

RESEARCH PAPER

Generation of human MHC (HLA-A11/DR1) transgenic mice for vaccine evaluation

Yang Zeng^{a,b}, Tongtong Gao^c, Guangyu Zhao^a, Yuting Jiang^a, Yi Yang^a, Hong Yu^a, Zhihua Kou^a, Yuchun Lone^b, Shihui Sun^a, and Yusen Zhou^{a,c}

^aState Key Laboratory of Pathogen and Biosecurity, Beijing Institute of Microbiology and Epidemiology, Beijing, China; ^bINSERM U1197 (ex U1014), University of Paris-Sud, Hospital Paul Brousse, Villejuif, France; ^cWenzhou Medical University, Zhejiang, China

ABSTRACT

The rapid occurrence of emerging infectious diseases demonstrates an urgent need for a new preclinical experimental model that reliably replicates human immune responses. Here, a new homozygous humanized human leukocyte antigen (HLA)-A11/DR1 transgenic mouse (HLA-A11^{+/+}/DR1^{+/+}/H-2- $\beta_{2m}^{-/-}$ /IA $\beta^{-/-}$) was generated by crossing HLA-A11 transgenic (Tg) mice with HLA-A2^{+/+}/DR1^{+/+}/H-2- $\beta_{2m}^{-/-}$ /IA $\beta^{-/-}$ mice. The HLA-A11-restricted immune response of this mouse model was then examined. HLA-A11 Tg mice expressing a chimeric major histocompatibility complex (MHC) molecule comprising the $\alpha 1$, $\alpha 2$, and β_{2m} domains of human HLA-A11 and the $\alpha 3$ transmembrane and cytoplasmic domains of murine H-2D^b were generated. The correct integration of HLA-A11 and HLA-DR1 into the genome of the HLA-A11/DR1 Tg mice (which lacked the expression of endogenous H-2-I/II molecules) was then confirmed. Immunizing mice with a recombinant HBV vaccine or a recombinant HIV-1 protein resulted in the generation of IFN- γ -producing cytotoxic T lymphocyte (CTL) and antigen-specific antibodies. The HLA-A11-restricted CTL response was directed at HLA immunodominant epitopes. These mice represent a versatile animal model for studying the immunogenicity of HLA CTL epitopes in the absence of a murine MHC response. The established animal model will also be useful for evaluating and optimizing T cell-based vaccines and for studying differences in antigen processing between mice and humans.

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Introduction

Major histocompatibility complex (MHC) molecules play a vital role in activating the adaptive immune system. They do this by presenting antigens (usually peptides) to immune cells in a specific context and by participating in the differentiation and maturation of T lymphocytes in the thymus.^{1,2} The incidence of disease caused by newly emerged viruses is increasing worldwide; therefore, candidate epitope-based vaccines are vital for providing immunoprotection from pathogens via the elicitation of humoral and cellular responses.^{3–8} Specific cytotoxic T lymphocyte (CTL) responses would lead to a marked reduction in the viral load, and may even clear the virus and cure some autoimmune diseases.⁹ A protective adaptive immune response is based on the effective activation and mobilization of B cells, cytotoxic T cells, and helper T cells.¹⁰ Synergism between MHC class I and II molecules is a key component of an effective host immune response.

Human MHC molecules (known as HLA (human leukocyte antigen) molecules) are the most polymorphic genes in the human genome. MHC restriction differs markedly according to geographical region and ethnicity. Several studies used MHC-I or MHC-II transgenic mice to examine the molecular mechanism(s) underlying disease, evaluate candidate vaccines, and screen HLA-restricted epitopes.^{11,12} For example, Ishioka

et al.¹³ used HLA-A2 and HLA-A11 transgenic (Tg) mice to evaluate a minigene DNA vaccine encoding multiple HLA-restricted CTL epitopes derived from HIV and HBV, whereas Pajot et al.¹⁴ used HLA-DR1 Tg mice to identify novel HLA-DR1-restricted epitopes derived from the HBV envelope protein. T helper cells play a vital role in boosting CTL responses and humoral immune responses against pathogens; indeed, the collaboration between T helper and cytotoxic CD8⁺ T cells is critical for an effective CTL response.^{15,16} We previously generated HLA-Tg mice expressing both HLA class I and II molecules (e.g., HLA-A2/DR1 Tg mice¹⁷ and HLA-A2/DP4 Tg mice¹⁸) and used them to develop and evaluate vaccines.^{19–22}

HLA-A11 is one of the most common HLA class I genotypes in the world, with a phenotypic frequency of about 20–30% in the Chinese population and 10–15% in European and American populations.²³ The first HLA-A11/K^b mouse was generated by Alexander et al.²⁴ and was used to develop HLA-A11-restricted epitope-based vaccines.²⁵ However, the relevance of this model to the human immune system was overshadowed by the presence of murine H-2 class I or II molecules, which were used preferentially as restricting elements in response to antigens.^{26,27} Furthermore, HLA-A11/K^b mice did not accurately reflect human T helper cell responses because they lacked the HLA-II molecules. Thus, a transgenic mouse model expressing

both HLA-A11 and HLA-DR1 molecules, but lacking H-2 class I and II molecules, is needed.

Here, we constructed an HLA-A11/DR1 (A11^{+/+}/DR01^{+/+}/ $\beta_2m^{-/-}$ /IA $\beta^{-/-}$) Tg mouse model based on the A*1101/DR01 genotypes, which represented 10–15% of the Chinese population, and used it to evaluate novel candidate vaccines.

Results

Generation of homozygous HLA-A11/DR1 (HLA-A11^{+/+}/DR01^{+/+}/ $\beta_2m^{-/-}$ /IA $\beta^{-/-}$) Tg mice

The construction of the recombinant gene encoding the chimeric HLA-A11 monochains illustrated in Figure 1A and B. Transgenic mice expressing the integrated HLA-A11 fragment were identified by PCR (data not shown). HLA-A11/DR1 (HLA-A11^{+/+}/DR01^{+/+}/ $\beta_2m^{-/-}$ /IA $\beta^{-/-}$) Tg mice were obtained by crossing the parental HLA-A11^{+/+} (HLA-A11) Tg mice with HLA-A2^{+/+}/DR01^{+/+}/ $\beta_2m^{-/-}$ /IA $\beta^{-/-}$ (HLA-A2/DR1) Tg mice. The genotype of the HLA-A11^{+/+}/DR1^{+/+}/ $\beta_2m^{-/-}$ /IA $\beta^{-/-}$ Tg mice was confirmed by PCR, with offspring-derived genomic DNA as a template. The exogenous HLA-A11 and HLA-DR1 fragments were integrated into the mouse genome (Fig. 2A), and the endogenous β_2m and H-2 IA β (Fig. 2B) genes were knocked out.

Expression of HLA-A11 and HLA-DR1 in splenocytes isolated from HLA-A11/DR1 Tg mice was detected by flow cytometry analysis. HLA-A11/DR1 Tg mice expressed both HLA-A11 (Fig. 2C) and HLA-DR1 (Fig. 2D), whereas no HLA transgenes were expressed in wild-type C57BL/6 mice or H-2-I/II knockout mice.

Taken together, these results demonstrated that the HLA-A11 and HLA-DR1 transgenes had integrated into the mouse genome and were expressed by splenocytes from HLA-A11/DR1 Tg mice, while competitive inhibition by mouse endogenous H-2-I/II molecules had been eliminated. Thus, the mouse model could be used to examine cooperation between human HLA-I/II molecules without interference from mouse H-2 molecules.

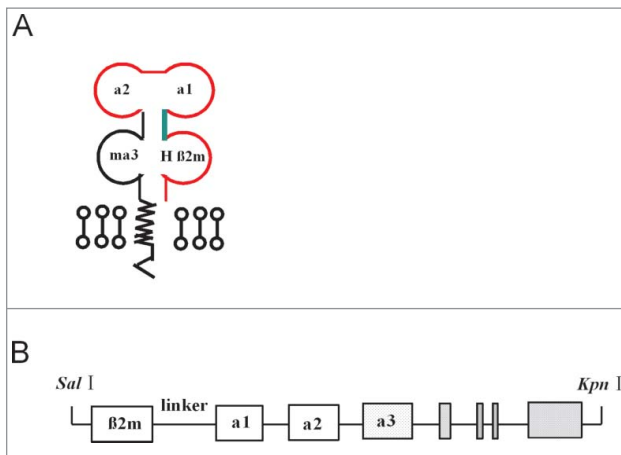


Figure 1. Schematic diagram of the chimeric human/mouse MHC class I gene. (A) Schematic diagram of the monochain chimeric human/mouse MHC class I gene showing the HHD structure of the chimeric human α_1 , α_2 , and β_2m HLA-A11/murine H-2 α_3 molecule. The murine H2 α_3 domain is covalently linked to human β_2m via a 15 amino acid linker. (B) Liner representation of the final construct.

Proportions of CD8⁺ and CD4⁺ T lymphocytes in the peripheral blood of HLA-A11/DR1 Tg mice

To test whether the expression of exogenous HLA transgenes would still allow positive and negative selection of CD4⁺ and CD8⁺ T lymphocytes in HLA-A11/DR1 Tg mice, we next examined the populations of HLA-regulated T lymphocytes in the periphery. The numbers of CD3⁺CD4⁺ and CD3⁺CD8⁺ T lymphocytes were counted by flow cytometry.

The isolated splenocytes were labeled with a PE-anti-mouse CD3 antibody, followed by a PEcy5-anti-mouse CD8 antibody. Figure 3A–D shows that 1.48% of the lymphocytes in HLA-A11/DR1 Tg mice, 3.32% of the lymphocytes in HLA-A11 Tg mice, and 1.16% of lymphocytes in HLA-DR1 Tg mice were CD8⁺ T cells. By contrast, 11.7% of lymphocytes in wild-type C57BL/6 mice were CD8⁺ T cells. Peripheral CD4⁺ T lymphocytes were labeled with a PE-anti-mouse CD3 antibody, followed by an APC-anti-mouse CD4 antibody. Figure 3E–H shows that 92.2% of the lymphocytes in HLA-A11/DR1 Tg mice, 62.6% in HLA-A11 Tg mice, and 94.8% in HLA-DR1 Tg mice were CD4⁺ T cells, compared with 64.6% in wild-type C57BL/6 mice.

These results showed that the percentage of CD4⁺ T cells in HLA-A11/DR1 Tg mice were similar to that in wild-type C57BL/6 mice, while the number of CD8⁺ T cells in HLA-A11/DR1 Tg mice was lower than that in wild-type C57BL/6 mice. However, the lower percentage of CD8⁺ T cell was in accordance with that in HLA-A2/DR1 Tg mice or HLA-A2/DP4 Tg mice in which efficient humoral and cellular immune responses could be developed.^{17,18} Thus, HLA transgenes enabled the differentiation and maturation of CD4⁺ and CD8⁺ T lymphocytes in transgenic mice.

HLA-A11/DR1 Tg mice mount effective humoral and cellular responses

To examine the humoral and cellular responses to vaccination in HLA-A11/DR1 Tg mice, the mice were immunized with a recombinant HBV vaccine (derived from the HBV envelope protein²⁸) or with a multi-epitope HIV-1-based candidate vaccine.²⁹

The amount of HBsAg-specific IgG antibodies in the serum of HLA-A11/DR1 Tg mice was similar to that in the serum of wild-type C57BL/6 mice (Fig. 4A). It is worth noting that HLA-A11/DR1 Tg mice generated 4–5-fold more IFN- γ than wild-type mice, indicating a satisfactory cellular immune response and an ability to mimic the human cytotoxic response (Fig. 4B).

HIV-1 protein vaccine, and the amount of HIV-specific IgG antibodies and the amount of IFN- γ secreted by CD8⁺ T lymphocytes was measured respectively. The results revealed an increase ($p < 0.05$) in the levels of specific IgG antibodies in HLA-A11/DR1 Tg mice (Fig. 5A). HLA-A11/DR1 Tg mice also secreted more IFN- γ than wild-type mice (Fig. 5B). These results suggest that HLA-A11/DR1 Tg mice show stronger cellular immune responses than C57BL/6 mice and are therefore a suitable model for evaluating vaccines.

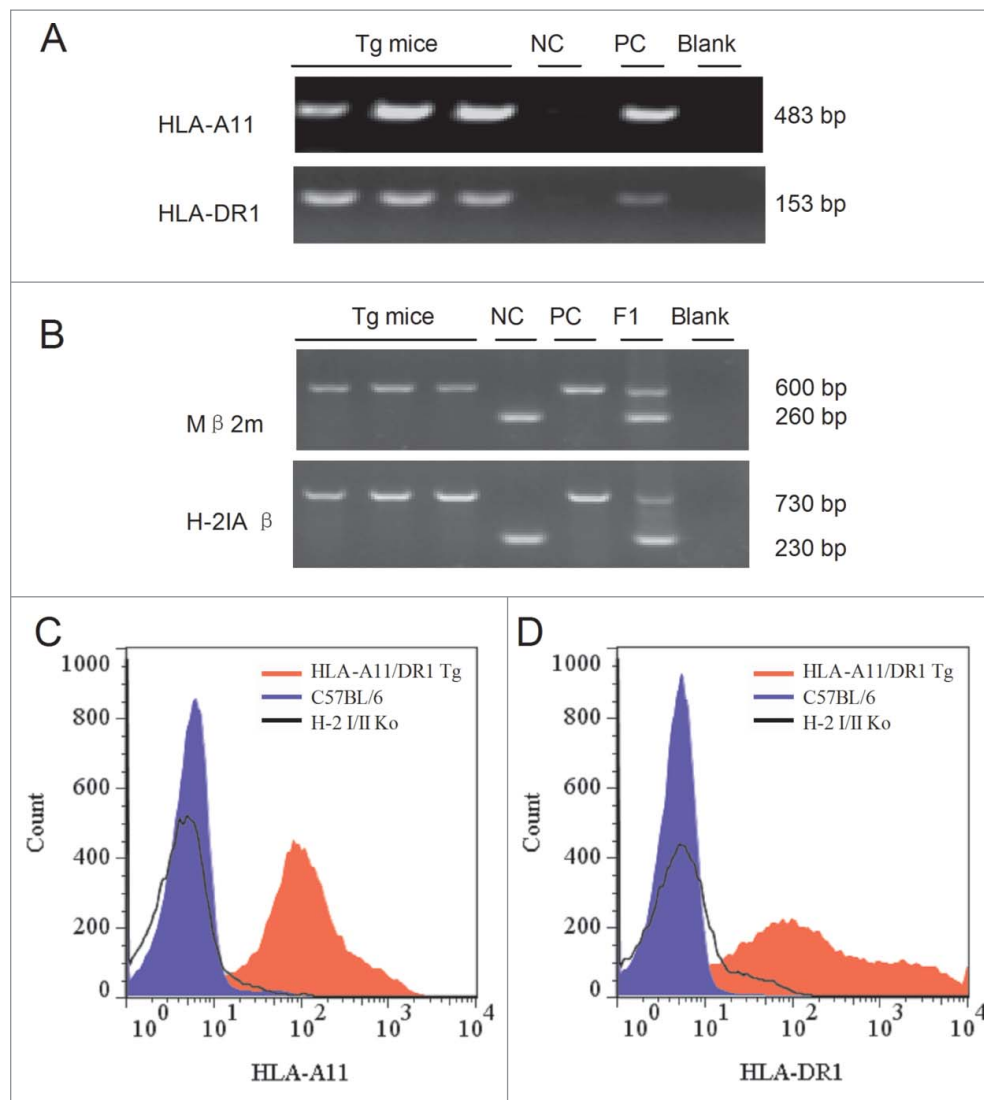


Figure 2. Genotype of HLA-A11/DR1 Tg mice and flow cytometric analysis of transgenic HLA molecules expressed on the surface of mouse immune cells. Genomic DNA purified from HLA-A11/DR1 Tg mice was analyzed by PCR using primers specific for HLA-A11, HLA-DR1, mouse β_2m , and H-2 IA β . (A) Identification of HLA-A11 (upper panel) and HLA-DR1 (lower panel). Tg mice, HLA-A11/DR1 Tg mice; NC, Negative control; PC, Positive control; Blank, dH₂O. (B) Identification of mouse β_2m (upper panel) and H-2 IA β (lower panel). Tg mice, HLA-A11/DR1 Tg mice; NC, Negative control; PC, Positive control; F1, Heterozygote control; Blank, dH₂O. (C and D) Splenocytes from HLA-A11/DR1 (red histogram), H-2/I/II knockout mice (black histogram), and wild-type C57BL/6 (blue histogram) mice were isolated and stained with a PE-anti-human-HLA-ABC antibody (C) or an FITC-anti-human-HLA-DR antibody (D) to detect HLA-A11 and HLA-DR1 expression, respectively.

HLA-A11/DR1 Tg mice show HLA-A11-restricted cytotoxic responses

To verify that HLA-A11/DR1 Tg mice mount a fully functional immune response, 2 known immunodominant HLA-A11-restricted epitopes, one derived from HIV-Pol₁₇₇₋₁₈₈ (QMAV-FIHNFKRK)³⁰ and the other derived from HIV-GP160₃₂₋₄₅ (KLWVTVYYGVVWR),³⁰ were used to measure IFN- γ production by CD8⁺ T lymphocytes from HLA-A11/DR1 Tg mice immunized with a recombinant HIV-1 protein vaccine.

Compared with those from wild-type C57BL/6 mice, CD8⁺T lymphocytes from HLA-A11/DR1 Tg mice were functionally restricted by the Tg human MHC class I molecules (HLA-A11). The HLA-A11-restricted HIV-specific CTL immune responses were specific for the HLA-A11-restricted-epitopes HIV-Pol₁₇₇₋₁₈₈ (Fig. 6A) and HIV-GP160₃₂₋₄₅ (Fig. 6B). Collectively, these data show that HLA-A11/DR1 Tg mice mount a functional HLA-A11 restricted immune response.

Discussion

Vaccines based on multiple epitopes can be used to target well-defined ethnic populations and prime robust immune responses.^{30,31} To facilitate the evaluation and development of novel candidate epitope-based vaccines, we constructed a novel humanized HLA-A11/DR1 Tg mouse model expressing both HLA-A11 and HLA-DR1 molecules in the absence of H-2 class I/II molecules. Different from previous HLA-A2/DR1¹⁷ and HLA-A2/DP4 Tg mouse models,¹⁸ HLA-A11/DR1 Tg mice showed combined expression of HLA-I allele A11 and HLA-II allele DR1, both of which are prevalent in the Chinese population. Because a quarter of the Chinese population express the HLA-A11 and HLA-DR1 genotypes,²³ this novel animal model could be used to predict and evaluate the cellular responses of this Chinese population more accurately than previous transgenic mouse models.

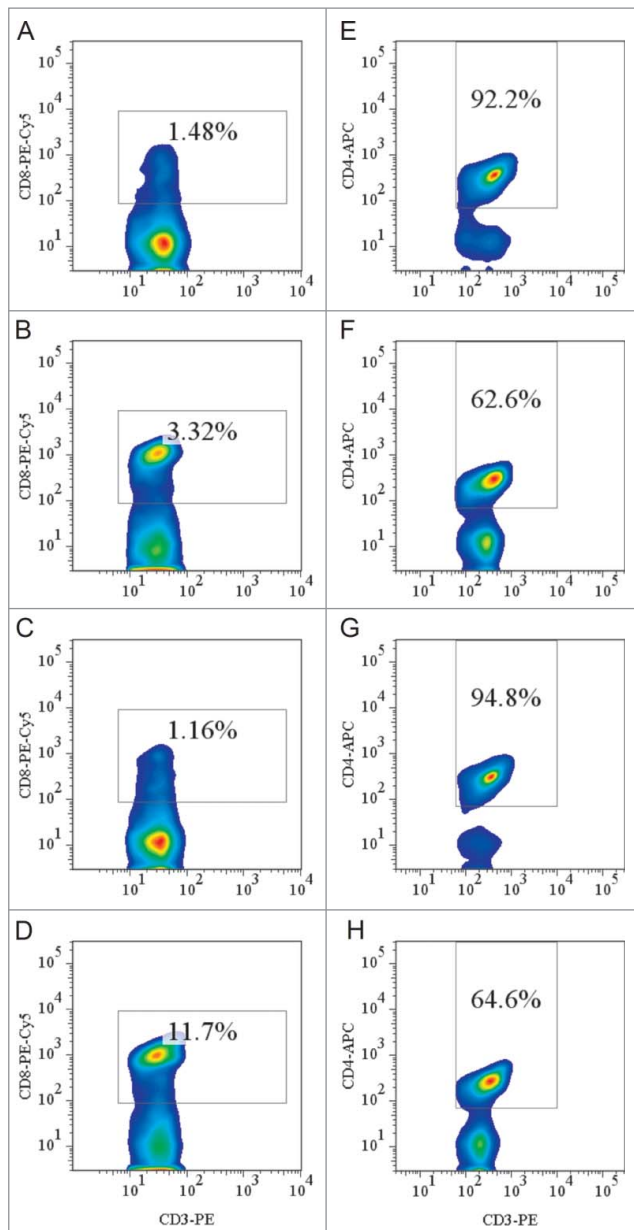


Figure 3. Flow cytometric analysis of peripheral CD8⁺ and CD4⁺ (T) lymphocytes. (A–D) Splenocytes from HLA-A11/DR1 mice (A), HLA-A11 mice (B), HLA-DR1 mice (C), and wild-type C57BL/6 mice (D) were isolated. CD3⁺ T lymphocytes were gated by staining with a FITC-anti-CD3 mAb and CD8⁺ T lymphocytes were gated by staining with a PEcy5-anti-CD8 mAb. (E–H) Splenocytes from HLA-A11/DR1 mice (E), HLA-A11 mice (F), HLA-DR1 mice (G), and wild-type C57BL/6 mice (H) were isolated. CD3⁺ T lymphocytes were gated by staining with a PE-anti-CD3 mAb, and CD4⁺ T lymphocytes were gated by staining with an APC-anti-CD4 mAb.

The HLA system is the most polymorphic gene cluster in human beings. HLA class I and II alleles are expressed in a codominant manner in any particular individual. Competition among different HLA molecules results in the selection of different immunodominant epitopes that recognize and present processed antigens to immune cells. Examining the function of a single HLA molecule in the absence of competitive inhibition is a real challenge. Humanized mice have allowed the investigation of individual HLA molecules due to the integration of specific HLA alleles into the mouse genome. Indeed, double-Tg mice (expressing human HLA class I and class II molecules) in the absence of murine H-2 molecules have been developed.^{17,18}

The phenotypic frequency of HLA-A11 in the Chinese population is about 20–30% compared with 10–15% in European, American, and Middle Eastern populations.²³ Thus, much research focused on HLA-A11 Tg mice for evaluating HLA-A11-restricted epitope-based vaccines.^{32,33}

To ensure that the transgenes functioned within the mouse genome, we designed a HLA-A11 HHD chimeric monochain expressing the $\alpha 1/\alpha 2/\beta_2m$ domains of HLA-A11 and the $\alpha 3$ domain of endogenous H-2. Replacing the human $\alpha 3$ domain with the mouse $\alpha 3$ domain preserved the species-specific binding affinity between the HLA-A11 chimeric monochain and the mouse CD8 molecule. Inactivating important components of the endogenous H-2 molecule prevented H-2 molecules from competing with the transgenic HLA molecules. Therefore, transgenic mice could only mount HLA-restricted immune responses.

Here, we confirmed the cell surface expression of HLA-I and HLA-II molecules by immune cells from HLA-A11 Tg mice. Although the percentage of CD8⁺ T lymphocytes in HLA-A11/DR1 mice was lower than that in the wild-type mice, the Tg HLA molecules efficiently presented antigens to immune cells to generate functional humoral and cellular immune responses. The low percentage of CD8⁺ T lymphocytes in HLA-A11/DR1 mice is in accordance with that in HLA-A2/DR1 Tg mice,¹⁷ HLA-A2/DP4 Tg mice,¹⁸ and other HLA class I transgenic mice models which also could develop efficient humoral and cellular immune responses similar to those in humans.^{34,35} Furthermore, studies show that the lower percentage of CD8⁺ T lymphocytes does not preclude the use of HLA-A2/DR1 Tg mice for studies of immune responses to pathogens such as HBV and HIV.^{34,36} Interesting, it is now well documented by several comparative studies that transgenic HLA class I molecules in chimeric monochain form are superior than traditional transgenic HLA class I molecule mice to induce more efficient T cell responses, even the number of CD8 T cell is lower.^{37,38}

We also confirmed that HLA-A11/DR1 Tg mice mount efficient and functional humoral and cellular immune responses to a recombinant HBV vaccine and a recombinant HIV-1 protein, and especially mount the specific A11-restricted cellular immune response to a recombinant HIV-1 protein. In our HLA-A11/DR1 transgenic mice, the HLA-DR1 genes were derived from the HLA-A2/DR1 transgenic mice in which the HLA-DR1-restricted CD4⁺T cell responses were described with the HLA-DR1-restricted epitope HBsAg_{179–194}.¹⁷ Furthermore, by using the HLA-A2/DR1-transgenic, H-2 class I/class II KO mice, Pajot et al.³⁶ identified 2 new HLA-DR1-restricted HIV-1 Gag p24-derived epitopes (Gag321–340 and Gag331–350) and confirmed the immunogenicity of 7 epitopes that had been described previously. In addition, it is well documented that in absence of MHC class II expression, there are no CD4⁺T cells and more interesting, there are also no humoral responses in the case of vaccination against HBV, as shown in HLA-A2 single transgenic H-2 class I/class II knock-out (KO) mice.³⁶ In the present HLA-A11/DR1 transgenic mice, we confirmed the HLA-DR1-restricted epitope HBsAg_{179–194} responses as reported in HLA-A2/DR1 transgenic mice (data not shown). Furthermore, we show an efficient specific antibodies responses (Fig. 4) which are reported highly protective against HBV viral infection. The results indicated that the novel HLA-A11/DR1

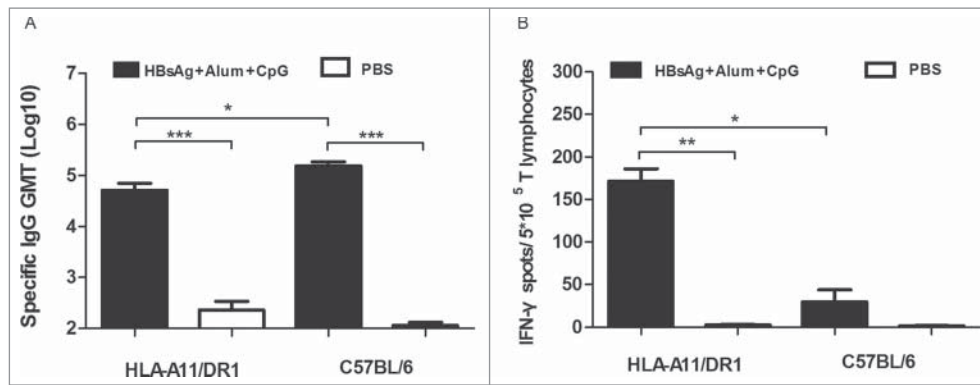


Figure 4. HBs-specific antibody- and cell-mediated responses after immunization with a recombinant HBsAg vaccine. (A) Sera were collected from immunized HLA-A11/DR1 Tg mice and wild-type C57BL/6 mice, and the titers of anti-HBs (IgG) antibodies were compared with those of PBS-immunized (hollow bar) mice in an ELISA test. (B) HBs epitope-specific IFN- γ production by cytotoxic T lymphocytes was examined by measuring the response of both HLA-A11/DR1 Tg mice and wild-type C57BL/6 mice to a recombinant HBs vaccine or PBS.

transgenic mice model has potential for evaluating both T cells (CD8 and CD4) and B cells activities as HLA-A2/DR1 transgenic mice, but covering different population. The two models are mutual important to evaluate available commercial vaccines and novel vaccine candidates development.

In summary, we generated a novel HLA-A11/DR1 Tg mouse strain that faithfully reproduced a human immune response and may be useful for identifying protective epitopes, designing human vaccine and for evaluating vaccination strategies. The humanized HLA-A11/DR1 Tg mouse is a promising and versatile preclinical model that will facilitate the study of human immune responses to a variety of antigens.

Materials and methods

Construction of the HHD chimeric HLA-A11 transgenic plasmid

A 2873bp chimeric HLA-A*1101 (GenBank accession number, D16841.1) monochain fragment was synthesized using GenScript and subcloned into the pET32a plasmid via the *SalI* and *KpnI* restriction sites to generate pET32a-A11, which encompassed the human HLA-A11 leader sequence, the $\alpha 1$ and $\alpha 2$ domains, murine H-2D^b(from the $\alpha 3$ domain to the COOH

terminus), and human β_2 -microglobulin (β_2m), which was covalently linked between the leader sequence and the 5' end of the HLA-A11 ($\alpha 1$ and $\alpha 2$) domains via a 15 amino acid linker encoding Gly₄Ser₃.³⁹ The mouse $\alpha 3$ domain within the chimeric HLA-A11 molecule facilitates interaction with mouse CD8 molecules.³⁹

Generation of HHD chimeric HLA-A11 Tg mice

The purified HLA-A11 fragment was released from pET32a-A11 by digestion with *SalI* and *KpnI* and microinjected into pronuclei from fertilized (C57BL/6J \times C57BL/6J) F1 mouse eggs to generate Tg embryos. The embryos were then reimplanted into pseudopregnant female mice. HLA-A11 Tg mice were identified by PCR of genomic DNA and by fluorescence activated cell sorting of T lymphocytes (see below). HLA-A11 Tg mice showing the highest expression of HLA-A11 were then crossed with HLA-A2/DR1 (HLA-A2^{+/+}/DR1^{+/+}/ β_2m ^{-/-}/IA β ^{-/-}) Tg mice on a C57BL/6 background (originally obtained from Dr. Yuchun Lone¹⁷). Offsprings with the HLA-A2^{-/-} A11^{+/+}/DR1^{+/+}/ β_2m ^{-/-}/IA β ^{-/-} genotype were identified by PCR.

Mice were bred in the animal facility at the Laboratory Animal Center, Chinese Academy of Medical Sciences, Beijing,

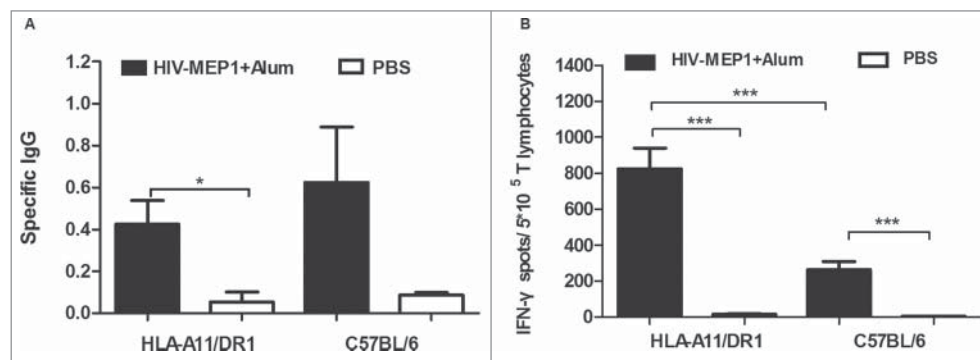


Figure 5. HIV-specific antibody and CD8⁺ (T)cell responses after immunization with a recombinant HIV-1 protein. (A) HIV-specific antibody titers in HLA-A11/DR1 Tg and C57BL/6 mice immunized with a recombinant HIV-1 protein or PBS. (B) HIV-specific IFN- γ production by cytotoxic T lymphocytes was examined by measuring responses to the recombinant HIV-1 protein in immunized mice (PBS-immunized mice were used as control).

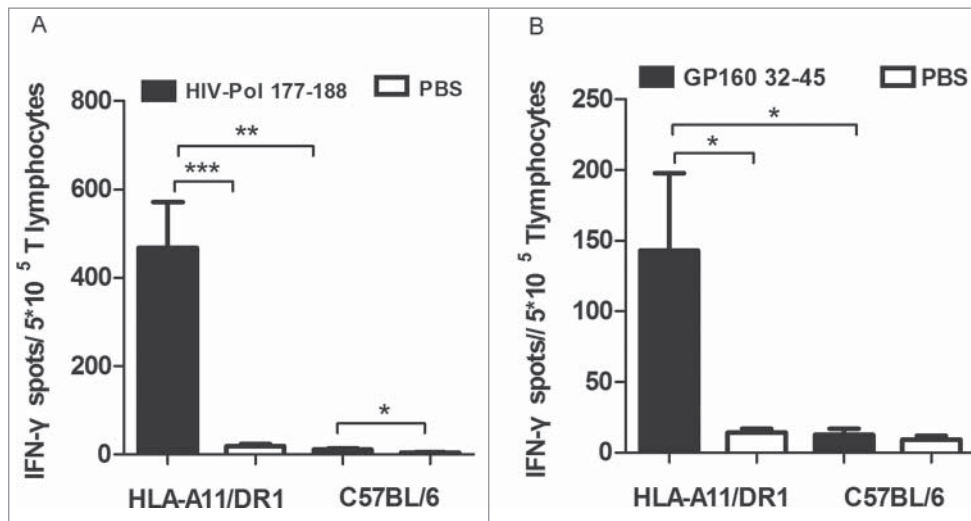


Figure 6. HLA-A11 restricted cytotoxic responses in HLA-A11 transgenic mice. Responses to (A) the HLA-A11-restricted epitope, HIV-Pol₁₇₇₋₁₈₈, and (B) the HLA-A11-restricted epitope, HIV-GP₃₂₋₄₅, were assessed in immunized HLA-A11 transgenic and wild-type C57BL/6 mice by counting the number of IFN- γ -secreting spots in ELISPOT assays.

China. All experiments were performed according to the approved protocols and guidelines of the animal facility at the Institutional Animal Care and Use Committees of the Laboratory Animal Center, State Key Laboratory of Pathogen and Biosecurity, Beijing Institute of Microbiology and Epidemiology (permit number, BIME 2014-0017), and the recommendations set out in the guide for the Care and Use of Laboratory Animals.

Genotype identification

HLA-A11 and HLA-DR1 transgenes and $\beta_2m/IA\beta$ knockout mice were identified by PCR. Murine genomic DNA was extracted as described previously,¹⁷ and PCR was conducted using the following primers: HLA-A11 forward, 5'-CTGGGT TTCATCCATCCG-3', and reverse, 5'-GATCGGTCTGGCT CTGAGC-3'; HLA-DR1 forward, 5'-GCTTCGAAATGGAAA ACCTG-3', and reverse, 5'-ATGTGCCTTACAGAGGCC-3'; mouse β_2m forward, 5'-CTGAGCTCTGTTTTCTGCTG-3', and reverse, 5'-CTTAAGTCTGCAGGCGTATG-3'; H-2 I α forward, 5'-TTCGTGTACCAGTTCATGGG-3', and reverse, 5'-TAGTTGTGCTGCACACCGT-3'; and Neo55a, 5'-CCTG CCGAGAAAGTATCCA-3'.

Flow cytometry analysis

The mice from experimental and control mice were euthanized and the splenocytes were isolated. The expression of HLA-A11 and HLA-DR1 was examined by flow cytometry analysis with PE-anti-human HLA-ABC and FITC-anti-human HLA-DR antibodies (anti-HLA class I and anti-HLA class II, respectively; BioLegend, San Diego, CA, USA). Splenocytes isolated from wild-type C57BL/6 and H-2-I/II knockout mice were used as controls. To determine the percentage of CD4⁺ and CD8⁺ cells, splenocytes were first labeled with a PE-anti-mouse CD3 antibody (BioLegend), followed by PE-cy5-anti-mouse CD8 or APC-anti-mouse CD4 (BioLegend). Splenocytes from wild-type C57BL/6 mice were used as the control.

Immunization of HLA-A11/DR1 Tg mice

Female HLA-A11/DR1 Tg mice or wild-type C57BL/6 mice (6–8 weeks old) received 3 intramuscular (i.m.) injections of recombinant HBV S antigen (1 μ g) adsorbed to 100 μ g of Al (OH)₃ (alum adjuvant) in combination with 10 μ g of CpG-ODNs (kindly provided by Dr. Honglin Xu²⁸) (total injected volume, 100 μ l). Each immunization was separated by an interval of 14 d. Sera were collected both before and after each immunization for serological analysis. Mice were sacrificed 10 d after the final boost, and the cytolytic responses to recombinant HBsAg vaccine were examined in an IFN- γ ELISPOT assay.

Female Tg and wild-type mice (6–8 weeks old) were anesthetized, and the tibialis anterior muscles were injected with 10 μ g of recombinant HIV-1 protein in alum adjuvant; mice received 3 injections at 14 day intervals.²⁹ The immunogenicity of 2 known HLA-A11-restricted epitopes derived from the HIV-Pol gene (aa177–188: QMAVFIHNFKRK)³⁰ and the HIV-GP160 gene (aa32–45: KLWVTVYYGVVWR)³⁰ was examined by measuring the amount of IFN- γ secreted by CD8⁺ T lymphocytes from HLA-A11/DR1 Tg mice and wild-type C57BL/6 mice. Mice were sacrificed 10 d after the final boost, and the cytolytic responses and HLA-A11-restricted CTL responses to the recombinant HIV-1 protein were determined in an ELISPOT assay.

Measurement of serum antibodies by ELISA

The serum levels of antibodies specific for the recombinant HBV vaccine and the recombinant HIV-1 protein in immunized HLA-A11/DR1 Tg mice and wild-type C57BL/6 mice were measured in ELISA. The plates coated with the HBV vaccine antigen or the recombinant HIV-1 protein were blocked with PBS supplemented with 0.1% Tween20 and 10% FCS and then washed 3 times. After the addition of mouse serum for 1 h, the plates were washed again, and bound antibodies were detected with HRP (horseradish peroxidase)-labeled

anti-mouse IgG (Santa Cruz Biotechnology, Inc.). Absorbance was then measured at 450_{nm} and 630_{nm} in a plate reader. The antibody titers (the mean of at least 3 determinations) were calculated using the serial end-point dilution method.⁴⁰

ELISPOT assay

An ELISPOT assay was performed to detect IFN- γ secreted by CD8⁺ T lymphocytes with BDTM ELISPOT kit (BD Biosciences, CA). Briefly, 96-well ELISPOT plates were coated with an anti-IFN- γ mAb overnight at 4°C and then blocked with blocking solution. Splenocytes (5×10^5) were added to each well and cultured in the presence of recombinant HBV vaccine, synthetic peptides, or recombinant HIV-1 protein (all at 10 μ g/ml) and incubated at 37°C/5% CO₂ for 48 h. IFN- γ -secreting cells were captured by an anti-IFN- γ mAb and detected by incubation with a biotinylated anti-IFN- γ mAb for 2 h at 37°C, followed by streptavidin-HRP for 1 h. The plates were then developed for 5–10 minutes in an ACE substrate solution (BD Biosciences, CA), washed 3 times, and air-dried at room temperature. Positive spots were counted using an ELISPOT plate reader (Cellular Technology Ltd).

Statistical analysis

Each experiment was repeated at least 3 times, and data were expressed as the mean \pm SEM. The means or geometric means of multiple groups were compared using Student's t-test. All statistical analysis was performed using GraphPad Prism version 5.0. A P value <0.05 was considered statistically significant.

Abbreviations

MHC	major histocompatibility complex
HLA	human leukocyte antigen
CTL	cytotoxic T lymphocyte
HIV	human immunodeficiency virus
HBV	hepatitis B virus

Disclosure of potential conflicts of interest

All authors agree to submit our manuscript to *Human Vaccines & Immunotherapeutics*. None of the authors have any conflicts of interest to declare.

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