Expression of a foreign epitope on the surface of the adenovirus hexon

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To present short protein sequences to the host immune system a foreign epitope has been expressed on the surface of the adenovirus virion as part of the hexon. As the trimeric hexon constitutes 240 out of the 252 capsomers of the virus, the foreign epitope is repeated 720 times on the virion surface. An eight amino acid sequence from the major antigenic site in the VP1 capsid protein of poliovirus type 3 was engineered into two regions of the adenovirus type 2 hexon. The two loop regions chosen to accommodate the foreign sequences are exposed on the surface of the virion, show sequence variation between serotypes and are the sites of interaction with neutralizing antibodies. Virus with substitutions in loop I had wild-type growth characteristics, whereas virus with substitutions in loop II grew poorly. Adenoviruses with poliovirus sequences in loop I were recognized and efficiently neutralized by antisera specific for the poliovirus sequence; an antiserum raised against the adenovirus with the poliovirus insert specifically recognized the VP1 capsid protein of poliovirus type 3. It is therefore feasible to alter the surface properties of the adenovirus virion and in doing so to manipulate the immune response to this virus.

Introduction

Human adenoviruses, of which 47 distinct serotypes have been identified (Hierholzer et al., 1988), replicate in the mucosal surfaces of respiratory, ocular and gastrointestinal tissues giving rise to a variety of acute infections. For over 30 years protection against infection by these viruses has been achieved by oral immunization which results in the generation of strong mucosal immunity. Live adenovirus types 4 and 7 (Ad4 and -7) in enteric coated capsules have been administered to military personnel to prevent acute respiratory disease with no indication of adverse reactions. Oral immunization with serotypes 1, 2 and 5 has also proved effective in clinical trials (reviewed in Ginsberg, 1984).

Adenoviruses are non-enveloped and contain a linear double-stranded DNA genome of approximately 36 kb that is packaged within an icosahedral capsid. Hexon is the major structural component of the virus capsid, forming the 20 facets of the icosahedron; the vertices are composed of a complex of penton base and fibre (Ginsberg, 1984). Type-specific antigenic determinants of hexon have been demonstrated on the surface of the virion (Willcox & Mautner, 1976a, b) but the antigenic structure is complex with monoclonal antibodies (MAbs) defining at least 19 different epitopes on the surface of hexon (Russell et al., 1981; Adam et al., 1986). Sequence comparison of human adenovirus hexons and alignment with the three-dimensional structure of the adenovirus type 2 (Ad2) hexon (Roberts et al., 1986) indicates that although the base of the hexon is very highly conserved the surface loops are highly variable (Kinloch et al., 1984; Toogood & Hay, 1988; Toogood et al., 1989). The prediction that these variable regions could represent type-specific neutralizing determinants was confirmed by the demonstration that antibodies raised against synthetic peptides derived from the loop regions could neutralize virus in a type-specific fashion (Toogood et al., 1992).

The variability in the length and sequence of these exposed loops suggested that they could be manipulated to express foreign antigenic determinants. This approach has been successfully used to generate a range of polioviruses with altered antigenic properties. Although the initial antigenic hybrids contained sequences from other poliovirus serotypes a wide range of epitopes from other pathogens have now been incorporated into poliovirus antigenic chimeras (Burke et al., 1988, 1989; Colbere-Garapin et al., 1988; Dedieu et al., 1992; Evans et al., 1989; Jenkins et al., 1990; Kitson et al., 1991; Lemon et al., 1992; Martin et al., 1988; Minor et al., 1990, 1991; Murdin & Wimmer, 1989; Murdin et al., 1991a, b, 1992; Murray et al., 1988a, b; Reimann et al., 1991). Although the relatively small size and ease of manipulation of polioviruses make them useful models for this type of work, the increased genetic stability of DNA viruses and the expanding use of adenoviruses as gene delivery vehicles make the construction of adeno-
viruses with altered surface properties an attractive proposition.

The objective of this work was to construct adenoviruses expressing foreign antigenic determinants on the surface of the virion. A type-specific antigenic determinant from poliovirus type 3 was engineered into the surface loops of the Ad2 major surface protein, hexon. One of the chimeric adenoviruses grew well in tissue culture and was neutralized by antisera specific for the inserted poliovirus epitope. An antisera raised against the adenovirus with the poliovirus insert specifically recognized the VP1 capsid protein of poliovirus type 3.

**Methods**

**Cells and viruses.** HeLa cells were grown in suspension in Glasgow-modified spinner Eagles’ medium (GS-MEM, Gibco) with 7% newborn calf serum. Adenovirus virions were prepared from infected HeLa cells by fluorocarbon extraction of freeze-fractured cells, followed by density gradient centrifugation (Hay et al., 1984). Virus was stored at −20 °C in 50% glycerol. The titre of the virus stock was in the region of 10^11 to 10^12 p.f.u. per ml. Adenovirus type 5 (Ad5) ts 2 was obtained from Dr V. Mautner, MRC Virology Unit, Glasgow, U.K. and was propagated at 32 °C. Virus titres at the permissive and non-permissive temperatures were determined by plaque assay on A549 cells as described previously (Williams, 1970). 293 cells (Graham et al., 1977) were grown as monolayers in G-MEM containing 10% fetal calf serum (FCS). Vaccine strains of poliovirus types 1, 2 and 3 were obtained from Dr M. Ferguson, National Institute for Biological Standards and Control, Potters Bar, U.K. and were propagated and titrated in HeLa cells. Monolayer cultures of A549 cells, a human cell line derived from lung carcinoma, were grown in G-MEM supplemented with 10% FCS.

**Construction of recombinant viruses.** The Ad2 HindIII A fragment cloned in pAT153 (Mautner & Boursnell, 1983) was obtained from Dr V. Mautner. Plasmid DNA was cleaved with Smal which has two recognition sites in the plasmid at positions 51.8 and 56.0 in the hexon gene. Recombination with the Ad2 hexon gene containing the poliovirus sequences should rescue infectivity of the ts mutant at the non-permissive temperature. DNA from Ad5 ts 2 virions and the modified HindIII A plasmid were cotransfected into 293 cells at the permissive temperature of 32 °C as described (Hay et al., 1984) and virus was harvested after 5 days. The virus yield was titrated at 32 °C and 38.5 °C. Plaques arising at 38.5 °C were inoculated into 24-well plates and viral DNA was tested for the presence of poliovirus sequences by PCR analysis. Isolates Ad5 PLI and Ad5 PLII contain the poliovirus sequences within the regions encoding hexon loops I or II.

**Antisera.** Two female Dutch rabbits were immunized subcutaneously with 50 μg of purified, heat-inactivated Ad5 PLI virions in Freund’s complete adjuvant. Rabbits were boosted after 2 weeks with 50 μg virions in Freund’s incomplete adjuvant, a test bleed being taken 2 weeks later and tested by ELISA for specific antibody. After a second boost serum was collected from each rabbit. Anti-poliovirus peptide antisera R77, R78, R126, R166, R222 and R235 raised in rabbits were obtained in freeze-dried form from Dr M. Ferguson and once rehydrated were used at the indicated dilutions. MAb R2/76 directed against the Ad2 hexon was obtained as an ascitic fluid from W. C. Russell (St Andrews).

**Western blots.** Proteins (20 μg) were separated in a 10% SDS-polyacrylamide gel and transferred to nitrocellulose using a semi-dry blotter. Non-specific binding sites on the membrane were blocked with 10% non-fat milk, the membrane was exposed to the primary rabbit antibody (1:3000 dilution), washed and exposed to a 1:3000 dilution of affinity-purified donkey anti-rabbit IgG conjugated to horseradish peroxidase (HRP) (Amersham). The blot was developed using enhanced chemiluminescence (Amersham).

**ELISA.** The reactivity of specific anti-peptide antibodies with Ad2 or Ad5 PLI was measured by ELISA using an HRP detection system. Microtitre plates (96-well) were coated overnight at 4 °C with 0.5 μg per well of purified adenovirus virions in PBS. After washing with PBS, non-specific binding sites were blocked by incubation with PBS containing 10% low-fat milk (Marvel) and 0.1% sodium azide for 3 h at 37 °C. Doubling dilutions of the antisera were allowed to react with the immobilized virus for 1 h at room temperature. After washing, plates were incubated with HRP-conjugated goat anti-rabbit IgG or a similarly conjugated anti-mouse IgG for 1 h at 37 °C, and the colour was subsequently developed with ortho-phenylenediamine and hydrogen peroxide for 30 min in the dark. Microtitre plates were read in a Tisiter Multiskan. Specificity was assessed by substituting the immune serum with the corresponding pre-immune serum.

**Neutralization test.** Appropriate dilutions of virus were determined that would cause a visible c.p.e. in a microtitre well of A549 cells following 3 days (poliovirus) or 5 days (adenovirus) infection. Thus for neutralization tests 50 μl of a dilution of virus stock was mixed with doubling dilutions (starting at 1:10) of each antisera in serum-free medium. Viruses and sera were incubated for 1 h at 37 °C and absorbed onto A549 cells in a microtitre plate for a further 1 h at 37 °C. GS-MEM containing 2% FCS was added and incubation continued for either 3 or 5 days at 37 °C. Monolayers were fixed with formaldehyde and stained with crystal violet.

**Results**

**Introduction of a poliovirus epitope into the adenovirus hexon**

The objective of this work was to present foreign epitopes to the host immune system by their expression as part of the adenovirus major capsid protein, hexon.
The adenovirus capsid contains 252 capsomers of which 240 are hexon and 12 are penton. The surface of the adenovirus virion is therefore largely composed of residues derived from hexon and as the molecule is a homotrimer any epitope expressed as part of the hexon will be repeated 720 times on the surface of the virion. Regions of hexon into which foreign sequences would be introduced were selected on the basis of knowledge of the three-dimensional structure of the Ad2 hexon (Roberts et al., 1986), comparative sequence analysis of hexons from different adenoviruses (Kinloch et al., 1984; Toogood & Hay, 1988; Toogood et al., 1989), and the location of neutralizing epitopes on the surface of the Ad2 hexon (Toogood et al., 1992). Regions in loop I (amino acids 284 to 291) and loop II (amino acids 442 to 449) were chosen as the points at which a foreign epitope could be introduced into hexon (Fig. 1). To test the feasibility of this approach part of the well characterized immunodominant antigenic site in the VP1 capsid protein (amino acids 93 to 100) of poliovirus type 3, P3/Leon/USA/37 strain (Minor et al., 1983; Evans et al., 1983; Ferguson et al., 1985) was independently introduced into either loop I or II of the Ad2 hexon.

This was accomplished by site-directed mutagenesis of an M13 subclone which was reinserted into the Ad2 HindIII A fragment. To aid in the selection of recombinants this plasmid was cotransfected with DNA prepared from Ad5 ts 2 which contains a temperature-sensitive lesion in the hexon gene (Boursnell & Mautner, 1981) and the virus yield was plated at the non-permissive temperature. The recombinant viruses obtained contain the poliovirus sequence Glu-Gln-Pro-Thr-Thr-Arg-Ala-Gln substituted in loop I (PLI) or loop II (PLII) and four additional amino acid changes that arise from insertion of the flanking XbaI sites that create a cassette into which can be inserted any sequences that are to be expressed (Fig. 1). The location of these substitutions on the surface of the hexon molecule can be determined from the three-dimensional structure of the Ad2 hexon.
Fig. 2 shows the sites of substitution although given the flexibility of the loop region no attempt has been made to model the substituted amino acids into the overall structure. In both loops I and II the substituted amino acids are in a solvent-exposed position which would be predicted to be accessible to recognition by antibodies.

Growth characteristics of adenoviruses containing poliovirus epitopes

Viable viruses with substitutions in loop I (Ad5 PLI) were obtained with relative ease whereas viruses with substitutions in loop II (Ad5 PLII) were very difficult to isolate as they gave rise to very small plaques and grew very poorly. It was therefore decided to focus attention on the properties of viruses bearing substitutions in loop I. Wild-type Ad2 and Ad5 PLI were therefore compared in a one-step growth curve (Fig. 3) which demonstrates that the viruses replicate with similar kinetics and produce virus stocks of similar titre. Examination of the plaques produced by Ad5 PLI and wild-type Ad2 (Fig. 3) reinforces the conclusion that the Ad5 PLI virus has essentially wild-type growth characteristics. Loop I was therefore identified as the preferred site for introduction of foreign sequences into the adenovirus hexon.

Table 1. Recognition of Ad5 PLI by poliovirus type 3-specific antisera

<table>
<thead>
<tr>
<th>Antiserum</th>
<th>Virus on solid phase*</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Ad2</td>
</tr>
<tr>
<td>Anti-hexon</td>
<td>10000</td>
</tr>
<tr>
<td>MAb R2/76</td>
<td>5000</td>
</tr>
<tr>
<td>Anti-Ad5 PLI</td>
<td>&lt; 20</td>
</tr>
<tr>
<td>R77</td>
<td>&lt; 20</td>
</tr>
<tr>
<td>R78</td>
<td>40</td>
</tr>
<tr>
<td>R126</td>
<td>&lt; 20</td>
</tr>
<tr>
<td>R166</td>
<td>40</td>
</tr>
<tr>
<td>R222</td>
<td>40</td>
</tr>
<tr>
<td>R235</td>
<td>40</td>
</tr>
</tbody>
</table>

* The ELISA titre is given as the reciprocal of the lowest dilution of antiserum that was at least twofold above the background reading. Poliovirus-specific antisera were raised against poliovirus type 3 VP1 amino acids 89 to 104 (EVDNEQPTTRAQKLFA; Ferguson et al., 1985).

Recognition of Ad5 PLI by poliovirus-specific antisera

To determine whether the poliovirus sequences introduced into the hexon were presented on the surface of the virion, purified virus was used as the solid phase in an ELISA with a variety of antibodies. It should be noted at this point that although PLI is in an Ad5 background it is likely, although in this case not determined, that the hexon is derived entirely from Ad2. A MAb directed against the Ad2 hexon, MAb R2/76, gave the same titre against Ad2 and Ad5 PLI, indicating that similar amounts of Ad2 and Ad5 PLI virus had coated the plates. A similar conclusion was reached when an antiserum raised against purified Ad5 PLI (anti-Ad5 PLI) also gave the same titre for both Ad2 and Ad5 PLI. A series of antibodies raised against synthetic peptides encompassing the octapeptide sequence from poliovirus type 3 inserted into Ad5 PLI (Ferguson et al., 1985) all reacted strongly with Ad5 PLI but failed to react with Ad2 (Table 1). The reactivity of the antipeptide antibodies to Ad5 PLI indicates that the poliovirus type 3 sequences are displayed on the surface of the adenovirus virion in a form that is fully accessible to antibody.
Table 2. Neutralization of poliovirus and Ad5 PLI by poliovirus type 3-specific peptide antisera

<table>
<thead>
<tr>
<th>Antiserum</th>
<th>Polio 1</th>
<th>Polio 2</th>
<th>Polio 3</th>
<th>Ad2</th>
<th>Ad5 PLI</th>
</tr>
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<tbody>
<tr>
<td>R77</td>
<td>&lt; 10</td>
<td>&lt; 10</td>
<td>160</td>
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<td>&lt; 10</td>
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<td>160</td>
<td>&lt; 10</td>
<td>&gt; 20000</td>
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<tr>
<td>R126</td>
<td>&lt; 10</td>
<td>&lt; 10</td>
<td>320</td>
<td>&lt; 10</td>
<td>5000</td>
</tr>
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<td>R166</td>
<td>&lt; 10</td>
<td>&lt; 10</td>
<td>20</td>
<td>&lt; 10</td>
<td>5000</td>
</tr>
<tr>
<td>R222</td>
<td>&lt; 10</td>
<td>&lt; 10</td>
<td>10</td>
<td>&lt; 10</td>
<td>&gt; 20000</td>
</tr>
<tr>
<td>R235</td>
<td>&lt; 10</td>
<td>&lt; 10</td>
<td>40</td>
<td>&lt; 10</td>
<td>&gt; 20000</td>
</tr>
</tbody>
</table>

* Neutralization titres are expressed as the reciprocal of the lowest dilution of antiserum that blocks infection.

Neutralization of Ad5 PLI by poliovirus-specific antisera

The antisera raised against the poliovirus type 3 peptide were capable of neutralizing that virus (Ferguson et al., 1985). It was therefore of interest to determine whether they could neutralize the adenovirus containing the poliovirus epitope. These antisera were therefore used in neutralization tests against poliovirus types 1 to 3, Ad2 and Ad5 PLI. None of the peptide antisera have neutralizing activity against poliovirus type 1, poliovirus type 2 or Ad2 but they have, as expected, neutralizing activity against poliovirus type 3 (Table 2). In addition all of the peptide antisera efficiently neutralized Ad5 PLI (Table 2). Comparison of the neutralization titres of the antisera for poliovirus type 3 and Ad5 PLI reveals that in almost all cases the sera are over 100-fold more active against Ad5 PLI than they are against poliovirus type 3.

Recognition of poliovirus type 3 by an antiserum raised against Ad5 PLI

To determine whether the epitope expressed on the surface of the Ad5 PLI hexon was capable of provoking an immune response, heat-inactivated purified virus was injected into rabbits and serum collected. The sera reacted vigorously in ELISA and neutralized both Ad2 and Ad5 PLI but failed to react with poliovirus type 3 in either of these tests. To determine whether antibodies had been produced against the poliovirus epitope, purified Ad2 virions and extracts from cells either mock-infected or infected with poliovirus types 1, 2 or 3 were separated by SDS–PAGE and subjected to Western blot analysis using the antisera raised against Ad5 PLI. As expected the antisera detected the structural protein of the Ad2 virion. No specific reaction was detected against uninfected cells or cells infected with poliovirus types 1 or 2 but a species corresponding to VP1 was detected in the extract of poliovirus type 3-infected cells (Fig. 4).

Discussion

We have demonstrated that viable adenoviruses can be constructed that contain a foreign sequence substituted into the exposed surface loops of the major structural protein, hexon. The regions of hexon into which these substitutions were made show type-specific variation (Toogood et al., 1989) and are the sites at which neutralizing antibodies interact (Toogood et al., 1992). Analysis of these regions by X-ray diffraction indicated that in addition to being located on the surface of the molecule they were disordered, displaying a considerable amount of conformational flexibility (Roberts et al., 1986). Substitution into loop I produced viruses that had wild-type growth characteristics, and substitution into loop II produced viruses with impaired growth. Although these viruses were efficiently neutralized by antipeptide antisera against the poliovirus sequence, poliovirus type 3 was not neutralized by antisera raised against the adenovirus expressing the poliovirus epitope. This may be a result of the heat inactivation process which could have modified the immunogenicity of the poliovirus sequences substituted into the hexon. A number of additional explanations could also account for this observation. It is likely that the poliovirus sequence in the adenovirus hexon may be held in a quite different conformation from that in its native environment. That this is the case is suggested by the observation that the sera raised against the synthetic peptide neutralize the
recombinant adenovirus at a much greater dilution than is required to neutralize poliovirus type 3. Another possibility is that the adenovirus hexon contains an immunodominant epitope that masks any immune response to the inserted poliovirus epitope. It is also relevant that the vaccine strain of poliovirus type 3 used in these experiments is only poorly neutralized by the sera raised against the poliovirus peptide (Ferguson et al., 1985 and Table 2, this report) and that the eight amino acid sequence inserted into adenovirus represents only a part of the 16 amino acid sequence that was used to generate the peptide antisera. Given the flexibility of the exposed surface loops of the adenovirus hexon it should be possible to insert considerably larger sequences of foreign material into the loop region. Comparison of the sequences of the surface loops of hexon from adenovirus types 2 and 40 reveals that relative to type 40 the type 2 hexon accommodates a 32 residue insertion of mainly acidic amino acids (Toogood et al., 1989).

Although the initial construction of the plasmid containing the poliovirus sequences was rather cumbersome, further insertion of other sequences should be considerably simplified as the poliovirus sequences are flanked by XbaI restriction sites that are unique in the plasmid. Thus any sequences with XbaI compatible ends can be inserted into the cleaved plasmid thereby providing a route for the construction of adenoviruses with hexons containing defined inserts of choice. These sequences still have to be inserted into the viral genome by homologous recombination in vivo which is rather inefficient, but a number of strategies are now available for manipulation of the entire adenovirus genome in a plasmid background that would avoid this step (Graham, 1990). Thus the adenovirus system described here may represent a good general method for presenting small antigenic epitopes to the immune system. The ability of orally administered adenovirus to elicit a strong and protective immune response suggests that the chimeric viruses described here may have the ability to induce a mucosal immune response to the foreign epitopes presented on the hexon surface.

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