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(54) **METHOD FOR REDUCING THE IMMUNOGENICITY OF ANTIBODY VARIABLE DOMAINS**

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(58) **Field of Search** **530/387.1, 387.3; 435/69.6, 70.21; 424/130.1, 133.1**

(56) **References Cited**

U.S. PATENT DOCUMENTS

4,816,567 A 3/1989 Cabilly et al.

FOREIGN PATENT DOCUMENTS

EP	239400	9/1987
EP	255694	2/1988
EP	274394	7/1988
EP	328404	6/1989
EP	323806	7/1989
EP	327000	8/1989
EP	332424	9/1989
EP	434257	6/1991
EP	438312	7/1991
GB	2 188 638	10/1987
WO	WO 87/02671	5/1987
WO	WO 89/00999	2/1989
WO	WO 91/08298	6/1991
WO	WO 91/09967	7/1991
WO	WO 91/10742	7/1991
WO	WO 91/11198	8/1991
WO	WO 91/11534	8/1991

OTHER PUBLICATIONS

Novotny et al., PNAS 83:226-30, 1986.*

Hunter, W.M. et al., Preparation of Iodine-131 Labelled

English, D. et al., Single Separation of Red Blood Cells: Granulocytes and Mononuclear Leukocytes on Discontinuous Density Gradients of Ficoll-Hypaque, 1974, J. Immunological Methods, 5, pp. 249-252 (cumulative).

Bernstein, F.C. et al., The Protein Data Bank: A Computer-based Archival File for Macromolecular Structures, 1977, J. Mol. Biol., 112, pp. 535-542 (cumulative).

Rodbard, D. et al., Improved Curve-Fitting, Parallelism Testing, Characterization Of Sensitivity And Specificity, Validation, And Optimization For Radioligand Assays, 1978, In: Radioimmunoassay & Related Procedures In Medicine, International Atomic Agency, Vienna, vol. 1, pp. 469-504 (cumulative).

Chirgwin, J. et al., Isolation of Biologically Active Ribonucleic Acid from Sources Enriched in Ribonuclease, 1979, Biochemistry, 18, pp. 5294-5299 (cumulative).

Flanagan, J.G. et al., Arrangement of human immunoglobulin heavy chain constant region genes implies evolutionary duplication of a segment containing γ , ϵ and α genes, 1982, Nature, 300, pp. 709-813 (cumulative).

Daugherty, B.L et al., Polymerase chain reaction facilitates the cloning, CDR-grafting and rapid expression of a murine monoclonal antibody directed against the CD18 component of leukocyte integrins, 1991, Nucleic Acids Research, 19, pp. 2471-2476 (cumulative).

Beatty, P.G. et al., Definition of a Common Leukocyte Cell-Surface Antigen (Lo95-150) Associated with Diverse Cell-Mediated Immune Functions, 1983, J. Immunol., 131, pp. 2913-1918 (cumulative).

Sanchez-Madrid, F. et al., A Human Leukocyte Differentiation Antigen Family With Distinct α -Subunits and a Common B-Subunit, 1983, J. Exp. Med. 158, pp. 1785-1803 (cumulative).

VanVoorhis, W.C. et al., Specific Antimononuclear Phagocyte Monoclonal Antibodies, 1983, J. Exp. Med., 158, pp. 126-145 (cumulative).

Wright, S.D. et al., Identification of the C3bi receptor of human monocytes and macrophages by using monoclonal antibodies (cumulative).

Gritz, L et al., Plasmid-encoded hygromycin B resistance: the sequence of hygromycin B phosphotransferase gene and its expression in *Escherichia coli* and *Saccharomyces cerevisiae*, 1983, Gene, 25, pp. 179-188 (cumulative).

(List continued on next page.)

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(57) **ABSTRACT**

A unique method is disclosed for identifying and replacing immunoglobulin surface amino acid residues which converts the antigenicity of a first mammalian species to that of a second mammalian species. The method will simultaneously change immunogenicity and strictly preserve ligand binding properties. The judicious replacement of exterior amino acid residues has no effect on the ligand binding properties but greatly alters immunogenicity.

OTHER PUBLICATIONS

- Berkner, K.L. et al., Effect of the tripartite leader on synthesis of a non-viral protein in an adenovirus 5 recombinant, 1985, Nucleic Acids, Res., 13, pp. 841–857 (cumulative).
- Riechmann, L. et al., Reshaping human antibodies for therapy, 1988, Nature, 332, pp. 323–327 (cumulative).
- LoBuglio, A.F. et al., Mouse/human chimeric monoclonal antibody in man: Kinetics and immune response, 1989, Proc. Nat'l Acad. Sci. USA, 86, pp. 4220–4224 (cumulative).
- Sheriff, S., Antibody–Antigen Complexes1, 1990, Annu. Rev. Biochem., 59, pp. 439–473 (cumulative).
- Padan, E., On the Nature of Antibody Combining Sites: Unusual Structural Features That May Confer on These Sites an Enhanced Capacity for Binding Ligands, 1990, Proteins: Struc., Funct., Genet., 7, pp. 112–124 (cumulative).
- Hakimi, J. et al., Reduce Immunogenicity And Improved Pharmacokinetics Of Humanized Anti-Tac In Cynomolgus Monkeys, 1991, J. Immunol., 147, pp. 1352–1359 (cumulative).
- Lo, S.K. et al., Transient Adhesion of Neutrophils To Endothelium, J. Exp. Med., 169, pp. 1779–1793 (cumulative).
- Sequences of Proteins of Immunol. Int., 4th Edition, U.S. Dept. of Health and Human Services (cumulative).
- Gorman, S.D. et al., Reshaping a therapeutic CD4 antibody, 1991, Proc. Nat'l. Acad. Sci., 88, pp. 4181–4185 (cumulative).
- Padlan, E. et al., Modeling of Antibody Combing Sites, 1991, Methods in Enzymology, 203, pp. 30–21 (cumulative).
- Padlan, E., A Possible Procedure for Reducing The Immunogenicity of Antibody Variable Domains While Preserving Their Ligand-Binding Properties, 1991, Molec. Immun., 28(4/5), pp. 489–498 (cumulative).
- Verhoeven, M. et al., Reshaping Human Antibodies: Grafting an Antilysozyme Activity, 1988, Science, 239, pp. 1534–1536 (cumulative).
- Queen, C. et al., A humanized antibody that binds to the interleukin 2 receptor, 1989, PNAS, 86, pp. 10029–10033 (cumulative).
- Bolger, M.B., et al., Computer Modeling of Combining Site Structure of Anti-hapten Monoclonal Antibodies, 1991, Methods in Enzy., 203, pp. 21–45 (cumulative).
- Lewin, ‘When Does Homology Mean Something else’, Science 237: 1570 (1987).
- Reeck et al. ‘Homology in Proteins and Nucleic Acids . . . out of it’, Cell, vol. 50, p 667, 1987.
- Creighton Proteins Structure and Molecular Principals, WH Freeman & Co.NY pp 93–94, 1983.
- Creighton, ‘Experimental Studies of Protein Folding and Unfolding’, Biophys. Mole/Bio, vol. 33, pp 231–233, 1975.
- Burgess et al. ‘Possible Dissociation of the Heparin-binding . . . by Site directed Mutagenesis of a Single Lysine Residue’, J. of Cell Biol., vol. 111, pp 2129–2138, 1990.
- Lazar et al. Transforming Growth Factor alpha: Mutation . . . Different . . . Biological Activities’, Mol Cell Biol. vol. 8 B(3), pp 1247–1252, 1988.
- Schwartz et al. ‘A superactive insulin [B10–Aspartic acid] insulin(human)’, Proc. Natl. Acad. Sci. USA, Vo.1, 84, pp. 6408–6411, 1987.
- Lin et al. ‘Structure Function Relationships in Glucagon: . . . (homoserine lactone 27)-glucagon’, Biochemistry, vol. 14, pp 1559–1563 1975.
- Rudikoff et al. ‘Functional antibody lacing a variable region disulfide bridge’, Proc. Natl. acad. Sci, USA, vol. 79, p 1979, 1982.
- Panka et al. ‘Variable region framework differences . . . antibodies’, Proc. Natl. Acad. Sci, USA, vol. 85, pp 3080–3084, 1988.
- Amit et al. ‘Three Dimensional Structure of an Antigen–Antibody Complex at 2.8A Resolution’, Science, vol. 233, p 747,753, 1986.
- Ehrlich et al. PCT Technology, Macmillan Publishers, pp 80–83, 1989.
- Paul, Fundamntal Immunology p 242, Third Edition 1993.
- Hunter et al., “Preparation of Iodine-131 Labeled Human Growth Hormone of High Specific Activity” 1962, Nature vol. 194, pp. 495–6.
- English et al., “Single Step Separation of Red Blood Cells, Granulocytes and Mononuclear Leukocytes on Discontinuous Density Gradients of Ficoll–Hypaque” 1974 J. Immunological Methods 5, pp. 349–252.
- Bernstein et al., “The Protein Data Bank: A Computer-based Archival File for Macromolecular Structures” 1997 J Mol Biol, 112, pp. 535–542.

* cited by examiner

Position	Fractional Accessibility		J539 Residue Exposure	Residues In Subgroup			
	KOL			I	II	III	
	Residue	Exposure					
1	E	1.00 Ex	E	1.00 Ex	Q	E	
2	V	0.23 mB	V	0.37 mB	VQ	VM	
3	Q	0.82 Ex	K	0.82 Ex	TQ	Q	
4	L	0.00 Bu	L	0.10 Bu	L	L	
5	V	0.87 Ex	L	1.00 Ex	RQKT	VL	
6	Q	0.00 Bu	E	0.09 Bu	E	E	
7	S	0.94 Ex	S	0.94 Ex	SG	SGG	
8	G	1.00 Ex	G	1.00 Ex	SG	GG	
9	G	0.00 Bu	G	0.00 Bu	G	G	
10	G	1.00 Ex	G	1.00 Ex	AGT	GA	
11	V	0.90 Ex	L	0.81 Ex	VL	LF	
12	V	0.25 mB	V	0.25 mB	V	V	
13	Q	0.71 mE	Q	0.87 Ex	K	Q	
14	P	0.59 PB	P	0.64 mE	KP	QP	
15	G	1.00 Ex	G	1.00 Ex	TS	GG	
16	R	0.73 mE	G	1.00 Ex	EQ	G	
17	S	0.66 mE	S	0.75 mE	TA	G	
18	L	0.28 mB	L	0.26mB	TL	L	
19	R	0.66 mE	K	0.75 mE	TS	RK	
20	L	0.00 Bu	C	0.00 Bu	LT	L	
21	S	0.71 mE	S	0.82 Ex	TC	SC	
22	C	0.00 Bu	C	0.00 Bu	CT	A	
23	S	1.00 Ex	A	1.00 Ex	TV	A	
24	S	0.00 Bu	A	0.00 Bu	FF	AS	
25	S	0.87 Ex	S	1.00 Ex	SG	G	
26	G	1.00 Ex	G	1.00 Ex	G	G	
27	F	0.10 Bu	F	0.10 Bu	GYD	FLG	
28	I	0.85 Ex	D	0.72 mE	TS	TN	
29	F	0.00 Bu	F	0.00 Bu	LI	F	
30	S	0.74 mE	S	0.83 Ex	SVI	S	
36	W	0.00 Bu	W	0.00 Bu	SW	W	
37	V	0.00 Bu	V	0.00 Bu	IV	V	
38	R	0.10 Bu	R	0.31 mB	IR	R	
39	Q	0.15 Bu	Q	0.28 mB	IQ	Q	
40	A	0.95 Ex	A	0.75 mE	PP	A	
41	P	0.90 Ex	P	0.73 mE	PG	PS	
42	G	1.00 Ex	G	1.00 Ex	PG	G	
43	K	0.86 Ex	K	0.86 Ex	KR	K	
44	G	1.00 Ex	G	1.00 Ex	AG	GS	
45	L	0.00 Bu	L	0.00 Bu	L	L	

FIG. 1a

Position	Fractional Accessibility			Residues in Subgroup				
	KOL		J539	I	II	III		
	Residue	Exposure	Residue	Exposure				
46	E	0.75 mE	E	0.73 mE	E	E		E
47	W	0.10 Bu	W	0.04 Bu	W	W		W
48	V	0.00 Bu	I	0.00 Bu	MV	LI		V
49	A	0.00 Bu	G	0.00 Bu	G	AG		GSA
66	R	0.36 mB	K	0.51 pB	R	R		R
67	F	0.00 Bu	F	0.00 Bu	V	LV		F
68	T	0.87 Ex	I	0.88 Ex	T	T		T
69	I	0.00 Bu	I	0.00 Bu	VMI	IV		I
70	S	0.78 mE	S	0.79 mE	TS	ST		S
71	R	0.11 Bu	R	0.00 Bu	RLA	KV		R
72	N	0.61 mE	D	0.55 pB	DK	D		DN
73	D	0.44 pB	N	0.43 pB	PETAS	T		DN
74	S	0.85 Ex	A	0.97 Ex	S	S		S
75	K	0.88 Ex	K	0.77 mE	TF	KR		K
76	N	0.69 mE	N	0.68 mE	NST	N		N
77	T	0.41 pB	S	0.33 mB	TQ	Q		T
78	L	0.00 Bu	L	0.00 Bu	AV	VF		LA
79	F	0.45 pB	Y	0.35 mB	Y	VS		YF
80	L	0.00 Bu	L	0.00 Bu	M	L		L
81	Q	0.53 pB	Q	0.69 mE	E	TKSIN		Q
82	M	0.00 Bu	M	0.00 Bu	L	ML		M
82a	D	0.73 mE	S	0.58 pB	SVRT	TSNIR		ND
82b	S	0.98 Ex	K	0.96 Ex	S	NS		S
82c	L	0.00 Bu	V	0.00 Bu	L	VM		L
83	R	0.73 mE	R	0.83 Ex	RFI	DT		RE
84	P	0.75 mE	S	0.90 Ex	S	PA		PA
85	E	0.82 Ex	E	0.90 Ex	E	VA		ED
86	D	0.00 Bu	D	0.11 Bu	D	D		D
87	T	0.54 pB	T	0.47 pB	T	T		T
88	G	1.00 Ex	A	0.00 Bu	A	A		A
89	V	0.58 PB	L	0.63 mE	V	TV		VL
90	Y	0.00 Bu	Y	0.00 Bu	Y	Y		YY
91	F	0.00 Bu	Y	0.08 Bu	Y	Y		Y
92	C	0.00 Bu	C	0.00 Bu	C	C		C
93	A	0.00 Bu	A	0.00 Bu	A	A		AT
94	R	0.17 Bu	R	0.15 Bu	R	RH		RP
				JHI	JH2	JH3	JH4	JH5
103	W	0.09 Bu	W	0.07 Bu	W	W	W	W
104	G	0.00 Bu	G	1.00 Ex	G	G	G	G
				JH6				

FIG. 1b

Position	Fractional Accessibility			J539	I	Residues In Subgroup					
	KOL	Exposure	Residue			II	III	JH1	JH2	JH3	JH4
Residue	Exposure	Residue	Exposure								JH5 JH6
105	Q	0.93 Ex	Q	0.99 Ex	Q	R	Q	Q	Q	Q	Q
106	G	0.00 Bu	G	0.00 Bu	G	G	G	G	G	G	G
107	T	0.22 mB	T	0.26 mB	T	T	T	T	T	T	T
108	P	0.99 Ex	L	0.67 mE	L	L	M	L	L	L	T
109	V	0.00 Bu	V	0.00 Bu	V	V	V	V	V	V	V
110	T	0.76 mE	T	0.69 mE	T	T	T	T	T	T	T
111	V	0.00 Bu	V	0.00 Bu	V	V	V	V	V	V	V
112	S	0.98 Ex	S	0.74 mE	S	S	S	S	S	S	S
113	S	0.94 Ex	A	0.84 Ex	S	S	S	S	S	S	S

FIG. 1c

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