

## Nonneutralizing Antibodies Are Able To Inhibit Human Immunodeficiency Virus Type 1 Replication in Macrophages and Immature Dendritic Cells

Vincent Holl,<sup>1</sup> Maryse Peressin,<sup>1</sup> Thomas Decoville,<sup>1</sup> Sylvie Schmidt,<sup>1</sup> Susan Zolla-Pazner,<sup>2</sup> Anne-Marie Aubertin,<sup>1</sup> and Christiane Moog<sup>1\*</sup>

EA 3770, ULP, Institut de Virologie, 3 rue Kœberlé, 67000 Strasbourg, France,<sup>1</sup> and Veterans Affairs and NYU Medical Centers, 423 East 23rd Street, Room 18124N, New York, New York 10010<sup>2</sup>

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**Only five monoclonal antibodies (MAbs) neutralizing a broad range of primary isolates (PI) have been identified up to now. We have found that some MAbs with no neutralizing activities according to the “conventional” neutralization assay, involving phytohemagglutinin-stimulated peripheral blood mononuclear cells as targets, efficiently inhibit the replication of human immunodeficiency virus type 1 (HIV-1) PI in macrophages and immature dendritic cells (iDC). The mechanism of inhibition is distinct from the neutralization of infectivity occurring via Fab fragments and involves the interaction of the F portion with the FcγRs present on macrophages and iDC. We propose that, if such nonneutralizing inhibitory antibodies limit mucosal HIV transmission, they should be induced by vaccination.**

It is now generally recognized that neutralizing antibodies (NAbs) constitute one of the elements of the adaptive immune response that must be induced by an effective vaccine against human immunodeficiency virus (HIV). Complete protection of macaques from experimental challenge was obtained after a passive transfer of NAbs (4, 12, 15). However, only five broadly NAbs have been identified and attempts to induce strong and broad NAb responses by immunization have failed. Broadly NAbs are also infrequently detected in sera from HIV type 1-infected individuals (13). NAbs are identified *in vitro* using “conventional” neutralization assays with phytohemagglutinin (PHA)-stimulated peripheral blood mononuclear cells (PBMC) as targets. This assay is based on the evaluation of HIV replication in CD4<sup>+</sup> T lymphocytes. Nevertheless, macrophages and dendritic cells (DC) are also permissive for HIV. Macrophages are considered a major reservoir of HIV *in vivo*, and immature DC (iDC) are among the first cells infected by HIV after mucosal transmission (1, 9). Thus, antibodies induced by immunization should also prevent HIV infection of these target cells.

To study the ability of MAbs to inhibit viral growth in various cells, the activity of anti-HIV MAbs was analyzed *in vitro* using a neutralization assay that measures, by flow cytometry, the percentage of infected cells by detecting intracellular p24 in lymphocytes obtained from PHA-PBMC (6, 11) and in macrophages (6) or iDC (7) generated by the differentiation of CD14<sup>+</sup> cells. Briefly, antibodies (Abs) were incubated for 1 h with viruses at concentrations of 2 to 10 μg/ml of p24, depending on the stain and the target cell (to reach 2 to 5% cells infected after a single round of infection) and then the mixture was added to target cells. This technique has the advantage of

allowing the characterization of infected cells by phenotyping. For the five previously described broadly NAbs, we found higher inhibitory activity when iDC instead of PHA-stimulated PBMC were used as target cells and an even higher activity when macrophages were the HIV targets (Table 1). When two subtype B primary isolates (PI), HIV-1BaL (obtained from the NIH) and Bx08 (courtesy of H. Fleury), were tested with each of the NAbs, activities were increased by 16- to 12,000-fold when macrophages and iDC were used as target cells compared to activities measured with PBMC. With a subtype C PI TV1 (obtained from S. Engelbrecht), a virus described as relatively “resistant” to neutralization (10), no neutralization was noted with MAbs IgG1b12 or 447-52D, but with the other three MAbs, increases of 8- to 2,000-fold in activity were found with macrophages and iDC compared to those with PBMC. Contrary to the change in inhibitory concentrations noted with the MAbs, the 90% inhibitory concentrations (IC<sub>90</sub>) for soluble CD4 and the T20 fusion inhibitor were similar on the three different target cells (Table 1), confirming previously published data for T20 (8). These results suggest that the increased inhibitory activity detected in iDC and macrophages is not due to a lower density of CD4 or to a reduction in virus fusion.

Next, the mechanism of HIV inhibition was investigated. The iDC and macrophages express FcγRs on their surfaces. These receptors are involved in immune complex capture, endocytosis, and degradation. To define their roles in HIV inhibition, we analyzed the inhibitory activity of the Fab fragments of MAb 447-52D (provided by R. Stanfield) (Fig. 1). Contrary to whole immunoglobulin G (IgG), 447-52D Fabs exhibit similar inhibition of HIV-1BaL infection in PBMC, macrophages, and iDC. This result strongly suggests a role for the Fc portion of the IgG in the increased inhibitory activity observed with macrophages and iDC.

Given these observations, we extended the studies by determining the IC<sub>90</sub> of additional anti-HIV MAbs (Table 2). Of the 45 MAbs to various epitopes of gp120 and gp41 tested, only

\* Corresponding author. Mailing address: EA 3770, Université Louis Pasteur, Institut de Virologie, 3 rue Kœberlé, F-67000 Strasbourg, France. Phone: 333 90 24 37 42. Fax: 333 90 24 37 23. E-mail: c.moog@viro-ulp.u-strasbg.fr.

TABLE 1. Inhibition of HIV-1BaL, Bx08, and TV1 by neutralizing MAbs, sCD4, and T20 when PHA-stimulated PBMC, macrophages, or iDC were used as target cells

MAb name	Epitope or target	IC <sub>90</sub> (μg/ml) <sup>a</sup>								
		PHA-stimulated cells			iDC			Macrophages		
		BaL	Bx08	TV1	BaL	Bx08	TV1	BaL	Bx08	TV1
2F5	gp41 (aa 662–667)	30 <sup>b</sup>	40 <sup>b</sup>	50 <sup>b</sup>	1 <sup>d</sup>	2 <sup>c</sup>	6 <sup>c</sup>	0.0025 <sup>f</sup>	0.01	0.2 <sup>c</sup>
IgG1 b12	CD4 binding site	25 <sup>b</sup>	50 <sup>b</sup>	>100	1 <sup>d</sup>	2 <sup>c</sup>	>50	0.5 <sup>d</sup>	5 <sup>c</sup>	>40
447-52D	Crown of the V3 loop	50 <sup>b</sup>	50 <sup>b</sup>	>100	2 <sup>c</sup>	5 <sup>c</sup>	>50	0.5 <sup>d</sup>	0.2 <sup>c</sup>	>50
4E10	gp41 (aa 671–676)	50 <sup>b</sup>	60 <sup>b</sup>	100 <sup>b</sup>	5 <sup>c</sup>	3 <sup>c</sup>	1.5 <sup>d</sup>	0.06 <sup>e</sup>	0.2 <sup>c</sup>	<0.1 <sup>e</sup>
2G12	gp120 (carbohydrate)	20 <sup>b</sup>	100 <sup>b</sup>	>100	1 <sup>d</sup>	2 <sup>c</sup>	5 <sup>c</sup>	0.05 <sup>e</sup>	0.1 <sup>c</sup>	1 <sup>d</sup>
sCD4	gp120 protein	15 <sup>b</sup>	40 <sup>b</sup>	10 <sup>c</sup>	10 <sup>b</sup>	40 <sup>b</sup>	10 <sup>c</sup>	5 <sup>c</sup>	20 <sup>b</sup>	5 <sup>c</sup>
T20	gp41 fusion domain	1 <sup>d</sup>	1 <sup>d</sup>	1 <sup>d</sup>	0.5 <sup>d</sup>	2 <sup>c</sup>	1 <sup>d</sup>	0.5 <sup>d</sup>	1 <sup>d</sup>	1 <sup>d</sup>

<sup>a</sup> IC<sub>90</sub> correspond to the concentrations (μg/ml) of MAbs or recombinant proteins that lead to a 90% reduction in the percentage of infected cells.

<sup>b</sup> Range, 11 to 100 μg/ml.

<sup>c</sup> Range, 2 to 10 μg/ml.

<sup>d</sup> Range, 0.3 to 1.9 μg/ml.

<sup>e</sup> Range, 0.05 to 0.2 μg/ml.

<sup>f</sup> Range, 0.005 to 0.04 μg/ml.

two, MAbs F425B4e8 and 391-D directed against the V3 loop, were able to inhibit HIV-1BaL infection of PHA-stimulated PBMC. In contrast, 13 of the 45 MAbs tested inhibit HIV-1BaL replication in macrophages. These 13 MAbs mainly recognize either a linear epitope on the V3 loop of gp120 or the principal immunodominant domain (PID) of gp41. In addition, one MAb that was active in the macrophage-based assay recognizes a conformational epitope on gp120 (anti-C5 MAb 221) and one is directed against amino acids (aa) 731 to 752 of gp41 (MAb 1577) (Table 2). Most of these MAbs also inhibit the replication of Bx08, and the three MAbs directed against the PID inhibit the “neutralization-resistant” strain TV1 (Table 3). These latter results with MAbs to the PID are particularly notable since MAbs to this region have not previously been shown to exhibit potent neutralizing activity (3, 5, 17). These results indicate that some MAbs exhibiting low or no inhibitory activity on T lymphocytes are able to successfully inhibit the replication of three PI in macrophages. Moreover, six of nine

MAbs tested were able to diminish HIV-1BaL replication in iDC (Table 3).

To determine whether the Fc portions of these IgGs bind to the FcγRs on the surface of macrophages and iDC, FcγRs were blocked by incubation for 30 min with 10 μg/ml of anti-FcγRI, anti-FcγRII, or anti-FcγRIII (BD-Pharmingen, San Diego, CA). The anti-FcγR Abs were added to cells before the addition of virus and MAb mixtures using conditions similar to those for neutralization assay. The inhibitory activity of MAb 240 on Bx08 replication was reversed after blockage of FcγRI in macrophages (Fig. 2A) and FcγRII in iDC (data not shown), demonstrating that binding of the Fc portion of IgG MAb 240-D to FcγR participates in HIV inhibition in macrophages and iDC. Moreover, the inhibitory activity of MAb 240-D targeting the PID was abolished in macrophages (Fig. 2B) and iDC (not shown) when 25 μg/ml of peptide corresponding to the gp41 PID (aa 593 to 616, ARP7022; obtained through the NIBSC) was added to the Ab-Bx08 mixture 1 h prior to incu-

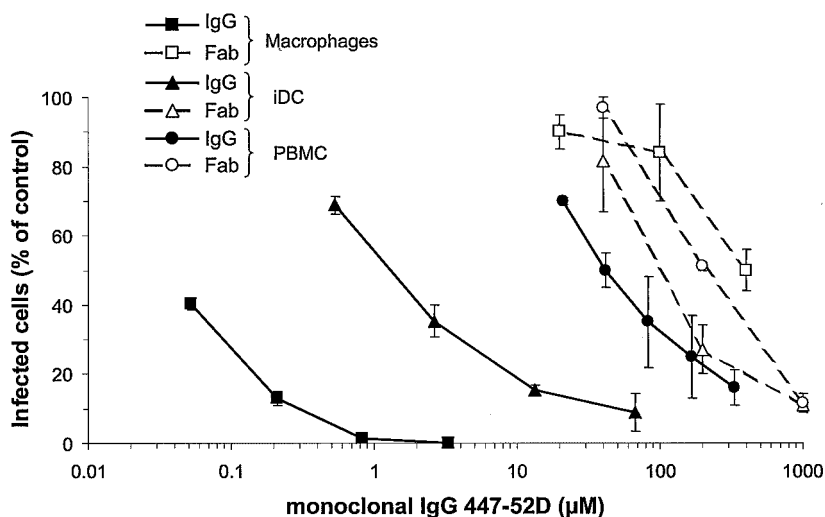


FIG. 1. Inhibition of HIV replication by MAb 447-52D (whole IgG) or its corresponding Fab fragment in PHA-stimulated PBMC, macrophages, and iDC. Values correspond to the percentage of infected cells in the presence of different dilutions of 447-52D versus infected cells without Ab (control infected cells). Mean and standard deviation of two independent wells obtained for one representative experiment are shown.

TABLE 2. Inhibition of HIV-1BaL replication in PHA-stimulated PBMC and macrophages by MAbs directed against various HIV epitopes<sup>a</sup>

Protein	Reference no.	MAb name	Ig	Epitope	IC <sub>90</sub> (μg/ml or dilution) <sup>b</sup>	
					PBMC	Macrophages
gp120	ARP 301	221	IgG1	gp160/gp120 (aa 482–495) C term of gp120	>50	20 <sup>c</sup>
	NIH 857	F105	IgG1k	Conformational gp120	>100	>50
	NIH 7369	654-30D	IgG1I	Tertiary gp120	ND	Ni
	ARP 3119	CA13	IgG1	Cross reactive to env	ND	Ni
	ARP 390	ICR39.13	IgG2b	Conformational (gp120)	Ni	Ni
	ARP 3041	11/68b	IgG1	gp120 (V1, V2+C4)	Ni	Ni
V3 loop	ARP 3036	8/64b	IgM	V3 (aa 300–315)	ND	Ni
	ARP 3038	10/540.w	IgG1	V3 (aa 311–321)	Ni	Ni
	ARP 3039	8/38	IgG2a	V3 (aa 300–315)	ND	Ni
	EVA 331	178.1.1	IgG2ak	V3 (to KSIRI sequence)	Ni	Ni
	EVA 3047	IIIB-V3-13	IgG1	V3 (IRIQRGPGRAFTIGC sequence)	>50	<b>3<sup>c</sup></b>
	ARP 3023	257-D IV	IgG1λ	V3 (KRIHI sequence)	>9.4	<b>0.05<sup>c</sup></b>
	ARP 3024	268-D IV	IgG1λ	V3 (HIGPGR sequence)	>14	<b>1<sup>c</sup></b>
	EVA 3056	MN215	IgG1	V3 KS/GIHIGPGKAFYTTGEI sequence)	>125	<b>10<sup>c</sup></b>
	NIH 7625	F425B4a1	IgG1λ	V3	>33	<b>0.4<sup>c</sup></b>
		391-D	IgG1k	V3	<b>20<sup>c</sup></b>	<b>1<sup>c</sup></b>
		NIH 7626	F425B4e8	IgG1k	Base of V3 loop	<b>25<sup>c</sup></b>
	NIH 2534	4G10	IgG1	V3 (RIQRGPGRAFVTGK)	ND	Ni
V2	ARP 324	CRA3	IgG2a	Conformational (V2 and C1)	Ni	Ni
	ARP 325	CRA4	IgG1	Conformational (V2)	Ni	Ni
	ARP 3075	62c	IgG1	Conformational (V2)	Ni	Ni
	ARP 3218	697D	IgG1λ	Conformational (V2, region 164–194)	>43	>20
C2, C4, C5		847-30	IgG1λ	C2	ND	>50
	ARP 388	ICR38	IgG2b	C4 (aa 427–436)	Ni	Ni
		858-D	IgG3λ	C5 (aa 495–516)	>100	>50
		1331A	IgG3λ	C5 (aa 495–516)	>100	>50
		450-D	IgG1λ	C5 (aa 503–509)	>100	>50
		722-D	IgG1k	C5 (aa 503–509)	ND	>50
	ARP 3221	670D	IgG1λ	C5 (aa 503–509)	>100	>50
CD4b5	EVA 3055	GP68	IgG1	CD4 binding site	Ni	Ni
	ARP 3220	654D	IgG1λ	CD4 binding site (discontinuous)	>25	>20
		570-D	IgG1λ	CD4 binding site	>100	>50
		654-D	IgG1λ	CD4 binding site	>100	>50
	ARP 3078	1.7B	IgG1	CD4 induced	>50	>50
	ARP 3079	4.8D	IgG1	CD4 induced	>50	>50
gp41	NIH 6882	5F3	IgG1λ	gp41 (526–543)	ND	>100
		181-D	IgG1k	gp141 (I) <sup>d</sup>	>100	>50
		240-D	IgG1k	gp41 (aa 579–604) (I)	>35	<b>1.8<sup>c</sup></b>
		246-D	IgG1k	gp41 (aa 579–604) (I)	>100	<b>0.8<sup>c</sup></b>
		50-69	IgG1λ	gp41 (aa 579–613 conformational) (I)	>100	<b>0.4<sup>c</sup></b>
	NIH 7623	F240	IgG1k	gp41 (aa 592–606)	>100	<b>0.5<sup>c</sup></b>
		98-6D	IgG1k	gp41 (aa 644–663) (II) <sup>e</sup>	>100	>50
		126-6	IgG1k	gp41 (aa 644–663) (II)	>100	>50
		167-D	IgG1λ	gp41 (aa 644–663) (II)	>100	>50
	NIH 1172	1577	IgG3	gp41 (731–752)	>100	<b>25<sup>c</sup></b>

<sup>a</sup> MAbs, selected because they either bind to native env or exhibit a restricted neutralizing activity, were obtained through the NIBSC, the NIH, or our laboratory (SZP).

<sup>b</sup> Values correspond to the ID<sub>90</sub> (μg/ml). Ni, no inhibitory activity detected at the dilution half of the supernatant provided; ND, not done.

<sup>c</sup> Boldface type indicates detection of inhibitory activity.

<sup>d</sup> Immunodominant domain of gp41 (amino acids 598 to 604, cluster I).

<sup>e</sup> Immunodominant domain of gp41 (amino acids 644 to 663, cluster II).

bation with target cells, whereas the addition of rgp160ΔPID (gp160 of Bx08 deleted from PID; kind gift from R. ElHabib, Aventis Pasteur) at 12.5 μg/ml had no effect. This competition experiment shows that the binding of the Fab portion of the MAb is necessary for HIV inhibition.

We have demonstrated that for macrophages and iDC, the inhibition of HIV replication by Abs can occur by two distinct

mechanisms, the first consisting of the neutralization of infectivity (which involves only the Fab part of the IgG) and the second depending on IgG-FcγR interaction, probably leading to endocytosis and the degradation of opsonized HIV particles. Abs that act only via the second mechanism in macrophages and iDC could be referred as nonneutralizing inhibitory Abs (NNiAbs) to distinguish them from NABs displaying both

TABLE 3. Inhibitory activity of MAbs against HIV-1BaL, Bx08, and TV1 replication in macrophages

Protein	MAb name	Epitope (sequence or domain)	IC <sub>90</sub> (μg/ml) <sup>a</sup>					
			PBMC		iDC	Macrophages		
			BaL	Bx08	BaL	BaL	Bx08	TV1
gp120	221	gp160/gp120 (aa 482–495)	>50	>100	ND	20 <sup>b</sup>	50 <sup>b</sup>	>100
V3 loop	IIIB-V3-13	V3 (IRIQRGPGRAFTIGC)	>50	>100	ND	3	ND	ND
	257-D IV	V3 (KRIHI sequence)	>9.4	ND	10 <sup>c</sup>	0.05 <sup>e</sup>	0.3 <sup>d</sup>	>9
	268-D IV	V3 (HIGPGR sequence)	>14	>14	15 <sup>b</sup>	1 <sup>d</sup>	0.5 <sup>d</sup>	>15
	MN215	V3 (KS/GIHIGPGKAFYTTGEI)	>125	>125	>50	10 <sup>c</sup>	40 <sup>b</sup>	20 <sup>b</sup>
	391-D	V3	20 <sup>b</sup>	50 <sup>b</sup>	ND	1 <sup>d</sup>	0.8 <sup>d</sup>	20 <sup>b</sup>
	F425B4a1	V3	>33	ND	ND	0.4 <sup>d</sup>	ND	ND
	F425B4e8	Base of V3 loop	25 <sup>b</sup>	100 <sup>b</sup>	4 <sup>c</sup>	1 <sup>d</sup>	5 <sup>c</sup>	>30
gp41	240-D	gp41 (aa 579–604) (I) <sup>f</sup>	>35	>35	20 <sup>b</sup>	1.8 <sup>d</sup>	1.9 <sup>d</sup>	0.5 <sup>d</sup>
	246-D	gp41 (aa 579–604) (I)	>100	>100	45 <sup>b</sup>	0.8 <sup>d</sup>	0.5 <sup>d</sup>	0.06 <sup>e</sup>
	50-69	gp41 (aa 579–613 conformational)	>100	>100	>50	0.4 <sup>d</sup>	0.8 <sup>d</sup>	0.6 <sup>d</sup>
	F240	gp41 (aa 592–606)	>100	>100	50 <sup>b</sup>	0.5 <sup>d</sup>	0.6 <sup>d</sup>	15 <sup>b</sup>
	1577	gp41 (aa 731–752)	>100	>100	>100	25 <sup>b</sup>	100 <sup>b</sup>	>100

<sup>a</sup> Values correspond to the ID<sub>90</sub> (μg/ml) determined when PHA-stimulated PBMC, macrophages, or iDC were used as target cells. ND, not done.

<sup>b</sup> Range, 11 to 100 μg/ml.

<sup>c</sup> Range, 2 to 10 μg/ml.

<sup>d</sup> Range, 0.3 to 1.9 μg/ml.

<sup>e</sup> Range, 0.05 to 0.2 μg/ml.

<sup>f</sup> Immunodominant domain of gp 41 (cluster I).

mechanism of inhibition and neutralizing PBMC infection. The NNiAbs are directed against epitopes distinct from those recognized by NABs and will not impair virus entry into cells. These Abs may not necessarily recognize functional envelope spikes, but they will link infectious virus particles to the target cell by efficient binding of the Fc region of IgG to FcγR and by binding of Fab regions to the HIV envelope. Binding of Ab to HIV is, however, apparently not sufficient for the inhibition of virus replication, as anti-C5 MAb 670D, for example, which binds to virus particles (16), was not able to inhibit HIV replication in macrophages (Table 2). The inhibition of HIV by Ab may also depend on the affinity of the Fc part of the Ab for FcγR and on an occupancy threshold below which MAb binds but does not inhibit. These parameters may also be involved in

virus inhibition. We showed that NNiAbs were mainly directed against the V3 loop and the PID. These two regions are accessible and highly immunogenic domains of HIV. Indeed, Abs directed to V3 loop and PID are detected in sera from HIV-infected individuals (2, 14) and the inhibition of HIV replication in macrophages (6) and iDC (7) has been evidenced with such sera.

We propose that NNiAbs that are not detected by using conventional assays using PBMC as target cells could participate in vivo in the protection of mucosal HIV transmission by preventing the infection of macrophages and iDC. These results, demonstrating new categories of protective Abs, may open new perspectives in the development and design of vaccines.

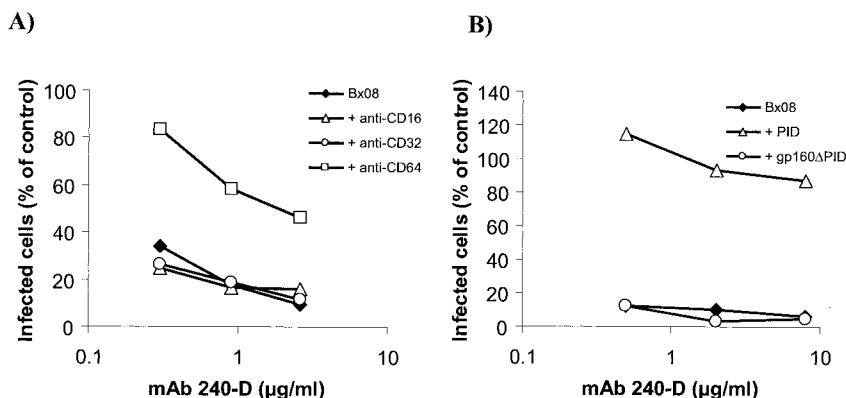


FIG. 2. Involvement of MAb 240-D paratope and Fc domains in its inhibitory activity of HIV replication in macrophages. (A) FcγRs were blocked by the addition of 10 μg of anti-FcγRI (CD64), anti-FcγRII (CD32), or anti-FcγRIII (CD16) to macrophages for 30 min before the addition of the virus-MAB 240 mixture using the conditions of the neutralization assay. (B) Competition experiments were performed by mixing, in conditions of neutralization assays, MAb 240-D and Bx08 with peptide PID (gp41, aa 593 to 616) at 25 μg/ml or rgp160ΔPID at 12.5 μg/ml. Then, 1 h later, the mixture was added to macrophages. The percentages of infected cells were determined by flow cytometry.

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