

	2. SEW2871 (5 mg/kg; Cayman Chemical) dissolved in a vehicle (PBS containing 5% acidified DMSO and 3% fatty acid-free BSA) or vehicle (23).	123 124 125
	3. JTE013 (3 mg/kg; Tocris Bioscience) dissolved in a vehicle (PBS containing 5% acidified DMSO and 3% fatty acid-free BSA) or vehicle (24).	126 127 128
	4. GST-RANKL (2 mg/kg, dissolved in PBS) (28).	129
	5. 29- or 30-G insulin syringes (Becton Dickinson).	130
	6. An indwelling needle: a 30-G needle attached to PE-10 tubing (Becton Dickinson).	131 132
2.6. Image Analysis		
	1. Imaging analysis software: Imaris (Bitplane) or Volocity (PerkinElmer).	133 134
	2. After Effects (Adobe).	135
2.7. In Vitro Migration Assay		
	1. EZ-TAXIScan (Effector Cell Institute, GE HealthCare; see Note 10) containing “41 Glass,” a small O-shaped ring, an EZ-TAXIScan chip, a rubber gasket, a holder base, wafer housing, a wafer clamp, a syringe guide, and a large O-shaped ring.	136 137 138 139 140
	2. Sample loading tip attached to a plastic syringe (Becton Dickinson).	141 142
	3. Microsyringe (10 μ L; MS-E10MIC; Exmire).	143
	4. 25-mm Thermanox plastic coverslip (Nunc; see Note 11).	144
	5. Scissors and tweezers.	145
	6. Serum-free Dulbecco’s modified Eagle’s medium (DMEM).	146
	7. Serum-free alpha-minimum essential medium (α MEM) and α MEM with 10% fetal calf serum (FCS), containing 1% penicillin and streptomycin.	147 148 149
	8. PBS, pH 7.4.	150
	9. 0.02% EDTA in PBS.	151
	10. Raw 264.7 cells (American Type Culture Collection, ATCC).	152
	11. Male or female C57BL/6 mice (6-9 weeks old).	153
	12. Syringe with 26-G needle.	154
	13. Mouse recombinant macrophage colony-stimulating factor (M-CSF; 100 ng/mL; PeproTech).	155 156
	14. Chemoattractant: S1P (10^{-6} , 10^{-7} , and 10^{-8} M; Enzo Life Sciences), dissolved in a vehicle (DMEM; see Note 12).	157 158
	15. ImageJ software (National Institutes of Health, NIH), equipped with an add-on program, MT Track J.	159 160

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3. Methods

3.1. *Intravital Two-Photon Imaging*

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1. Start up the two-photon microscope and turn on the heater in the environmental chamber (see Note 13).
2. All the procedures on mice are performed under anesthesia (see Note 14). Shave the hair and apply hair-removal lotion on top of the mouse's head (see Note 15). Cut the skin minimally with iris scissors for insertion of the O-ring. Fix the O-ring on the parietal bone with adhesive and petrolatum or difloil grease (see Note 16), which prevents leakage of PBS and fills the O-ring.
3. Insert an indwelling needle into the tail vein to treat the mouse with reagents during observation.
4. Intravenously inject 100 μ L of 2 mg/mL 70-kDa Texas Red-conjugated dextran in PBS (see Note 17).
5. Immobilize the mouse on the custom-made stereotactic holder as tightly as possible to avoid drift owing to respiration and pulsation (Fig. 1; see Note 18).
6. Focus on the bone marrow cavity at an appropriate depth and look through ocular lenses with the help of a mercury lamp. Change the light source from the mercury lamp to the Ti-Sapphire laser and the optical path to the NDD. Set the zoom ratio, z-positions, the interval time, and the duration time using observation software attached to the microscope (Fig. 2).
7. Inject 100 μ L of each reagent per mouse.
8. Analyze images by measuring cellular velocities, migration lengths, and contact times using image processing and analysis software.

3.2. *In Vitro Migration Assay*

3.2.1. *Bone Marrow-Derived M-CSF-Dependent Mononuclear Cells*

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1. Harvest bone marrow cells from C57BL/6 mice by flushing serum-free α MEM medium using a 26-G syringe.
2. Wash the harvested cells twice with α MEM containing FCS and culture the cells in α MEM containing FCS and 100 ng/mL M-CSF for 3 days.
3. Wash cells once with PBS and treat them with 0.02% EDTA in PBS.
4. Collect suspended cells and culture them in α MEM containing FCS and 100 ng/mL M-CSF for another 3 days.

3.2.2. *Setting of EZ-TAXIScan*

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1. Place "41 Glass" in holder base. Drop DMEM in the center of the glass and place a coverslip over the liquid (see Note 19). Wipe off excess fluid, making sure no air bubbles are trapped between the glasses.

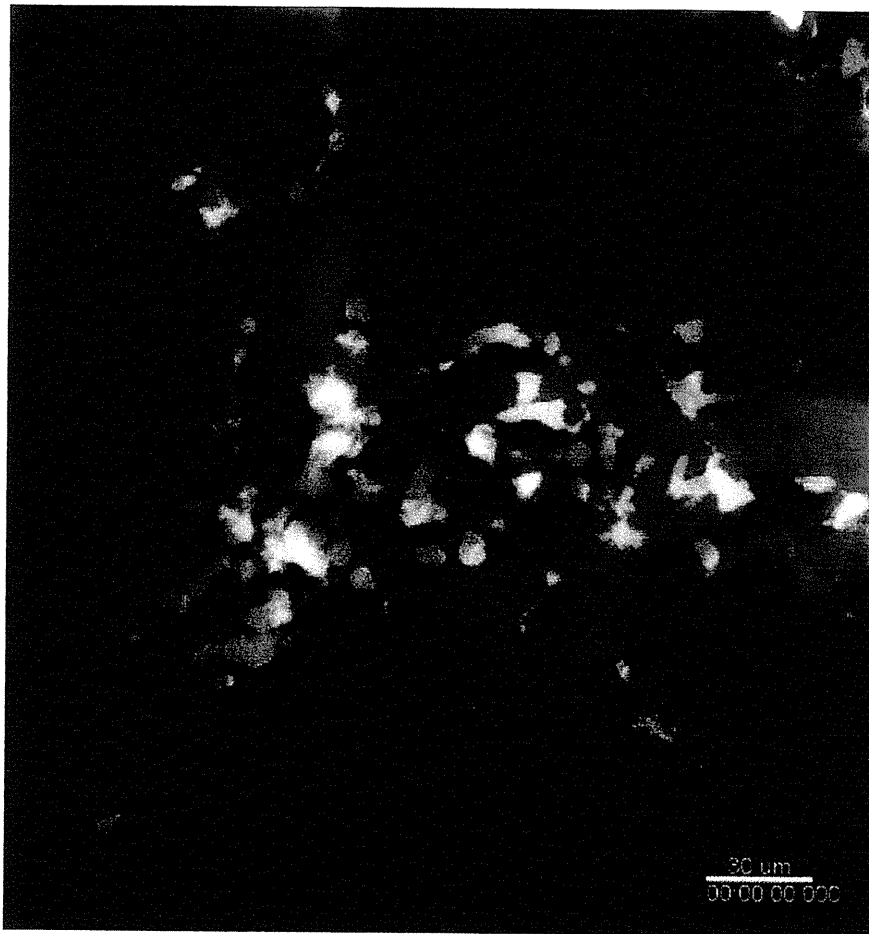


Fig. 2. Osteoclast precursors visualized by intravital two-photon imaging. Murine parietal bone of heterozygous Cx_3CR1 -EGFP knock-in mice is visualized. CX_3CR1 -EGFP-positive cells appear *green* in the bone marrow cavity. Collagen fibers in bone are detected by second-harmonic generation (*in blue*) and the blood vessels are visualized by 70-kDa dextran-conjugated Texas Red.

2. Place the small “O-shaped ring” on top of the wafer housing. 202
Gently close the inner lever and fix the wafer housing. 203
3. Fill inside the wafer housing with 4 mL of DMEM. 204
4. Place the EZ-TAXIScan chip gently into the wafer housing 205
(see Note 20), making sure no air bubbles are trapped. To 206
protect the chip, place the rubber gasket beneath the wafer 207
clamp. Place wafer clamp on wafer housing. Place the large 208
O-shaped ring upside of the wafer housing. Gently close the 209
outer lever (see Note 21). 210
5. Place the assembled device on top of the preheated 211
EZ-TAXIScan microscope. 212

3.2.3. Alignment of Cells

1. Remove 1 mL of medium from the upper side of the holder. 213
2. Add 1 μ L of cells (Raw 264.7 cells or bone-marrow derived 214
M-CSF-dependent mononuclear cells) to the second chamber 215
using a microsyringe with a syringe guide. 216

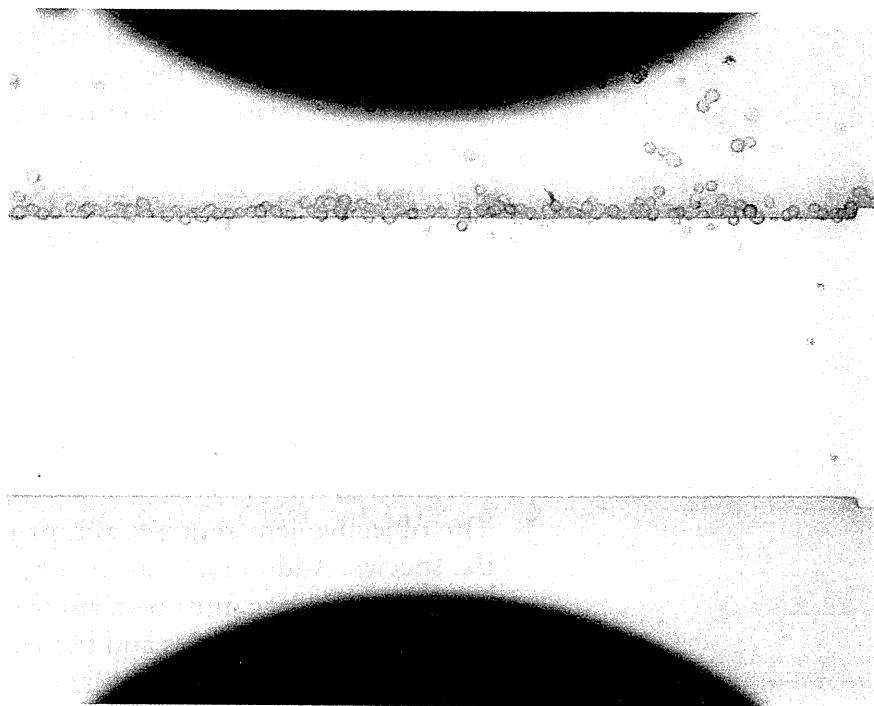


Fig. 3. Alignment of cells on one side of the EZ-TAXIScan chip before analysis. Before analysis of cell migration, cells should be aligned in the chamber on one side.

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3.2.4. Analysis of Chemotaxis

3. Remove 8–10 μL of medium from the third chamber using a microsyringe.
 4. Monitor alignment of the cells on one side (Fig. 3).
 5. Return the removed medium inside the holder.
1. Begin image acquisition using 1-min intervals.
 2. Add 1 ml of the chemoattractant (S1P: 10^{-6} , 10^{-7} , and 10^{-8} M) to the third chamber.
 3. Sequential image data are processed and migration speeds and tracking distances are calculated using analysis software.

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4. Notes

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1. The two-photon microscopy setup is also available from other microscope manufacturers (Zeiss, Nikon, and Olympus). Regarding objective lenses, higher NA and longer WD should be desirable. Bone marrow can be observed through an inverted microscope.
2. A femtosecond-pulsed infrared laser is also available from Coherent (Chameleon).
3. The more channels the NDD has, the more colors can be detected.

[AU2]

4. As sufficient space is necessary to place a living mouse between the objective lens and the stage, we replaced the normal stage for the section with a customized one. 236
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5. Because temperature is a critical factor for cell mobility, a decrease in body temperature of the animal must be prevented. Thus, we set up an environmental chamber, which can enclose the animal with the microscope stage and the objective lens. 239
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6. Global S1PR1 deficiency causes embryonic lethality at e12.5 to e14.5 due to defective blood vessel development (29). 243
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7. Although S1PR2-deficient mice suffer from auditory impairment due to vessel defects in the inner ear, they survive and reproduce (25). 245
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8. The objective lens requires almost the same WD as that from the species. Additionally, as the objective lens is water-immersible, the substance must be filled with the same refractive index as water between species and the lens. The O-ring works both as a spacer and a PBS reservoir. 248
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9. We used a red dye in this protocol because the target cells express EGFP. If the target cells are red, dextran-conjugated fluorescein isothiocyanate (FITC) can be used. Far-red dyes, such as Qdot-650, can be an alternative if the NDD has a channel that can detect long-wavelength signals. As blood vessels inside the bone marrow cavity have relatively high permeability, dextran with molecular weights over 70 kDa should be used. 253
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10. The EZ-TAXIscan is a visually accessible chemotactic chamber in which one compartment containing ligand (for example, SIP of various concentrations) and another compartment containing cells are connected by a microchannel. A stable concentration gradient of chemoattractant can be reproducibly formed and maintained through the channel without medium flow. Phase-contrast images of migrating cells are acquired at 1-min intervals. 261
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11. 0.2 mm-thick cover slips coated with other reagents (such as collagen and fibronectin) can also be used. 268
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12. Stock solutions of SIP are difficult to make. To accomplish this, SIP should be dissolved in BSA (4 mg/mL)-containing buffers or in other organic solvents with sonication or gentle warming (at 45–60°C). 270
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13. It takes some time for the laser and the temperature to stabilize. 274
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14. All animals must be handled according to institutional and national guidelines and regulations under an approved protocol. 276
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15. Remove hair as much as possible to avoid hair coming into the visual field because it produces strong background autofluorescence. 278
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16. Avoid glue contamination of the visual fields: some glues can produce autofluorescence.
 17. With excretion of the dextran into urine, fluorescence fades after several hours. If necessary, additional dosage of the dye should be added.
 18. Do not fasten too tightly because the animal can be hurt.
 19. The coated side should be set upward.
 20. The EZ-TAXIScan chip should be handled gently with tweezers. To protect it from drying, place it in a liquid at all times. After using, the EZ-TAXIScan chip should be sonicated and stored in 20–70% ethanol at room temperature. There are four types of EZ-TAXIScan chips with depths of 4, 5, 6, and 8 μm . The suitable size depends on the cell types. The 8- μm -depth chip should be used for Raw 264.7 cells and 5- μm -depth chip for bone marrow-derived M-CSF-dependent mononuclear cells.
 21. Before setting up the assembled device, make sure no air bubbles remain inside the chamber. If air bubbles are present, remove them with the sample-loading tip attached to the plastic syringe.

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Baseline anti-citrullinated peptide antibody (ACPA) titers and serum interleukin-6 (IL-6) levels possibly predict progression of bone destruction in early stages of rheumatoid arthritis (ERA)

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Abstract A prospective study was made to seek for a convenient biomarker to predict progression of bone destruction (PBD) in early stages of rheumatoid arthritis (ERA). All participated patients had definite RA and their radiographic stages were mild less than stage II of the Steinbrocker classification, naïve for treatment of any DMARDs or corticosteroids. After the entry, they were treated according to the 2002 ACR management guideline

for RA. The candidate biomarkers (RF-IgM, RF-IgG, CARF, ACPA, CRP, ESR, NTx, MMP-3, IL-6 and osteopontin) were measured at the entry. PBD was assessed radiographically by interval changes in the modified Sharp scores (Δ SHS) for 24 months. The associations between Δ SHS and baseline biomarkers were assessed statistically by multivariate regression analyses. Both the baseline ACPA and IL-6 levels correlated with PBD, suggesting that they could predict PBD in ERA.

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Keywords Rheumatoid arthritis · Bone/joint destruction · Biomarker · Prediction · Anti-citrullinated peptide antibody (ACPA) · Interleukin-6 (IL-6)

Introduction

Rheumatoid arthritis (RA) is a systemic autoimmune disease of unknown origin characterized by chronic destructive polyarthritis which leads to disability and increased mortality. Although its etiology is unknown, the disease is thought to develop through three main processes such as autoimmunity, inflammation and subsequent bone resorption. On the other hand recent cumulative evidence suggests that early diagnosis and therapeutic interventions at the early stages of RA are important [1]. However, clinical course or prognosis of RA is not uniform but various. Therefore, the individual patient needs an optimal therapeutic strategy according to their prognosis. In order for this, the individual clinical course or prognosis should be predicted accurately. Although several candidate predictive biomarkers have been reported, none has been definite [2].

The aim of this prospective study was to seek for a useful biomarker measurable in the blood samples to

predict progression of bone destruction (PBD) in mild radiographic stages of RA under the real clinical circumstances. The representative biomarkers were selected as a candidate according to the main disease processes such as autoimmunity, inflammation and bone resorption.

Materials and methods

Subjects

This study has been approved by the Ethical Committee of NHO Osaka Minami Medical Center and registered at the Japan Pharmaceutical Information center (JAPIC), No. Japic CTI-070480. Written informed consent was obtained from every patient enrolled in this study. A total of 52 patients with RA (12 males and 40 females, average age 56.5 years) were enrolled in this study. All the patients were fulfilled with the 1987 ACR classification criteria for RA [3] and their radiographical stages were relatively mild less than stage II of the Steinbrocker criteria [4]. They had not been used either any disease modifying anti-rheumatic drugs (DMARDs) or corticosteroids before the entry. After the entry, they were treated according to the 2002 ACR management guideline for RA [5]. At the endpoint (24 M), more than 90 % of the patients was administered DMARDs, (methotrexate, 76.5 %, 7.8 ± 2.6 mg/week; salicylazosulfapyridine, 19.1 %; leflunomide, tacrolimus and buccilamine, <5 %), and 85.5 % of the patients received a low dose of corticosteroids (prednisolone 4.7 ± 2.8 mg/day), and 10.6 % of the patients were treated with biologics (infliximab, etanercept, tocilizumab).

Biomarkers

The candidate biomarkers were selected according to the main disease processes such as autoimmunity, inflammation and bone resorption. They were measurable in blood samples and previously suggested to be associated with bone destruction in RA [6–11]. They included rheumatoid factor-IgM (RF-IgM), RF-IgG, anti-agalactosyl IgG antibody (CARF) and anti-citrullinated peptide antibody (ACPA) for autoimmunity, C-reactive protein (CRP), erythrocyte sedimentation rate (ESR), interleukin-6 (IL6) for inflammation, type I collagen cross-linked N-telopeptide (NTx) and matrix metalloproteinase 3 (MMP3) for bone resorption, and osteopontin (Op) for inflammation and bone resorption. They were measured at the entry according to the individual standard method. Anti-CCP2 antibody was measured using the DIASTATTH Anti-CCP kit (Axis-Shield Diagnostics Limited, United Kingdom) with the cutoff at 4.6 U/ml. IgM-RF was measured using RF-the latex X1 fixation test (DenkaSeiken, Tokyo, Japan)

(cutoff value: 18 IU/ml). CARF was measured using the Eitest CA-RF enzyme immunoassay (EIA) (Sanko Junyaku, Tokyo, Japan) (cutoff value: 6.0 AU/ml). MMP-3 was measured using Panakuria MMP-3 plate for ELISA (Daiichi Fine Chemical, Toyama, Japan) (cutoff value: male 121 ng/ml, female 59.7 ng/ml). CRP was measured using high sensitivity CRP-the latex fixation test (Denka Seiken, Tokyo, Japan) (cutoff value: 0.3 mg/dl). IL-6 was measured using a sandwich ELISA kit for human IL-6 (R&D Systems Inc, Minneapolis, USA) (cutoff value: 9.96 pg/ml). OPN was measured using a sandwich ELISA kit for human OPN (Immuno-Biological Laboratories, Gunma, Japan).

Assessment of radiographical progression of bone/joint destruction (PBD)

The conventional radiographs of both hands and feet were scored on joint space narrowing (JSN) and erosions (E) according to the van der Heijde modified Sharp scores (SHS) [12] at the entry of study (A) and 24 M later (B) by two trained observers. PBD was assessed by subtracting (A) from (B), defined as Δ SHS (Δ TSS, Δ E and Δ JSN).

Statistical analyses

All statistical analyses were undertaken using the statistical package for JMP ver. 8.0 (SAS Institute, Tokyo, Japan). Associations between biomarker levels and radiographic progression (Δ SHS) were assessed by multivariate linear regression analyses with variable step-wise (backward) selection technique (cut-off $p < 0.20$). All tests were two-sided and conducted at the 0.05 significant level.

Results

Correlations between the individual biomarkers and radiographical progression (Δ SHS)

The correlations between the baseline individual biomarker and radiographical progression were presented by Pearson's correlation coefficients (Table 1).

Radiographic progressions (Δ SHS: Δ TSS, Δ E, Δ JSN) were strongly correlated with the individual baseline score (TSS*, E*, JSN*). Significant correlations ($p < 0.05$) were also noted in IL-6 and ESR for Δ TSS, in ACPA for Δ E and in IL-6 and ESR for Δ JSN. Therefore, we decided that the baseline SHS (TSS*, E*, and JSN*) was added but CARF was omitted for further statistical analyses to avoid multicollinearity because of the strong correlation ($|r| > 0.9$) with RF-IgM. ESR was also omitted because of missing data.

Table 1 Pearson 's correlation coefficients and *p* values

	Objective variables			Explanatory variables					
	ΔE	ΔJSN	ΔTSS	Age	E*	JSN *	TSS*	RF-IgM	RF-IgG
ΔE	1								
ΔJSN	0.4354 0.0013	1							
ΔTSS	0.8174 <0.0001	0.8743 <0.0001	1						
Age	-0.0266 0.8515	-0.0099 0.9447	-0.0207 0.8845	1					
E*	0.6358 <0.0001	0.3865 0.0047	0.5900 <0.0001	0.0845 0.5515	1				
JSN*	0.4718 0.0004	0.3821 0.0052	0.4988 0.0002	0.2436 0.0819	0.6464 <0.0001	1			
TSS*	0.6073 <0.0001	0.4234 0.0018	0.5982 <0.0001	0.1838 0.1922	0.9005 <0.0001	0.9139 <0.0001	1		
RF-IgM	0.0053 0.9712	-0.0370 0.8010	-0.0205 0.8890	-0.0367 0.8024	-0.0725 0.6204	-0.0152 0.9174	-0.0475 0.7460	1	
RF-IgG	0.2200 0.1373	0.2474 0.0936	0.2768 0.0596	-0.1449 0.3311	0.1515 0.3092	-0.0076 0.9597	0.0775 0.6046	0.0244 0.8736	1
CRP	0.0912 0.5201	0.1698 0.2287	0.1578 0.2638	0.0773 0.5861	0.0863 0.5428	0.0277 0.8456	0.0617 0.6638	-0.0371 0.8004	0.1415 0.3428
ESR	0.2833 0.1441	0.5492 0.0025	0.4801 0.0097	0.1756 0.3713	0.1611 0.4128	0.1023 0.6046	0.1407 0.4751	-0.1506 0.4534	0.2175 0.2858
MMP3	0.2003 0.1771	-0.0207 0.8902	0.0955 0.5231	0.2420 0.1013	0.4288 0.0026	0.1315 0.3781	0.3049 0.0372	-0.0688 0.6534	0.0112 0.9406
NTx	-0.0963 0.5197	-0.0269 0.8574	-0.0694 0.6431	0.2730 0.0634	-0.0109 0.9420	-0.0964 0.5194	-0.0598 0.6898	-0.1142 0.4551	0.0089 0.9529
IL-6	0.2078 0.1520	0.4461 0.0013	0.3969 0.0047	0.0920 0.5294	0.1697 0.2438	0.1985 0.1715	0.2037 0.1603	-0.0915 0.5407	0.2672 0.0760
CCP	0.3252 0.0334	0.0405 0.7966	0.2059 0.1854	-0.1457 0.3511	0.0506 0.7473	0.1378 0.3781	0.1039 0.5075	0.0854 0.5909	0.3844 0.0109
CARF	-0.0482 0.7476	-0.0159 0.9155	-0.0363 0.8087	0.0863 0.5635	-0.0578 0.6995	0.1561 0.2946	0.0562 0.7077	0.9378 <0.0001	-0.0829 0.5795
OPN	-0.0297 0.8345	0.1298 0.3590	0.0671 0.6367	-0.1721 0.2224	-0.122 0.3887	-0.1172 0.4080	-0.1371 0.3519	0.1309 0.3702	0.1421 0.3408

Table 1 continued

	Explanatory variables							
	CRP	ESR	MMP3	NTx	IL-6	CCP	CARF	OPN
ΔE								
ΔJSN								
ΔTSS								
Age								
E*								
JSN*								
TSS*								
RF-IgM								
RF-IgG								
CRP	1							
ESR	0.6462 0.0002	1						
MMP3	0.0678 0.6505	0.2626 0.1949	1					
NTx	0.0604 0.6866	0.3458 0.0836	0.1213 0.4166	1				
IL-6	0.1688 0.2464	0.5170 0.0081	0.1524 0.3177	0.3622 0.0145	1			
CCP	-0.1274 0.4155	0.1804 0.4102	-0.0986 0.5295	-0.2524 0.1025	-0.1390 0.3800	1		
CARF	-0.1090 0.4659	-0.1523 0.4576	-0.0427 0.7755	-0.1382 0.3541	-0.1694 0.2658	0.1963 0.2071	1	
OPN	0.2664 0.0563	0.2633 0.1759	-0.1070 0.4743	0.1158 0.4381	0.2981 0.0375	-0.0968 0.5367	-0.1686 0.2572	1

The correlations between the individual biomarker and radiographical progression were presented by Pearson's correlation coefficients. Radiographic progression (ΔSHS: ΔTSS, ΔE, ΔJSN) were strongly correlated with the individual baseline score (TSS*, E*, JSN*). In addition, significant correlations ($p < 0.05$) were noted in IL-6 and ESR for ΔTSS, in ACPA for ΔE and in IL-6 and ESR for ΔJSN. The correlation between RF-IgM and CARF was strong ($|r| > 0.9$)

Upper line: correlation coefficients; lower line: p values

Assessment of associations between the baseline biomarkers and Δ SHS by multivariate regression analyses with step-wise selection

At the first step, multivariate regression analyses were performed in all variables except ESR and CARF. Then, the variables were deleted one by one (cutoff $p < 0.2$) and the analyses were repeated. Finally, for Δ TSS, IL-6 and ACPA as well as TSS* were selected. For Δ E, ACPA and E* were selected. For Δ JSN, IL-6 and JSN* score were selected. Then, the final analysis was performed using these selected variables. As a result shown in Table 2, the associations remained significant in IL-6 ($p = 0.0091$) as well as in TSS* ($p = 0.0001$) but not in ACPA ($p = 0.1176$) for Δ TSS. For Δ E, ACPA showed significant associations ($p = 0.0129$) as well as E* ($p < 0.0001$). For Δ JSN, IL-6 still showed significant associations ($p = 0.0039$) as well as JSN* ($p = 0.0262$).

Discussion

The aim of this prospective study was to seek for a useful and convenient biomarker to predict PBD in mild

Table 2 Multivariate analysis with selected variables

Variable	Regression coefficient	<i>p</i> value
(A) Δ TSS		
TSS*	0.1819	0.0001
IL-6	0.1358	0.0091
ACPA	0.0051	0.1176
(B) Δ E		
E*	0.2298	<0.0001
ACPA	0.0044	0.0129
(C) Δ JSN		
JSN*	0.1165	0.0262
IL-6	0.1010	0.0039

Associations between biomarkers levels and radiographic progression (Δ SHS) were assessed by multivariate linear regression analyses with variable step-wise (backward) selection technique. At the first step, multivariate regression analyses were undertaken in all variables except ESR and CARF. Then, the variables were deleted one by one (cutoff $p < 0.2$) and the analyses were repeated. Finally, in Δ TSS, IL-6 and ACPA as well as the baseline TSS (TSS*) were selected. In Δ E, ACPA and the baseline E score (E*) were selected. In Δ JSN, IL-6 and the baseline JSN score (JSN*) were selected. Then, the final analysis was performed using these selected variables. The associations remained significant for IL-6 ($p = 0.0091$) as well as the baseline TSS ($p = 0.0001$) but not for ACPA ($p = 0.1176$) in Δ TSS. In Δ E, ACPA showed significant associations ($p = 0.0129$) as well as the baseline E score ($p < 0.0001$). In Δ JSN, IL-6 still showed significant associations ($p = 0.0039$) as well as the baseline JSN score ($p = 0.0262$). All tests were two-sided and conducted at the 0.05 significant level

radiographic stages of RA under the real clinical circumstances. The candidate biomarkers in this study were selected by taking consideration into the main pathogenic disease processes such as autoimmunity, inflammation and bone resorption. There were two main findings. Firstly, the baseline SHS (TSS*, E*, JSN*) significantly correlated with PBD (Δ TSS, Δ E and Δ JSN). All patients enrolled in this study had short disease duration (<2 years) and with mild bone destruction (less than stage II of the Steinbrocker classification). They still stayed at so-called “the Window of Opportunity” stage. This strongly suggests and supports the importance of earlier therapeutic interventions in RA as emphasized in lots of previous clinical studies [1].

Secondly, among the candidate biomarkers, the baseline serum ACPA titers and IL-6 levels showed significant associations with PBD. It is not surprising that ACPA is a significant predictor for PBD and their titer is important [7, 13]. We also previously reported that the usefulness of prediction for on-set of RA in undifferentiated polyarthritis [14]. These seem to be reasonable because ACPA is categorized into a biomarker for autoimmunity which is an earlier event in the pathogenesis of RA compared to the other subsequent events such as inflammation and tissue damages (bone/cartilage resorption). Another reason is that ACPA is more specific for autoimmunity in RA than other biomarkers such as RF-IgG (RF-IgM and RF-IgG) [8, 14]. On the other hand, there have been lots of supportive reports which suggest that MMP-3 is useful for predicting PBD in RA [10], they were not significantly associated with Δ SHS in this prospective study. The reason for this discrepancy is not clear. However, this suggests that MMP-3 might reflect on ongoing inflammation and bone/cartilage destruction in the affected joints rather than the trigger [15] because proinflammatory cytokines induce MMPs, including MMP-3 [16].

In this study, we also showed IL-6 was a significant predictor for PBD in early stages of RA. IL-6 is well known as one of the major proinflammatory cytokines in the pathogenesis of RA as well as TNF and IL-1 [17]. In addition, it has been revealed as an important therapeutic target [17]. In fact, recent clinical trials have shown an outstanding efficacy of anti-IL-6 therapy in RA [18]. Besides IL-6, other proinflammatory cytokines, such as TNF α and IL-1, especially TNF α is interesting as a candidate marker as well as IL-6, because it is well known that TNF α is one of the key cytokines in the pathogenesis of RA, and TNF inhibitors are major biologics in the treatment of RA. We tried to measure serum TNF α levels in the RA patients by using a sandwich ELISA kit (QTA00B) for human TNF α (R&D System Inc, Minneapolis, USA), however, all of the serum TNF α levels of the patients were within the range of the healthy volunteers (not detectable ~ 9.03 pg/ml). So, TNF α was not included as a candidate marker in this study.

On the other hand, as mentioned above, anti-cytokine therapies especially using biologics is now well known like a master of treatment of RA. However, they have some critical concerns such as severe adverse effects and high costs. In addition, we are not able to know which cytokine is the most optimal target before the start of the treatment. The finding suggests that the early stages of RA patients showing high serum IL-6 levels might be good candidates for anti-IL6 therapy. Another advantage of IL-6 is that it reflects on the actual degree of inflammation and it can be measured easily and accurately in sera. It also changes more dynamically in sera compared to TNF and IL-1 [19].

In summary, both the baseline ACPA titers and serum IL-6 levels as well as the baseline SHS (TSS*, E*, JSN*) correlated with PBD, suggesting that ACPA and serum IL-6 levels have potentials for predicting PBD in early stages of RA and earlier therapeutic intervention is also important.

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Conflict of interest All authors have no conflicts of interest.

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二光子励起顕微鏡による生きた組織の観察

Visualization of live tissues and organs by using intravital two-photon microscopy

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Key Words: live imaging, two-photon microscopy, immune system, bone homeostasis, chemokine

■ Abstract ■

動物の本質は「動き」にあるが、これは生体のみが備え、死体にはない。近年、蛍光顕微鏡技術の長足の進歩により、生体を「生きたまま」で観察することができるようになった。この技術によって、特にそれまで見えなかった細胞や分子の「動き」を直接見ることができるようになり、生命科学の領域にパラダイムシフトをもたらされつつある。本稿では、二光子励起顕微鏡を用いた「生体イメージング」の方法論と今後の応用について概説する。

■二光子励起顕微鏡の原理と応用

蛍光観察では、注目する細胞や分子などを蛍光分子で標識する。蛍光分子は一般に、エネルギー的に低い状態（基底状態）と高い状態（励起状態）があるが、普段は基底状態にある。このエネルギー差に相当する光（光子）を当てると、蛍光分子はこのエネルギーを吸収して励起状態になるが、自然にまた励起状態へと戻っていく（図上）。この際、そのエネルギーに相当する光を放出し、これを観察しているのが蛍光観察である。

二光子励起とは、蛍光観察の際に、光子1個ではなく、2個の光子を蛍光分子に同時に当てることにより励起させることである。光子1個 対 蛍光分子1個 による1対1反応に比べて、複数の光子による励起（多（二）光子励起）は極めて起こりにくい現象であるが、光子密度を非常に高くすれば非線形的に起こり得る（Göppert-Mayer, 1931）。

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この「二光子励起」の現象は、光子密度が異常に高い場所でのみ起こる。蛍光顕微鏡でいえば、光が一点に凝集される点、すなわち「焦点」のみで起こり得る現象である。これを利用して「焦点のみで励起が起こるような顕微鏡」として「二光子励起顕微鏡」が作られた。

まとめると、この顕微鏡には以下のような長所がある。

① 高い空間（z軸）解像度

焦点平面のみでしか励起が起こらない（その他の平面では励起に必要なエネルギーに満たない光子が当たっているものの励起には至らない）ため、非観察面からの蛍光シグナルがない（非観察面からの蛍光シグナルは「ピンボケ」の原因となる）。

② 高い組織透過性（深部組織の観察に威力を発揮）

2個の光子を同時に当てて蛍光分子を励起するため、当てる光子1個分のエネルギーは小さくて済む（約半分）。エネルギーが半分ということは、光子の波長が2倍になることであり、実際二光子励起で用いるレーザーは近赤外域にある（波長が780~1000 nm）。波長が長い赤外光は、短い可視光や紫外光よりも浸透性が高く、深い組織まで励起・観察することが可能となる。

これらの特長は、「組織・臓器を生かしたままで観察」するために極めて有用である。固定した（もはや生きていない）組織や臓器は、パラフィンなどで包埋して薄切すれば、どんな場所でも観察できるが、生きた組織（特に生きた個体内）では、観察したい場所が対物レンズでアプローチできる場所よりも深いことがある。このような場合、二光子励起顕微鏡が威力を発揮する。

■生きた組織・個体の観察の実際

「生体イメージング」は大きく2つに分類される。「tissue explant imaging」では、実験動物を屠殺して注目する組織・臓器を取り出し、酸素化した培養液中で生かして観察する。一方、「intravital imaging」では、動物を麻酔下で生かしたまま、観察したい組織・臓器を露出して観察する。Intravitalの方法論はより困難であるが多くの利点があり、特に、動物を生かしているので循環血流が保たれる点は非常に大きい。これは、特に免疫・血液系のように、血流を介した細胞の動態が重要なシステムの解析に有用である。

■生きた骨組織・骨髄内の二光子励起観察

筆者はintravital two-photon imagingによる、生きた骨組織・骨髄内の高解像度イメージング法を世界に先駆けて開発した。硬い石灰質に囲まれた骨組織の内部は、生きたままでの観察が特に困難であると考えられていた。骨基質に含まれるリン酸カルシウム結晶は、励起光を容易に散乱させるため、二光子励起に用いる近赤外線レーザーを用いても深部まで到達させることは難しかった。筆者はシステムを改良し、骨基質が比較的薄いマウス頭頂骨を用いて、生きた骨髄内を外部から非侵襲的に高解像度で観察できる実験系を確立した(図下)1)。これを用いて、骨髄内の細胞が血中から遊走したり逆に再還流する様子をリアルタイムで可視化し、この動態を制御するケモカインを同定したり、

骨髄には雑多な血液系・間葉系細胞が、所狭しと詰め込まれている。多様な血液系細胞はそれぞれ決まった場所(ニッチ)に存在し、また互いに複雑な静的・動的ネットワークを形成しているが、これらについては未だ不明な点が数多く残されており、二光子励起顕微鏡による“非破壊検査”を用いた今後の解明が期待される。

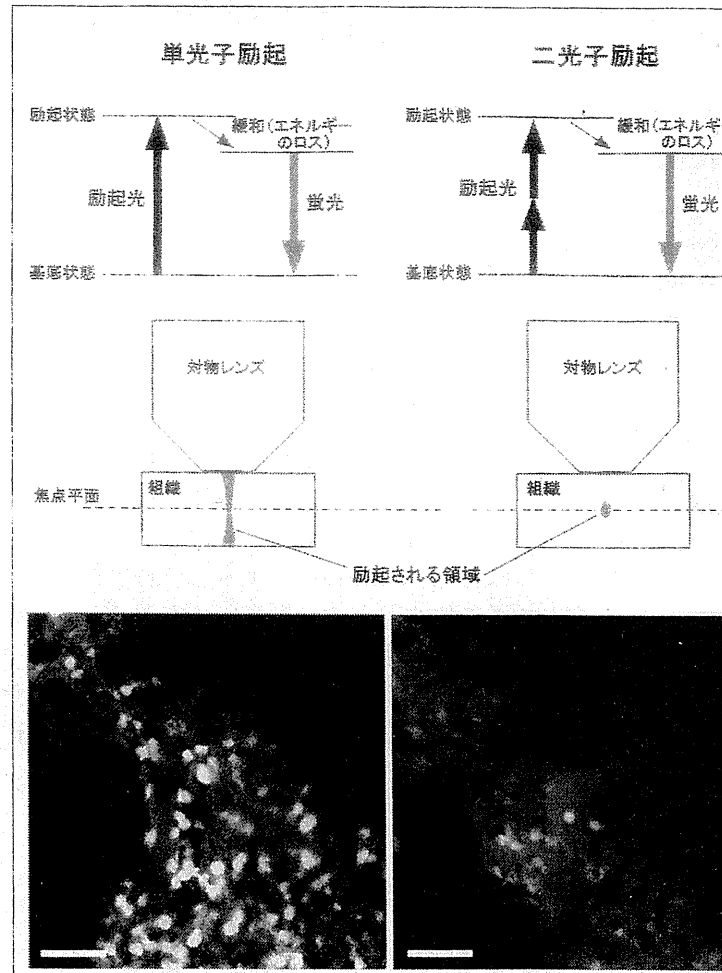


図 二光子励起顕微鏡の原理とその生体観察

上図：通常の蛍光観察では、1個の蛍光分子を1個の光子で励起するが(左図)、多(二)光子励起では複数(2個)の光子で励起する(右図)。このような現象は非常に起こりにくく、光子密度が極大となる焦点平面のみで起こる(下図)。このため、観察したい部位のみ蛍光することになるので高い空間解像度が得られ、非観察部位が励起されないため光毒性が低く退色が少ない。

下図：顆粒球系(LysM⁺;左側)および単球系(CX,CR1⁺;右側)をそれぞれGFP標識したトランスジェニックマウスの骨髄腔の生体二光子励起イメージング。骨髄内の血管構造をTexas Redを結合した高分子デキストランを静脈注射にて可視化している。実際にはこれを一定時間間隔で撮影し、動画を作成する(文献より一部改変;動画については筆者HP<<http://bioimaging.ifrec.osaka-u.ac.jp>>を参照)。スケールバー:30 μm。

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破骨細胞のイメージングと RANKL シグナル

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破骨細胞は、単球系血液細胞から分化・成熟する多核巨細胞であり、「骨吸収」という特殊な機能を持つ唯一の細胞である。しかし、生きた骨組織内で成熟破骨細胞がどのように骨吸収を行っているのか、生体内における成熟破骨細胞の動態についてはこれまで明らかにされてこなかった。我々は最近、二光子励起顕微鏡を駆使することでマウスを生かしたまま、骨表面上での生きた成熟破骨細胞の動態を可視化することに成功し、成熟破骨細胞の骨吸収が RANKL (receptor activator of NF- κ B ligand) によって動的に制御されていることを明らかにした。

本稿ではこの研究成果に加え、我々が開発した骨のライブイメージングの方法論やその応用について概説する。

RANKL signaling and bone diseases.

In vivo imaging of mature osteoclasts and RANKL signaling.

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Osteoclasts are 'bone-resorbing' giant polykaryons that differentiate from mononuclear macrophage/monocyte-lineage hematopoietic precursors. However how the activity of mature osteoclasts is regulated *in vivo* remains unclear. To answer the question, we utilized an advanced imaging system for visualizing live bone tissues with intravital multiphoton microscopy that we have recently established. By means of the system we have recently succeeded in visualization of mature osteoclasts in live bones and revealed that RANKL regulates bone-resorptive functions of mature osteoclasts *in vivo*. Here we show the latest data and the detailed methodology of intravital imaging of bone tissues, and also discuss its further application.

はじめに

破骨細胞は単球系血液細胞から分化・成熟する多核巨細胞であり、「骨吸収」という特殊な機能を

持つ唯一の細胞である。骨髄から採取した単球系細胞を M-CSF (macrophage colony-stimulating factor) と RANKL (receptor activator of

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NF- κ B ligand)の刺激下で *in vitro* で培養すると、破骨細胞様の多核巨細胞が形成され、中には100核以上の巨細胞も観察される。しかし「このような多核巨細胞が *in vivo* でも本当に形成されるのか」、「生きた骨組織内で成熟破骨細胞はどのように骨吸収を行うのか」など、生体内での成熟破骨細胞の動態についてはこれまで明らかにされてこなかった。

硬い石灰質に囲まれた骨組織の内部は従来生きたままでの観察が困難であったが、我々は、組織深部の観察が可能な「二光子励起顕微鏡」を駆使して、マウスを生かしたままで骨組織内の細胞動態を観察するイメージング方法を確立した¹⁾。この方法を用いて、我々は最近、骨表面上での生きた成熟破骨細胞の動態を可視化することに成功し、成熟破骨細胞の骨吸収がRANKLによって動的に制御されていることを明らかにした。

本稿では、これらの研究成果の解説に加えて骨組織内の生体二光子励起イメージングの方法論や、その今後の応用と将来性について、実際の画像を紹介しながら概説する。

骨組織内の生体二光子励起イメージング

1. 二光子励起顕微鏡のメリット

二光子励起顕微鏡では、通常の蛍光顕微鏡観察(共焦点レーザー顕微鏡も含む)で用いる励起光の半分のエネルギー(2倍の波長)をもったレーザー光を、細かいパルス状に放出したものを励起光源に用いる。二光子励起は、レンズで集約された光子が集まる1点の焦点面にしか起こらないため、非常にクリアな画像が得られ(高い空間解像度)、観察対象となる組織・臓器への光毒性や蛍光の退色もきわめて小さく抑えることができる(低い組織侵襲性)。また、励起光として通常の半分のエネ

ルギー(2倍の波長)の近赤外光(波長が780~1,000 nm)を用いるため、組織の深部まで励起光を到達させることができる(高い組織透過性)。以上の特長から、二光子励起顕微鏡は、「組織・臓器を生かしたままで観察」するためにきわめて有用である^{2,3)}。

2. 骨組織内の“生体(= intravital)”イメージングの実際

骨基質に含まれるリン酸カルシウム結晶は、励起光を容易に散乱させるため、二光子励起に用いる近赤外線レーザーを用いても深部まで到達させることは難しい。現在の近赤外線レーザーでは、軟部組織であれば表面から800~1,000 μ mまで到達が可能であるが、骨組織の場合は、150~200 μ mが限界である。このため、我々は、骨基質が薄くて骨表面から骨髓腔まで80~120 μ mで到達できるマウスの頭頂骨を用いて、骨組織内の生体イメージングに取り組んでいる^{1,4,5)}(図1)。実際には、麻酔したマウスの頭頂骨の皮膚を切開し、露出させた後、顕微鏡用のステージにマウスを固定して観察する。この方法では、骨髓腔内を流れる豊富な血流が得られているため、骨組織に定着している細胞の動きのみならず、血管から骨髓内へ細胞が流入したり、逆に血中へ還流していく様子を観察することができる。さらには、薬剤を尾静脈などから全身投与すると、血流を通して速やかに観察部位に到達させることができる。

この「生体二光子励起イメージング」を用いて、我々はこれまで、破骨前駆細胞の骨吸収面への遊走・接着が、脂質メディエーターの一種であるスフィンゴシン1リン酸(S1P)によって動的に制御されていることを解明した⁶⁾。しかしながら、

RANKL: receptor activator of nuclear factor- κ B ligand (破骨細胞分化因子)

M-CSF: macrophage colony-stimulating factor (マクロファージコロニー刺激因子)

S1P: sphingosine-1-phosphate (スフィンゴシン1リン酸)

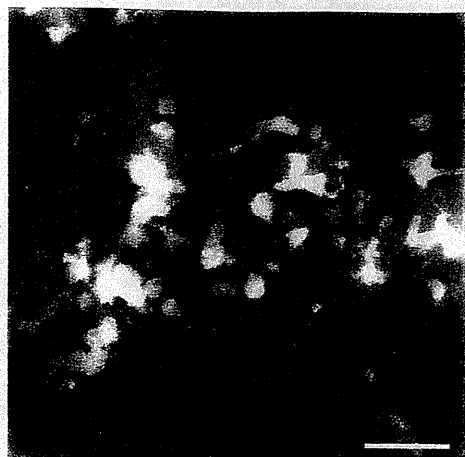


図1 骨組織内での破骨前駆細胞の生体二光子励起イメージング

破骨前駆細胞を含む単球系細胞を緑色にラベルしたマウスの骨髄腔の生体二光子励起イメージング。骨髄腔内の血管構造は、赤色蛍光 (Texas Red) を結合させた高分子テクニストランを静脈注射して可視化している。青色は骨組織を示す。実際の実験ではこれを一定時間間隔で撮影し、動画を作成する。

(スケールバー: 30 μm)

(カラーグラフィック5頁参照) (筆者作成)

「破骨前駆細胞が骨表面に到達して成熟破骨細胞に分化した後、骨吸収が生体内でどのように調節されているのか」という点については、これまで明らかにされてない。そこで、我々は次に「生体二光子励起イメージング」技術を駆使して、生体骨組織内での成熟破骨細胞の動態を可視化することに取り組んだ。

生体イメージングによる成熟破骨細胞の動態解明

1. 生きた成熟破骨細胞の動態の可視化

我々は、成熟破骨細胞に特異的に緑色蛍光蛋白質 (GFP) を発現させたマウスの骨組織内を二光子励起顕微鏡で観察することにより、生きた成熟破骨細胞の動態の可視化に成功した (図 2a)。その結果、骨表面で骨吸収を行っている成熟破骨細胞

には、「①動きがなく、今まさに骨吸収をしている状態の細胞 (図 2b)」と、「②動いていて、骨吸収をしていない状態の細胞 (図 2c)」の少なくとも2種類が存在することが明らかとなった (論文投稿中)。

次に我々は、これらのマウスに各種薬剤を投与して、成熟破骨細胞の動態の変化を検討した。まず、成熟破骨細胞を蛍光標識したマウスに RANKL を腹腔内投与して破骨細胞の機能を亢進させ、投与2日後にマウスの骨組織内を生体二光子励起顕微鏡で観察した。その結果、成熟破骨細胞数が増加し、そのほとんどが「①動きがなく、今まさに骨吸収をしている状態の細胞」であった。一方、骨粗鬆症の治療薬であるビスホスホネートを投与して破骨細胞の機能を抑制させると、骨表面での成熟破骨細胞数が減少し、また残った細胞のほとんどが、「②動いていて、骨吸収をしていない状態の細胞」であった。このように、二光子励起イメージングを用いることにより、成熟破骨細胞の絶対数のみならず、成熟破骨細胞の機能 (骨吸収をしているのか、あるいは骨吸収をしていないのか) も同時に評価することができるようになった。

2. RANKL による成熟破骨細胞の骨吸収制御機構の解明

次に我々は、上述の成熟破骨細胞を蛍光標識したマウスに RANKL を急速に静脈内投与した。その結果、投与後 30 分という比較的短い時間に、成熟破骨細胞が「②骨吸収をしていない状態」から「①今まさに骨吸収をしている状態」へと変わっていく様子が観察された。従来、RANKL は破骨前駆細胞が成熟破骨細胞へ分化するための必須の因子であることが知られていた。今回我々が行ったイメージングの結果により、RANKL は破

GFP: green fluorescence protein, green fluorescent protein ともいう。(緑色蛍光タンパク質)

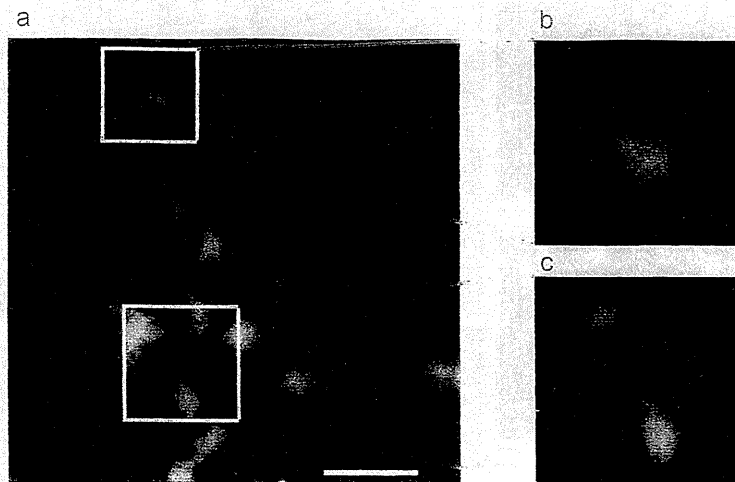


図2 骨組織内での成熟破骨細胞の生体二光子励起イメージング

成熟破骨細胞を緑色にラベルしたマウスの骨髄腔の生体二光子励起イメージング (a)。骨髄腔内の血管構造は、赤色蛍光 (Texas Red) を結合させた高分子デキストランを静脈注射して可視化している。青色は骨組織を示す。骨表面で骨吸収を行っている成熟破骨細胞には、「動きがなく、今まさに骨吸収をしている状態の細胞 (b)」と、「動いていて、骨吸収をしていない状態の細胞 (c)」の少なくとも2種類が存在する。(スケールバー: 40 μm) (カラーグラフィック5頁参照) (筆者作成)

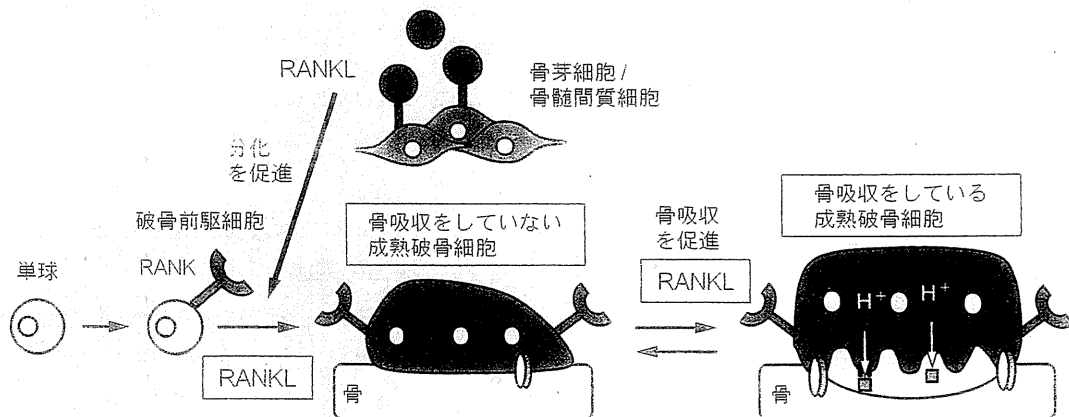


図3 生体内における成熟破骨細胞の骨吸収メカニズム

成熟破骨細胞には、「骨吸収をしていない状態の細胞」と、「今まさに骨吸収している状態の細胞」との2種類が存在する。また、RANKL は破骨細胞分化を促進するだけでなく、成熟した破骨細胞にも作用し、骨吸収を促進する役割も担っている。(筆者作成)

骨細胞の分化を促進するだけでなく、成熟した破骨細胞にも作用し、骨吸収を促進する役割も担っていることが明らかとなった (図3)。

これまでの成熟破骨細胞の骨吸収に関する研究のほとんどは、固定した骨組織を切り出して行われたものであるため、細胞の動的な情報を得るこ

とが困難であった。しかし、今回我々が行ったイメージング系は、マウスを生かしたまま、骨表面での成熟破骨細胞の骨吸収の動態をリアルタイムで観察することができるため、今後、骨吸収性疾患の病態解明や新規薬剤の開発においても強力な手段となり得ると考えられる。

今後の展開

1. 頭頂骨以外の骨組織のイメージング系の開発

現時点では、十分な解像度で可視化できる骨組織は、骨梁が薄くレーザー光を透過させやすい頭頂骨に限られている。基本的には、どこの部分の骨であっても、骨代謝や骨髄細胞の動態などには変化がないと考えられるが、それらを実証するためには、やはり長管骨など一般に広く研究に用いられている骨組織をライブイメージングにより解析する必要がある。現在我々は、長管骨のライブイメージング系の確立に取り組んでいる。

2. 長時間のライブイメージング系の開発

ガス麻酔下でマウスを生かしたまま、骨組織を手術的に露出してイメージングに当たっている現在の方法では、連続した撮影時間は4~5時間程度が限界である。細胞の動きや細胞間の接触時間などをイメージングするのであれば、この観察時間で十分であるが、それより長い時間のかかる現象(例えば、破骨細胞の分化など)をイメージングするためには、別の測定系を構築する必要がある。(マウスを長期間にわたり麻酔管理するか、手術野を閉じて経日的観察を可能にするなど)。現在我々は、長時間のライブイメージング系の確立にも取り組んでいる。

おわりに

近年の蛍光イメージング技術の急速な進歩のおかげで、さまざまな臓器・組織での生きた細胞の

観察が活発に行われている。しかしながら、これまでの生体イメージング研究の多くは、生体内での細胞の動き(動く速さ)や細胞間相互作用(細胞同士の接触時間)を観察するにすぎなかった。これからの生体イメージング研究では、「細胞が動いて接触した後に何が起こるのか」、すなわち生体内での細胞の「機能」や「分化」をイメージングすることが重要になってくると思われる。特に骨組織は、破骨細胞や骨芽細胞による骨代謝制御の場であるばかりでなく、リンパ球を始めとして顆粒球・単球など多種多様な血液細胞の発生・機能分化にとって重要な部位である。我々が開発した骨組織の生体二光子励起イメージングの方法論は、骨髓腔内での各細胞の挙動・位置決めとその分化制御や、血液系幹細胞が多能性を維持する特殊な環境(ニッチ)の解明において強力な手段となることが強く期待される。

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