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Nucleotide sequences: Sequences of nucleotides or amino acids longer than 50 bases/residues should not be presented in the text or in table form, but rather should be submitted as a publication quality figure. Original nucleotide sequences and determined nucleotide sequences encoding reported amino acid sequences described in the manuscript must be submitted to GenBank (http://www.ncbi.nlm.nih.gov/Genbank/) or European Nucleotide Archive (http://www.ebi.ac.uk/ena/) at the time of manuscript submission. An accession number and sequence availability are required at the time of publication. The accession number should be accompanied by the Web site address of the databank.

Microarray data: The JI will not publish descriptive manuscripts that report microarray data, unless such information can be considered of unusual immunological significance and/or include functional experiments that provide novel insight into mechanism. As with other scientific approaches, current experimental, quantitation,

verification, and statistical analyses are expected. Microarray experiments should be Minimum Information About a Microarray Experiment (MIAME) compliant (for guidelines, see http://www.mged.org/Workgroups/MIAME/miame.html). Whereas limited online space may be available for supplemental tables associated with the manuscript, complete microarray data must be deposited in the appropriate public database (e.g., GEO [http://www.ncbi.nlm.nih.gov/geo/], ArrayExpress [http://www.ebi.ac.uk/microarray-as/ae/], or CIBEX [http://cibex.nig.ac.jp/index.jsp]), and must be accessible without restriction from the date of publication. An entry name or accession number must be included in the paper before publication. The accession number should be accompanied by the Web site address of the databank.

STYLE GUIDE

General style conventions: In general, *The JI* follows *Scientific Style and Format: The CSE Manual for Authors, Editors, and Publishers, 7th Edition*, published by the Council of Science Editors, Inc., in instances where style issues are not directly addressed.

Abbreviations for references: PubMed (http://www.ncbi.nlm.nih. gov/journals) is the primary source for journal name abbreviations.

Nomenclature: The most current links for nomenclature guidelines are posted online.

Allergen nomenclature: Nomenclature for allergens should be assigned in cooperation with the International Union of Immunological Societies (IUIS) Allergen Sub-Committee. Authors of accepted manuscripts that describe novel allergens will be requested to complete a brief standard form available at IUIS Allergen Nomenclature (http://www.allergen.org/Allergen.aspx).

CD nomenclature: For the purpose of consistency, *The JI* will follow CD nomenclature. For murine molecules, *The JI* will follow the nomenclature previously published (*J. Immunol.* 160: 3861–3868, 1998). For human molecules, standard CD nomenclature will be followed as updated (*J. Immunol.* 168: 2083–2086, 2002). See also http://www.HCDM.org.

Chemical names: The JI uses *The Merck Index* (http://library.dialog.com/bluesheets/html/bl0304.html) and the *IUPAC-IUB Commission on Biochemical Nomenclature-Chemical Abstracts* (http://www.chem.qmul.ac.uk/iupac/bibliog/white.html) as the primary references for proper spelling and style of chemical names.

Chemokine/chemokine receptor nomenclature: The systematic name for chemokines and chemokine receptors should be used. The original name may be given in parentheses if desired. See *Cytokine 21:48–9, 2003*.

Enzyme nomenclature: Enzyme Nomenclature (http://www.chem.qmul.ac.uk/iubmb/enzyme/) is *The JI* source for style and spelling of enzyme names.

Gene nomenclature: The HUGO guidelines for gene nomenclature (http://www.genenames.org/) may be used for naming human genes. Mouse Genome Informatics (http://www.informatics.jax.org/) is a reference source for naming mouse genes.

Genetic nomenclature for mice: The JI uses the revisions for standardized genetic nomenclature for mice published periodically in Mouse Genome. A current listing of inbred strains of mice and rats is available at Mouse Genome Informatics. Authors are encouraged to deposit their mapping data with the Mouse Genome Database (MGD) (http://www.informatics.jax.org/) before publication and to include the assigned MGD accession numbers in their manuscripts. Data may be submitted electronically by e-mail. Information about electronic submission of datasets can be obtained at the Data and Nomenclature Submissions page. Gene symbols should be reserved with MGD in advance of publication. An electronic nomenclature submission form is available from the MGD Web site.

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HLA nomenclature: HLA nomenclature is updated periodically by the World Health Organization Nomenclature Committee for Factors of the HLA System. A recent reference is *Hum. Immunol.* 64: 919–20, 2003. Annual comprehensive revisions are published in *Human Immunology*, usually in the spring. See also: http://www.ebi.ac.uk/imgt/hla/.

SUPPLEMENTAL MATERIALS

Supplemental Data:

- Supporting data that are not essential to understanding the material presented in the manuscript may be submitted with the original paper for peer review; however, the print version of the paper must stand on its own without the supplemental material.
- Supplemental material is primarily intended for short videos, large tables, large sequence alignments, or large data sets.
 Additional supplemental figures and tables that support the interpretation and conclusions drawn in the manuscript may, however, also be submitted for review with the manuscript.
- Supplemental material must be submitted as separate files from the rest of the manuscript during the online submission; select "Supplemental Data" as the "File Type" when uploading the files.
- Apart from videos, all files must be either PDF or Excel file format; multiple PDF files should be combined into a single PDF file.
- For Cutting Edge manuscripts, no more than two supplemental figures and/or tables may be submitted. One of the two allowable Cutting Edge supplemental items can be a description, no more than one page in length, of computational or bioinformatics methods. Methods that are normally found in the body of the article may not be included in this Cutting Edge supplemental item.
- For Full-Length manuscripts, no more than four supplemental figures and/or tables may be submitted.
- Each supplemental figure should comprise no more than a single 8.5" × 11" PDF page, and be large enough to be legible when that page is opened.
- Legends or short explanations must accompany all supplemental figures and videos; no other supplementary text is permitted in full-length manuscripts. Legends should be placed below the corresponding figure in the PDF. Legends for videos should be submitted as a single PDF. Table legends must be prepared as footnotes to the table; all tables must have a title.

*Note: Excel files will be converted to PDFs for the review process only. At publication, the file(s) will be uploaded in the original Excel format.

Videos must be 320×480 pixels or smaller for best viewing within a browser. Videos must be no longer than 30 seconds and under 10MB, with no sound or voice-over. Submit videos in MPG or QuickTime format. Change QuickTime file extensions to ".mov" so that Web browsers will recognize the file type and play the movie. Compress videos as much as possible to help control file size. Name videos by order of citation appearance (e.g., video1.mov). Select "Video" as the "File Type" when uploading the files during online submission. Authors will be notified if problems exist with videos as submitted and will be asked to take responsibility for modifications. No editing will be done to videos at the Editorial Office. A legend or short explanation must accompany the video.

Links to the supplemental material will appear in two places in the online journal: in the Table of Contents and in the information box associated with the first page of the full-text article. There will not be any links in the body of the article. In the printed paper, supplemental material will be footnoted the first time mentioned: "The online version of this article contains supplemental material." Supplemental Materials are posted online as provided by the author. There is a publication charge of \$75 per supplemental figure or table.

CUTTING EDGE MANUSCRIPT PREPARATION

Manuscripts submitted to the *Cutting Edge* section should conform to the *General Guidelines* for full-length manuscripts as well as the additional guidelines below:

- 1. Cutting Edge articles, including figures and references, must fit within four journal pages. Authors should estimate the size of figures and tables and limit the text accordingly. One printed page in The JI contains approximately 8,000 characters, including spaces. Thus, a four-page Cutting Edge article would contain approximately 32,000 characters. The formula for estimating the number of pages is provided in Manuscript Preparation.
- 2. The Abstract is limited to 150 words.
- The Materials and Methods section may be sharply limited, but should be sufficient to allow the evaluation of results and conclusions.
- 4. Authors may combine the *Results* and *Discussion* sections.

PREPARATION OF THE REVISED MANUSCRIPT

Follow *The JI* Editorial Office instructions contained in the previous decision letter carefully and thoroughly. A revised manuscript not returned within 9 months of the date of the decision letter will be considered a new manuscript and subject to a new, complete review.

Individual manuscript files, files for each figure and table (even if they are unchanged from the previous submission), and a point-by-point reply to all referee comments must be uploaded to the system. The revised manuscript text must be marked to show changes using yellow highlighting (Microsoft Word files preferred). Do not show deletions, because if the manuscript is accepted, this version will be immediately sent for publication. High-resolution figure files must be submitted. Figures must be in **TIFF**, **EPS**, or **PDF** format and prepared as described under *Figures*.

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Submit online at ji.msubmit.net.

- 1. For the initial submission of Full-Length manuscripts:
 - Either a PDF of the entire manuscript (text, figures, and tables), or individual manuscript, figure, and table files may be uploaded to the system. If individual files are uploaded, the system then creates a single PDF for review purposes.
- 2. For Cutting Edge submissions and all revised manuscripts:
 - Individual manuscript, high resolution figure, and table files must be uploaded to the system. Authors should save copies for themselves of all the files in their original formats.
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- 1. Ensuring the scientific integrity of the submission;
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- 5. Making unique materials available to qualified investigators;
- Complying with requirements to deposit microarray and other similar datasets in public venues;
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- 8. Ensuring that all conflicts of interest or financial interests are listed;
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All potential reviewers are contacted individually to determine availability. Manuscript files are sent to at least two expert reviewers. Reviewers are asked to complete the review of the manuscript within 2 weeks and to return a short review form. Based on the reviewers' comments, the Section Editor recommends a course of action and communicates the reviews and recommendations to the Deputy Editor for a final decision.

The Deputy Editor considers the comments made by the reviewers and the recommendation of the Section Editor, selects those comments to be shared with the authors, makes a final decision concerning the manuscript, and prepares the decision letter for signature by the Editor-in-Chief. If revisions of the manuscript are suggested, the Deputy Editor also recommends who should review the revised paper when resubmitted. Authors are informed of the decision by e-mail; appropriate comments from reviewers and editors are appended.

Decisions: There are four categories for initial decisions: accept, accept with minor revision, return for revision, and reject. Some manuscripts are accepted provisionally, pending relatively minor revisions. In this case, the Deputy Editor may conduct the rereview. For many manuscripts, authors are invited to resubmit if revision or additional experimentation can address major cri-

ticisms. Typically, one or more reviewers will then be asked to consider the adequacy of the revisions. *Cutting Edge* papers are allowed only minor revisions because of time constraints. All revised manuscripts are carefully reexamined, and ultimate acceptability is not guaranteed. *The JI* does not provide for an advance determination of the acceptability of a particular manuscript for publication, nor does it promise expedited review of selected manuscripts.

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STANDARD ABBREVIATIONS

The abbreviations listed here are used without definition in articles published in The JI. The form may be used for both singular and plural, or made plural with "s" at the author's option.

Å, angstrom

aa, amino acid (only with numbers)

Ab, antibody

ABTS, 2,2'-azinobis(3-ethylbenzthiazoline-6-sulfonic acid)

ADP, adenosine 5'-diphosphate

Ag, antigen

AIDS, acquired immunodeficiency syndrome

AMP, adenosine 5'-monophosphate

ANOVA, analysis of variance

AP-1, activator protein 1

APC, Ag-presenting cell

ATP, adenosine triphosphate

BALB/c, a mouse strain

BALT, bronchus-associated lymphoid tissue

BAPTA-AM, 1,2-bis(2-aminophenoxy)ethane-N,N,N',N'-

tetraacetic acid acetoxymethyl ester

BCR, B cell receptor

bp, base pair (only with numbers)

BrdU, 5-bromo-2'-deoxyuridine

BSA, bovine serum albumin

C, complement

C region, constant region of Ig

cAMP, cyclic AMP

C terminus, carboxyl terminus

C-terminal, carboxyl-terminal

CCL, CC chemokine ligand

CCR, CC chemokine receptor

CD40L, CD40 ligand

cDNA, complementary DNA

CDP, cytidine 5'-diphosphate

CDR, complementarity determining region

C/EBP, CCAAT/enhancer-binding protein

CFA, complete Freund's adjuvant

CFSE, 5-(and 6-)-carboxyfluorescein diacetate succinimidyl ester

CFU, colony-forming unit

cGMP, guanosine 3',5'-cyclic monophosphate

CHAPS, 3-[(3-cholamidopropyl)dimethylammonio]-1-

propanesulfonate

Ci. curie

CIITA, class II transactivator

CLIP, class II-associated invariant-chain peptide

cM, centiMorgan(s)

CMP, cytidine 5'-monophosphate

CMV, cytomegalovirus

CNS, central nervous system

CoA, coenzyme A

Con A, concanavalin A

CpG, cytosine guanine dinucleotide

cpm, counts per minute

CREB, cAMP response binding protein

cRNA, complementary RNA

CSF, colony-stimulating factor

CTL, cytotoxic T lymphocyte

CTLA, cytolytic T lymphocyte-associated Ag

CTP, cytidine 5'-triphosphate

CXCL, CXC chemokine ligand

CXCR, CXC chemokine receptor

d, deoxy; distilled (as in dH2O)

D region, diversity region of Ig or T cell receptor for Ag

Da, dalton (only with numbers)

DAPI, 4',6'-diamidino-2-phenylindole

DEAE, diethylaminoethyl

df, degrees of freedom

DMEM, Dulbecco's modified Eagle's medium

DMSO, dimethylsulfoxide

DNA, deoxyribonucleic acid

DNase, deoxyribonuclease

DNP, dinitrophenyl

dpm, disintegrations per minute

ds, double-stranded (as dsDNA)

DTT, dithiothreitol

E, erythrocyte

EBV, Epstein-Barr virus

EC₅₀, 50% effective concentration

ECL, enhanced chemiluminescence

ED₅₀, 50% effective dose

EDTA, ethylenediaminetetraacetic acid

EGTA, ethylene glycol-bis(β-aminoethyl ester)-N,N,N',N'tetraacetic acid

ELISA, enzyme-linked immunosorbent assay

ELISPOT, enzyme-linked immunospot

EMSA, electrophoretic mobility shift assay

ERK, extracellular signal-regulated kinase

E:T ratio, effector to target ratio

Fab, Ag-binding fragment

F(ab')2

F-actin, filamentous actin

FACS, fluorescence-activated cell sorting

FAM, 6-carboxyfluorescein

FBS, fetal bovine serum

FcR, Fc receptors (e.g., FcγRI)

FCS, fetal calf serum

FITC, fluorescein isothiocyanate

FLICE, Fas-associated death domain-like IL-1β-converting

enzyme

FLIP, FLICE inhibitory protein FLT3, *fms*-related tyrosine kinase 3

fMLF, formyl-methionyl-leucyl-phenylalanine

fura 2-AM, fura 2-acetoxymethyl ester

g, gram (only with numbers)

GALT, gut-associated lymphoid tissue

GAPDH or G3PDH, glyceraldehyde-3-phosphate

dehydrogenase

G-CSF, granulocyte CSF GDP, guanosine 5'-diphosphate GFP, green fluorescent protein

GM-CSF, granulocyte-macrophage CSF GMP, guanosine 5'-monophosphate gp, glycoprotein (e.g., gp100) GPI, glycosylphosphatidylinositol GST, glutathione *S*-transferase GTP, guanosine 5'-triphosphate h, hour (only with numbers)

H chain, heavy chain

H&E, hematoxylin and eosin

HBSS, Hanks' balanced salt solution

HEPES, N-2-hydroxyethylpiperazine-N'-2-ethanesulfonic acid

HIV, human immunodeficiency virus

HLA, human histocompatibility leukocyte Ag HPLC, high performance liquid chromatography

HRP, horseradish peroxidase HSV, herpes simplex virus

HUVEC, human umbilical vein endothelial cell IC₅₀, 50% inhibition/inhibitory concentration ICAM, intercellular adhesion molecule

ICOS, inducible costimulator Id, idiotype; idiotypic determinant

ID₅₀, 50% infective dose or 50% inhibiting dose

IDO, indoleamine 2,3-dioxygenase IFA, incomplete Freund's adjuvant IFN, interferon (e.g., IFN-γ) Ig, immunoglobulin IgH, Ig heavy chain

IκB or I-κB, inhibitory NF-κB IL, interleukin (e.g., IL-2) i.m., intramuscular

IMDM, Iscove's modified Dulbecco's medium IMEM, Iscove's minimal essential medium

i.p., intraperitoneal

ITAM, immunoreceptor tyrosine-based activation motif ITIM, immunoreceptor tyrosine-based inhibitory motif

IU, international unit i.v., intravenous

J region, joining region of Ig or T cell receptor for Ag

JAK or Jak, Janus kinase JNK, c-Jun N-terminal kinase kb, kilobase (only with numbers) kbp, kilobase pair (only with numbers)

 $K_{\rm a}$, association constant

 $K_{\rm d}$, distribution coefficient; dissociation constant

 $K_{\rm D}$, affinity constant

kDa, kilodalton (only with numbers)

L chain, light chain LD₅₀, 50% lethal dose

LFA, leukocyte (lymphocyte) function-associated Ag

LIF, leukemia inhibitory factor LPS, lipopolysaccharide

LU, lytic unit

2-ME, 2-mercaptoethanol mAb, monoclonal Ab 2-ME, 2-mercaptoethanol

MACS, magnetic-activated cell sorting

MALDI, matrix-assisted laser desorption ionization

MALDI-TOF, matrix-assisted laser desorption ionization-time

of flight

MALT, mucosa-associated lymphoid tissue MAPK, mitogen-activated protein kinase MCP, monocyte chemoattractant protein

M-CSF, macrophage CSF 2-ME, 2-mercaptoethanol

MEK, mitogen-activated protein kinase kinase

MEM, minimum essential medium
MES, 2-(*N*-morpholino)ethanesulfonic acid
mg, milligram (only with numbers)
MHC, major histocompatibility complex
min, minute (only with numbers)
MIP, macrophage-inflammatory protein
ml, milliliter (only with numbers)
MLC, mixed lymphocyte culture
MLR, mixed leukocyte reaction

mo, month(s) (only with numbers)
MOPS, 4-morpholinepropanesulfonic acid

*M*_r, relative molecular mass mRNA, messenger RNA

MTT, 3-(4,5-dimethylthiazol-2-yl)-2,5-dimethyltetrazolium

bromide

 μg , microgram (only with numbers) μl , microliter (only with numbers)

m.w., molecular weight

MyD88, myeloid differentiating factor 88

n, number in study or group

NAD, nicotinamide adenine dinucleotide

NADH, reduced NAD

NaDodSO₄, sodium dodecyl sulfate

NADP, NAD phosphate

NADPH, reduced NAD phosphate NBT, nitroblue tetrazolium ND, not determined

NDP, nucleoside 5'-diphosphate

NF, nuclear factor

NFAT or NF-AT, nuclear factor of activated T cells

NF-κB, nuclear factor κB

Ni-NTA, nickel-nitrilotriacetic acid

NK cell, natural killer cell

NMP, nucleoside 5'-monophosphate

NO, nitric oxide NOD, nonobese diabetic NS, not significant

nt, nucleotide (only with numbers)

N-terminal, NH2-terminal or amino-terminal N terminus, NH2 terminus or amino terminus

NTP, nucleoside 5'-triphosphate OCT, octamer-binding factor

OD, optical density OVA, ovalbumin *p*, probability

PAGE, polyacrylamide gel electrophoresis PBL, peripheral blood lymphocyte

PBMC, peripheral blood mononuclear cell

PBS, phosphate-buffered saline PCR, polymerase chain reaction

PE, phycoerythrin

PECAM-1, platelet endothelial cell adhesion molecule-1

PerCP, peridinin chlorophyll protein

PFU, plaque-forming unit PG, prostaglandin PHA, phytohemagglutinin

PI3K, phosphatidylinositol 3-kinase

PIPES, piperazine-N,N'-bis(2-ethane sulfonic acid)

PMA, phorbol myristate acetate PMSF, phenylmethylsulfonyl fluoride

PWM, pokeweed mitogen r, recombinant (e.g., rIFN-γ) R, receptor (e.g., IL-2R)

RACE, rapid amplification of cDNA end RAG, recombination-activating gene

RANTES, regulated upon activation, normal T cell expressed and

secreted

RBC, red blood cell

RFLP, restriction fragment length polymorphism

RIA, radioimmunoassay RNA, ribonucleic acid RNase, ribonuclease rpm, revolutions per minute RPMI, (usually RPMI 1640) rRNA, ribosomal RNA

RT-PCR, reverse transcriptase polymerase chain reaction

s, second (use only with numbers)

s.c., subcutaneous

SCID, severe combined immunodeficiency

SD, standard deviation SDS, sodium dodecyl sulfate

SDS-PAGE, sodium dodecyl sulfate-polyacrylamide gel

electrophoresis SE, standard error

SEM, standard error of the mean

SHIP, src homology 2-containing inositol 5'-phosphatase

SIV, simian immunodeficiency virus

sp. act., specific activity SRBC, sheep red blood cell ss, single-stranded (e.g., ssDNA) SSC, standard saline citrate

STAT, signal transducer and activator of transcription

SV40, simian virus 40 $t_{1/2}$, half-life, half-time

TAMRA, 5-(and 6)-carboxytetramethylrhodamine TAP, transporter associated with Ag processing Tat, terminal deoxynucleotidyltransferase

TBS, Tris-buffered saline TBST, TBS with Tween 20 TCA, trichloroacetic acid TCR, T cell receptor for Ag TDP, thymidine 5'-diphosphate

TdT, terminal deoxynucleotidyltransferase

TGF, transforming growth factor

Th cell, T helper cell

TLC, thin layer chromatography

TLR, Toll-like receptor

TMP, thymidine 5'-monophosphate

TNF, tumor necrosis factor

TNP, trinitrophenyl

TRAIL, TNF-related apoptosis-inducing ligand Tris, tris(hydroxymethyl)aminomethane

tRNA, transfer RNA

TTP, thymidine 5'-triphosphate

TUNEL, Tdt-mediated dUTP nick end labeling

U, unit (only with numbers) UDP, uridine 5'-diphosphate UMP, uridine 5'-monophosphate UTP, uridine 5'-triphosphate

UV, ultraviolet

v/v, volume to volume ratio (%) v/w, volume to weight ratio (%) V region, variable region of Ig

VCAM, vascular cell adhesion molecule V(D)J or VDJ, variable diversity joining

VLA, very late activation Ag W, watt (only with numbers) WBC, white blood cell WEHI medium

wk, week (only with numbers) xid, X-linked immunodeficiency

Zap70, ζ -associated protein 70 (or ζ -chain-associated protein

Keywords

Animals Human Other Animals

Cells B Cells Dendritic Cells Endothelial Cells Eosinophils Mast Cells/Basophils Monocytes/Macrophages Natural Killer Cells Neutrophils Stem Cells Stromal Cells T Cells T Cells, Cytotoxic

Diseases Autoimmunity Diabetes EAE/MS

Th1/Th2 Cells

Endotoxin Shock Graft Versus Host Disease Immunodeficiency Diseases Rheumatoid Arthritis Systemic Lupus Erythematosus

Infections AIDS Bacterial Fungal Parasitic-Helminth Parasitic-Protozoan

Molecules

Cytokines

AcutePhase Reactants Adhesion Molecules Antibodies Antigens/Peptides/Epitopes Autoantibodies Cell Surface Molecules Chemokines Complement Cytokine Receptors

Fc Receptors Lipid Mediators Lipopolysaccharide MHC Nitric Oxide

Protein Kinases/Phosphatases Superantigens T Cell Receptors Transcription Factors

Processes

Gene Regulation

Allergy Antigen Presentation/Processing Apoptosis Cell Activation
Cell Differentiation Cell Proliferation Cell Trafficking

Chemotaxis Comparative Immunology/Evolution Costimulation Cytotoxicity Gene Rearrangement

Tissues Lung Mucosa Skin

Techniques Gene Therapy Molecular Biology Transgenic/Knockout Mice

Spleen & Lymph Nodes

Hematopoiesis

Neuroimmunology

Signal Transduction

Repertoire Development

Reproductive Immunology

Tolerance/Suppression/Anergy

Inflammation

Phagocytosis

Transplantation

Vaccination

Tumor Immunity

Memory



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