

Description**FIELD OF THE INVENTION**

5 The invention relates generally to methods and compositions for treating diseases associated with excessive interferon- γ (IFN- γ) production, and more particularly to methods and compositions employing the Epstein-Barr virus (EBV) protein BCRF1 for effectively reducing levels of IFN- γ .

BACKGROUND

10 The immune system comprises a highly interactive complex of tissues, cell types, and soluble factors. Recently, it has been suggested that several diseases and immune disorders may be associated with imbalances among certain components of the immune system, particularly cytokines: e.g. Mosmann et al., *Ann. Rev. Immunol.*, Vol. 7, pgs. 145-173 (1989); Cher et al., *J. Immunol.*, Vol. 138, pgs. 3688-3694 (1987);
 15 Mosmann et al., *Immunol. Today*, Vol. 8, pgs. 223-227 (1987); and Heinzel et al., *J. Exp. Med.*, Vol. 169, pgs. 59-72 (1989).

For example, a large body of evidence suggests that excessive production of gamma interferon (IFN- γ) is responsible for autoimmune diseases associated with the major histocompatibility complex (MHC): Hooks et al., *New England J. Med.*, Vol. 301, pgs. 5-8 (1979) (elevated serum levels of IFN- γ correlated with
 20 autoimmunity); Basham et al., *J. Immunol.*, Vol. 130, pgs. 1492-1494 (1983) (IFN- γ can increase MHC gene product expression); Battazzo et al., *Lancet*, pgs. 1115-1119 (11/12/83) (aberrant MHC gene product expression correlated with some forms of autoimmunity); Hooks et al., *Ann. N.Y. Acad. Sci.*, Vol. 301, pgs. 21-32 (1980) (higher IFN- γ levels correlated to greater severity of disease in SLE patients, and histamine-release enhancing activity of interferon can be inhibited by anti-interferon sera); Jacob et al., *J. Exp. Med.*,
 25 Vol. 166, pgs. 798-803 (1987) (amelioration and delay of onset of disease conditions in mouse models of systemic lupus erythematosus by blocking anti-IFN- γ monoclonal antibodies); and Iwatani et al., *J. Clin. Endocrin. and Metabol.*, Vol. 63, pgs. 695-708 (1986) (anti-IFN- γ monoclonal antibody eliminated the ability of leucoagglutinin-stimulated T cells to induce HLA-DR expression). It has been hypothesized that excess IFN- γ causes the inappropriate expression of MHC gene products which, in turn, causes autoimmune
 30 reactions against the tissues whose cells are inappropriately expressing the MHC products and displaying autoantigens in the context of the products. Thus McDevit, [*Clin. Res.*, Vol. 34, pgs. 163-175 (1985)] has suggested that reducing IFN- γ levels in autoimmune patients, e.g. by administering IFN- γ antagonists, could have beneficial effects.

In addition to the above evidence, IFN- γ may also play a role in allergy by its ability to increase the
 35 number and density of Fc ϵ receptors on monocytes; it has been implicated in the pathogenesis of sarcoidosis and psoriasis; and it is believed to augment cell-mediated immunity, which plays a major role in tissue rejection in allogeneic transplant patients.

In view of the above, the availability of compounds capable of reducing IFN- γ levels would be highly advantageous for treatment of diseases associated with inappropriate immune responses, such as some
 40 parasitic diseases, allergy, and MHC-associated immune disorders, including rheumatoid arthritis, systemic lupus erythematosus (SLE), myasthenia gravis, insulin-dependent diabetes mellitus, thyroiditis, and the like.

SUMMARY OF THE INVENTION

45 The invention relates to methods and compositions for treating diseases associated with high levels of IFN- γ production. The method of the invention comprises the step of administering a disease-controlling amount of a BCRF1, a protein derived from the Epstein-Barr virus. The invention further includes expression vectors for producing recombinant BCRF1, purified BCRF1, and pharmaceutical compositions for use with the method. Preferably, the BCRF1 used with the invention is selected from the group of mature
 50 polypeptides of the open reading frame defined by the amino acid sequence shown in SEQ ID NO: 1, wherein the abbreviations indicate the L forms of the amino acids; and the amino acids are listed starting from the N-terminus.

BRIEF DESCRIPTION OF THE FIGURES

55 Figure 1 is a diagrammatic illustration of a mammalian expression vector useful in the production of BCRF1.

Figure 2 is a diagrammatic illustration of a bacterial expression vector useful in the production of BCRF1.

DETAILED DESCRIPTION OF THE INVENTION

The invention relates to methods and compositions for treating diseases associated with excessive IFN- γ production. The invention is based in part on the discovery that the nucleic acid sequence encoding a recently discovered protein, designated cytokine synthesis inhibitory factor (CSIF), possesses a high degree of homology with the EBV BCRF1 open reading frame. EBV is a human herpesvirus endemic in all human populations, and has been associated with several diseases: e.g. Dillner et al., Adv. Cancer Res., Vol 50, pgs. 95-158 (1988); Thorley-Lawson, Biochim. Biophys. Acta, Vol. 948, pgs. 263-286 (1988); and Tasato, Adv. Cancer Res., Vol. 49, pgs. 75-125 (1987). EBV has a double-stranded DNA genome of about 172 kilobases [Baer et al., Nature, Vol. 310, pgs. 207-211 (1984)]. The genome contains many open reading frames apparently corresponding to proteins produced by EBV, one of which is BCRF1.

The invention includes mature polypeptides, or proteins, of the BCRF1 open reading frame. For secreted proteins, an open reading frame usually encodes a polypeptide that consists of a mature or secreted product covalently linked at its N-terminus to a signal peptide. The signal peptide is cleaved prior to secretion of the mature, or active, polypeptide. The cleavage site can be predicted with a high degree of accuracy from empirical rules [e.g., von Heijne, Nucleic Acids Research, Vol. 14, pgs. 4683-4690 (1986)], and the precise amino acid composition of the signal peptide does not appear to be critical to its function [e.g. Randall et al., Science, Vol. 243, pgs. 1156-1159 (1989); Kaiser et al., Science, Vol. 235, pgs. 312-317 (1987)]. Consequently, mature proteins are readily expressed by vectors encoding signal peptides quite different from that encoded by the open reading frame defined by Formula I.

A wide range of expression systems (i.e. host-expression vector combinations) can be used to produce the proteins of the invention. Possible types of host cells include, but are not limited to, bacterial, yeast, insect, mammalian, and the like. Many reviews are available which provide guidance for making choices and/or modifications of specific expression systems: e.g. (to name a few), de Boer and Shepard, "Strategies for Optimizing Foreign Gene Expression in *Escherichia coli*," pgs. 205-247, in Kroon, ed. Genes: Structure and Expression (John Wiley & Sons, New York, 1983), review several *E. coli* expression systems; Kucheralapati et al., Critical Reviews in Biochemistry, Vol. 16, Issue 4, pgs. 349-379 (1984), and Banerji et al., Genetic Engineering, Vol. 5, pgs. 19-31 (1983) review methods for transfecting and transforming mammalian cells; Reznikoff and Gold, eds., Maximizing Gene Expression (Butterworths, Boston, 1986) review selected topics in gene expression in *E. coli*, yeast, and mammalian cells; and Thilly, Mammalian Cell Technology (Butterworths, Boston, 1986) reviews mammalian expression systems. Likewise, many reviews are available which describe techniques and conditions for linking and/or manipulating specific cDNAs and expression control sequences to create and/or modify expression vectors suitable for use with the present invention, e.g. Sambrook et al., Molecular Cloning: A Laboratory Manual, 2nd Ed. (Cold Spring Harbor Laboratory, N.Y., 1989).

An *E. coli* expression system is disclosed by Riggs in U.S. Patent 4,431,739, which is incorporated by reference. Particularly useful prokaryotic promoters for high expression in *E. coli* are the tac promoter, disclosed by de Boer in U.S. Patent 4,551,433, which is incorporated herein by reference, and the pL promoter, disclosed by Remaut et al., Gene, Vol 15, pgs. 81-93 (1981), which is incorporated by reference. Secretion expression vectors are also available for *E. coli* hosts. Particularly useful are the pIn-III-ompA vectors, disclosed by Ghayeb et al., in EMBO J., Vol. 3, pgs. 2437-2442 (1984), in which the cDNA to be transcribed is fused to the portion of the *E. coli* OmpA gene encoding the signal peptide of the ompA protein which, in turn, causes the mature protein to be secreted into the periplasmic space of the bacteria. U.S. Patents 4,336,336, 4,411,994, 4,332,892 and 4,338,397 also disclose secretion expression vectors for prokaryotes. Accordingly, these references are incorporated by reference. Numerous strains of bacteria are suitable hosts for prokaryotic expression vectors, including: strains of *E. coli*, such as W3110 (ATCC No. 27325), JA221, C600, ED767, DH1, LE392, HB101, X1776 (ATCC No. 31244), X2282, RR1 (ATCC No. 31343) MRCI; strains of *Bacillus subtilis*; and other enterobacteriaceae such as *Salmonella typhimurium* or *Serratia marcescens*, and various species of *Pseudomonas*. General methods for deriving bacterial strains, such as *E. coli* K12 X1776, useful in the expression of eukaryotic proteins, are disclosed by Curtis III in U.S. Patent 4,190,495. Accordingly this patent is incorporated by reference. In addition to prokaryotic and eukaryotic microorganisms, expression systems comprising cells derived from multicellular organism may also be used to produce proteins of the invention. Of particular interest are mammalian expression systems because their post-translational processing machinery is more likely to produce biologically active mammalian proteins.

Several DNA tumor viruses have been used as vectors for mammalian hosts. Particularly important are the numerous vectors which comprise SV40 replication, transcription, and/or translation control sequences coupled to bacterial replication control sequences, e.g. the pcD vectors developed by Okayama and Berg, disclosed in *Mol. Cell. Biol.*, Vol. 2, pgs. 161-170 (1982) and *Mol. Cell. Biol.*, Vol. 3, pgs. 280-289 (1983), and improved by Takebe et al., *Mol. Cell. Biol.*, Vol. 8, pgs. 466-472 (1988). Accordingly, these references are incorporated herein by reference. Other SV40-based mammalian expression vectors include those containing adenovirus regulatory elements, disclosed by Kaufman and Sharp, in *Mol. Cell. Biol.*, Vol. 2, pgs. 1304-1319 (1982), and by Clark et al., in U.S. patent 4,675,285, both of which are incorporated herein by reference. Monkey cells are usually the preferred hosts for the above vectors. Such vectors containing the SV40 ori sequences and an intact A gene can replicate autonomously in monkey cells (to give higher copy numbers and/or more stable copy numbers than nonautonomously replicating plasmids). Moreover, vectors containing the SV40 ori sequences without an intact A gene can replicate autonomously to high copy numbers (but not stably) in COS7 monkey cells, described by Gluzman, *Cell*, Vol. 23, pgs. 175-182 (1981) and available from the ATCC (accession no. CRL 1651). The above SV40-based vectors are also capable of transforming other mammalian cells, such as mouse L cells, by integration into the host cell DNA.

The biological activity of the BCRF1s of the invention is readily determined in IFN- γ inhibition assays. Such assays require a cell line or cell population that synthesizes IFN- γ . Conveniently, peripheral blood lymphocytes (PBLs) that have been stimulated with a mitogen such as phytohemagglutinin (PHA) can serve as such a cell population. Roughly, the assay works as follows: The PHA-stimulated PBLs are divided into two equal parts. To one part, a sample containing a BCRF1 is added. The other part serves as a control. After several days the supernatants of both cultures are tested for IFN- γ . This is conveniently done with a standard ELISA assay using commercially available monoclonal and polyclonal antibodies for IFN- γ , e.g. Genzyme, Inc. (Boston, MA). Alternatively, the readout of the assay can be the amount of IFN- γ mRNA transcribed, for example, as measured by RNA-blotting, PCR, or like methodology. PBLs are obtained using standard techniques, e.g. Mishell et al., eds., *Selected Methods in Cellular Immunology* (Freeman, New York, 1980).

When polypeptides of the present invention are expressed in soluble form, for example as a secreted product of transformed yeast or mammalian cells, they can be purified according to standard procedures of the art, including steps of ammonium sulfate precipitation, ion exchange chromatography, gel filtration, electrophoresis, affinity chromatography, and/or the like; e.g. *Enzyme Purification and Related Techniques, Methods in Enzymology*, 22:233-577 (1977), and *Scopes, Protein Purification: Principles and Practice* (Springer-Verlag, New York, 1982) provide guidance in such purifications. Likewise, when polypeptides of the invention are expressed in insoluble form, for example as aggregates, inclusion bodies or the like, they can be purified by standard procedures in the art, including separating the inclusion bodies from disrupted host cells by centrifugation, solubilizing the inclusion bodies with chaotropic agents and with reducing agents, diluting the solubilized mixture, and lowering the concentration of chaotropic agent and reducing agent so that the polypeptide takes on a biologically active conformation. These last procedures are disclosed in the following references, which are incorporated by reference: Winkler et al., *Biochemistry*, 25: 4041-4045 (1986); Winkler et al., *Biotechnology*, 3: 992-998 (1985); Kothe et al., U.S. patent 4,569,790; and European patent applications 86306917.5 and 86306353.3.

As used herein, "effective amount" means an amount sufficient to ameliorate a symptom of a disease condition mediated by excessive IFN- γ . The effective amount for a particular patient may vary depending on such factors as the state of the disease condition being treated, the overall health of the patient, the method of administration, the severity of side-effects, and the like. Generally, BCRF1 is administered as a pharmaceutical composition comprising an effective amount of BCRF1 and a pharmaceutical carrier or excipient. A pharmaceutical carrier can be any compatible, non-toxic substance suitable for delivering the compositions of the invention to a patient. Generally, compositions useful for parenteral administration of such drugs are well known, e.g. *Remington's Pharmaceutical Science*, 15th Ed. (Mack Publishing Company, Easton, PA 1980). Alternatively, compositions of the invention may be introduced into a patient's body by implantable drug delivery system: e.g. Urquhart et al., *Ann. Rev. Pharmacol. Toxicol.*, Vol. 24, pgs. 199-236 (1984); Lewis, ed. *Controlled Release of Pesticides and Pharmaceuticals* (Plenum Press, New York, 1981); U.S. patent 3,773,919; U.S. patent 3,270,960; and the like.

When administered parenterally, the BCRF1 is formulated in a unit dosage injectable form (e.g., a solution, suspension, or emulsion) in association with a pharmaceutical carrier. Such carriers are inherently nontoxic and nontherapeutic. Examples of such carriers are normal saline, Ringer's solution, dextrose solution, and Hank's solution. Nonaqueous carriers such as fixed oils and ethyl oleate may also be used. A preferred carrier is 5% dextrose/saline. The carrier may contain minor amounts of additives such as substances that enhance isotonicity and chemical stability, e.g., buffers and preservatives. BCRF1 is

preferably formulated in purified form substantially free of aggregates and other proteins at a concentration in the range of about 5 to 20 $\mu\text{g/ml}$. Preferably, BCRF1 is administered by continuous infusion so that an amount in the range of about 50-800 μg is delivered per day (i.e. about 1-16 $\mu\text{g/kg/day}$). The daily infusion rate may be varied based on monitoring of side effects, blood cell counts, and the like.

EXAMPLES

Example 1. Expression of BCRF1 in COS 7 Monkey cells

A gene encoding the open reading frame for BCRF1 was amplified by polymerase chain reaction using primers that allowed later insertion of the amplified fragment into an *EcoRI*-digested pcD(SR α) vector (Figure 1). The coding strand of the inserted fragment is shown in SEQ ID NO. 2.

Clones carrying the insert in the proper orientation were identified by expression of BCRF1 and/or the electrophoretic pattern of restriction digests. One such vector carrying the BCRF1 gene was designated pBCRF1(SR α) and was deposited with the ATCC under accession number 68193. pBCRF1(SR α) was amplified in *E. coli* MC1061, isolated by standard techniques, and used to transfect COS 7 monkey cells as follows: One day prior to transfection, approximately 1.5×10^6 COS 7 monkey cells were seeded onto individual 100 mm plates in Dulbecco's modified Eagle medium (DME) containing 5% fetal calf serum (FCS) and 2 mM glutamine. To perform the transfection, COS 7 cells were removed from the dishes by incubation with trypsin, washed twice in serum-free DME, and suspended to 10^7 cells/ml in serum-free DME. A 0.75 ml aliquot was mixed with 20 μg DNA and transferred to a sterile 0.4 cm electroporation cuvette. After 10 minutes, the cells were pulsed at 200 volts, 960 μF in a BioRad Gene Pulser unit. After another 10 minutes, the cells were removed from the cuvette and added to 20 ml of DME containing 5% FCS, 2mM glutamine, penicillin, streptomycin, and gentamycin. The mixture was aliquoted to four 100 mm tissue culture dishes. After 12-24 hours at 37°C, 5% CO₂, the medium was replaced with similar medium containing only 1% FCS and the incubation continued for an additional 72 hours at 37°C, 5% CO₂, after which the medium was collected and assayed for its ability to inhibit IFN- γ synthesis.

10 ml aliquots of freshly isolated PBLs (about 2×10^6 cells/ml) were incubated at 37°C with PHA (100 ng/ml) in medium consisting of (i) 90% DME supplemented with 5% FCS and 2 mM glutamine, and (ii) 10% supernatant from COS 7 cells previously transfected with pBCRF1(SR α). After 24 hours the cells and supernatants were harvested to assay for the presence of either IFN- γ mRNA or IFN- γ protein, respectively. Controls were treated identically, except that the 10% supernatant was from COS 7 cultures previously transfected with a plasmid carrying an unrelated cDNA insert. The BCRF1-treated samples exhibited about a 50% inhibition of IFN- γ synthesis relative to the controls.

Example 2. Expression of BCRF1 in *Escherichia coli*

A gene encoding a mature BCRF1 of the sequence given in SEQ ID NO. 3 may be expressed in *E. coli*:

The cDNA insert of pBCRF1(SR α) is recloned into an M13 plasmid where it is altered twice by site-directed mutagenesis: first to form a *ClaI* site at the 5'-end of the coding region for the mature BCRF1 polypeptide, and second to form a *BamHI* site at the 3'-end of the coding region for the mature BCRF1 polypeptide. The mutated sequence is then readily inserted into the TRPC11 expression vector described below.

The TRPC11 vector was constructed by ligating a synthetic consensus RBS fragment to *ClaI* linkers (ATGCAT) and by cloning the resulting fragments into *ClaI*-restricted pMT11hc (which had been previously modified to contain the *ClaI* site). pMT11hc is a small (2.3 kilobase) high copy, AMP^R, TET^S derivative of pBR322 that bears the πVX plasmid *EcoRI*-*HindIII* polylinker region. (πVX is described by Maniatis et al., Molecular Cloning: A Laboratory Manual, Cold Spring Harbor Laboratory, 1982.) This was modified to contain the *ClaI* site by restricting pMT11hc with *EcoRI* and *BamHI*, filling in the resulting 5-sticky ends and ligating with *ClaI* linker (CATCGATG), thereby restoring the *EcoRI* and *BamHI* sites and replacing the *SmaI* site with a *ClaI* site. One transformant from the TRPC11 construction had a tandem RBS sequence flanked by *ClaI* sites. One of the *ClaI* sites and part of the second copy of the RBS sequence were removed by digesting this plasmid with *PstI*, treating with *BaI*31 nuclease, restricting with *EcoRI* and treating with T4 DNA polymerase in the presence of all four deoxynucleotide triphosphates. The resulting 30-40 bp fragments were recovered by PAGE and cloned into *SmaI*-restricted pUC12. A 248 bp *E. coli* trpP-bearing *EcoRI* fragment derived from pKC101 (described by Nichols et al. in *Methods in Enzymology*, Vol. 101, pg. 155 (Academic Press, N.Y. 1983)) was then cloned into the *EcoRI* site to complete the TRPC11

construction, which is illustrated in Figure 2. TRPC11 is employed as a vector for BCRF1 by first digesting it with *Cla*I and *Bam*HI, purifying it, and then mixing it in a standard ligation solution with the *Cla*I-*Bam*HI fragment of the M13 containing the nucleotide sequence coding for the mature BCRF1. The insert-containing TRPC11, referred to as TRPC11-BCRF1, is propagated in *E. coli* K12 strain JM101, e.g.

5 available from the ATCC under accession number 33876.

Applicants have deposited *E. coli* MC1061 carrying pBCRF1(SR α) with the American Type Culture Collection, Rockville, MD, USA (ATCC), under accession number 68193. This deposit was made on December 20th 1989 under conditions as provided under ATCC's agreement for Culture Deposit for Patent Purposes, which assures that the deposit will be made available to the US Commissioner of Patents and

10 Trademarks pursuant to 35 USC 122 and 37 CFR 1.14, and will be made available to the public upon issue of a U.S. patent, which requires that the deposit be maintained. Availability of the deposited strain is not to be construed as a license to practice the invention in contravention of the rights granted under the authority of any government in accordance with its patent laws.

The Deposit has been modified to conform to the requirements of the Budapest Treaty on the Deposit

15 of Microorganisms.

The descriptions of the foregoing embodiments of the invention have been presented for purpose of illustration and description. They are not intended to be exhaustive or to limit the invention to the precise forms disclosed, and obviously many modifications and variations are possible in light of the above teaching. The embodiments were chosen and described in order to best explain the principles of the

20 invention to thereby enable others skilled in the art to best utilize the invention in various embodiments and with various modifications as are suited to the particular use contemplated. It is intended that the scope of the invention be defined by the claims appended hereto.

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SEQUENCE LISTING

5 SEQ ID NO: 1
 SEQUENCE TYPE: Amino acid
 SEQUENCE LENGTH: 170 amino acid residues
 STRANDEDNESS: single
 10 TOPOLOGY: linear
 MOLECULE TYPE: protein/polypeptide
 ORIGINAL SOURCE ORGANISM: Epstein-Barr virus
 15 PROPERTIES: BCRF-1

	Met	Glu	Arg	Arg	Leu	Val	Val	Thr	Leu	Gln	Cys	Leu	Val	Leu	Leu	
					5					10					15	
20	Tyr	Leu	Ala	Pro	Glu	Cys	Gly	Gly	Thr	Asp	Gln	Cys	Asp	Asn	Phe	
					20					25					30	
	Pro	Gln	Met	Leu	Arg	Asp	Leu	Arg	Asp	Ala	Phe	Ser	Arg	Val	Lys	
					35					40					45	
25	Thr	Phe	Phe	Gln	Thr	Lys	Asp	Glu	Val	Asp	Asn	Leu	Leu	Leu	Lys	
					50					55					60	
	Glu	Ser	Leu	Leu	Glu	Asp	Phe	Lys	Gly	Tyr	Leu	Gly	Cys	Gln	Ala	
					65					70					75	
30	Leu	Ser	Glu	Met	Ile	Gln	Phe	Tyr	Leu	Glu	Glu	Val	Met	Pro	Gln	
					80					85					90	
	Ala	Glu	Asn	Gln	Asp	Pro	Glu	Ala	Lys	Asp	His	Val	Asn	Ser	Leu	
					95					100					105	
35	Gly	Glu	Asn	Leu	Lys	Thr	Leu	Arg	Leu	Arg	Leu	Arg	Arg	Cys	His	
					110					115					120	
	Arg	Phe	Leu	Pro	Cys	Glu	Asn	Lys	Ser	Lys	Ala	Val	Glu	Gln	Ile	
					125					130					135	
40	Lys	Asn	Ala	Phe	Asn	Lys	Leu	Gln	Glu	Lys	Gly	Ile	Tyr	Lys	Ala	
					140					145					150	
	Met	Ser	Glu	Phe	Asp	Ile	Phe	Ile	Asn	Tyr	Ile	Glu	Ala	Tyr	Met	
					155					160					165	
45	Thr	Ile	Lys	Ala	Arg											
					170											

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SEQ ID NO: 2

SEQUENCE TYPE: Nucleotide

SEQUENCE LENGTH: 498 bases

5 STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: DNA sequence

ORIGINAL SOURCE ORGANISM: Epstein-Barr virus

10 PROPERTIES: BCRF-1

AATTC ATG GAG CGA AGG TTA GTG GTC ACT CTG CAG TGC CTG GTG CTG 47
 CTT TAC CTG GCA CCT GAG TGT GGA GGT ACA GAC CAA TGT GAC AAT 92
 15 TTT CCC CAG ACC TAA GAG ATG CCT TCA GTC GTG TTA AAA CCT TTT 137
 TCC AGA CAA AGG ACG AGG TAG ATA ACC TTT TGC TCA AGG AGT CTC 182
 TGC TAG AGG ACT TTA AGG ATG CCA GGC CCT GTC AGA AAT GAT CCA 227
 ATT CTA CCT GGA GGA AGT CAT GCC ACA GGC TGA AAC CAG GAC CCT 272
 20 GAA GCC AAA GAC CAT GTC AAT TCT TTG GGT GAA AAT CTA AAG ACC 317
 CTA CGG CTC CGC CTG CGC AGG TGC CAC AGG TTC CTG CCG TGT GAG 362
 AAC AAG AGT AAA GCT GTG GAA CAG ATA AAA AAT GCC TTT AAC AAG 407
 CTG CAG GAA AAA GGA ATT TAC AAA GCC ATG AGT GAA TTT GAC ATT 452
 25 TTT ATT AAC TAC ATA GAA GCA TAC ATG ACA ATT AAA GCC AGG TGA G 498

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SEQ ID NO: 3

SEQUENCE TYPE: Amino acid

SEQUENCE LENGTH: 147 amino acid residues

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: protein/polypeptide

ORIGINAL SOURCE ORGANISM: Epstein-Barr virus

PROPERTIES: BCRF-1

15	Thr	Asp	Gln	Cys	Asp	Asn	Phe	Pro	Gln	Met	Leu	Arg	Asp	Leu	Arg	5	10	15
	Asp	Ala	Phe	Ser	Arg	Val	Lys	Thr	Phe	Phe	Gln	Thr	Lys	Asp	Glu	20	25	30
20	Val	Asp	Asn	Leu	Leu	Leu	Lys	Glu	Ser	Leu	Leu	Glu	Asp	Phe	Lys	35	40	45
	Gly	Tyr	Leu	Gly	Cys	Gln	Ala	Leu	Ser	Glu	Met	Ile	Gln	Phe	Tyr	50	55	60
25	Leu	Glu	Glu	Val	Met	Pro	Gln	Ala	Glu	Asn	Gln	Asp	Pro	Glu	Ala	65	70	75
	Lys	Asp	His	Val	Asn	Ser	Leu	Gly	Glu	Asn	Leu	Lys	Thr	Leu	Arg	80	85	90
30	Leu	Arg	Leu	Arg	Arg	Cys	His	Arg	Phe	Leu	Pro	Cys	Glu	Asn	Lys	95	100	105
	Ser	Lys	Ala	Val	Glu	Gln	Ile	Lys	Asn	Ala	Phe	Asn	Lys	Leu	Gln	110	115	120
35	Glu	Lys	Gly	Ile	Tyr	Lys	Ala	Met	Ser	Glu	Phe	Asp	Ile	Phe	Ile	125	130	135
	Asn	Tyr	Ile	Glu	Ala	Tyr	Met	Thr	Ile	Lys	Ala	Arg				140	145	

Claims

Claims for the following Contracting States : AT, BE, CH, DE, DK, FR, GB, IT, LU, NL, SE

- 45 1. An Epstein-Barr virus BCRF1 protein capable of inhibiting the synthesis of interferon- γ , defined by the amino acid sequence shown in SEQ ID NO.3.
2. A recombinant expression vector capable of expressing an Epstein-Barr virus BCRF1 protein in a host, wherein said BCRF1 protein is capable of inhibiting the synthesis of interferon- γ .
- 50 3. The expression vector of Claim 2 wherein said BCRF1 protein is defined by the amino acid sequence shown in SEQ ID NO.3.
4. The expression vector of Claim 2 or Claim 3 wherein said host is mammalian.
- 55 5. The expression vector of Claim 4 consisting of pBCRF1(SR α), deposited with the American Type Culture Collection, Rockville, MD, USA, under accession number ATCC 68193.



SEQ ID NO: 3

SEQUENCE TYPE: Amino acid

SEQUENCE LENGTH: 147 amino acid residues

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STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: protein/polypeptide

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ORIGINAL SOURCE ORGANISM: Epstein-Barr virus

PROPERTIES: BCRF-1

15	Thr	Asp	Gln	Cys	Asp	Asn	Phe	Pro	Gln	Met	Leu	Arg	Asp	Leu	Arg	5	10	15
	Asp	Ala	Phe	Ser	Arg	Val	Lys	Thr	Phe	Phe	Gln	Thr	Lys	Asp	Glu	20	25	30
20	Val	Asp	Asn	Leu	Leu	Leu	Lys	Glu	Ser	Leu	Leu	Glu	Asp	Phe	Lys	35	40	45
	Gly	Tyr	Leu	Gly	Cys	Gln	Ala	Leu	Ser	Glu	Met	Ile	Gln	Phe	Tyr	50	55	60
25	Leu	Glu	Glu	Val	Met	Pro	Gln	Ala	Glu	Asn	Gln	Asp	Pro	Glu	Ala	65	70	75
	Lys	Asp	His	Val	Asn	Ser	Leu	Gly	Glu	Asn	Leu	Lys	Thr	Leu	Arg	80	85	90
30	Leu	Arg	Leu	Arg	Arg	Cys	His	Arg	Phe	Leu	Pro	Cys	Glu	Asn	Lys	95	100	105
	Ser	Lys	Ala	Val	Glu	Gln	Ile	Lys	Asn	Ala	Phe	Asn	Lys	Leu	Gln	110	115	120
35	Glu	Lys	Gly	Ile	Tyr	Lys	Ala	Met	Ser	Glu	Phe	Asp	Ile	Phe	Ile	125	130	135
	Asn	Tyr	Ile	Glu	Ala	Tyr	Met	Thr	Ile	Lys	Ala	Arg				140	145	

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Claims

Claims for the following Contracting States : AT, BE, CH, DE, DK, FR, GB, IT, LU, NL, SE

- | | | |
|----|----|--|
| 45 | 1. | An Epstein-Barr virus BCRF1 protein capable of inhibiting the synthesis of interferon- γ , defined by the amino acid sequence shown in SEQ ID NO.3. |
| | 2. | A recombinant expression vector capable of expressing an Epstein-Barr virus BCRF1 protein in a host, wherein said BCRF1 protein is capable of inhibiting the synthesis of interferon- γ . |
| 50 | 3. | The expression vector of Claim 2 wherein said BCRF1 protein is defined by the amino acid sequence shown in SEQ ID NO.3. |
| | 4. | The expression vector of Claim 2 or Claim 3 wherein said host is mammalian. |
| 55 | 5. | The expression vector of Claim 4 consisting of pBCRF1(SR α), deposited with the American Type Culture Collection, Rockville, MD, USA, under accession number ATCC 68193. |



6. The expression vector of Claim 2 wherein said host is bacterial.
7. A pharmaceutical composition for treating a disease associated with excessive interferon- γ production, comprising an Epstein-Barr virus BCRF1 protein in association with a pharmaceutically acceptable carrier or excipient, wherein said BCRF1 protein is capable of inhibiting the synthesis of interferon- γ .
8. A pharmaceutical composition according to Claim 7 wherein said BCRF1 protein is selected from the group of mature proteins of the open reading frame defined by the amino acid sequence shown in SEQ ID NO.1.
9. A pharmaceutical composition according to Claim 8 wherein said mature protein is defined by the amino acid sequence shown in SEQ ID NO.3.
10. A method of manufacturing a pharmaceutical composition for treating a disease associated with excessive interferon- γ production, comprising the admixture of an Epstein-Barr virus BCRF1 protein with a pharmaceutically acceptable carrier or excipient, wherein said BCRF1 protein is capable of inhibiting the synthesis of interferon- γ .
11. The use of an Epstein-Barr virus BCRF1 protein for the manufacture of a medicament for the treatment of a disease associated with excessive interferon- γ production.
12. The use according to Claim 11 wherein said medicament is adapted for intravenous delivery of an amount of said BCRF1 protein in the range of about 1-16 $\mu\text{g/kg}$ body weight of said patient per day.

25 Claims for the following Contracting States : ES, GR

1. A method of producing an Epstein-Barr virus BCRF1 protein capable of inhibiting the synthesis of interferon- γ , the method comprising the steps of:
 constructing a vector comprising a nucleotide sequence coding for said protein, wherein the nucleotide sequence is capable of being expressed by a host containing the vector and wherein the nucleotide sequence is selected from the group encoding BCRF1 proteins;
 incorporating the vector into the host; and
 maintaining the host in a culture medium under conditions suitable for expression of the nucleotide sequence into said protein.
2. The process of Claim 2 wherein the protein is defined by the amino acid sequence shown in SEQ ID NO.3.
3. The process of Claim 2 wherein the nucleotide sequence has the DNA sequence shown in SEQ ID NO.2.
4. A recombinant expression vector capable of expressing an Epstein-Barr virus BCRF1 protein in a host, wherein said BCRF1 protein is capable of inhibiting the synthesis of interferon- γ .
5. The expression vector of Claim 4 wherein said BCRF1 protein is defined by the amino acid sequence shown in SEQ ID NO.3.
6. The expression vector of Claim 4 or Claim 5 wherein said host is mammalian.
7. The expression vector of Claim 6 consisting of pBCRF1(SR α), deposited with the American Type Culture Collection, Rockville, MD, USA, under accession number ATCC 68193.
8. The expression vector of Claim 4 wherein said host is bacterial.
9. A method of manufacturing a pharmaceutical composition for treating a disease associated with excessive interferon- γ production, comprising the admixture of an Epstein-Barr virus BCRF1 protein with a pharmaceutically acceptable carrier or excipient, wherein said BCRF1 protein is capable of inhibiting the synthesis of interferon- γ .



10. A method according to Claim 9 wherein said BCRF1 protein is selected from the group of mature proteins of the open reading frame defined by the amino acid sequence shown in SEQ ID NO.1.
- 5 11. A method according to Claim 10 wherein said mature protein is defined by the amino acid sequence shown in SEQ ID NO.3.
12. The use of an Epstein-Barr virus BCRF1 protein for the manufacture of a medicament for the treatment of a disease associated with excessive interferon- γ production.
- 10 13. The use according to Claim 12 wherein said medicament is adapted for intravenous delivery of an amount of said BCRF1 protein in the range of about 1-16 μ g/kg body weight of said patient per day.

Patentansprüche

Patentansprüche für folgende Vertragsstaaten : AT, BE, CH, DE, DK, FR, GB, IT, LU, NL, SE

- 15 1. Zum Hemmen der Synthese von Interferon- γ befähigtes Epstein-Barr-Virus-BCRF1-Protein, das durch die in SEQ ID Nr. 3 dargestellte Aminosäuresequenz definiert ist.
- 20 2. Zum Expressieren eines Epstein-Barr-Virus-BCRF1-Proteins in einem Wirt befähigter, rekombinanter Expressionsvektor, wobei das BCRF1-Protein zum Hemmen der Synthese von Interferon- γ befähigt ist.
3. Expressionsvektor von Anspruch 2, wobei das BCRF1-Protein durch die in SEQ ID Nr. 3 dargestellte Aminosäuresequenz definiert ist.
- 25 4. Expressionsvektor von Anspruch 2 oder 3, wobei der Wirt ein Säuger ist.
5. Expressionsvektor von Anspruch 4, der aus pBCRF1(SR α) besteht, der bei der American Type Culture Collection, Rockville, MD, USA, unter der Zugangsnummer ATCC 68193 hinterlegt ist.
- 30 6. Expressionsvektor von Anspruch 2, wobei der Wirt ein Bakterium ist.
7. Pharmazeutische Zusammensetzung zum Behandeln einer Erkrankung, die mit überschüssiger Interferon- γ -Produktion verbunden ist, welche ein Epstein-Barr-Virus-BCRF1-Protein zusammen mit einem pharmazeutisch annehmbaren Träger oder Vehikel umfaßt, wobei das BCRF1-Protein zum Hemmen der Synthese von Interferon- γ befähigt ist.
- 35 8. Pharmazeutische Zusammensetzung gemäß Anspruch 7, wobei das BCRF1-Protein aus der Gruppe reifer Proteine mit dem offenen Leserahmen ausgewählt ist, der durch die in SEQ ID Nr. 1 dargestellte Aminosäuresequenz definiert ist.
- 40 9. Pharmazeutische Zusammensetzung gemäß Anspruch 8, wobei das reife Protein durch die in SEQ ID Nr. 3 dargestellte Aminosäuresequenz definiert ist.
- 45 10. Verfahren zu Herstellen einer pharmazeutischen Zusammensetzung zum Behandeln einer Erkrankung, die mit übermäßiger Interferon- γ -Produktion verbunden ist, welches das Vermischen eines Epstein-Barr-Virus-BCRF1-Proteins mit einem pharmazeutisch annehmbaren Träger oder Vehikel umfaßt, wobei das BCRF1-Protein zum Hemmen der Synthese von Interferon- γ befähigt ist.
- 50 11. Verwendung eines Epstein-Barr-Virus-BCRF1-Proteins zur Herstellung eines Arzneimittels zur Behandlung einer mit übermäßiger Interferon- γ -Produktion verbundenen Erkrankung.
12. Verwendung gemäß Anspruch 11, wobei das Arzneimittel an die intravenöse Abgabe einer Menge des BCRF1-Proteins im Bereich von etwa 1-16 μ g/kg Körpergewicht des Patienten je Tag angepaßt ist.

55 Patentansprüche für folgende Vertragsstaaten : ES, GR

1. Verfahren zum Herstellen eines zum Hemmen der Synthese von Interferon- γ befähigten Epstein-Barr-Virus-BCRF1-Proteins, wobei das Verfahren die Schritte des:



- Aufbauens eines Vektors, der eine für das Protein kodierende Nukleotidsequenz umfaßt, wobei die Nukleotidsequenz durch einen den Vektor enthaltenden Wirt exprimiert werden kann und wobei die Nukleotidsequenz aus der BCRF1-Proteine kodierenden Gruppe ausgewählt ist;
Einbringens des Vektors in den Wirt und
5 Erhaltens des Wirts in einem Kulturmedium unter zur Expression der Nukleotidsequenz in das Protein geeigneten Bedingungen umfaßt.
2. Verfahren von Anspruch 2, wobei das Protein durch die in SEQ ID Nr. 3 dargestellte Aminosäuresequenz definiert ist.
 - 10 3. Verfahren von Anspruch 2, wobei die Nukleotidsequenz die in SEQ ID Nr. 2 dargestellte DNA-Sequenz besitzt.
 4. Zum Exprimieren eines Epstein-Barr-Virus-BCRF1-Proteins in einem Wirt befähigter, rekombinanter Expressionsvektor, wobei das BCRF1-Protein zum Hemmen der Synthese von Interferon- γ befähigt ist.
 - 15 5. Expressionsvektor von Anspruch 4, wobei das BCRF1-Protein durch die in SEQ ID Nr. 3 dargestellte Aminosäuresequenz definiert ist.
 - 20 6. Expressionsvektor von Anspruch 4 oder 5, wobei der Wirt ein Säuger ist.
 7. Expressionsvektor von Anspruch 6, der aus PBCRF1(SR α) besteht, der bei der American Type Culture Collection, Rockville, MD, USA, unter der Zugangsnummer ATCC 68193 hinterlegt ist.
 - 25 8. Expressionsvektor von Anspruch 4, wobei der Wirt ein Bakterium ist.
 9. Verfahren zum Herstellen einer pharmazeutischen Zusammensetzung zum Behandeln einer Erkrankung, die mit überschüssiger Interferon- γ -Produktion verbunden ist, welches das Vermischen ein Epstein-Barr-Virus-BCRF1-Proteins zusammen mit einem pharmazeutisch annehmbaren Träger oder
30 Vehikel umfaßt, wobei das BCRF1-Protein zum Hemmen der Synthese von Interferon- γ befähigt ist.
 10. Verfahren gemäß Anspruch 9, wobei das BCRF1-Protein aus der Gruppe reifer Proteine mit dem offenen Leserahmen ausgewählt ist, der durch die in SEQ ID Nr. 1 dargestellte Aminosäuresequenz definiert ist.
 - 35 11. Verfahren gemäß Anspruch 10, wobei das reife Protein durch die in SEQ ID Nr. 3 dargestellte Aminosäuresequenz definiert ist.
 12. Verwendung eines Epstein-Barr-Virus-BCRF1-Proteins zur Herstellung eines Arzneimittels zur Behandlung einer mit übermäßiger Interferon- γ -Produktion verbundenen Erkrankung.
 - 40 13. Verwendung gemäß Anspruch 12, wobei das Arzneimittel an die intravenöse Abgabe einer Menge des BCRF1-Proteins im Bereich von etwa 1-16 μ g/kg Körpergewicht des Patienten je Tag angepaßt ist.
 - 45 **Revendications**
Revendications pour les Etats contractants suivants : AT, BE, CH, DE, DK, FR, GB, IT, LU, NL, SE
 1. Protéine BCRF1 du virus de Epstein-Barr capable d'inhiber la synthèse de l'interféron- γ , définie par la séquence d'acides aminés montrée dans SEQ ID N° 3.
 - 50 2. Vecteur d'expression recombinant capable d'exprimer une protéine BCRF1 du virus de Epstein-Barr chez un hôte, où ladite protéine BCRF1 est capable d'inhiber la synthèse de l'interféron- γ .
 3. Vecteur d'expression de la revendication 2, où ladite protéine BCRF1 est définie par la séquence d'acides aminés montrée dans SEQ ID N° 3.
 - 55 4. Vecteur d'expression de la revendication 2 ou la revendication 3 où ledit hôte est mammifère.



5. Vecteur d'expression de la revendication 4 consistant en pBCRF1 (SR α), déposé au American Type Culture Collection, Rockville, MD, USA, sous le numéro d'accèsion ATCC 68193.
6. Vecteur d'expression de la revendication 2, où ledit hôte est bactérien.
7. Composition pharmaceutique pour le traitement d'une maladie associée à une production excessive de l'interféron- γ , comprenant la protéine BCRF1 du virus de Epstein-Barr en association avec un support ou excipient acceptable en pharmacie, où ladite protéine BCRF1 est capable d'inhiber la synthèse de l'interféron- γ .
8. Composition pharmaceutique selon la revendication 7, où ladite protéine BCRF1 est choisie dans le groupe consistant en protéines mûres du cadre de lecture ouvert défini par la séquence d'acides aminés montrée à SEQ ID N° 1.
9. Composition pharmaceutique selon la revendication 8, où ladite protéine mûre est définie par la séquence d'acides aminés montrée à SEQ ID N° 3.
10. Méthode de fabrication d'une composition pharmaceutique pour le traitement d'une maladie associée à une production excessive de l'interféron- γ , comprenant le mélange d'une protéine BCRF1 du virus d'Epstein-Barr avec un support ou excipient acceptable en pharmacie, où ladite protéine BCRF1 peut inhiber la synthèse de l'interféron- γ .
11. Utilisation d'une protéine BCRF1 du virus de Epstein-Barr pour la fabrication d'un médicament pour le traitement d'une maladie associée à une production excessive de l'interféron- γ .
12. Utilisation selon la revendication 11, où ledit médicament est adapté à une administration intraveineuse d'une quantité de ladite protéine BCRF1 comprise entre environ 1-16 μ g/kg de poids corporel dudit patient par jour.
- 30 **Revendications pour les Etats contractants suivants : ES, GR**
 1. Méthode de production d'une protéine BCRF1 du virus de Epstein-Barr capable d'inhiber la synthèse de l'interféron- γ , la méthode comprenant les étapes de :
construire un vecteur comprenant une séquence de nucléotide codant pour ladite protéine, où la
séquence de nucléotides peut être exprimée par un hôte Contenant le vecteur et où la séquence de
nucléotides est choisie dans le groupe codant les protéines BCRF1;
incorporer le vecteur dans l'hôte; et
maintenir l'hôte dans un milieu de culture dans des conditions appropriées à l'expression de la
séquence de nucléotides dans ladite protéine.
 2. Procédé de la revendication 1, où la protéine est définie par la séquence d'acides aminés montrée dans SEQ ID N° 3.
 3. Procédé de la revendication 2, où la séquence de nucléotides a la séquence d'ADN montrée dans SEQ ID N° 2.
 4. Vecteur d'expression recombinant capable d'exprimer une protéine BCRF1 du virus de Epstein-Barr chez un hôte, où ladite protéine BCRF1 est capable d'inhiber la synthèse de l'interféron- γ .
 5. Vecteur d'expression de la revendication 4, où ladite protéine BCRF1 est définie par la séquence d'acides aminés montrée dans SEQ ID N° 3.
 6. Vecteur d'expression de la revendication 4 ou la revendication 5 où ledit hôte est mammifère.
 7. Vecteur d'expression de la revendication 6 consistant en pBCRF1 (SR α), déposé au American Type Culture Collection, Rockville, MD, USA, sous le numéro d'accèsion ATCC 68193.
 8. Vecteur d'expression de la revendication 4, où ledit hôte est bactérien.



9. Méthode de fabrication d'une composition pharmaceutique pour le traitement d'une maladie associée à une production excessive de l'interféron- γ , comprenant le mélange d'une protéine BCRF1 du virus d'Epstein-Barr avec un support ou excipient acceptables en pharmacie, où ladite protéine BCRF1 peut inhiber la synthèse de l'interféron- γ .
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10. Méthode selon la revendication 9, où ladite protéine BCRF1 est choisie dans le groupe des protéines mûres du cadre de lecture ouvert défini par la séquence d'acides aminés montrée par SEQ ID N°1.
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11. Méthode selon la revendication 10, où ladite protéine mûre est définie par la séquence d'acides aminés montrée dans SEQ ID N°3.
12. Utilisation d'une protéine BCRF1 du virus de Epstein-Barr pour la fabrication d'un médicament pour le traitement d'une maladie associée à une production excessive de l'interféron- γ .
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13. Utilisation selon la revendication 11, où ledit médicament est adapté à une administration intraveineuse d'une quantité de ladite protéine BCRF1 comprise entre environ 1-16 μ g/kg de poids corporel dudit patient par jour.

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