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Usage of T Cell Receptor Repertoire is Restricted in Synovial Lymphocytes in Rheumatoid Arthritis

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= Abstract =

Background: Rheumatoid arthritis is an autoimmune disease characterized by a chronic inflammatory process, primarily involving the synovial membrane of peripheral joints, where T cell activation is found. To address the superantigen stimulation in rheumatoid arthritis, T cell clonality and the expression of activation markers were analyzed. **Methods:** To detect *TCRBV* usage, inverse PCR and sequencing were done. Monoclonal antibodies were used for flow cytometric analysis of *TCRBV8* or *TCRBV5*. As results, a restricted usage of *TCRBV3* gene was detected in synovial lymphocytes from one rheumatoid arthritis patient. However, preferential usage for *TCRBV8*, which may be one indicator for stimulation by staphylococcal superantigen, was not obvious although general activation of T cells was found as high DR+ percentage in synovial T cells. These data show specific antigen rather than superantigen might involve the pathogenesis of rheumatoid arthritis.

Key Words: , . T Chain

, 가 (2).

T

1% TCR (T cell

(1). 80% 가 receptor) 가 .

가 sublining layer

가 . 40-50%가 T

CD4+ 가 CD4:CD8 4:1 14:1

CD4+ 가

lymphoid aggregate

lymphoid aggregate

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가 CD4+ (3). 1990 T 가 TCR V β family V gene repertoire (4) T TCR 가 T 가 가 T 가 superantigen T 가 (6-8)가 (9,10) T 가 superantigen TCR repertoire T TCR repertoire가 (11). 가 가 RNA RNeasy kit (Qiagen, Santa Claris, CA) First strand cDNA random primer (pd(N)₆, Pharmacia Fine Chemicals, Uppsala, Sweden) RNA 1 μ g random primer 0.1IU, dNTP 200 mM, DTT 800mM reverse transcriptase (Gibco BRL, Gaithersburg, MD) 400U 42 60 cDNA DNA polymerase 가 16 2 cDNA T4 DNA ligase 가 second stranded cDNA circularization PCR PCR degenerate primer forward primer 5'-GGG TCG ACC TGT GCA CCT CCT TCC CAT T-3' , inverse primer 5'-GCA TGC GGC CGC ATG GCC ATG GTC AAG AGA-3' primer 20pM, dNTP 200mM, MgCl₂ 2mM, KCl 50mM 0.001% gelatin 10mM Tris-HCl (pH; 8.3) Taq polymerase (Bioneer,) 2.5unit 가 PCR 94 1 , 55 1 , 72 2 30 . 30 72

10 elongation . PCR 1.5% agarose gel
 PCR 10µl . Inverse PCR
 Fig. 1

Gene cloning

50ng pT7Blue T-vector (Novagen, Madison, WI)
 PCR 0.2 pM DNA ligase (Gibco BRL,
 Gaithersburg, MD) 2U 가 16 2
 ligation . Competent cell (genotype : endA1
 hsdR17 (rk-, mk+) supE44 thi-1 gyrA96 relA1 lac [F',
 proAB, lacqZDM15, Tn10(tetr) recA] transformation
 50µg/MØ ampicillin, 15mg/MØ tetracycline, X-gal
 IPTG (Gibco BRL, Gaithersburg, MD)가 가
 LB agar plate clone
 . clones Sal I Not I
 insert (300 bp)가 clone automatic
 sequencer (; ALFexpress, Amersham Pharmacia
 Biotech AB, Uppsala, Sweden)

National Center for Biotechnology Information
 GenBank chain family

CD3, CD4, CD8 DR
 (Becton Dickinson, San Jose, CA) TCR β
 TCRBV5 TCRBV8 (T Cell Diagnostics, Cambridge,
 MA) . 5µ
 1 10µl 가 30 4
 0.1% bovine serum albumin PBS 2

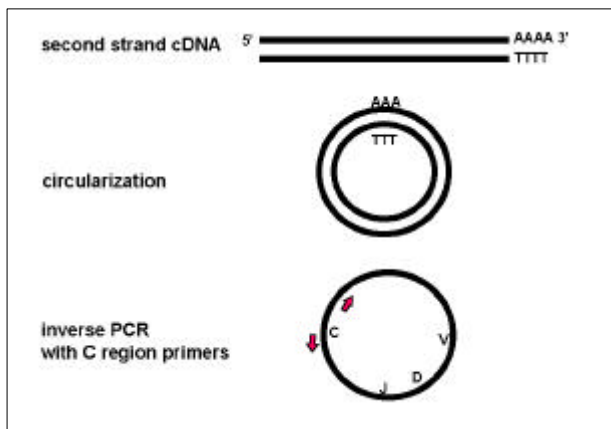


Fig. 1. Scheme of inverse PCR

(FACstar, Becton Dickinson, San
 Diego, CA)

1. TCRB V

(RA-1 RA-2)
 inverse PCR 300 bp
 PCR (Fig. 2). RA-1
 18
 TCRBV3
 10
 TCRBV3

T
 (Table 1).

2. TCRB V8 TCRB V5

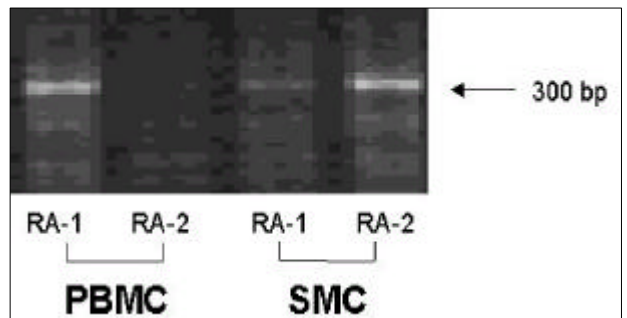


Fig. 2. Inverse PCR product. PCR was done with degenerate primers of TCRBC. PCR products were loaded on 1% agarose gel. PBMC; peripheral blood mononuclear cells. SMC; synovial mononuclear cells.

Table 1. Distribution of TCRBV family among T cell cDNA clones

| | TCRB V | No. of positive clones |
|-------------------|----------|------------------------|
| Patient (RA-1) | TCRB V1 | 1 |
| | TCRB V3 | 10 |
| | TCRB V5 | 4 |
| | TCRB V7 | 1 |
| | TCRB V21 | 2 |

Inverse PCR using C region primers was done to identify V region genes. Total 18 clones were tained and sequencing was done.

| | | T | T |
|-----------------------------------|------------------------|-----------------------------|-----------------------------|
| Staphylococcal toxin superantigen | | 7.8% | 1.9% 4.6% |
| <i>TCRB V8</i> T | | (Table 3). | CD3+ DR+ |
| <i>TCRB V8</i> | | RA-1 | 가 RA-2 |
| T (4.7 ± 1.7%) | T (2.3 ± | (Table 3). | 가 |
| 1.8%) | (Table | | |
| 2). | (RA-2) | <i>TCRB V5</i> | |
| | 6.0% | | |
| superantigen | <i>TCRB V5</i> | | |
| T (2.9 ± 2.4%) | T (4.1 ± 1.9%) | TCR β chain family | α chain family β |
| (Table 2). | | chain family | 가 |
| 3. | | , α chain allelic exclusion | |
| 2 | | (17) allele | |
| 50 U/ml | rIL-2 | α chain | β chain |
| 가 | | T | (18)가 |
| <i>TCRB V</i> | | PCR | semi-quantitation RNA |
| CD8+ 가 | CD4+ | | |
| 가 | (Table 3). | | |
| | anti- <i>TCRB V5</i> | 17% (| HLA-DQ2 가) |
| | anti- <i>TCRB V8</i> | CD8+ | 20-40%가 <i>TCRA V12.1</i> 가 |
| | RA-1 | 가 | |
| <i>TCRB V5+</i> 가 0.8%, | <i>TCRB V8+</i> 가 1.2% | | |
| , <i>TCRB V5+</i> 가 1.8%, | <i>TCRB V8+</i> | (19, 20) | |
| 가 1.0% | RA-2, <i>TCRB V5+</i> | <i>TCRA V12.1+</i> CD8+ | |
| <i>TCRB V8+</i> | 6.0% | <i>TCRA V12.1+</i> 가 | |

Table 2. *TCRB V5* or *TCRB V8* positive T cells in rheumatoid arthritis patients

| Lymphocytes | | <i>TCRB V5</i> positive % | <i>TCRB V8</i> positive % |
|----------------------|------|---------------------------|---------------------------|
| Rheumatoid arthritis | PBMC | 2.9 +2.4* | 2.3 +1.8*** |
| | SMC | 4.1 +1.9** | 4.7 +1.7**** |
| Healthy control | PMBC | 3.7 +1.0 | 3.7 +0.7 |

Cells were reacted with anti-*TCRB V5* or anti-*TCRB V8* monoclonal antibodies and a total of 10,000 to 20,000 cells were analyzed by FACstar. N= 21 for rheumatoid arthristis and 15 for healthy control. Data are expressed as mean + S.D.. PBMC; peripheral blood mononuclear cells. SMC; synovial mononuclear cells. Student's T test was done. *, **, ***, **** ; not significant at the p>0.01 level versus PBMC from healthy control group.

Table 3 Surface markers of synovial T cells of rheumatoid arthritis after *in vitro* culture

| positive % | Fresh | | | | | 3 months after culture | | | | |
|------------|-------|------|--------|----------------|----------------|------------------------|------|--------|----------------|----------------|
| | CD8 | CD4 | DR/CD3 | <i>TCRB V5</i> | <i>TCRB V8</i> | CD8 | CD4 | DR/CD3 | <i>TCRB V5</i> | <i>TCRB V8</i> |
| RA-1 | 50.3 | 27.9 | 14.3 | 0.8 | 1.2 | 11.3 | 64.5 | 29.8 | 1.8 | 1.0 |
| RA-2 | 48.6 | 30.1 | 26.4 | 6.0 | 7.8 | 16.8 | 55.9 | 35.9 | 1.9 | 4.6 |

Cells were stained with each monoclonal antibodies and a total of 5,000 to 20,000 cells were analyzed by FACstar.

가 T
 inverse PCR T 2
TCRB V recombinant IL-2 가 3
TCRB V3 CDR3 CD4+ 가 가 DR+
 가 TCR
TCRBV3 T T
TCRB V3 T recombinant IL-2 가
 IL-2
*TCRB V3*가 가 가
 T chain
 family CD4+ T 가 (3)
*TCR VB3*가 superantigen CD8+
 가
 가 T , 가
 T 가 3 CD4+ T
 T (11). CD8+ T CD4+ T 가
 Superantigen staphylococcal toxin 가 IL-2
 superantigen CD4+ T 가 IL-2
TCRB V8 T 가
TCRB V8 T T DR 가
 가
 superantigen T 2.2 ± 1.0 (11)
TCRB V5 T DR+ T
 T 가 T
 가
 T 가 DR+ 가 CD4+ T CD8+ T
 T CD4+
 T 가
 (20) T TCR repertoire T
 TCR (20). 가
 T 40% IL-2
 TCR chain 가 T IL-15 T
 (21) cartilage proteoglycan 가
 (22, 23) type II collagen(24) 가 가 (25).
 T
 T
 가 가
 TCR repertoire

- 가 가 broad spectrum
가 T (26)
- T
- T
- 가 T T
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