

INTERLEUKIN 1:

A MOLECULAR BASIS FOR SICK PEOPLE FEELING BAD

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INTRODUCTION

The acute phase response refers to the characteristic constellation of events induced by infection, inflammation, immunologic reactions, injury and neoplasia. Some of the features of this response include fever, somnolence, anorexia, lassitude, myalgias, leukocytosis, and an elevated erythrocyte sedimentation rate. For many years it was suspected that the acute phase response might be caused by products of tissue damage. For example, Homburger showed in 1945 that pus from sterile chemically-induced abscesses could increase the level of plasma fibrinogen in experimental animals (1). Subsequently, three separate lines of investigation coalesced to provide insight into the nature of the mediator causing the acute phase response. Paul Beeson found that a partially purified extract obtained from rabbit leukocytes (Leukocytic Pyrogen) could induce fever (2). Later experiments described material found in the plasma of animals during experimental fever (Endogenous Pyrogen) that had all the properties of Leukocytic Pyrogen (3,4). Kampschmidt and colleagues studied the effect of extracts of leukocytes on serum iron and zinc levels and published a series of experimental results indicating that changes in acute phase reactants could be accounted for by the action of a Leukocytic Endogenous Mediator (5,6). Although it was initially thought that polymorphonuclear leukocytes produced these endogenous factors, subsequent work clearly indicated that mononuclear phagocytes were their major source (7). At about the same time, Gery and co-workers demonstrated that a factor produced by mononuclear phagocytes, Lymphocyte Activating Factor, had the capacity to enhance responsiveness of T lymphocytes (8,9). This factor, which was found to exert a number of activities on various lymphocyte populations, was given the name Interleukin 1 to delineate it from other cytokines active on lymphoid cells (10). It quickly became apparent that the activities of Endogenous Pyrogen, Leukocytic Endogenous Mediator and Lymphocyte Activating Factor could be accounted for by the same or a very similar molecule (11-13). Over

the past 5 years there has been a rapid expansion in knowledge concerning the role of IL 1 and related molecules in the mediation of various aspects of the acute phase response, culminating in the isolation of the cDNA's encoding the IL 1 genes (14-17) and the identification of IL 1 receptors (18,19). It has become clear that the IL 1 family of molecules exerts a wide range of biologic activities that play a central role not only in host defense against a variety of challenges but also as an important element in the pathogenesis of a number of chronic diseases.

1975

Chen

IL 1 mRNA levels in macrophages

macrophage products and IL 1 in rheumatoid synovial fluid

1979

International Lymphocyte Workshop

Interleukin-1 (IL-1)

1984,85

Lomedico, Aaron March, Parham
encoding IL 1 genes 14-17

Molecular cloning of the cDNA

1989

Figini

Identification of interleukin 1 receptors (18,19)

TABLE I

A BRIEF HISTORY OF INTERLEUKIN 1

- 1948 - Beeson: Granulocyte pyrogen (endogenous pyrogen)
 - extracts of leukocytes cause fever (2)
- 1969 - Kampschmidt: Leukocyte endogenous mediator
 - phagocyte product induces the acute phase response (5)
- 1972 - Gery: Lymphocyte activating factor
 - macrophage product enhances T lymphocyte responses (8,9)
- 1979 - International Lymphokine Workshop: Interleukin 1 (10)
- 1984,85 - Lomedico, Auron, March, Furutani: Molecular cloning of the cDNA encoding interleukin 1 (14-17)
- 1985 - Dower: Identification of interleukin 1 receptors (18,19)

TABLE II
ACRONYMS, SYNONYMS AND PSEUDONYMS OF
MEMBERS OF THE INTERLEUKIN 1 FAMILY

<u>FACTOR</u>	<u>ACTIVITY</u>	<u>REFERENCES</u>
EP - Endogenous pyrogen	A monocyte/macrophage product that induces fever in vivo.	(20-23)
LEM - Leukocyte endogenous mediator	A monocyte/macrophage product that induces neutrophilia, stimulates the production of acute phase reactants and depresses serum zinc and iron in vivo.	(24)
LAF - Lymphocyte activating factor	A monocyte/macrophage product that augments mitogen induced proliferation of murine thymocytes. This factor also augments antibody production from T cell depleted B lymphocytes.	(8,9)
BAF:	B cell activating factor (25-28)	
MP:	Mitogenic protein (29)	
HP-1:	Helper-peak 1 (30)	
TRF-III:	T cell replacing factor III (31)	
TRF _M :	T cell replacing factor _M (32)	
BDF:	B cell differentiation factor (33)	
MCF - Mononuclear cell factor	A monocyte product that augments the production of PGE ₂ and collagenase from synovial fibroblasts.	(34-38)
PIF - Proteolysis inducing factor	A factor (M _r 4200) found in plasma of febrile, burned or traumatized patients that stimulates muscle cells to produce prostaglandins leading to proteolysis, release of amino acids and muscle wasting. It is also pyrogenic, causes elevation in acute phase reactants and has thymocyte stimulatory activity.	(39-40)
ETAF- Epidermal cell-derived thymocyte activating factor	A factor produced by keratinocytes with IL-1 activity	(41,42)
Catabolin	A factor produced by porcine synovial cells and monocytes that induces proteoglycan matrix degradation of cartilage.	(43-45)

Table III

INTERLEUKIN 1

1. 14,000 - 17,000 dalton protein (46)
 - sensitive to heating, acid pH
 - pronase sensitive
 - murine IL 1 - trypsin resistant
 - human IL 1 - trypsin, chymotrysin sensitive
 - inhibited by the arginine specific inhibitor, phenyl glyoxal (47)
2. Charge heterogeneity:
 - pI 5.0
 - pI 7.0
3. Not glycosylated (48)
4. Low molecular weight fragments of 2 and 4kD found in urine and plasma are active (49-51)

TABLE IV
ACTIVITIES OF INTERLEUKIN 1

T Lymphocytes

1. Co-mitogenic with lectins (8,9)
2. ↑ Lymphokine production (IL 2, IFN γ) (52,53)
3. ↑ Activity of cytotoxic T cells (54)
4. ↑ Receptors for IL-2 (55,56)
5. Chemotaxis (57)
6. Radioprotection (58)
7. ↑ Cell cycle progression of activated cells (59)
8. ↑ Membrane viscosity and (?) antigen binding (60)

B Lymphocytes

1. Maturation of pre-B cells (61,62)
2. Cofactor in clonal expansion (63,64)
3. Promote differentiation of antibody forming cells (65,66)
4. Chemotaxis (57)

NK Cells

1. Enhance killing and synergistically enhance killing with IFN- γ (67,68)

Macrophages

1. ↑ Production of prostaglandin (69)
2. ↑ Tumoricidal activity (70)

Polymorphonuclear Leukocytes

1. ↑ Release from bone marrow (71)
2. ↑ Margination (72)
3. ↑ Extravasation (73)
4. Chemotaxis (74,75)
5. Activation of glucose metabolism (74)
6. Stimulate NBT reduction (74)
7. Stimulate release of lysosomal enzymes (76,77)
8. Stimulate adherence to endothelial cells (78,79)

Endothelial Cells

1. ↑ Proliferation (80)
2. ↑ Prostacyclin production (81)
3. ↑ Adhesiveness (82,83)
4. ↑ Procoagulant activity (84,85)
5. Induction of new surface antigens (86)

Synovial Cells and Fibroblasts

1. PGE₂ production (34-38)
2. Collagenase production (34-38,87)
3. Proliferation (88,89)
4. Production of β interferon (90)

TABLE IV (Cont)

Chondrocytes

1. ↑ Production of collagenase (91)
Neutral proteases (92), plasminogen activator (93)
2. ↑ Production of PGE₂ (94)

Central Nervous System

1. Stimulation of hypothalamic PGE₂ production → fever (20)
2. Somnolence (95)
3. Slow wave sleep (96,97)
4. Anorexia (98)
5. Proliferation of glial cells (99)

Hepatocytes

1. ↑ Synthesis of acute phase proteins (24)
2. ↓ Synthesis of albumin and prealbumin (24)

Muscle Cells

1. Proteolysis (PIF) (39,40)

Mesangial Cells

1. Proliferation (100)

Osteoblasts

1. ↑ Proliferation (101)
2. ↑ Alkaline phosphatase production (101)

Osteoclasts

1. ↑ Bone resorption (102)

Epidermal Cells

1. ↑ Synthesis of collagen type IV (103)

Adipocytes

1. ↓ Lipoprotein lipase (104)

Tumor Cells

1. ↑ Cytocidal activity (105,106)

TABLE V
MULTIPLE BIOLOGIC ACTIVITIES ATTRIBUTED TO
INTERLEUKIN 1

Fever*
Hypoferremia
Hypozincemia
Hypercupremia
Increased
Blood neutrophils*
Hepatic acute-phase proteins*
Bone resorption*
Cartilage breakdown
Muscle proteolysis
Slow-wave sleep
Endothelial procoagulant
Chondrocyte proteases
Synovial collagenase*
Endothelial neutrophil adherence
Neutrophil degranulation*
Neutrophil superoxide
Interferon production
Proliferation of
Fibroblasts
Glial cells
Mesangial cells
Synovial fibroblasts
EBV B-cell lines
Chemotaxis of
Monocytes
Neutrophils*
Lymphocytes*
Stimulation of PGE ₂ in
Hypothalamus
Cortex
Skeletal muscle
Dermal fibroblast*
Synovial fibroblast*
Chondrocyte
Macrophage/monocyte*
Endothelium (PGI ₂)
Decreased
Hepatic albumin synthesis*
Appetite*
Brain binding of opioids
Lipoprotein lipase activity*
Augmentation of
T-cell responses*
B-cell responses
NK activity
IL-2 production*
Lymphokine production

* Activity confirmed by recombinant materials

TABLE VI
INTERLEUKIN 1 AND THE ACUTE PHASE RESPONSE

I. Increased hepatic synthesis of:

- a) serum amyloid A protein (107-114)
- b) fibrinogen (115-124)
- c) C reactive protein (118,121,125,126)
- d) haptoglobin (118-120,123,126-128)
- e) ceruloplasmin (118,123)
- f) complement factor B (129)

II. Decreased hepatic synthesis of:

- a) albumin (128)
- b) pre-albumin (128)

III. Decrease in serum

- a) iron - sequestration in lactoferrin complexes (126,130-134)
- b) zinc - induction of hepatic metallothionein (120,131,132,135)

TABLE VII

INDUCERS OF IL-1 PRODUCTION FROM MONOCYTE/MACROPHAGES

Microorganisms	
Viruses	
Bacteria	
Spirochetes	
Yeasts	
Microbial products	
Endotoxins from gram-negative bacteria	
Peptidoglycans from all bacteria	
Exotoxins from pathogenic strains of <i>Staphylococci</i> and <i>Streptococci</i>	
Yeast polysaccharides	
Inflammatory agents	
Bile salts	
Etiocolanolone	
C5a	
Silica	
Urate crystals	
Plant lectins	
Phytohemagglutinin	
Concanavalin A	
Cytokines	
Colony-stimulating factor	
Tumor necrosis factor	
Other inducers	
Bleomycin	
PolyI:C	
Muramyl dipeptide	

TABLE VIII
AGENTS THAT INHIBIT THE ACTIVITY OF INTERLEUKIN-1

Inhibitors of Release from Monocytes and Macrophages

Corticosteroids (136-138)
Prostaglandins E₂ (139)
Chloroquine (140,141)
Ultraviolet light (142)

Inhibitors of Activity

Cyclooxygenase inhibitors (fever, muscle proteolysis) (21)
Urinary inhibitors: 20-30 kDa - urine of febrile people (143)
Uromodulin: 85 kDa - urine of pregnant women, normal and febrile plasma (144)
Products of macrophages (22kDa [145,146]), EBV transformed B cell lines (95kDa [147])
Inhibitors in synovial fluid (148,149)
Inhibitors in serum (150,151)

TABLE IX
CELL SOURCES OF IL-1 AND IL-1 LIKE ACTIVITIES (152)

Cell Type	Stimulants	Ref.
Monocyte, macrophages, and macrophage lines	LPS, particles, PMA MDP,	(153)
	C containing Immune complexes,	(145,154)
	Cytokines (CSF, IFN- γ , TNF)	(155-160)
	Activated T cells	(161-165)
	C5a	(166)
	Cell injury	(167)
Keratinocytes	PMA,MDP,LPS	(41,42,74,75,168-170)
Dendritic cells	LPS	(171)
Langerhans cells	LPS, <i>Staph. albus</i>	(172)
Large granular lymphocytes	LPS	(173)
B cell lines	None	(147,174,175)
B lymphoblasts	LPS, anti-IgM	(176)
Endothelial cells	LPS, thrombin	(177)
Mesangial cells	Cell cycle dependent	(178)
Astrocytes	LPS	(179)
Glioma cell lines	Cell cycle dependent	(180,181)
Microglial cells	LPS, <i>Staph. aureus</i> , injury	(182)
Neutrophils	Aluminum hydroxide	(183)
Fibroblasts	MDP	(184)
Epithelial cells	PMA, LPS, UV irradiation	(185,186)
Corneal epithelium	-	(187)

Abbreviations: LPS = lipopolysaccharide; PMA = phorbol myristate acetate; MDP = muramyl dipeptide; CSF = colony stimulating factor; IFN- γ = interferon γ ; TNF = tumor necrosis factor

TABLE X

**DETECTION OF IL-1-LIKE ACTIVITY IN
HUMAN TISSUES AND FLUIDS**

1. Synovial fluid (148,149)
2. Gingival fluid (188)
3. Plasma
 - a) febrile animals (189)
 - b) marathon runners (190)
 - c) following ovulation (150)
 - d) burned, traumatized, febrile or cirrhotic subjects (PIF) (40,40a)
 - e) patients with Crohn's disease (191)
4. Urine (192)
5. Stratum corneum from the human heel (193)

TABLE XI

DEFECTIVE INTERLEUKIN 1 PRODUCTION FROM
PERIPHERAL BLOOD MONOCYTES IN VIVO

1. Patients with some cancers (194,195)
2. Patients with systemic lupus erythematosus (196)
3. Some neonates (197)
4. Some elderly persons (198)
5. Malnutrition (199,200)

TABLE XII

FUNCTIONAL COMPARISON OF INTERLEUKIN 1
AND TUMOR NECROSIS FACTOR (TNF, CACHECTIN)

Shared Activities

1. Pyrogenic (157)
2. Stimulation of prostaglandin E₂ and collagenase production by synovial cells and dermal fibroblasts (34-38,201)
3. Activation of osteoclasts → bone resorption (101,202)
4. Stimulation of procoagulant activity by vascular endothelial cells (84,85,203)
5. Stimulation of neutrophil adherence to endothelial cells (78,79,82,83,204)
6. Stimulation of neutrophil phagocytosis, respiratory burst and degranulation (74,76,77,205)
7. Inhibition of lipoprotein lipase (104,206)
8. Induction of new surface antigens on endothelial cells (86)

Unique features of TNF

1. Cytolytic activity (206)
2. Induction of IL-1 biosynthesis (157)
3. Shock (206)

Unique features of IL-1

1. Thymocyte stimulatory activity
2. Induction of acute phase reactants

TABLE XIII
CHARACTERISTICS OF MURINE AND HUMAN
INTERLEUKIN 1 PRECURSOR COMPLEMENTARY DNA
(14-17)

	Murine (pI 5)	Human IL-1 α (pI 5)	Human IL-2 β (pI 7)
Source	P388D1	Human macrophages	Human monocytes
mRNA	2.0 Kb	2.0 - 2.4 Kb	1.6 Kb
Abundance of mRNA (% total polyA + RNA)	0.005%	0.01%	0.1%
Amino acids	270	271	269
Predicted M _r of primary translation product	31,026	30,606	30,747*
Molecular weight of secreted product with biologic activity	17,992 (C terminal 156 amino acids)	17,500 (C terminal 159 amino acids)	17,500 (C terminal 153 amino acids)
N terminal hydrophobic signal sequence	No	No	No
Membrane spanning region	No	No	No
Homology**			

* Not biologically active

** Over 50% of the identical positions between IL-1 proteins occur in the M_r 17,500 C terminal biologically active portion of the molecule and a large proportion of these occur in the middle and C terminal regions suggesting that these regions are critical for biologic activity. However, 47% of all common residues occur in the portion of the primary translation product that precedes the active IL-1 segment. This degree of homology is not expected in a non-functional portion of a protein and therefore suggests that the N-terminal portion of the molecule may have some unknown biologic function.

FIGURE 1: Nucleotide Sequence and Predicted Amino Acid Sequence of Human IL-1 β Primary Translation Product (16)

5' TTCGAGGCACAAGGCACAACAGGCTGCTCTGGGATTCTCTTCAGCCAATCTTCATTGCTCAAGTGCTGAAGCAGCC	-1
ATG GCA GAA GTA CCT GAG CTC GCC AGT GAA ATG ATG GCT TAT TAC AGT GGC AAT GAG GAT	60
Met Ala Glu Val Pro Glu Leu Ala Ser Glu Met Met Ala Tyr Tyr Ser Gly Asn Glu Asp	20
GAC TTG TTC TTT GAA GCT GAT GGC CCT AAA CAG ATG AAG TGC TCC TTC CAG GAC CTG GAC	120
Asp Leu Phe Phe Glu Ala Asp Gly Pro Lys Gln Met Lys Cys Ser Phe Gln Asp Leu Asp	40
CTC TGC CCT CTG GAT GGC GGC ATC CAG CTA CGA ATC TCC GAC CAC CAC TAC AGC AAG GGC	180
Leu Cys Pro Leu Asp Gly Gly Ile Gln Leu Arg Ile Ser Asp His His Tyr Ser Lys Gly	60
TTC AGG CAG GCC GCG TCA GTT GTT GTG GCC ATG GAC AAG CTG AGG AAG ATG CTG GTT CCC	240
Phe Arg Gln Ala Ala Ser Val Val Val Ala Met Asp Lys Leu Arg Lys Met Leu Val Pro	80
TGC CCA CAG ACC TTC CAG GAG AAT GAC CTG AGC ACC TTC TTT CCC TTC ATC TTT GAA GAA	300
Cys Pro Gln Thr Phe Gln Glu Asn Asp Leu Ser Thr Phe Phe Pro Phe Ile Phe Glu Glu	100
GAA CCT ATC TTC TTT GAC ACA TGG GAT AAC GAG GCT TAT GTG CAC GAT GCA CCT GTA CGA	360
Glu Pro Ile Phe Phe Asp Thr Trp Asp Asn Glu Ala Tyr Val His Asp ALA PRO VAL ARG	120
TCA CTG AAC TGC ACG CTC CGG GAC TCA CAG CAA AAA AGC TTG GTG ATG TCT GGT CCA TAT	420
SER LEU ASN Cys THR LEU ARG ASP SER GLN GLN LYS SER LEU VAL MET SER GLY PRO TYR	140
GAA CTG AAA GCT CTC CAC CTC CAG GGA CAG GAT ATG GAG CAA CAA GTG GTG TTC TCC ATG	480
GLU LEU LYS ALA LEU HIS LEU GLN GLY GLN ASP MET GLU GLN GLN VAL VEL PHE Ser Met	160
TCC TTT GTA CAA GGA GAA GAA AGT AAT GAC AAA ATA CCT GTG GCC TTG GGC CTC AAG GAA	540
SER PHE VAL GLN GLY GLU GLU Ser ASN ASP LYS Ile Pro Val Ala Leu Gly Leu Lys Glu	180
AAG AAT CTG TAC CTG TCC TGC GTG TTG AAA GAT GAT AAG CCC ACT CTA CAG CTG GAG ACT	600
LYS ASN LEU TYR LEU SER Cys Val Leu Lys Asp Asp Lys Pro Thr Leu Gln Leu Glu SER	200
GTA GAT CCC AAA AAT TAC CCA AAG AAG AAG ATG GAA AAG CGA TTT GTC TTC AAC AAG ATA	660
VAL ASP PRO Lys Asn Tyr Pro Lys Lys Lys Met GLU LYS ARG PHE VAL PHE ASN Lys ILE	220
GAA ATC AAT AAC AAG CTG GAA TTT GAG TCT GCC CAG TTC CCC AAC TGG TAC ATC AGC ACC	720
GLU ILE ASN ASN Lys LEU GLU Phe Glu SER ALA GLN PHE PRO ASN TRP TYR ILE Ser Thr	240
TCT CAA GCA GAA AAC ATG CCC GTC TTC CTG GGA GGG ACC AAA GGC GGC CAG GAT ATA ACT	780
Ser Gln Ala Glu Asn Met PRO VAL PHE LEU GLY GLY Thr Lys Gly Gly Gln Asp Ile Thr	260
GAC TTC ACC ATG CAA TTT GTG TCT TCC TAA AGAGAGCTGTACCCAGAGAGTCCTGTGCTGAATGTCGAC	849
Asp Phe Thr Met GLN PHE VAL SER SER End	
TCAATCCCTAGGGCTGGCAGAAAGGGAACAGAAAGGTTTTTGTAGTACGGCTATAGCCTGGACTTTCCTGTTGTCTACAC	928
CAATGCCCAACTGCCTTAGGGTAGTGCTAAGAGGATCCTGTCCATCAGCCAGGACAGTCAGCTCTCTCCTTTC	1007
AGGGCCAATCCAGCCCTTTTGTGAGCCAGGCCTCTCT - 3'	1047

Amino acids in capitals indicate those residues confirmed by protein sequencing analysis.

FIGURE 2: Nucleotide Sequence and Predicted Amino Acid Sequence of Human IL-1 α Primary Translation Product (16)

5' -- TGAGGGAGTCATTTCATTTGCGYTTGAGTCAGCAAGAAGTCAAG																				-1
ATG	GCC	AAA	GTT	CCA	GAC	ATG	TTT	GAA	GAC	CTG	AAG	AAC	TGT	TAC	AGT	GAA	AAT	GAA	GAA	60
Met	Ala	Lys	Val	Pro	Asp	Met	Phe	Glu	Asp	Leu	Lys	Aan	Cys	Tyr	Ser	Glu	Asn	Glu	Glu	20
GAC	AGT	TCC	TCC	ATT	GAT	CAT	CTG	TCT	CTG	AAT	CAG	AAA	TCC	TTC	TAT	CAT	GTA	AGC	TAT	120
Asp	Ser	Ser	Ser	Ile	Asp	His	Leu	Ser	Leu	Asn	Gln	Lys	Ser	Phe	Tyr	His	Val	Ser	Tyr	40
GGC	CCA	CTC	CAT	GAA	GGC	TGC	ATG	GAT	CAA	TCT	GTG	TCT	CTG	AGT	ATC	TCT	GAA	ACC	TCT	180
Gly	Pro	Leu	His	Glu	Gly	Cys	Met	Asp	Gln	Ser	Val	Ser	Leu	Ser	Ile	Ser	Glu	Thr	Ser	60
AAA	ACA	TCC	AAG	CTT	ACC	TTC	AAG	GAG	AGC	ATG	GTG	GTA	GTA	GCA	ACC	AAC	GGG	AAG	GTT	240
Lys	Thr	Ser	Lys	Leu	Thr	Phe	Lys	Glu	Ser	Met	Val	Val	Val	Ala	Thr	Asn	Gly	Lys	Val	80
CTG	AAG	AAG	AGA	CGG	TTG	AGT	TTA	AGC	CAA	TCC	ATC	ACT	GAT	GAT	CAG	CTG	GAG	GCC	ATC	300
Leu	Lys	Lys	Arg	Arg	Leu	Ser	Leu	Ser	Gln	Ser	Ile	Thr	Asp	Asp	Asp	Leu	Glu	Ala	Ile	100
GCC	AAT	GAC	TCA	GAG	GAA	GAA	ATC	ATC	AAG	CCT	AGG	TCA	GCA	CCT	TTT	AGG	TTC	CTG	AGC	360
Ala	Asn	Asp	Ser	Glu	Glu	Glu	Ile	Ile	Lys	Pro	Arg	Ser	Ala	Pro	Phe	Ser	Phe	Leu	Ser	120
AAT	GTG	AAA	TAC	AAC	TTT	ATG	AGG	ATC	ATC	AAA	TAC	GAA	TTC	ATC	CTG	AAT	GAC	GCC	CTC	420
Asn	Val	Lys	Tyr	Asn	Phe	Met	Arg	Ile	Ile	Lys	Tyr	Glu	Phe	Ile	Leu	Asn	Asp	Ala	Leu	140
AAT	CAA	AGT	ATA	ATT	GGA	GCC	AAT	GAT	CAG	TAC	CTC	AGC	GCT	GCT	GCA	TTA	CAT	AAT	CTG	480
Asn	Gln	Ser	Ile	Ile	Arg	Ala	Asn	Asp	Gln	Tyr	Leu	Thr	Ala	Ala	Ala	Leu	His	Asn	Leu	160
GAT	GAA	GCA	GTG	AAA	TTT	GAC	ATG	GGT	GCT	TAT	AAG	TCA	TCA	AAG	GAT	GAT	GCT	AAA	ATT	540
Asp	Glu	Ala	Val	Lys	Phe	Asp	Met	Gly	Ala	Tyr	Lys	Ser	Ser	Lys	Asp	Asp	Ala	Lys	Ile	180
ACC	GTG	ATT	CTA	AGA	ATC	TCA	AAA	ACT	CAA	TTG	TAT	GTG	ACT	GCC	CAA	GAT	GAA	GAC	CAA	600
Thr	Val	Ile	Leu	Arg	Ile	Ser	Lys	Thr	Gln	Leu	Tyr	Val	Thr	Ala	Gln	Asp	Glu	Asp	Gln	200
CCA	GTG	CTG	CTG	AAG	GAG	ATG	CCT	GAG	ATA	CCC	AAA	ACC	ATC	ACA	GGT	AGT	GAG	ACC	AAC	660
Pro	Val	Leu	Leu	Lys	Glu	Met	Pro	Glu	Ile	Pro	Lys	Thr	Ile	Thr	Gly	Ser	Glu	Thr	Asn	220
CTC	CTC	TTC	TTC	TGG	GAA	ACT	CAG	GGC	ACT	AAG	AAC	TAT	TTC	ACA	TCA	GTT	GCC	CAT	CCA	720
Leu	Leu	Phe	Phe	Trp	Glu	Thr	His	Gly	Thr	Lys	Asn	Tyr	Phe	Thr	Ser	Val	Ala	His	Pro	240
AAC	TTG	TTT	ATT	GCC	ACA	AAG	CAA	GAC	TAC	TGG	GTG	TGC	TTG	GCA	GGG	GGG	CCA	CCC	TCT	780
Asn	Leu	Phe	Ile	Ala	Thr	Lys	Gln	Asp	Tyr	Trp	Val	Cys	Leu	Ala	Gly	Gly	Pro	Pro	Ser	260
ATC	ACT	GAC	TTT	CAG	ATA	CTG	GAA	AAC	CAG	GCG	TAG	GTCTGGAGTCTCAGTTGTCTCACTTGTGCAG								847
Ile	Thr	Asp	Phe	Gln	Ile	Leu	Glu	Asn	Gln	Ala	End								271	
TGTTGACAGTTTCATATGTACCATGTACATGAAGAAGCTAAATCCTTTACTGTTAGTCATTTGCTGAGCATGTACTGAGC																				926
CTTGTAATTCTAAATGAATGTTTACACTCTTTGTAAGAGTGAACCAACACTAACATATAATGTTGTTATTTAAAGAAC																				1005
ACCCTATATTTTGCATAGTACCAATCATTTTAATTATTATCTTCATAACAATTTTAGGAGGACCAGAGCTACTGACTA																				1084
TGGCTACCAAAAAGACTCTACCCATATTACAGATGGGCAAATTAAGGCATAAGAAAACTAAGAAATATGCACAATAGCA																				1163
ATTGAAACAAGAAGCCACAGACCTAGGATTTTCATGATTTTCATTTCAACTGTTTGCCCTTCTGCTTTTAAGTTGCTGATGA																				1242
ACTCTTAATCAAATAGCATAAGTTTCTGGGACCTCAGTTTTATCATTTTCAAAATGGAGGGAATAATACCTAAGCCCTTC																				1321
CTGCCGCAACAGTTTTTTATGCTAATCAGGGAGGTCATTTTGGTAAAATACTTCTCGAAGCCGAGCCTCAAGATGAAGG																				1400
CAAAGCAGGAAATGTTATTTTTTAATTATTATTATATATGATATTATATAAATATATTTAAGATAAATTATAATATACTAT																				1479
ATTTATGGGAACCCCTTCATCCTCTGAGTGTGACCAAGGCTCCTCCACAATAGCAGACAGTGTTTTCTGGGATAAGTAA																				1558
GTTTGATTTTCATTAAATACAGGGCATTTTGGTCCAAGTTGTGCTTATCCCATAGCCAGGAAACTCTGCATCTAGTACTT																				1637
GGGAGACCTGTAATCATATAATAAATGTACATTAATTACCTTGAGCCAGTAATTTGGTCCGATCTTTGACTCTTTTGCCA																				1716
TTAAACTTACCTGGGCATTCCTGTTTCATTCAATTCACCTGCAATCAAGTCTACAAGCTAAAATTAGATGAACCTCAA																				1795
CTTTGACAACCATAGACCACTGTATACAAAATTTCTTTCTGGAATGTAATCAATGTTTCTCTAGGTTCTAAAAATT																				1874
GTGATCAGACCATAATGTTACATTATTATCAACAATAGTGATTGATAGAGTGTTATCAGTCATAACTAAATAAAGCTTG																				1953
CAAGAAAAAATAAATAAAAAA - 3'																				1979

FIGURE 3: Alignment of Human IL-1 α , Murine IL-1 and Human IL-1 β Amino Acid Sequences (16)

α	Met	Ala	Lys	Val	Pro	Asp	Met	Phe	Glu	Asp	Leu	Lys	Asn	Cys	Tyr	Ser	Glu	Asn	Glu	Glu
β	Met	Ala	Lys	Val	Pro	Asp	Leu	Phe	Glu	Asp	Leu	Lys	Asn	Cys	Tyr	Ser	Glu	Asn	Glu	Asp
	Met	Ala	Glu	Val	Pro	Glu	Leu	Ala	Ser	Glu	Met	Met	Ala	Tyr	Ser	Ser	Gly	Asn	Glu	Asp
	Asp	Ser	Ser	Ser	Ile	Asp	His	Leu	Ser	Leu	Asn	Gln	Lys	-	Ser	Phe	Tyr	His	Val	Ser
	Tyr	Ser	Ser	Ala	Ile	Asp	His	Leu	Ser	Leu	Asn	Gln	Lys	-	Ser	Phe	Tyr	Asp	Ala	Ser
	Asp	Leu	Phe	Phe	Glu	Ala	Asp	Gly	Pro	Leu	Gln	Met	Lys	Cys	Ser	Phe	Gln	Asp	Leu	Asp
	Tyr	Gly	Pro	Leu	His	Glu	Gly	Cys	Met	Asp	Gln	Ser	Val	Ser	Leu	Ser	Ile	Ser	Glu	Thr
	Tyr	Gly	Ser	Leu	His	Glu	Thr	Cys	Thr	Asp	Gln	Phe	Val	Ser	Leu	Arg	Thr	Ser	Glu	Thr
	Leu	Cys	Pro	Leu	-	-	-	-	-	Asp	Gly	Gly	Ile	Gln	Leu	Arg	Ile	Ser	Asp	His
	Ser	Lys	Thr	Ser	Lys	Leu	Thr	Phe	Lys	Glu	Ser	Met	-	Val	Val	Val	-	Ala	Thr	-
	Ser	Lys	Met	Ser	Asn	Phe	Thr	Phe	Lys	Glu	Ser	Arg	-	Val	Thr	Val	Ser	Ala	Thr	Ser
	His	-	Tyr	Ser	Lys	Gly	-	Phe	Arg	Gln	Ala	Ala	Ser	Val	Val	Val	-	Ala	Met	-
	-	Asn	Gly	Lys	Val	Leu	Lys	Lys	Arg	Arg	Leu	Ser	Leu	Ser	Gln	Ser	Ile	Thr	Asp	Asp
	Ser	Asn	Gly	Lys	Ile	Leu	Lys	Lys	Arg	Arg	Leu	Ser	Phe	Ser	Glu	Thr	Phe	Thr	Glu	Asp
	-	-	Asp	Lys	-	Leu	Arg	Lys	Met	Leu	Val	Pro	Cys	Pro	Gln	Thr	Phe	Gln	Glu	Asn
	Asp	Leu	Glu	Ala	Ile	Ala	Asn	Asp	Ser	-	Glu	Glu	Glu	Ile	Ile	-	-	-	-	-
	Asp	Leu	Gln	Ser	Ile	Thr	His	Asp	Leu	-	-	Glu	Glu	Thr	Ile	-	-	-	-	-
	Asp	Leu	Ser	Thr	Phe	Phe	Pro	Phe	Ile	Phe	Glu	Glu	Glu	Pro	Ile	Phe	Phe	Asp	Thr	Trp
	-	-	-	-	Lys	Pro	Arg	Ser	Ala	Pro	Phe	Ser	Phe	Leu	Ser	Asn	Val	Lys	Tyr	Asn
	-	-	-	-	Gln	Pro	Arg	Ser	Ala	Pro	Tyr	Thr	Tyr	Gln	Ser	Asp	Leu	Arg	Tyr	Lys
	Asp	Asn	Glu	Ala	Tyr	Val	His	Asp	Ala	Pro	-	-	-	Val	Arg	Ser	Leu	Asn	Cys	Thr
	Phe	Met	Arg	Ile	Ile	Lys	Tyr	Glu	Phe	Ile	Leu	Asn	Asp	Ala	Leu	Asn	Gln	Ser	Ile	Ile
	Leu	Met	Lys	Leu	Val	Arg	Gln	Lys	Phe	Val	Met	Asn	Asp	Ser	Leu	Asn	Gln	Thr	Ile	Tyr
	Leu	Arg	Asp	Ser	Gln	Gln	Lys	Ser	Leu	Val	Met	Ser	-	-	-	-	-	Gly	Pro	Tyr
	Arg	Ala	Asn	Asp	Gln	-	Tyr	Leu	Thr	Ala	Ala	Ala	Leu	His	Asn	Leu	Asp	Glu	Ala	Val
	Gln	Asp	Val	Asp	Lys	His	Tyr	Leu	Ser	Thr	Thr	Trp	Leu	Asn	Asp	Leu	Gln	Gln	Glu	Val
	Glu	Leu	Lys	Ala	Leu	His	-	Leu	Gln	Gly	-	-	-	Gln	Asp	Met	Glu	Gln	Gln	Val
	Lys	Phe	Asp	Met	Gly	Ala	Tyr	Lys	Ser	Ser	Lys	-	Asp	Asp	Ala	Lys	Ile	Thr	Val	Ile
	Lys	Phe	Asp	Met	Tyr	Ala	Tyr	Ser	Ser	Gly	Gly	-	Asp	Asp	Ser	Lys	Tyr	Pro	Val	Thr
	Val	Phe	Ser	Met	Ser	Phe	Val	Gln	Gly	Glu	Glu	Ser	Asn	Asp	-	Lys	Ile	Pro	Val	Ala
	Leu	Arg	Ile	Ser	Lys	Thr	Gln	Leu	Tyr	Val	Thr	Ala	Gln	-	Asp	Glu	Asp	Gln	Pro	Val
	Leu	Lys	Ile	Ser	Asp	Ser	Gln	Leu	Phe	Val	Ser	Ala	Gln	-	Gly	Glu	Asp	Gln	Pro	Val
	Leu	Gly	Leu	Lys	Glu	Lys	Asn	Leu	Tyr	Leu	Ser	Cys	Val	Leu	Lys	Asp	Asp	Lys	Pro	Thr
	Leu	Leu	Lys	Glu	Met	Pro	Glu	Ile	Pro	Lys	Thr	Ile	Thr	-	-	Gly	Ser	Glu	Thr	Asn
	Leu	Leu	Lys	Glu	Leu	Pro	Glu	Thr	Pro	Lys	Leu	Ile	Thr	-	-	Gly	Ser	Glu	Thr	Asp
	Leu	Gln	Leu	Glu	Ser	Val	Asp	-	Pro	Lys	Asn	Tyr	Pro	Lys	Lys	Lys	Met	Glu	Lys	Arg
	Leu	Leu	Phe	Phe	Trp	Glu	Thr	His	Gly	Thr	Lys	Asn	Tyr	Phe	Thr	Ser	Val	Ala	His	Pro
	Leu	Ile	Phe	Phe	Trp	Lys	Ser	Ile	Asn	Ser	Lys	Asn	Tyr	Phe	Thr	Ser	Ala	Ala	Tyr	Pro
	Phe	Val	Phe	Asn	Lys	Ile	Glu	Ile	Asn	Asn	Lys	Leu	Glu	Phe	Glu	Ser	Ala	Gln	Phe	Pro
	Asn	Leu	Phe	Ile	Ala	Thr	Lys	Gln	-	-	Asp	Tyr	Trp	Val	Cys	Leu	-	-	Ala	Gly
	Glu	Leu	Phe	Ile	Ala	Thr	Lys	-	-	Glu	Gln	Ser	Arg	Val	His	Leu	-	-	Ala	Arg
	Asn	Trp	Tyr	Ile	Ser	Thr	Ser	Gln	Ala	Glu	Asn	Met	Pro	Val	Phe	Leu	Gly	Gly	Thr	Lys
	Gly	Pro	Pro	Ser	Ile	Thr	Asp	Phe	Gln	Ile	Leu	Glu	Asn	Gln	Ala	-	-	-	-	-
	Gly	Leu	Pro	Ser	Met	Thr	Asp	Phe	Gln	Ile	Ser	-	-	-	-	-	-	-	-	-
	Gly	Gly	Gln	Asp	Ile	Thr	Asp	Phe	Thr	Met	Gln	Phe	Val	Ser	Ser	-	-	-	-	-

TABLE XIV
GENOMIC ORGANIZATION OF THE
HUMAN IL-1 α GENE
(207)

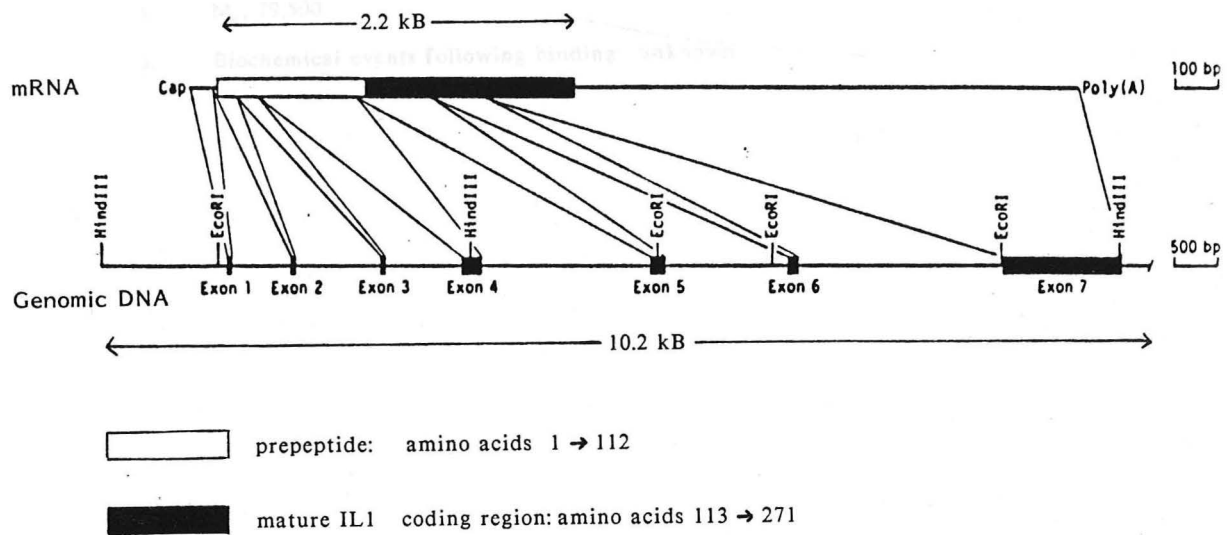


TABLE XV

INTERLEUKIN-1 RECEPTORS (18,19)

1. Widely distributed in low numbers
2. K_a : $2.6 - 3.6 \times 10^9/M$
- [Preliminary evidence for both high (K_d 8-15 pM) and lower (K_d 1nM) affinity receptors (208)]
3. Bind both IL-1 α and IL1 β but not other polypeptide hormones
4. M_r , 79,500
5. Biochemical events following binding: unknown

Species	Cell Type	IL-1 Receptor	Ref.
Mouse	T lymphoma	1.0 x 10 ⁵ / 10 ⁶	450
Mouse	T lymphoma	1.0 x 10 ⁵ / 10 ⁶	451
Mouse	T lymphoma	1.0 x 10 ⁵	452
Mouse	Fibroblast	1.0 x 10 ⁵	453
Mouse	T lymphoma	1.0 x 10 ⁵ / 10 ⁶	454
Human	T lymphoma	1.0 x 10 ⁵	455
Human	Monocyte	1.0 x 10 ⁵	456
Human	Myeloid leukemia	1.0 x 10 ⁵	457
Human	Endothelial cell	1.0 x 10 ⁵	458
Human	Prothrombocyte	1.0 x 10 ⁵	459
Human	Myeloma	1.0 x 10 ⁵	460
Human	B lymphoma	1.0 x 10 ⁵	461
Human	Neutrophil	1.0 x 10 ⁵	462
Human	Macrophage	1.0 x 10 ⁵	463
Human	Epithelial cell	1.0 x 10 ⁵	464
Human	Endothelial cell	1.0 x 10 ⁵	465

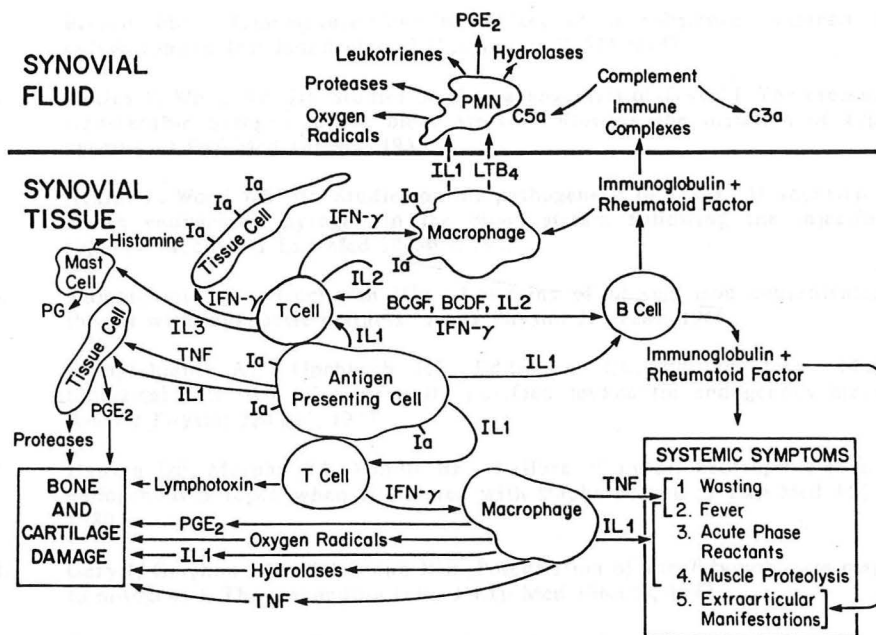
Dowling, S.K. et al. J. Exp. Med. 157: 521 (1983)

TABLE XVI
CELLULAR DISTRIBUTION OF IL-1 RECEPTORS (18)

Primary cells			
Species	Cell Type		Receptor sites per cell
Mouse	Thymocytes		<10
Mouse	PNA+ fraction		<10
Mouse	PNA- fraction		27
Mouse	Spleen cells		<10
Mouse	Lymph node cells		<10
Human	Peripheral blood mononuclear cells		27
Human	T cell line		100
Human	Gingival fibroblasts		4x10 ³
In vitro cell lines			
Species	Cell Type	Designation	
Mouse	T lymphoma	LBRM-33-1A4	550
Mouse	T lymphoma	LBRM-33-5A4	<10
Mouse	T lymphoma	EL-4	185
Mouse	Fibroblast	L929	144
Mouse	Fibroblast	SC-1	1850
Human	T lymphoma	Jurkat-FHCRC	<10
Human	T lymphoma	HSB-2	79
Human	T leukemia	PEER	<10
Human	Monocyte	U937	<10
Human	Myelogenous leukemia	KG-1	<10
Human	Erythroleukemia	K562	<10
Human	Promyelocyte	HL-60	<10
Human	Myeloma	ARH77	330
Human	B lymphoma	BMB	540
Human	Melanoma	A375	465
Human	Hepatoma	SKHEP-2	<10
Rat	Fibroblast	XC	450
Rat	Hepatoma	HEP-2	540
Rat	Epithelial cell	HTC	510
Bovine	Endothelial cell	CPAE	750

Dower, S.K. et al J. Exp. Med. 162:501, 1985

FIGURE 4: The role of interleukin 1 in the pathogenesis of rheumatoid arthritis



Abbreviations: BCDF, B cell differentiation factor; BCGF, B cell growth factor; Ia, class II molecules encoded by genes of the major histocompatibility complex; IFN γ , interferon γ ; IL-1, interleukin 1; IL-2, interleukin-2; IL-3, interleukin-3; PGE $_2$, prostaglandin E $_2$; TNF, tumor necrosis factor.

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