

Suppression of NF- κ B Activation in Normal T Cells by Supernatant Fluid from Human Renal Cell Carcinomas

T lymphocytes from patients with renal cell carcinoma (RCC) show reduced immune function and impaired activation of the transcription factor, NF- κ B. We determined the mechanism of NF- κ B suppression in T cells of RCC patient and determined whether supernatant fluid from RCC explants (RCC-S) induced the same phenotype of NF- κ B suppression in normal T cells that is observed in patient T cells. The pattern of κ B-binding activity in T cells of RCC patient was altered as compared to that seen in T cells obtained from normal volunteers. In some patients, no activation of RelA/NF κ B1-binding activity was detectable, while in others κ B-binding activity was modestly induced but the duration was reduced. I κ B α was degraded normally following stimulation in both normal controls and T cells from RCC patients. RCC-S did not alter the cytoplasmic levels of RelA and NF- κ B1 but did suppress their nuclear localization and inhibited the activation of RelA/NF- κ B1 binding complexes. These results show that RCC-S can induce in normal T cells the same phenotype of impaired NF- κ B activation that is detected in T cells of RCC patient. It also appears that NF- κ B suppression by RCC-S may contribute to the immunosuppression of host immunity.

Key Words: Transcription factors; T-lymphocytes; Kidney neoplasms; Proteins, NF- κ B

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INTRODUCTION

Among renal cell carcinoma (RCC) patients, 30% have metastases at the time of diagnosis. Two-year survival rate of RCC patient is 10-20% (1, 2). Metastatic RCC is ordinarily resistant to both radiation therapy and chemotherapy (3). Although the development of immunotherapy, including interferon- α , interleukin-2 and interleukin-12 has resulted in improved response rates in metastatic RCC, the treatment of metastatic RCC remains difficult with less than 25% of patients responding favorably (4). Several mechanisms contribute to the immune unresponsiveness of cancer patients, including immune suppression directly induced by the cancer through its production of soluble immune suppressive factors (5). T cells represent an important component of the host immune response against human tumor cells. The impairment of T cell function and signaling pathways observed in cancer patients may hinder the development of T cell immunity and reduce the effectiveness of immunotherapy. NF- κ B is important to T cell activation and regulates the inducible expression of a large number of genes necessary for the development of T cell

immunity. The κ B/Rel family of proteins is composed of p105/p50 (NF- κ B1), p65 (RelA), p100/p52 (NF- κ B2), c-Rel and RelB which bind the κ B sequence motif in homo or heterodimeric form (6-10). NF- κ B is present in an inactive form in the cytoplasm of T cells through interaction with inhibitory proteins, collectively termed I κ Bs. I κ B α inhibits both DNA binding activity and nuclear localization of RelA and c-Rel proteins (6, 11). The principle steps in the activation of NF- κ B involve the dissociation of dimeric forms containing RelA, c-Rel and NF- κ B1 from I κ B α after phosphorylation and subsequent degradation of I κ B α (6, 11, 12). We determined the mechanism of NF- κ B suppression in T cells of RCC patient and determined RCC-S induced suppression of NF- κ B activation in normal T cells is similar to that observed in T cells of RCC patient.

MATERIALS AND METHODS

Cell culture and isolation of T cells

Peripheral blood was obtained from patients with con-

firmed diagnosis of renal cell carcinoma and healthy volunteers. Tumor tissue was obtained from RCC patients and used to make tumor supernatants. Normal T cells were used as control and compared to patient T cells. Normal T cells from healthy donors were used for culturing with tumor supernatants. Peripheral blood lymphocytes (PBL) were isolated from healthy volunteers and RCC patients. T cells were purified as previously described (13, 14): PBL were subjected to Ficoll-Hypaque (Pharmacia, Piscataway, NJ, U.S.A.) density gradient centrifugation and then depleted of macrophages, B cells and NK cells by negative selection using magnetic separation with microbeads coated with antibodies to CD14, CD16 and CD19 (Miltenyi Biotec Inc., Sunnyvale, CA, U.S.A.). To obtain activated T cells, T cells (5×10^5 /mL) were cultured in the absence and presence of stimulus for various lengths of times. The medium used was RPMI1640 supplemented with 5% fetal calf serum. PMA (20 ng/mL) (Sigma, St. Louis, MO, U.S.A.) plus ionomycin (0.75 μ g/mL) (Sigma) or cross-linked anti-CD3 (10 μ g/mL) (Ortho Biotech, Rariton, NJ, U.S.A.) plus IL-2 (1000 IU) (Hoffman LaRoche, Nutley, NJ, U.S.A.) were used as stimulus.

Preparation of tumor supernatants

To generate tumor supernatant fluid, tumor tissues from primary renal cell carcinoma were obtained from patients undergoing radical nephrectomy. 3×3 mm explants were made from tumors using a No. 10 Bard-Parker scalpel and incubated overnight in RPMI1640 medium. Thereafter, 1 g of 3×3 mm explants were placed into a T-75 flask with 15 mL of Dulbecco's Modified Eagles Medium without additional supplements. After three days of culture at 37°C with 95% air and 5% CO₂ the supernatant fluid was harvested, filtered and stored at -70°C until use. To determine the effect of tumor-derived soluble products on normal T cells, T cells were cultured in complete RPMI1640 with and without RCC-S. The supernatant fluid was added to T cell cultures, using 20-50% dilution.

Preparation of cytoplasmic and nuclear extracts

Cytoplasmic and nuclear extracts were prepared according to Schreiber et al. (15) with minor modifications. T cells (10×10^6 /mL) were harvested and washed with cold PBS, and then sedimented by centrifugation. The cell pellet was resuspended in 150 μ L of buffer A (10 mM HEPES, pH 7.9, 10 mM KCl, 0.1 mM EDTA, 0.1 mM EGTA, 1 mM DTT, 100 μ g/mL PMSF, 2 μ g/mL aprotinin, 2 μ g/mL leupeptin, 100 μ g/mL Pefabloc and 100 μ g/mL chymostatin) by gentle pipetting. The cells

were incubated on ice for 15 min and then 10 μ L of 10% Nonidet P-40 solution (Sigma) was added and centrifuged. The supernatant fluid containing the cytoplasmic extracts was aliquoted. Pelleted nuclei were resuspended in 50 μ L of buffer C (25% glycerol, 20 mM HEPES, pH 7.9, 0.4 M NaCl, 1 mM EDTA, 1 mM EGTA, 1 mM DTT, 100 μ g/mL PMSF, 2 μ g/mL aprotinin, 2 μ g/mL leupeptin, 100 μ g/mL Pefabloc and 100 μ g/mL chymostatin). After mixing at 4°C for 20 min and centrifuge, supernatants containing the nuclear proteins were taken and stored at -70°C. Protein concentration was measured with a commercial kit (Pierce BCA kit, Rockford, IL, U.S.A.).

Western blot analysis

Protein samples (5-10 μ g) were mixed with equal volume of 2X Laemmli sample buffer, boiled and resolved by electrophoresis in 10% sodium dodecyl sulfate-polyacrylamide gels. The proteins were transferred from the gel to nitrocellulose membranes. The membranes were blocked by incubating in 5% nonfat dry milk in TBST overnight. The membranes were sequentially incubated with specific antibody (two hr for anti-I κ B α and anti-c-Rel, one hr for anti-RelA and anti-NF- κ B1) and then with horseradish peroxidase-conjugated donkey anti-rabbit IgG (Amersham, Chicago, IL, U.S.A.; 1:2000 dilution in TBST). The membranes were washed with TBST and specific immune complexes were detected by enhanced chemiluminescence (Dupont, Boston, MA, U.S.A.).

Electrophoretic mobility shift assay (EMSA)

EMSA was performed using nuclear extracts obtained at various times after stimulation with either PMA plus ionomycin or cross-linked anti-CD3 plus IL-2. Nuclear extracts (5-10 μ g) were incubated in a 25 μ L total reaction volume containing 20 mM HEPES, pH 7.9, 80 mM NaCl, 0.1 mM EDTA, 1 mM DTT, 8% glycerol and 2 μ L of poly (dI-dC) (Pharmacia, Piscataway, NJ, U.S.A.) for 15 min at 4°C. The reaction mixture was then incubated with radiolabeled oligonucleotide (2×10^5 cpm) for 20 min at room temperature. The samples were analyzed by electrophoresis in 6% non-denaturing polyacrylamide gel with 0.25X TBE buffer (22.3 mM Tris, 22.2 mM boric acid, 0.5 mM EDTA). The gels were dried and analyzed by autoradiography. An oligonucleotide corresponding to κ B element from the IL-2R α gene was prepared by using an Applied Biosystems oligonucleotide synthesizer (model 381A). The sequence of the oligonucleotide was: 5'-CAACGGCAGGGGAATCTCCCTCTCCTT-3'. The underlined sequence represents the κ B motif.

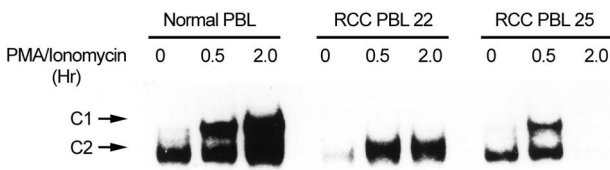


Fig. 1. Analysis of stimulus-induced κ B binding activity in normal and patient T cells by EMSA. Normal PBL, normal T cells; RCC PBL, T cells of RCC patient.

RESULTS

κ B-binding activity in T cells from RCC patients

T cells from normal donors and RCC patients were stimulated with PMA/ionomycin and κ B-binding activity of those cells were analyzed by EMSA. In normal T cells, stimulation with PMA/ionomycin resulted in the appearance of two distinct κ B motif-specific DNA-binding activities (designated C1 and C2 in Fig. 1). Both complexes were observed within 30 min of stimulation and remained elevated for at least two hr. In six of patient-derived T cells (n=10), the C1 complex was not detectable. In T cells obtained from the other four patients, the C1 activity was observed after 30 min of stimulation but returned to basal levels in two hr (Fig. 1) (Table 1). Nuclear extracts from patient T cells showed constitutive expression of the C2 complex and the abundance of this activity was increased by stimulation. Similar pat-

Table 1. κ B/Rel protein expression and NF- κ B activity in N-PBL, P-PBL and N-PBL with RCC-S

Expression & activity	N-PBL (n=5)	P-PBL (n=10)	N-PBL+RCC-S (n=10)
+	5		
-		6	6
+/-		4	4

N-PBL, normal T cells; P-PBL, T cells of RCC patient; RCC-S, supernatant from RCC explants

terns of response were observed when normal and patient T cells were stimulated with anti-CD3/IL-2 (data not shown).

Expression of NF- κ B and I κ B α in T cells from RCC patients

The nuclear localization of RelA, c-Rel and NF- κ B1 in the different cell populations was examined by Western analysis of cytoplasmic and nuclear extracts at various times following stimulation. In normal T cells, expression of c-Rel, RelA and NF- κ B1 was increased in the nucleus after 30 min and two hr of stimulation. There were decreased expression of RelA and c-Rel in the nuclei of patient T cells suggesting impaired translocation (Fig. 2). Alterations in levels of nuclear RelA were the most dramatic. In four of 10 RCC patient T cells modest but significant levels of RelA were detected within the nucleus after 30 min of stimulation but faded into the background in two hr (Fig. 2A). In six other samples nuclear RelA was undetectable at any time (Fig. 2B) (Table 1). Levels of NF- κ B1 and c-Rel in the nucleus were increased modestly following stimulation. Interestingly, no C1 κ B-binding activity was seen under these conditions. T cells from normal donors and RCC patients were stimulated for various times with PMA/ionomycin to determine the change in I κ B α , an inhibitor of NF- κ B activity, by Western analysis. A characteristic decrease in I κ B α expression was observed within 30 min in normal T cells (Fig. 3) which coincides with the nuclear appearance of RelA (Fig. 2). In patient-derived T cells (n=10), the magnitude and kinetics of I κ B α degradation were similar to that of normal T cells (Table 2). These observations demonstrate that early events in NF- κ B acti-

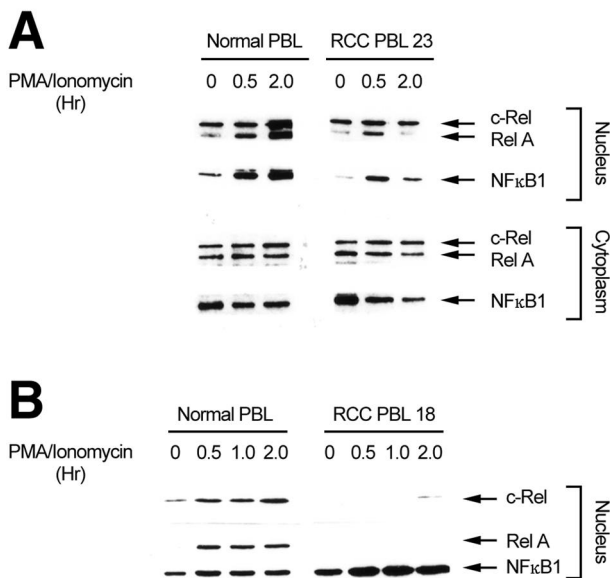


Fig. 2. Altered nuclear translocation of RelA and c-Rel showed in patient T cells.

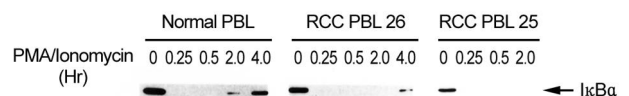


Fig. 3. Normal degradation of I κ B α following stimulation is observed in normal T cells and patient T cells.

Table 2. Degradation of I κ B α in N-PBL, P-PBL and N-PBL with RCC-S

Degradation	N-PBL (n=5)	P-PBL (n=10)	N-PBL+RCC-S (n=10)
+	5	10	8
-			2

Abbreviations are as noted in Table 1

vation are normal in patient T cells.

Effect of RCC-S in NF- κ B activation

Normal T cells were cultured in RCC-S overnight and then stimulated with PMA/ionomycin to determine whether RCC-S could alter activation of NF- κ B in normal T cells. NF- κ B proteins remained in the cytoplasm but failed to undergo nuclear translocation after stimulation (Fig. 4A). κ B-binding activity was also suppressed by RCC-S as determined by EMSA (Fig. 4B) (Table 1). In most cases (n=8), RCC-S did not alter the normal degradation of I κ B α . In a few experiments (n=2), the degradation of I κ B α was reduced by RCC-S (Table 2). This frequency is similar to that observed in T cells from RCC patient (n=10).

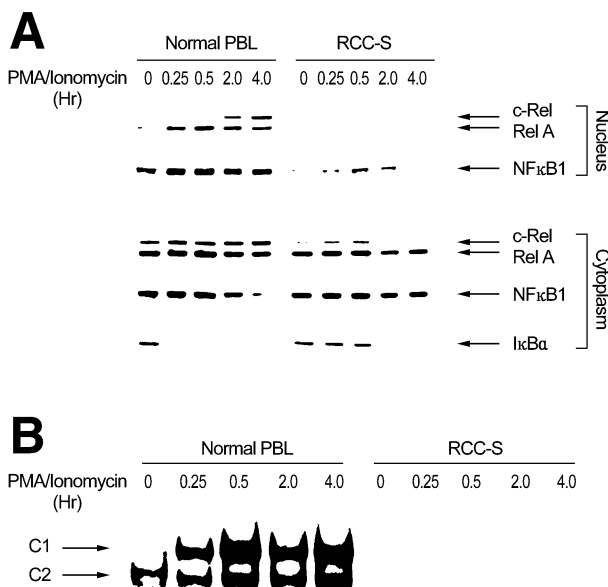


Fig. 4. RCC-S suppresses the activation of NF κ B in normal T cells. (A) Western blot: RCC-S inhibits the nuclear translocation of RelA, c-Rel and NF κ B1. The degradation of I κ B α occurs normally in normal T cells exposed to RCC-S in some experiments. (B) EMSA: RCC-S suppressed κ B DNA binding activity.

DISCUSSION

The development of cellular immunity is dependent on the activation of T cells. Activation-dependent stimulation of new gene transcription is now well documented, involving transacting factors such as NF- κ B which regulates the expression of many genes important to T cell function and growth (6, 16, 17).

T cells from cancer patients or tumor-bearing animals are impaired in their response to extracellular stimuli including the activation of κ B-specific DNA-binding activity (14, 18). Our results show that there are alterations in both the expression and DNA-binding activity of NF- κ B family members of RCC patient T cells. The major inducible κ B-specific binding complex (C1) either did not appear in the nucleus or was modestly activated and exhibited a markedly reduced time course. The failure of c-Rel and RelA to accumulate normally in the nuclei of patient T cells was due to an event occurring after phosphorylation and degradation of the inhibitor I κ B α since this latter process occurred normally in RCC patient T cells.

Although the biological nature of the soluble products in tumor supernatants is not known, the culture supernatant of tumor cells may produce a variety of immunosuppressive molecules such as IL-10, TGF- β 1 and prostaglandin E2 (5). They may contribute to immune unresponsiveness of cancer patients and inhibit the activation of NF- κ B. Our results demonstrate that RCC-S can induce in normal T cells the same phenotype of impaired NF- κ B activation that is detected in T cells of RCC patient. These alterations in NF- κ B activity may be due to a product(s) of renal tumor cells. Stimulus-dependent degradation of I κ B α in the cytoplasm was not impaired by RCC-S in most experiments. However, the degradation of I κ B α was reduced by RCC-S in two out of 10 cases. Thus, there appears to be at least two different mechanisms by which RCC-S can suppress NF- κ B activation. First, NF- κ B proteins do not translocate to the nucleus at all. Second, NF- κ B proteins degrade in the nucleus after translocation.

In conclusion, two patterns of altered NF- κ B activation and alterations in NF- κ B activity induced by RCC-S may contribute to the lack of antitumor immunity in RCC patients.

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