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Acute effect of infection by adipogenic human adenovirus Ad36

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Abstract

Human adenovirus Ad36 is causally and correlatively associated in animals and humans, respectively, with increased adiposity and altered metabolic profile. We inoculated rats with Ad36, UV-inactivated Ad36 or mock-infected. Four-days later, Ad36-infected rats showed 23% greater epididymal fat pad weight and viral mRNA, the viral DNA could also be detected in tissues viz. the liver, brain, and adipose tissue. Intranasal or intra-peritoneal routes of viral inoculations showed similar tissue affinity. Serum cytokine response was remarkably down regulated. Ad36 acutely suppresses systemic immune response and spreads widely. This information will help to determine Ad36 tissue tropism and its metabolic consequences.

Keywords

obesity; adiposity; adipose tissue; cytokine; immune response; MCP1; IL6; IL18; IL12

Although obesity is associated with several health risks[9], not all obese individuals are adversely affected by it[24]. To better utilize health care resources, it would help immensely to selectively target prevention and treatment efforts by identifying "at risk" individuals from those who are relatively protected from the adverse effects of obesity. Recent emerging evidence from animal models[18] and human studies[17] indicates that some forms of obesity may indeed carry relatively low health risk. We reported that human adenovirus Ad36 infection appears to be one such marker of "low risk" obesity. Experimental infection of animals with Ad36 induces obesity without increasing food intake, and induces relative-hypolipidemia[6-8] and better insulin sensitivity[20]. In-vitro, Ad36 induces adipogenic commitment, differentiation and lipid accumulation in adipocyte progenitors[21,22,25], and increases cellular glucose uptake[26].[23] These causal effects of Ad36 infection are mirrored in humans naturally infected with Ad36, who have greater body weight and body fat[1], but relative hypolipidemia[1], greater potential of preadipocyte differentiation[21], better glycemic control and lower inflammatory response of adipose tissue[23], compared to uninfected individuals. Thus, Ad36 induces obesity with metabolically favorable profile in animals and is associated with such obesity in humans. Strong association of Ad36 infection with human obesity and metabolic phenotype makes it an important candidate for evaluating its causative role in human obesity. Also, Ad36 proteins are promising novel candidates[22] as a tool to reveal regulatory controls which could be therapeutically manipulated to alleviate cardiovascular risk factors in humans. Despite the increasing data and significance of Ad36, the fundamental information about the acute spread of Ad36 in a host is unknown, which is critical for determining the hostvirus interaction in future.

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Ad36 belongs to serotype D of 51 human adenoviruses identified and is antigenically unique [27]. Ad36 was isolated from fecal sample of a girl suffering from enteritis[27] and shows about 15 - 17% prevalence in adult population in the United States[1]. About 30% of the obese, but only 11% of the non-obese adults are naturally infected with the virus[1]. While Ad36 DNA can be isolated from human adipose tissue[21], Ad36 tissue tropism or the host immune response to experimental infection in humans cannot be studied due to ethical considerations. Since Ad36 can successfully infect rats, we used this animal model to study acute host response to the viral infection as described in the following two experiments.

For experiment 1, twenty-two male Wistar rats (4-weeks old) were obtained from Harlan (Madison, WI). After one week of acclimatization to the new surroundings, rats were randomized into 3 weight-matched groups (control-N=8, Ad-36 inactivated-N=7 and Ad-36-N=7) and housed individually in micro-isolator cages under biosafety level-two containment. Rats were inoculated intranasally with media (mock), 8×10^5 particles of either UV inactivated Ad-36 or Ad-36, respectively. *Ad-libitum* access to water and rat chow was offered. Protective clothing, gloves, shoes, hairnets and masks were used to enter the rooms and care was taken to prevent cross contamination. Four days post inoculation, animals were sacrificed after an overnight fast. Trunk blood was collected and serum was separated. Lung, pancreas, heart, spleen, brain, stomach, epididymal fat, liver and spleen were carefully separated, and flash frozen in liquid nitrogen. Various techniques used and assays that were conducted are described below.

Second experiment determined the effect of a different route of infection on spread of Ad36, 4 wk old male Wistar rats were intraperitoneally (i.p.) inoculated with either 200 μ L cell culture media (N = 6) or 10⁷ PFU of Ad36 (N=6). Four days post inoculation, the animals were sacrificed and epididymal, visceral and retroperitoneal fat tissue, lung, liver spleen and skeletal muscle were harvested. DNA was isolated and a nested PCR was conducted as described below.

Ad-36 was obtained from American Type Culture Collection (catalog no. VR-913; American Type Culture Collection, Manassas, VA). The virus was plaque-purified as previously described[6].

Purified virus was UV irradiated with 250 mJ/cm² using a Stratalinker UV crosslinker (cat# 400079, Stratagene).

RNA was extracted using the RNeasy Mini Kit, as per the manufacturer's instructions (cat # 74104, Qiagen). Residual DNA was eliminated by using Amplification Grade Deoxyribonuclease I (cat # 18068-015, Invitrogen). One µg of total RNA was reverse-transcribed to cDNA using iScript[™] cDNA Synthesis Kit (cat # 170-8890, Bio-Rad) as perthe manufacturer's protocol. Samples were stored at -80°C untilused for Real Time PCR.

A standard was generated using DNA pooled from the experimental samples. At least 3 data points, representing 10-fold dilutions, were required for generating a standard curve. Non-template control reaction mixtures contained water instead of sample. Real-time quantitative PCR was carried out in optical 96 or 384 well reaction plates using ABI PRISM 7700 sequence detector (Applied Biosystems, Branchburg, NJ) using a SYBER Green detection system (cat # 170-8880, Bio-Rad). Both, samples and standards were run at least in duplicate, and each transcript level was adjusted to the housekeeping gene used (β -actin). The reactions were performed using the following parameters: one cycle of 48°C for 30 min, then 95°C for 10 min, followed by 40 cycles at 95°Cfor 15 s and 60°C for 1 min. For the detection of Ad-36 DNA the following primers were used: forward 5'-GGCATACTAACCCAGTCCGATG-3' and reverse 5'-TTGCCAGAATCCCACCCATAC-3'.

PCR core system II (cat # M7665, Promega) was used for the amplification of cDNA, obtained as described above. Water was used a negative PCR control. Positive PCR control was DNA from Ad36 infected A549 cells. DNA was denatured for 2 min at 95°C and subjected to 35 cycles of PCR (94°C for 1 min, 58°C for 1 min, 72°C for 2 min) followed by extension at 72° C for 5 min. PCR products were visualized on a 1.2 % agarose gel with a 100 bp DNA ladder (cat # G-2101, Promega). The following primers were used to detect Ad-36 E4Orf1: forward 5'-GGCATACTAACCCAGTCCGATG -3' and reverse 5'-

AATCACTCTCCCAGCAGCAGG -3'. The expected product size was 138 bp.

Total DNA was extracted from the tissue using DNeasy Tissue Mini Kit (cat # 69504, Qiagen). DNA samples were stored at -80° C until used for amplification. PCR core system II (cat # M7665, Promega) was used for the amplification of DNA. Negative PCR controls were water and DNA from uninfected rats. Positive PCR control was DNA from Ad36 infected A549 cells. DNA was denatured for 2 min at 95°C and subjected to 35 cycles of PCR (94°C for 1 min, 58° C for 1 min, 72°C for 2 min) followed by extension at 72°C for 5 min. PCR products were visualized on a 1.2 % agarose gel with a 100 bp DNA ladder (cat # G-2101, Promega).

The following primers from the fiber protein region were used.

Ad-36 Outer forward primer (5'-GTCTGGAAAACTGAGTGTGGATA),

Ad-36 Outer reverse primer (5'-ATCCAAAATCAAATGTAATAGAGT),

Ad-36 Inner forward primer (5'-TTAACTGGAAAAGGAATAGGTA),

Ad-36 Inner reverse primer (5'-GGTGTTGTTGGTTGGCTTAGGATA).

Expected product size: 650bp

A 14-plex assay using Luminex xMAP technology was used to determine serum cytokine levels (Millipore corporation, cat # RCYTO-80K-PMX). Serum samples from 5 randomly selected animals from each group were used. The lowest detection limit for all assays was 24.5pg/mL.

Upon termination of the experiments, we observed that the body weight of rats from all three groups remained almost identical 4 d post inoculation (Figure 1A), but interestingly, epididymal fat pad weight of the Ad36 infected group was 23% heavier (p=.01) compared to that of the uninfected control group (Figure 1B). As expected, UV-inactivated virus did not influence the fat pad weight.

Two experiments determined the spread of Ad36 to body tissues in rats inoculated via two separate routes. Samples from all harvested tissues showed the presence of viral DNA (Figures 2A–B), but not the uninfected control animals (a representative sample shown in Figure 2B). Epididymal fat pad showed Ad36 E4orf1 mRNA, indicative of viral replication process (Figure 2C).

The following cytokines were below the detection level (24.5pg/mL) for all animals in the group: Granulocyte-monocyte-colony-stimulating factor (GMCSF), IL1, IL4, IL18, IL10, IL12, IL5, TNF α and were excluded from the analysis. The rest of the cytokines were analyzed and if values were below the detection levels, they were considered as 24.5 pg/mL. Overall, not the UV-inactivated Ad36, but the wild type virus appears to reduce cytokine response 2d and 4d post inoculation (Figure 3). Although the reduction in cytokine response to Ad36 infected reached statistical significance for only macrophage chemoattractant protein 1 (MCP-1) and IL-18, perhaps due a limited number of animals per group (n = 5 per group), many of the cytokine levels were reduced by 50% to 95% by the virus.

Pasarica et al.

In humans, adenovirus infections are common and cause acute upper respiratory tract infections, enteritis or conjunctivitis[10]. Adenoviruses may persist asymptomatically in adult humans[13]. Although adenoviruses are not thought to integrate with host DNA, recent reports suggest this to be a possibility[28]. There are 6 major subgroups (A–F) among the 50 human adenoviruses. Each subgroup has a number of specific serotypes. Ad36 belongs to subgroup D, serotype 36. Ad36 is serologically different compared to other human adenoviruses [27]. Adenoviruses are non-enveloped DNA viruses that replicate in host cell nucleus

Several medium to long-term studies about the adipogenic effect of adenoviruses are published [5–8,20]. However, surprisingly little is known about the acute host response to these adipogenic adenoviruses, which is fundamental to understanding the role of these viruses in human obesity and for devising treatment or prevention strategies. While Ad36 is a strong candidate for determining its role in human obesity and is relatively most studied virus, sketchy information is available about it's target tissue and the immediate response of its host. Changes in food intake or activity do not explain the metabolic effects of Ad36 and the animals do not show overt signs of infection or premature mortality even up to 7 mo post infection [6-8,20]. Animals could be experimentally infected by intra nasal or intra peritoneal inoculation of Ad36. Also, blood-borne, air- or fecal-oral transmission of Ad36 was observed[6-8,20]. Ad36 DNA appears in the blood of experimentally infected chickens and their uninfected cagemates in about 12 h post inoculation and lasts in the blood for up to 4 wks[7] Also, Ad36 DNA could be detected in the blood and adipose tissue of animals 5 wks post infection[6], by capillary electrophoresis method. Ad36 infected marmosets excreted active virus in feces up to 2 months post infection and harbored viral DNA in adipose tissue up to 7 months[8], and in hamsters up to 4 wks post infection when the experiments were terminated.

In the present study, an increase in epididymal fat pad weight in just 4 d post Ad36 infection was surprising. No change in the fat pad weight of the UV-inactivated group further supports the role of active virus in increasing the adipose tissue. Also, mRNA expression of Ad36 E4orf1, the adipogenic gene of the virus[22] in these fat pads is particularly noteworthy. This is the first demonstration of Ad36 gene expression in adipose tissue of an experimentally infected animal host, suggesting active viral replication and possible adipogenic effect on the tissue.

As determined by the presence of viral DNA, intra-nasally or intra-peritoneally inoculated Ad36 spreads rapidly to several tissues, including some known for adenoviral tropism, such as the liver. Although the spread of the virus to the brain may be somewhat unconventional, it may explain the changes in neurotransmitter levels observed in Ad36 infected rats[20]. Spread of the virus to adipose tissue and skeletal muscle is consistent with our previous findings[6, 8] and important due to our recent in-vitro findings in these tissues[26] [21,23]. Given the effect of the virus on adipose tissue derived stem cell commitment and differentiation[21], spread of Ad36 to adipose tissue is particularly intriguing. About 27% of the Pima Indians screened harbored Ad36 DNA in their subcutaneous adipose tissue and showed greater adipogenic potential compared to their antibody negative counterparts[21]. Functional effect, if any, of Ad36 viral DNA presence on these tissues remain to be determined.

Many viruses subvert the host immune system for successful infection, spread or persistence and understanding these immune modulations may provide key information for defense against the microbe. Overall, Ad36 appears to down-regulate the host immune response in just 2 to 4 days post inoculation. In response to infection, IL-12 induces IFN γ and along with IL-18 host mounts a Th1 type of immune response. In early stages of infection, many viruses downregulate the Th1 response by suppressing the production of these cytokines [4]. Down regulation of this cytotoxic immune response may allow Ad36-infected cells to survive, which may be of particular relevance to cells from the adipose tissue, which expands in response to

Ad36 infection. A long-term and tissue-specific assessment of cytokine response to Ad36 is required to further understand the interaction.

Inflammatory cytokines are positively associated with obesity and insulin resistance [15,29] and approaches to transgenically knockout inflammatory cytokines such as MCP1, TNF α (Tumor necrosis factor alpha) show improvement in insulin resistance[14] and knockout of cytokines such as IL-18 or receptor of IL-1 increase adiposity[11,19]. In-vitro infection of human adipose tissue by Ad36 down regulates inflammatory cytokines (unpublished observations). Thus, the down regulation of inflammatory cytokines by Ad36 may contribute to its adipogenic effect and the paradoxical improvement in glycemic control. Long-term studies are required to test this hypothesis.

Although Ad36 appears to acutely down regulate response of the inflammatory cytokines investigated, other intact human adenoviruses or replication deficient adenoviral vectors are known to elicit a strong and acute inflammatory cytokine response[3,12], which in fact is a serious limitation of otherwise efficient application of human adenoviruses as vectors[2]. Extensive research has been undertaken to overcome this shortcoming of human adenoviral vector use[2]. One of the alternative strategies is to use an adenovirus that does not elicit strong inflammatory cytokine response. In response to such concerns, Iacobelli-Martinez et al [16] reported human adenoviruses Ad16, Ad35 and Ad37, which down regulate acute inflammatory response of cytokines such as IL-12, IFN γ and IL-6, and suggested the use of these adenoviruses for gene delivery and vaccine development as implications. A pronounced anti-inflammatory cytokine response by Ad36, coupled with its ability to infect several host tissues also suggests its possible role as vector.

In conclusion, acute effects of Ad36 in rats include greater fat pad weight, and attenuation of host immune response, which may help in the widespread tissue distribution observed. This study provides the basis to study tissue tropism and long-term modulation of host immune response and its metabolic consequences. The data also provide candidate host tissues for the virus in humans naturally infected with Ad36.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

Acknowledgments

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Pasarica et al.



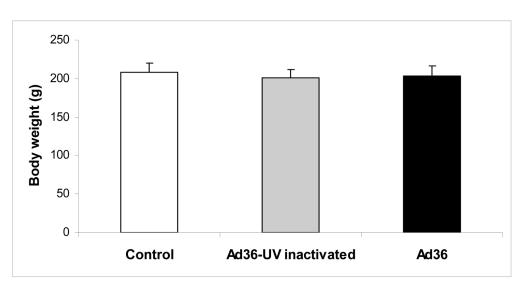


Figure 1B: Epididymal fat pad weight 4 d post inoculation

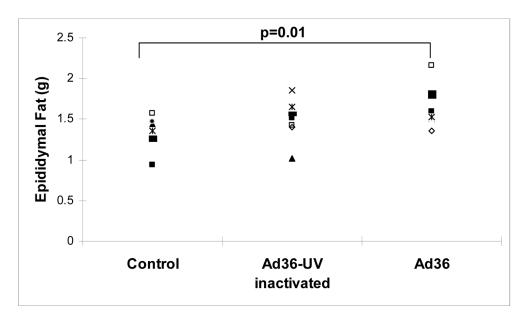


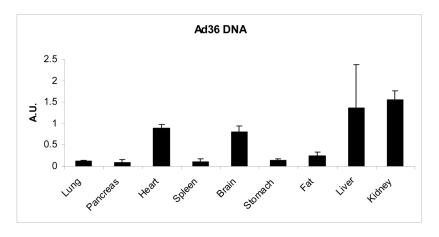
Figure 1.

Figure 1A: Body weight of rats in three groups on day 4 post inoculation was not significantly different.

Figure 1B: Epididymal fat pad weights of rats on day 4 post inoculation.

Pasarica et al.

Figure 2A: Ad36 DNA in tissues, 4 d post i.n. inoculation





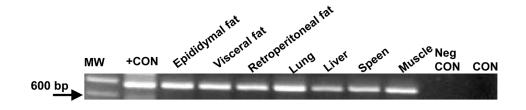


Figure 2C: Ad36 E4orf1 mRNA in epididymal adipose tissue

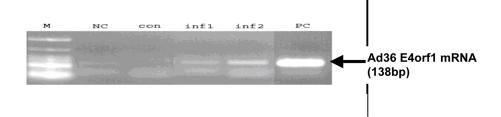
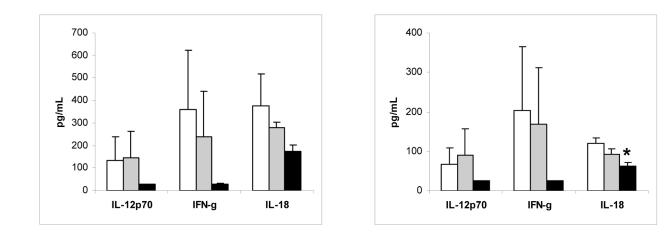


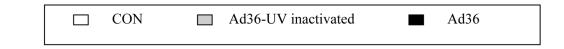
Figure 2.

Figure 2A: Presence of Ad36 DNA in various tissues of rats infected by an intra-nasal inoculation. The quantities were determined by quantitative real time PCR. Figure 2B: Presence of Ad36 DNA in various tissues of rats infected by intraperitoneal inoculation. The quantities were determined by nested PCR. MW: molecular weight ladder. Neg. CON: Water template, CON: Tissue from uninfected control animal. Figure 2C: Ad36 E4orf1 mRNA in epididymal fat pad of two representative animals (inf 1 and inf2) from the Ad36 infected group. PC: Positive control mRNA from Ad36 infected A549 cells. NC: negative control water template. M: molecular weight ladder.

Figure 3A: 2 d post inoculation







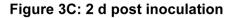


Figure 3D: 4 d post inoculation

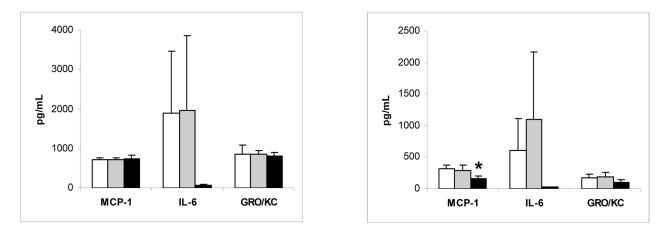


Figure 3.

Figures 3A–D: Serum cytokine levels in 5 randomly selected animals from each group, 2 and 4 days post inoculation. * p < 0.05. GRO/KC: growth regulated oncogene.