Comparative Antirhinoviral Activities of Soluble Intercellular Adhesion Molecule-1 (sICAM-1) and Chimeric ICAM-1/Immunoglobulin A Molecule

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We conducted a comparative study of the antirhinovirus activities of soluble intercellular adhesion molecule-1 (sICAM-1) and a chimeric ICAM-1/immunoglobulin A (IgA) molecule (IC1-5D/IgA) for nine major receptor group human rhinovirus (HRV) serotypes and for a variant of HRV-39 relatively resistant to inhibition by sICAM-1. IC1-5D/IgA inhibited the infectivity of eight of the nine wild-type HRVs and the resistant HRV-39 variant and was 60 to 170 times more potent than sICAM-1 on a molar basis. In contrast to sICAM-1, IC1-5D/IgA directly neutralized the infectivity of the representative HRVs by $\sim 1 \log_{10}$. These results expand on the antirhinovirus spectrum of IC1-5D/IgA, confirm that dimeric forms of sICAM-1 have a higher antirhinoviral potency than monomeric sICAM-1, and indicate that cross-linking of two adjacent receptor binding sites on the virus capsid by a divalent receptor enhances the direct inactivation of viral infectivity.

Intercellular adhesion molecule-1 (ICAM-1) has been identified as the cell receptor for the major group of human rhinoviruses (HRVs) (4, 9, 10). Major HRVs bind to the N-terminal immunoglobulin-like domain (D1) of ICAM-1 through residues located in the central portions of the floor of the canyon that surrounds the fivefold vertex of the virus capsid (8). Recombinant soluble forms of ICAM-1 (sICAM-1) have antirhinoviral activity in vitro, which is mediated through three characterized mechanisms of action: competition for the receptor-binding sites on the virus, hindrance of an early infection event such as entry or uncoating, and, to a substantially lesser extent, direct inactivation with formation of empty capsids (2, 5–7).

One such sICAM-1 (6) was shown to have antirhinoviral activity in vitro against 88 of the 90 numbered HRVs belonging to the major receptor group, with 50% effective inhibitory concentrations (EC₅₀) ranging from 0.1 to 41.1 μ g/ml in WI-38 human embryonic lung fibroblast cells (3). Chimeric immunoadhesin molecules have been constructed from either the two most distal domains of ICAM-1 coupled with the heavy chain of immunoglobulin A1 (IgA1), IgG, or IgM, or from all five extracellular domains of ICAM-1 coupled with the heavy chains of IgA1 or IgG (7). The most active of these molecules, a chimera composed of the five extracellular domains of ICAM-1 coupled with the heavy chain of IgA (IC1-5D/IgA), was shown by plaque reduction assay to inhibit the infectivity of HRV-3 approximately 200-fold more effectively on a molar basis than monomeric sICAM-1. IC1-5D/IgA also inhibited the binding of HRV-3 to HeLa cells approximately 7-fold more effectively than monomeric sICAM-1 and was approximately 12-fold more efficient than sICAM-1 in inducing formation of empty capsids of HRV-3 (7).

We have done a comparative study of the inhibitory activities of sICAM-1 (provided by Steven D. Marlin, Boheringer Ingelheim Pharmaceuticals, Ridgefield, Conn.) and the chimultiple HRV serotypes compared with that of the monomeric sICAM-1 is in keeping with the hypothesis that multivalent binding results in higher virus-receptor affinity (7). We have previously isolated a variant of HRV-39 moderately resistant to sICAM-1 by serial passages in HeLa cells in the presence of 100 μ g of sICAM-1 per ml, a concentration ~100 times the EC₅₀ for the wild-type virus (1). The EC₅₀ value for this variant was ~30-fold higher than the value for the wild-type virus, an increase that is within the ~400-fold range of values reported for the numbered major HRV serotypes (3). Direct and indirect evidence suggested that the sICAM-1 resistance was a preexisting phenotype selected from the pool of wild-type virus presumably because of an altered binding phenotype with reduced virus-receptor affinity (1). In

mera IC1-5D/IgA (provided by Timothy A. Springer, Harvard

Medical School, Boston, Mass.) for nine major HRV serotypes

(serotypes 3, 13, 14, 16, 23, 39, 68, 73, and 80) and for a variant

of HRV-39 selected for moderate resistance to sICAM-1 (1).

All of the wild-type major HRV serotypes were originally

obtained from the American Type Culture Collection. The

 EC_{50} s were determined by cytopathic effect inhibition assay in

WI-38 cells as described previously (2). The chimera IC1-5D/

IgA inhibited the infectivity of eight wild-type major HRV

serotypes tested with EC₅₀s between 0.02 and 0.48 μ g/ml

(equivalent to binding site concentrations between 0.2 and 4.8

nM) (Table 1). IC1-5D/IgA was more potent than monomeric

sICAM-1 by 50 to 143 times on a weight basis and by 60 to 170

times on a molar basis. These results are consistent with and

expand on the previously reported inhibition of one HRV

serotype by IC1-5D/IgA (7). Moreover, the more potent

activity of the divalent IC1-5D/IgA that we observed against

binding phenotype with reduced virus-receptor affinity (1). In the present study, the moderately resistant variant designated HRV-39/7p6 had a sICAM-1 EC₅₀ value 38-fold higher than that of the wild-type HRV-39 (Table 1). In contrast, the EC₅₀ value of IC1-5D/IgA was only modestly (approximately fivefold) increased for the sICAM-1-resistant variant compared with the wild-type HRV-39 (Table 1). This observation may suggest that the reduction in receptor-binding affinity of the resistant variant, which may result from a slight canyon conformational variation(s), can be partially compensated by the

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TABLE 1. Inhibitory effect of sICAM-1 and chimeric ICAM-1/IgA molecule (IC1-5D/IgA) on HRV cytopathic effect in human embryonic lung fibroblasts (WI-38 strain)

HRV serotype	$EC_{50} \ (\mu g/ml)^a$		Ratio of sICAM-1 EC ₅₀ to IC1-5D/IgA
	sICAM-1	IC1-5D/IgA	EC ₅₀ to ICI-5D/IgA EC ₅₀
3	5.7 ± 0.6	0.04 ± 0.01	143
13	20.6 ± 7.1	0.27 ± 0.12	76
14	6.8 ± 0.8	0.06 ± 0.01	113
16	1.2 ± 0.1	0.02 ± 0.01	60
23	>100 ^b	>32.0	
39	1.0 ± 0.3	0.02 ± 0.01	50
39/7p6 ^c	38.8 ± 22.6	0.10 ± 0.09	388
68	7.9 ± 2.3	0.07 ± 0.04	113
73	30.4 ± 4.9	0.48 ± 0.47	63
80	52.1 ± 27.1	0.42 ± 0.40	124

^{*a*} Values are means \pm standard deviations from two to three independent assays.

^b Result taken from previously published assays (3).

^c HRV-39 variant moderately resistant to inhibition by sICAM-1 (1).

higher virus-receptor affinity consequent to multivalency. Martin and colleagues postulated that virus escape from inhibition by multivalent immunoadhesins would be expected to occur at a lower frequency than that to monomeric soluble receptor (7). Our data indicate that the IC1-5D/IgA molecule retains a greater relative inhibitory effect for a virus selected for relative resistance to sICAM-1 and is consistent with this hypothesis.

In a previous study (3), we found that HRV serotypes 23 and 25, previously classified as belonging to the major receptor group (11), were not susceptible to inhibition by 100 μ g of sICAM-1 per ml (~1.2 μ M). In addition, receptor specificity studies using receptor blocking with an excess of anti-ICAM-1 monoclonal antibody in two cell lines suggested that those two serotypes used a cell receptor different from ICAM-1 (3). In the present study, IC1-5D/IgA also had no effect at the highest concentration tested on the infectivity of HRV-23 (Table 1). This suggests that the valency of the soluble receptor does not alter the susceptibility of HRV-23 and provides additional evidence that HRV-23 may use a different cell receptor.

We have previously shown by infectivity reduction assays that HRV-39 was not directly inactivated to a significant extent $(<0.5 \log_{10} \text{ reduction in infectivity})$ by incubation with monomeric sICAM-1 for up to 24 h (2). In the present study, we assessed the neutralizing effect of IC1-5D/IgA on the infectivity of two major HRV serotypes, HRV-39 and HRV-13, and the variant of HRV-39 resistant to sICAM-1, as described previously (2). Briefly, approximately 10⁶ 50% tissue culture infective doses of each virus were incubated in medium containing a concentration of sICAM-1 or IC1-5D/IgA equal to ~ 10 times the IC₅₀ of each molecule for the respective virus, or in plain medium, for 1 h at 33°C on a rocker platform. Each virus-drug or virus-medium mixture was then serially diluted in 10-fold dilutions, and the infectivity was determined on quadruplicate monolayers of WI-38 cells in 96-well plates. Guinea pig neutralizing antibody to HRV-39 (American Type Culture Collection) was used as a positive control for the HRV-39 assays. Confirming previous observations, there was no significant reduction of infectivity of HRV-39, HRV-13, or HRV-39/7p6 after incubation with sICAM-1 (Fig. 1). A reduction in infectivity of HRV-39 and HRV-13 of approximately 1.0 log₁₀ was observed after incubation with IC1-5D/IgA. This observation is in keeping with data previously published by Martin et al. (7), who found that IC1-5D/IgA was roughly 12 times more

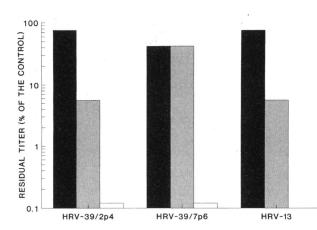


FIG. 1. Effect of incubation with sICAM-1 or IC1-5D/IgA on the infectivity of plaque purified wild-type HRV-39/2p4, HRV-39/7p6 (a variant of HRV-39 selected for resistance to sICAM-1), and pooled culture supernatant of HRV-13. Residual virus titers obtained after 1 h of incubation of 10⁶ 50% tissue culture infective doses with concentrations equal to 10 times the EC₅₀ value of each molecule for the respective virus are expressed as percentages of the titer of a control incubated with plain medium. Concentrations of sICAM-1 were 10 µg/ml for HRV-39/2p4, 390 µg/ml for HRV-39/7p6, and 210 µg/ml for HRV-39/2p4, 1 µg/ml for HRV-39/7p6, and 3 µg/ml for HRV-39. (2000) 2p4, 1 µg/ml for HRV-39/7p6, and 3 µg/ml for HRV-39. (2000) site are the average of two independent experiments. Symbols: solid bars, sICAM-1; hatched bars, IC1-5D/IgA; open bars, anti-HRV-39 guinea pig neutralizing antibody used as a positive control.

efficient that sICAM-1 in inducing conformational changes in the virus capsid. In contrast to wild-type HRV-39, no significant reduction in the infectivity of HRV-39/7p6 was observed after incubation with IC1-5D/IgA (Fig. 1). This result indicates that cross-linking adjacent receptor binding sites by means of multivalency of the soluble receptor does not affect its ability to directly inactivate a HRV-39 variant that binds the receptor with lower affinity.

The enhanced antirhinoviral potency of this multimeric molecule of sICAM-1 and its direct inactivation of some HRV serotypes make it an interesting one for further study and potential clinical development.

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