Dunnnett’s test or the Student-Newman-Keuls test. Tumor volume differences between the RTX and RTX-IFNγ groups were statistically analyzed using unpaired Student’s t-test. The threshold for significance was P < 0.05 in all statistical analyses.

Results

Cell proliferation

- Raji cells

  RTX-IFNγ treatment was more effective at suppressing cell proliferation than RTX treatment on Day 2 and Day 3 after drug treatment. On Day 3, proliferation of the RTX-IFNγ group cells was 60% of the control group, whereas that of the RTX group was 89% of the control group. Cell counts were significantly decreased in the RTX-IFNγ treatment group compared to the control group, whereas the RTX and IFNγ groups showed no significant difference compared to the control group. Indeed, the IFNγ group showed an increased number of Raji cells (Fig. 1a).

- Ramos cells

  The same in vitro experiment using RAMOS cells showed that cell proliferation in the RTX-IFNγ group was 69% of the control group on Day 3, whereas proliferation of the RTX group was 88% of the control group on Day 3. Also on Day 3, the RTX-IFNγ group showed significant suppression of cell proliferation compared to the control group, whereas treatment with RTX or IFNγ alone showed no significant difference compared to the control group (Fig. 1b).
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Cell apoptosis assay

The percentage of apoptotic cells was not significantly different among the four groups, as shown by flow cytometry (Fig. 2).

Cell cycle assay

We subjected Raji cells to a cell cycle assay to investigate why RTX-IFNγ treatment suppressed cell proliferation without affecting apoptosis.

The percentage of cells in G0/G1 phase was significantly increased in the RTX-IFNγ group compared with that in the other three groups (Fig. 3), and there was no significant difference in the percentage of cells in S phase and G2/M phase among the four groups (data not shown).

Cells treated with RTX or IFNγ alone showed no significant difference in the percentage of cells in G0/G1 phase compared with the control group, and RTX induced G1/S checkpoint cell arrest only when combined with IFNγ.

Western blotting

We then investigated the cell arrest findings at the G1/S checkpoint in cells treated with RTX-IFNγ using Western blot analysis. The expression level of p27kip1 protein was increased and
that of both Cdk 2 and Cyclin E was decreased in the RTX-IFN\(\gamma\) group in the Raji cell line compared to the other groups (Fig. 4).

**Antitumor effect in vivo**

We conducted *in vivo* experiments in mice to investigate the antitumor effect of RTX-IFN\(\gamma\) administration and the effect of ADCC and CDC.
Rituximab and RTX-IFNγ significantly suppressed tumor volumes in mice compared to the control after Day 9. Treatment with IFNγ alone showed no antitumor effect in the mouse xenograft model (Fig. 5).

**Fig. 4.** Western blot analysis of G1/S checkpoint-related proteins

We investigated the protein expression levels of p27kip1, cyclin E, and Cdk 2, which are related to G1/S checkpoint arrest, using Western blotting. Raji cells (1×10⁶) were seeded in 6-well plates and cultured with each drug for 48 hours. The expression level of p27kip1 increased and that of cyclin E and Cdk 2 decreased in the RTX-IFNγ group compared to the other groups.

**Fig. 5.** Antitumor effect of RTX-IFNγ in the C.B-17 SCID mouse xenograft model

C.B-17 SCID mice were used. Raji cells (1×10⁷) were injected subcutaneously into both sides of the back of each mouse. We divided the mice into 4 groups: control, PBS 100 µl; RTX 100 mg/100 µl PBS; IFNγ 100 IU/100 µl PBS; RTX-IFNγ (100 mg/100 µl PBS + 100 IU/100 µl PBS) and started drug treatment at Day 14 post-injection (Day 0 in the figure). Tumor volumes were calculated as length×width×height. Measurements and drug treatments were performed three times a week. The curve shows the mean ± standard deviation of the volume of the tumors; the numbers of tumors shown as below (Cont : N = 4, RTX : N = 6, IFNγ : N = 5, RTX-IFNγ : N = 6).

Treatment with RTX and RTX-IFNγ significantly suppressed tumor volumes compared with the control after Day 9. Mice treated with RTX-IFNγ showed complete tumor resolution by Day 14, whereas treatment with RTX alone caused tumor regression. RTX-IFNγ significantly decreased tumor volumes compared with RTX alone after Day 7 (Dunnett’s test: *P < 0.05 and **P < 0.01 vs. control, unpaired Student’s t-test †P < 0.05 and ‡P < 0.01 vs. RTX alone).
Discussion

Tumor-derived cells were treated with RTX alone, IFNγ alone, or RTX-IFNγ. Untreated cells served as controls. We experimentally examined in vitro cell proliferation, conducted apoptosis and cell cycle assays, performed Western blotting to identify changes in the levels of proteins related to the cell cycle, and investigated tumor growth in a mice xenograft experiment.

In cell proliferation experiments, treatment with RTX or IFNγ alone induced no significant suppression compared with the control, whereas treatment with RTX-IFNγ significantly suppressed cell proliferation. In the in vivo mouse experiments, the administration of RTX-IFNγ significantly suppressed tumor growth compared with the single therapies. Cell cycle assays showed that the ratio of cells in G0/G1 phase significantly increased following exposure to RTX-IFNγ compared with the other groups. Furthermore, RTX-IFNγ treatment increased the level of p27kip1 protein and decreased the levels of Cyclin E and Cdk 2 protein in the single treatment groups.

Treatment of lymphomas with RTX

RTX is an anti-human CD20 antibody used to treat various diseases because normal B cells express CD20 on the surface from the Pro B cell phase. B-cell non-Hodgkin lymphoma is treated by administering RTX combined with chemotherapeutic agents such as cyclophosphamide, hydroxydaunorubicin, oncovan and prednisolone (CHOP) is used as a first-line treatment. RTX can enhance the antitumor effect of chemotherapeutics and decrease the required drug dose and associated side effects. Therefore, RTX may allow the safer use of chemotherapeutics by patients unable to tolerate full-dose chemotherapy.

A new mechanism for an antitumor effect: arresting the cell cycle

IFNγ alone showed no antitumor effect in either in vitro or in vivo experiments, whereas treatment with RTX-IFNγ showed a higher antitumor effect than RTX alone. This is consistent with a previous study showing that trastuzumab and IFNγ combined therapy extended tumor-free survival. Indeed, some tumors in the present study were completely resolved following treatment with RTX-IFNγ. RTX-IFNγ combined therapy may therefore strongly enhance the antitumor effects of clinical chemotherapeutics, allowing their dose to be reduced. Therefore, treatment with RTX-IFNγ might extend the survival rate of patients with B cell lymphoma.

RTX is thought to exhibit ADCC and CDC, resulting in an antitumor effect; however, ADCC and CDC require secondary signals. In the presented in vitro experiments, the medium lacked effector cells or active complement and RTX single therapy did not significantly reduce cell numbers compared with the control. In contrast, treatment with RTX-IFNγ reduced cell numbers in the absence of effector cells or active complement in the presented in vitro experiments.

Trastuzumab can induce antibody internalization within 24 hours after binding to erbB2 receptors on the cell membrane, resulting in apoptosis. Trastuzumab also can induce signal transduction and phenotypic changes in cells. In contrast, RTX does not induce antibody-
internalization and thus cannot induce cell apoptosis. We observed no significant differences in apoptosis among the four groups in the current study, thereby demonstrating that treatment with RTX-IFNγ has an antitumor effect without causing apoptosis and necrosis.

Suppressing the malignant potential of cancer by arresting the cell cycle

There are several reports that treatment of HER2-positive breast cancer with trastuzumab can change the malignant phenotype. The cell cycle assay used in this study showed a significant increase in the number of cells in the G0/G1 phase following treatment with RTX-IFNγ compared with the other three groups. We therefore investigated several proteins related to the G1/S checkpoint using Western blotting and found that p27\(^{kip1}\) increased and cyclin E and Cdk 2 decreased in cells treated with RTX-IFNγ. These changes might reflect cell cycle arrest at the G1/S checkpoint and suggest that RTX-IFNγ can change the malignant phenotype to arrest the cell cycle.

Cells undergoing abnormal cell proliferation tend to accumulate genetic mutations, which can in turn increase the malignant potential of cancers and remarkably increase patient mortality, because the mutations promote adaptation to the microenvironment, metastatic ability, and treatment resistance. RTX-IFNγ can affect the G1/S checkpoint and arrest the cell cycle leading to suppression of mutagenesis, thereby preventing the accumulation of gene mutation and subsequent enhancement of carcinogenesis.

RTX treatment for autoimmune diseases

RTX reduces the number of plasma cells with ADCC and CDC, and is used for the treatment of many diseases such as idiopathic thrombocytopenic purpura, rheumatoid arthritis, and systemic lupus erythematosus. In this study, we demonstrated that RTX-IFNγ reduced the number of B-cells, suggesting that treatment with RTX-IFNγ could be more effective for the treatment of these diseases compared to RTX alone.

Limitations

We did not investigate the antitumor effect of RTX-IFNγ combined with chemotherapies.

We also did not investigate markers related to the epithelial-mesenchymal transition, such as Snail, Slug, and TWIST, or the relationship between cell cycle arrest and epithelial-mesenchymal transition by measuring TGF-β.

Conclusions

RTX-IFNγ combined therapy may induce a synergistic antitumor effect superior to RTX single therapy. RTX-IFNγ acts via a new antitumor mechanism, affecting cancer cells by arresting the cell cycle. Our study suggests that RTX-IFNγ treatment may enhance clinical chemotherapies, reduce the dose of chemotherapeutics, and extend tumor-free survival. RTX-IFNγ is expected to arrest the cell cycle and therefore prevent increase in malignancy.
Conflict of interest disclosure

The authors have no conflict of interest to declare.

References


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