Immune Tolerance Split between Hepatitis B Virus Precore and Core Proteins

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The function of the hepatitis B virus (HBV) precore or HBeAg is largely unknown because it is not required for viral assembly, infection, or replication. However, the HBeAg does appear to play a role in viral persistence. It has been suggested that the HBeAg may promote HBV chronicity by functioning as an immunoregulatory protein. As a model of chronic HBeAg exposure and to examine the tolerogenic potential of the HBV precore and core (HBcAg) proteins, HBe/HBeAg-transgenic (Tg) mice crossed with T cell receptor (TCR)-Tg mice expressing receptors for the HBc/HBeAgs (i.e., TCR-antigen double-Tg pairs) were produced. This study revealed three phenotypes of HBe/HBeAg-specific T-cell tolerance: (i) profound T-cell tolerance most likely mediated by clonal deletion, (ii) T-cell clonal ignorance, and (iii) nondeletional T-cell tolerance mediated by clonal anergy and dependent on the structure, location, and concentration of the tolerogen. The secreted HBeAg is significantly more efficient than the intracellular HBcAg at eliciting T-cell tolerance. The split T-cell tolerance between the HBeAg and the HBcAg and the clonal heterogeneity of HBe/HBeAg-specific T-cell tolerance may have significant implications for natural HBV infection and especially for precore-negative chronic hepatitis.

The nucleoprotein of the hepatitis B virus (HBV) exists in two structural forms. The nucleocapsid, designated the hepatitis core antigen (HBcAg), is an intracellular 21-kDa protein that self-assembles into particles that encapsidate the viral genome and polymerase and is essential to the function and maturation of the virion. A unique feature of the HBV is the production of a secreted, nonparticulate second form of the nucleoprotein designated the precore or hepatitis B e antigen (HBeAg). The precore and core proteins are translated from 2 distinct RNA species that have different 5' initiation sites (14). The precore mRNA encodes a hydrophobic signal sequence that directs the precore protein to the endoplasmic reticulum, where it undergoes N- and C-terminal cleavage within the secretory pathway and is secreted as an 18-kDa monomeric protein (26, 38, 42, 53). The HBcAg and HBeAg are distinctly recognized by antibodies (22) but, due to their extensive amino acid homology, are highly cross-reactive at the T-cell level (2, 36, 50). In contrast to the HBcAg, the function of secretory HBeAg in the viral life cycle is unknown inasmuch as it is not required for assembly, infection, or replication (6, 10, 44). It has been proposed that the HBeAg may have an immunoregulatory function in promoting viral persistence. For example, HBeAg-specific Th2 cells may cross-regulate HBcAg-specific Th1 cells or the secreted HBeAg may preferentially behave as a tolerogen and inactivate HBcAg-specific T cells through deletion or clonal anergy in the periphery (35, 36).

In the most successful chronic infections, viral antigens either evade the host immune response or tolerate the host immune system. During chronic viral infections such as HBV it may be useful to consider the viral antigens as neoself-antigens due to their long-term coexistence with the host immune system. Immune tolerance to self- or neoself-antigens can be mediated by a variety of mechanisms. For example, strongly stimulating self-antigens can delete autoreactive T cells during development in the thymus or in peripheral lymphoid tissues (15, 23). Conversely, antigen receptor stimulation in the absence of costimulatory molecules or stimulation with weak antigens can elicit a nondeletional program of anergy (47). T-cell anergy is a tolerance mechanism in which the T cell is functionally inactivated following an initial antigen encounter but remains alive in a hypoactivated state. Two broad categories of T-cell anergy have been defined as clonal anergy, principally a growth arrest state, and adaptive tolerance, or in vivo deletion or clonal anergy in the periphery (35, 36).

In this report we have compared the tolerogenic potential of the HBeAg and HBcAg in mice transgenic (Tg) for the two proteins and in HBe/HBeAg-Tg mice bred to T-cell receptor (TCR)-Tg mice expressing receptors specific for the HBe/HBeAgs (i.e., TCR-antigen [Ag] double-Tg pairs). This study revealed three phenotypes of HBe/HBeAg-specific T-cell tolerance: (i) profound T-cell tolerance most likely mediated by...
clonal deletion, (ii) nondeletional T-cell tolerance dependent on the structure, location, and/or concentration of the tolerogen mediated by in vivo anergy, and (iii) T-cell clonal ignorance. Importantly, the HBeAg appears more efficient at eliciting T-cell tolerance than the HBeAg, and this split immune tolerance may have significant implications during a natural HBV infection.

MATERIALS AND METHODS

Mice. C57BL/10 (B10) (H-2\textsuperscript{d}), H-2 congenic B10.S (H-2\textsuperscript{s}), and B10.S MRL-Fas\textsuperscript{sky} mice were obtained from the breeding colony of the Vaccine Research Institute of Stanford University. The HBV-Tg lineages used in this study expressed HBV proteins of the ayw subtype, as produced as described previously, and are listed in Table 1 (16–18, 31, 33). For convenience, we designate the HBeAg-Tg mice as HBc(lo) and HBe(bi) and the HBcAg-Tgs as HBc(lo) and HBc(bi) based on protein expression levels. The entire HBV-Tg represents the 1.3.3.23 lineage that contains a greater-than-genome-length copy of the HBV (ayw) genome and replicates HBV at high levels as described previously (17). The TCR-Tg lineages used in this study are listed in Table 2. The TCR-a chain V3 fragments and beta-chain V(D)J fragments were derived from HBeAg-specific T-cell hybridomas, and characterization of VJ, V\textsubscript{\gamma}, V\textsubscript{\delta}, and J\textsubscript{\gamma} usage and the V(D)J junctional region sequence of \beta chain and VJ junctional regions of \alpha chain from the 2B2, 4E4, and 1E9 T-cell hybridomas have been published previously (8). Modified \alpha chain V3 fragments were inserted into an Xho-NotI-excised TCR a chain shuttle vector which contains a rearranged TCR a chain genomic DNA and the endogenous \beta chain enhancer. Modified \beta chain V(D)J fragments were inserted into a ClaI-NotI-excised TCR \beta chain shuttle vector that contains a rearranged TCR \beta \textchain genomic DNA and the endogenous \beta enhancer as described previously (8). The shuttle vectors (20) were kindly provided by M. M. Davis, Stanford University. The 15.4-kb \alpha chain and 19.8-kb \beta chain TCR DNA fragments were comicroinjected into fertilized mouse (C57BL/10, B10 or B10.s) embryos. Progeny mice were screened for the presence of the TCR transgenes in peripheral blood lymphocytes by PCR analysis with primers located on the V and the CDR3 region for each TCR transgene. The expression of the TCR transgenes in peripheral blood lymphocytes and lymphoid tissues was confirmed by immunofluorescence and reverse transcription-PCR by using monoclonal antibodies (MAbs) and oligonucleotide primers specific for the transgene TCRs. The expression of the TCR transgenes in peripheral blood lymphocytes and lymphoid tissues was confirmed by immunofluorescence and reverse transcription-PCR by using monoclonal antibodies (MAbs) and oligonucleotide primers specific for the transgene TCRs.

<table>
<thead>
<tr>
<th>Transgene</th>
<th>Promoter</th>
<th>Expression of \textsuperscript{b}</th>
<th>Reference</th>
</tr>
</thead>
<tbody>
<tr>
<td>HBeAg</td>
<td>MT</td>
<td>Low (10–20 ng/ml)</td>
<td>31</td>
</tr>
<tr>
<td>HBeAg</td>
<td>MT</td>
<td>Low (0.25 ng/mg)</td>
<td>33</td>
</tr>
<tr>
<td>HBeAg</td>
<td>MUP</td>
<td>High (4–10 \mu g/ml)</td>
<td>18</td>
</tr>
<tr>
<td>HBeAg</td>
<td>MUP</td>
<td>High (1–2 \mu g/mg)</td>
<td>16</td>
</tr>
<tr>
<td>Entire HBV</td>
<td>HBV</td>
<td>Intermediate (50 ng/ml)</td>
<td>17</td>
</tr>
</tbody>
</table>

\textsuperscript{a} MT, metallothionein I; MUP, major urinary protein.

TABLE 2. Summary of HBc/HBeAg-specific TCR-Tg lineages used in this study

<table>
<thead>
<tr>
<th>Lineage</th>
<th>Source</th>
<th>T-cell hybridoma\textsuperscript{a}</th>
<th>TCR (V\textsubscript{\alpha}/V\textsubscript{\beta})</th>
<th>Specificity</th>
<th>TCR\textsuperscript{a} frequency (splenic CD4\textsuperscript{+}) (%)</th>
<th>Avidity</th>
<th>Tolerance\textsuperscript{c} (dbl-Tgs)</th>
</tr>
</thead>
<tbody>
<tr>
<td>11/4-12</td>
<td>B10/eAg-Tg</td>
<td>2B2</td>
<td>11.1/4</td>
<td>129–140/A\textsuperscript{b}</td>
<td>67</td>
<td>Low</td>
<td>Ignorance</td>
</tr>
<tr>
<td>7/16-5</td>
<td>B10/+</td>
<td>4E4</td>
<td>5/11</td>
<td>129–140/A\textsuperscript{b}</td>
<td>53</td>
<td>Intermediate</td>
<td>Anergy</td>
</tr>
<tr>
<td>8/12-2</td>
<td>B10.S/+</td>
<td>1E9</td>
<td>--/4</td>
<td>120–131/A\textsuperscript{b}</td>
<td>37</td>
<td>High</td>
<td>Deletion (?)</td>
</tr>
</tbody>
</table>

\textsuperscript{a} See reference 8 for details of these hybridomas.

\textsuperscript{b} Determined by breeding TCR-Tg mice with HBV-Tg lineages which express either high or low levels of HBeAg or HBeAg.

\textsuperscript{c} Recombinant proteins and synthetic peptides. Recombinant HBeAg of the ayw subtype was produced in Escherichia coli and purified as described previously (45). A recombinant HBeAg corresponding in sequence to serum-derived HBeAg encompassing the 10 precore amino acids remaining after cleavage of the precursor and residues 1 to 149 of HBeAg was produced as described previously (46). The presence of the 10 precore amino acids prevents particle assembly, and HBeAg is recognized efficiently by HBeAg-specific MAbs but displays little HBe antigenicity. The following HBeAg-derived synthetic peptides representing Th cell recognition sites were used and designated by amino acid position from the N terminus of HBeAg ayw: 120 to 131 (I\textalpha\textsuperscript{a}), VSFGWIRTI PPA; 129 to 140 (I\textalpha\textsuperscript{b}), PPAYRTTPNAPIL; and 120 to 140, a 21-mer comprising both T-cell sites. Peptides were synthesized by the simultaneous peptide synthesis method as previously described (43). Serology. HBeAg was measured in diluted Tg mouse sera by a commercial enzyme-linked immunosorbent assay (ELISA), and recombinant (rHBeAg) was used as a standard. Anti-HBe and anti-HBe immunoglobulin G (IgG) antibodies were measured in murine sera by an indirect solid-phase ELISA by using rHBeAg or rHBeAg as the solid-phase ligand as described previously (29). Serial dilutions of both sera sera and preimmunization sera were made, and the data are expressed as antibody titers representing the reciprocals of the highest dilutions of sera required to yield an optical density at 492 nm (OD\textsubscript{492}) three times an equal dilution of preimmunization sera. IgG isotype-specific ELISAs were performed with IgG1-, IgG2a-, IgG2b-, and IgG3-specific secondary antibodies (Southern Biotechnology, Birmingham, Ala.). In vitro cytokine analysis. Spleen cells from either unprimed or primed TCR-Tg, TCR-Ag double- or triple-Tg, or wild-type mice were cultured (5 \times 10\textsuperscript{6}/ml) with various concentrations of a series of antigens. Culture supernatants were harvested at 48 h for interleukin-2 (IL-2) determination and at 96 h for gamma interferon (IFN-\gamma) determinations. Cytokines were measured by two-site ELISA with pairs of cytokine-specific MAbs. One unlabelled MAb was adsorbed to the microtiter plate well and used as a capture antibody, and the other labeled MAb served as the probe. To determine T-cell proliferation, TCR-Tg spleen T cells were labeled in vitro with the intracellular dye carboxy-fluorescein diacetate succinimidyl ester (CFSE) by using the Vybrant CFSE SE tracer kit (Molecular Probes, Eugene, Ore.,). CFSE (0.5 mM) was added to the cell suspensions, and the mixture was incubated for 10 min at 37°C. The labeling reaction was stopped by repetitive washing with ice-cold RPMI medium–10% fetal calf serum. Labeled cells (5 \times 10\textsuperscript{6}/ml) were cultured with various concentrations of antigen for 4 or 7 days. Cultured cells were harvested, washed, and labeled with anti-TCR V\textsubscript{\beta}11 antibodies. The number of cell divisions are determined by dilution of the intracellular dye CFSE in TCR V\textsubscript{\beta}11+ gated T cells by flow cytometry.

Flow cytometry. Single-cell suspensions of thymus or spleen were prepared. Before staining, cells were incubated with an anti-Fe MAb (2B4G2) to block nonspecific Fe receptor uptake. For staining with directly labeled antibodies, 10\textsuperscript{6}
cells were incubated with antibodies at 4°C for 15 min. Cells were washed three times and analyzed in a FACScan (Becton Dickinson). Gates were set only on viable cells, and usually >10^6 cells were analyzed with LYSIS II (Becton Dickinson). The murine antibodies used for two- and three-color staining were as follows: anti-TCR Vβ4 (KT4), anti-TCR Vβ11 (RR8-15), anti-TCR Vα11 (RR8-1), anti-CD4 (H129.19), and anti-CD8a (53–6.7) (BD Bioscience, Palo Alto, Calif.).

Liver injury model. 7/16-5 TCR × HBe/HBC double-Tg, or TCR × HBe × HBC triple-Tg mice were injected with the 7/16-5 TCR-specific peptide 129 to 140 (50 μg) emulsified in complete Freund’s adjuvant (CFA). At multiple time points, the extent of hepatocellular injury was monitored by measuring serum alanine aminotransferase (sALT) activity. sALT activity was measured in a Kodak EtaChem DTSC II chemistry analyzer (Ortho Clinical Diagnostics, Raritan, N.J.). Normal sALT levels in mice in our colony range from 20 to 40 U/liter; therefore, we designated sALT levels greater than 100 U/liter elevated.

Adoptive transfer. Donor spleen cells (30 × 10^6) derived from 7/16-5 TCR-Tg mice were injected intravenously into CD4/CD8-depleted HBcAg or HBeAg single-Tg recipient mice. The spleen cell inocula contained ~3 × 10^6 TCR+ CD4+ T cells. Whole-spleen inocula were used for convenience and yielded similar results to transferred, purified CD4+ T-cell populations.

**RESULTS**

Differing phenotypes of HBc/HBeAg-specific T-cell tolerance. Three TCR-Tg lineages (Table 2) were selected that represent three phenotypes of antigen responsiveness but more importantly of tolerance induction by the HBc/HBeAgs in TCR/Ag double-Tg pairs. Naïve splenic T cells from 8/12–2 single TCR-Tg mice are efficiently activated by culture with relatively low concentrations of the HBc/HBeAgs as measured by production of IL-2 (Fig. 1, top row), IFN-γ, and proliferation (data not shown). This is especially noteworthy because the frequency of HBc/HBeAg-specific T cells is quite low (<5%) among Vβ4+ cells due to pairing with endogenous Vα chains in this Vβ4+-only TCR-Tg lineage. The estimate of <5% is based on the percentage of CFSE-labeled, 8/12–2 Vβ4+ T cells that undergo cell division in the presence of the HBcAg (data not shown). Naïve splenic T cells from 7/16-5 single TCR-Tg mice demonstrate intermediate HBc/HBeAg-specific IL-2 production in terms of levels of IL-2 produced and minimum antigen dose required compared to the 8/12–2 TCR-Tg lineage. We have previously classified 11/4–12 TCR-Tg T cells as low avidity, consistent with the right-shifted dose-response curve (8). It is also noteworthy that 11/4–12 TCR-Tg splenic T cells preferentially recognize the HBcAg compared to the particular HBcAg, which is a very unusual pattern (Fig. 1, top row).

The three TCR-Tg lineages representing relatively high, intermediate, and low avidity for the HBc/HBeAgs were bred with antigen-Tg mice (Table 1) expressing either high (hi) or relatively low (lo) levels of the HBc/HBeAgs. As shown in Fig. 1 (second row), IL-2 production by naïve splenic T cells of 8/12–2 × HBc(lo) double-Tg mice is significantly reduced (≤300 U/ml) compared to 8/12–2 single TCR-Tg mice at all in vitro HBc/HBeAg concentrations. Predictably, naïve splenic T cells of 8/12–2 × TCR-Tg mice bred with high-serum HBcAg expressers (4 to 10 μg/ml) (Fig. 1, third row) produced absolutely no IL-2 upon culture with the HBc/HBeAgs. An intermediate HBcAg serum concentration of 50 ng/ml, as present in HBV-Tg mice, also eliminated in vitro IL-2 production in 8/12–2 × HBV double-Tg mice (Fig. 1, row 5). This result indicated the tolerogenic potential of serum HBeAg in the context of the other viral proteins and in an in vivo model of HBV replication.

A completely different phenotype is represented by 11/4–12 × HBc/HBeAg double-Tg mice. Regardless of the level of HBc/HBeAg expression in these double-Tg mice, 11/4–12 TCR-Tg T cells were unaffected and produced equivalent levels of HBc/HBeAg-specific IL-2 in vitro, as did single TCR-Tg 11/4–12 T cells (Fig. 1, column 3). We have previously reported that 11/4–12 TCR-Tg T cells are not tolerized in HBc(lo) double-Tg mice (8); however, the observation that 11/4–12 TCR-Tg T cells also ignore the HBc/HBeAgs in high-expresser double-Tg mice is interesting because the serum concentration of HBcAg in the high-expresser Tg mouse (4 to 10 μg/ml) is 10-fold higher than the minimal HBcAg concentration required to activate the 11/4–12 T cells in vitro (0.4 μg/ml).

A more complex phenotype is illustrated by 7/16–5 TCR-Tg T cells. The 7/16–5 T cells appear to ignore low in vivo concentrations of HBcAg (and HBeAg) (data not shown) in terms of in vitro IL-2 production by 7/16–5 × HBe(lo) double-Tg T cells, which is equivalent to IL-2 production in single 7/16–5
Unprimed naive spleen cells (5 TCRTg mice did not spontaneously produce anti-HBc antibody. The OD492 reading of undiluted supernatant in the antigen-specific ELISA. The level of in vitro antibody production is expressed as an OD492 reading of undiluted supernatant in the antigen-specific ELISA. Titers are expressed as the reciprocal of the highest dilution of serum sera to yield an OD492 value three times that of pre-p120-131 injection sera. wks, weeks.

TCR-Tg T cells. However, at the higher in vivo HBcAg concentration present in 7/16–5 × HBc(hi) double-Tg mice, naive T cell HBc/HBeAg-specific in vitro IL-2 production is significantly reduced compared to 7/16–5 single TCR-Tg mice. It is notable that the tolerance exhibited in 7/16–5 × HBc(hi) double-Tg mice is not due to clonal deletion of 7/16–5 TCR-Tg T cells either in the thymus or in the spleen, as determined by fluorescence-activated cell sorter (FACS) analysis (data not shown). It is also of interest that 7/16–5 TCR-Tg T cells are not significantly tolerized in 7/16–5 × HBc(hi) double-Tg mice, as demonstrated by relatively efficient HBc/HBeAg-specific in vitro IL-2 production. In fact, 7/16–5 × HBc(hi) double-Tg mice undergo spontaneous anti-HBc seroconversion between 4 and 6 weeks of age and remain anti-HBc antibody positive (Fig. 2A). Male 7/16–5 × HBc(hi) double-Tg mice spontaneously seroconvert to anti-HBc positivity somewhat earlier than females, which is consistent with higher HBcAg expression in the livers of male mice (data not shown). In contrast, 7/16–5 × HBc(hi) double-Tg mice do not spontaneously produce anti-HBe antibody in vivo. Whereas spleen cells of all 7/16–5 TCR-Tg mice produce IgM anti-HBc antibody in vitro upon culture with the HBcAg, only 7/16–5 × HBc(hi) double-Tg spleen cells produce IgG anti-HBc antibody in vitro (Fig. 2B). The lack of IgG anti-HBe antibody production in vitro by 7/16–5 × HBc(hi) double-Tg spleen cells indicates that the failure to detect anti-HBe in vivo was not due to masking or immune complexing with excess serum HBcAg. Therefore, in vivo tolerance induction of 7/16–5 TCR-Tg T cells in double-Tg mice is nondeletional, HBcAg concentration dependent, and preferentially induced in HBcAg double-Tg but not HBcAg double-Tg mice (i.e., T-cell tolerance is split between the HBcAg and the HBcAg).

Mechanisms of tolerance in double-Tg mice. (i) 8/12–2 × HBcAg double-Tg mice. The fact that 8/12–2 TCR-Tg T cells are of relatively high avidity and are very sensitive to tolerance induction by even low concentrations of serum HBcAg predicts that clonal deletion is a likely mechanism. However, because the frequency of Vβ4+ TCR-Tg T cells specific for the HBc/HBeAgs is so low (<5%) in this Vβ-only TCR-Tg lineage, it is not possible to obtain conclusive evidence of deletion by FACS analysis. Although a decrease in Vβ4+ T-cell frequency is observed in the thymuses and spleens of 8/12–2 × HBc(lo) double-Tg mice compared to 8/12–2 single TCR-Tg mice, the differences are inconclusive. Low serum HBcAg concentrations are sufficient to elicit fatty acid synthase (FAS)-mediated apoptosis in the periphery as shown by comparing B10.S HBc(lo)-Tg mice with FAS-deficient (lpr/lpr) HBc(lo)-Tg mice in terms of T-cell peptide 120 to 131-induced anti-HBe autoantibody production (Fig. 3). The HBc/HBeAg 120 to 131 sequence is the recognition site for 8/12–2 TCR-Tg T
cells and also the predominant T-cell recognition site in wild-type B10.S (H-2b) mice. For example, neonatal tolerance induction with the 120 to 131 peptide renders B10.S mice non-responsive to the entire HBc/HBeAg at the T-cell level (30). Although B10.S/HBe(lo)-Tg mice are highly tolerant to the HBcAg, injection with p120-131 elicits low levels and transient anti-HBe IgG antibody production of primarily the IgG2b isotype. This indicates that at least low-level residual T helper cell function persists in B10.S/HBe(lo)-Tg mice, perhaps reflecting T cells that avoided deletion in the thymus (Fig. 3). B10.S HBe(lo)-Tg mice on a homozygous lpr background (elpr) produce significantly more anti-HBe antibody with a wider IgG isotype distribution, indicating that FAS-mediated apoptosis in the periphery may at least contribute to maintaining T-cell tolerance to the HBeAg in the B10.S strain. We suspect that both thymic (central) and peripheral deletion mechanisms play a role in HBeAg-specific tolerance among high-avidity T cells, as illustrated by the 8/12-2 TCR-Tg lineage and polyclonal HBeAg-specific T cells in B10.S HBeAg-Tg mice.

(ii) 11/4-12 × HBe/HBeAg double-Tg mice. We have previously shown that 11/4-12 TCR-Tg T cells are not deleted in the thymus or spleen of double-Tg mice expressing low levels of the HBe/HBeAgs (8). Further, FACS analysis of high-expresser HBe/HBeAg × 11/4-12 double-Tg mice also revealed no clonal deletion in the thymus or the spleen (data not shown). Furthermore, the function of 11/4-12 TCR-Tg T cells in terms of in vitro cytokine production is not altered by the coexpression of high levels of the HBe/HBeAgs in double-Tg mice. Therefore, low-avidity 11/4-12 TCR-Tg T cells are neither tolerized nor activated by coexpression of even high levels of the HBe/HBeAg and coexist with these viral antigens in vivo by virtue of clonal ignorance.

(iii) 7/16-5 × HBe/HBeAg double-Tg mice. FACS analysis of 7/16-5 × HBe and 7/16-5 × HBe double-Tg mice expressing low and even high levels of the HBe/HBeAgs revealed no clonal deletion in the thymus or the spleen of 7/16-5 TCR-Tg T cells (data not shown). However, functional analysis of 7/16-5 T cells derived from HBe/HBeAg-expressing double-Tg mice revealed apparent clonal ignorance in low-HBe/HBeAg-expressing environments, whereas high serum concentrations of HBeAg in vivo significantly altered 7/16-5 T-cell function (Fig. 1). The tolerogenic effects of secreted HBeAg observed in 7/16-5 × HBe(hi) double-Tg mice were not observed for the HBeAg in 7/16-5 × HBe(hi) double-Tg mice. For example, comparison of the kinetics of HBe/HBeAg-specific in vitro IL-2 and IFN-γ production by naive splenic T cells revealed low-level and transient IL-2 and IFN-γ production by 7/16-5 T cells derived from 7/16-5 × HBe(hi) double-Tg mice compared to 7/16-5 single TCR-Tg and 7/16-5 × HBe(hi) double-Tg mice (Fig. 4). Note that IL-2 production peaked in all cultures at day 2; however, it was significantly lower at day 2 in the 7/16-5 × HBe(hi) T-cell culture and was undetectable by day 4. Similarly, IFN-γ levels were rising in the 7/16-5 and 7/16-5 × HBe(hi) T-cell cultures at day 4 but were declining in the 7/16-5 × HBe(hi) T-cell culture. The in vitro proliferative capacities of 7/16-5 T cells derived from 7/16-5 single TCR-Tg mice and 7/16-5 × HBe(hi) double-Tg mice were also compared. Naive splenic T cells were labeled with the intracellular fluorescent dye CFSE and cultured with p120-140 (0.5 μg/ml) for 4 or 7 days, and the harvested Vβ11+ T cells were analyzed by FACS (Fig. 5). Note that 7/16-5 TCR-Tg T cells derived from 7/16-5 × HBe(hi) double-Tg mice were able to proliferate (as demonstrated by dilution of the CFSE dye) upon p120-140 peptide stimulation, which indicates an absence of a state of profound or absolute anergy in these T cells. Rather, the 7/16-5, Vβ11+ T cells derived from 7/16-5 × HBe(hi) double-Tg mice were substantially slower to enter cell division than 7/16-5 single TCR-Tg T cells after 4 days in culture (i.e., 37.8% nondivided cells compared to 10.7%, respectively). Furthermore, after 7 days in culture, 14.0% of 7/16-5 × HBe(hi)-derived Vβ11+ T cells failed to divide compared to only 2.4% of 7/16-5-derived T cells. Similarly, T cells derived from 7/16-5 × HBe(hi) double-Tg mice underwent fewer total cell divisions than single TCR-Tg T cells (the same for ≥4 and ≥8 cell division windows at days 4 and 7, respectively) (Fig. 5). Therefore, 7/16-5 TCR-Tg T cells are not deleted but are rendered anergic in the presence of high serum levels of HBeAg, as determined by reduced proliferation and cytokine production upon antigen-specific recall in vitro. Two broad categories of T-cell anergy designated either clonal anergy or adaptive tolerance (in vivo anergy) have been described previously (47). The 7/16-5 TCR-Tg T cells in HBe(hi) double-Tg mice appear to exemplify adaptive tolerance because IFN-γ, as well as IL-2, production is inhibited (Fig. 4) and not reversed by exogenous IL-2 (data not shown) and antigen persistence is required to maintain the anergic state in vivo (9). In adaptive tolerance model systems, a phase of clonal activation and expansion.
repertoire in B10.S mice. For this purpose B10.S/HBeAg is relevant to the polyclonal HBeAg-specific T-cell interest to determine whether the tolerogenic potential of the T-cell hybridoma produced in the B10.S strain. It was of highly sensitive to HBeAg-induced tolerance, was derived from Tg mice compared to B10.S/H9252/H11001 antibody production is profoundly reduced in B10.S/HBe(hi)-Tg reactive at the T-cell level. As shown in Fig. 6A, IgG anti-HBc response to the HBcAg, since the two antigens are cross-reactive at the T-cell level. The HBcAg was used as an immunogen because and HBe(hi)-Tg [B10.S/HBe(hi)] mice were immunized with rHBcAg. The HBcAg was used as an immunogen because it affects the HBcAg-specific immune response. This is relevant because both forms of the antigen are present during chronic HBV infection and tolerance split between the HBeAg and the HBeAg is likely. For example, B10.S/HBe(hi)-Tg mice do not spontaneously produce anti-HBe in vivo and B10.S/HBc(hi)-Tg mice do spontaneously seroconvert to anti-HBc (Fig. 6A, inset), suggesting that the HBeAg is less tolerogenic in this polyclonal T-cell model, as previously indicated in TCR clonal T-cell repertoire. Furthermore, the T-cell response to HBcAg, as measured by splenic IFN-γ production recalled by the HBeAg in vitro, was severely reduced in B10.S/HBe(hi)-Tg mice compared to B10.S/+ mice immunized with the HBeAg (Fig. 6B). This indicates that the polyclonal HBC/HBeAg-specific T-cell repertoire in B10.S mice behaves similarly to the 8/12–2 TCR-Tg lineage in terms of its sensitivity to tolerance induction by circulating HBeAg and that tolerance elicited by the HBeAg affects the HBeAg-specific immune response. This is relevant because both forms of the antigen are present during chronic HBV infection and tolerance split between the HBeAg and the HBeAg is likely. For example, B10.S/HBe(hi)-Tg mice do not spontaneously produce anti-HBe in vivo and B10.S/HBc(hi)-Tg mice do spontaneously seroconvert to anti-HBc (Fig. 6A, inset), suggesting that the HBeAg is less tolerogenic in this polyclonal T-cell model, as previously indicated in TCR × HBc double-Tg mice (Fig. 2).

Implications of HBeAg-specific T-cell tolerance for natural HBV infection. (i) Secreted HBeAg is tolerogenic for a polyclonal T-cell repertoire. The 8/12–2 TCR-Tg lineage, which is highly sensitive to HBeAg-induced tolerance, was derived from a T-cell hybridoma produced in the B10.S strain. It was of interest to determine whether the tolerogenic potential of the HBeAg is relevant to the polyclonal HBeAg-specific T-cell repertoire in B10.S mice. For this purpose B10.S/+ wild-type and HBe(hi)-Tg [B10.S/HBe(hi)] mice were immunized with rHBcAg. The HBcAg was used as an immunogen because serum HBeAg present in B10.S/HBe(hi)-Tg mice will not mask or immune complex with anti-HBe antibody to prevent its detection because the two antigens are not cross-reactive at the B-cell level. Further, it was of interest to determine whether potential tolerance to the HBeAg could affect the immune response to the HBcAg, since the two antigens are cross-reactive at the T-cell level. As shown in Fig. 6A, IgG anti-HBc antibody production is profoundly reduced in B10.S/HBe(hi)-Tg mice compared to B10.S/+ mice after HBeAg immunization. Moreover, the T-cell response to HBcAg, as measured by splenic IFN-γ production recalled by the HBeAg in vitro, was severely reduced in B10.S/HBe(hi)-Tg mice compared to B10.S/+ mice immunized with the HBeAg (Fig. 6B). This indicates that the polyclonal HBC/HBeAg-specific T-cell repertoire in B10.S mice behaves similarly to the 8/12–2 TCR-Tg lineage in terms of its sensitivity to tolerance induction by circulating HBeAg and that tolerance elicited by the HBeAg affects the HBeAg-specific immune response. This is relevant because both forms of the antigen are present during chronic HBV infection and tolerance split between the HBeAg and the HBeAg is likely. For example, B10.S/HBe(hi)-Tg mice do not spontaneously produce anti-HBe in vivo and B10.S/HBc(hi)-Tg mice do spontaneously seroconvert to anti-HBc (Fig. 6A, inset), suggesting that the HBeAg is less tolerogenic in this polyclonal T-cell model, as previously indicated in TCR × HBc double-Tg mice (Fig. 2).

(ii) Perinatal exposure to HBeAg is tolerogenic. A number of studies indicate that the HBeAg, either free or complexed to immunoglobulin, can cross the human placenta (1, 21, 24, 51, 54). Previous studies suggested that in utero exposure to HBeAg through placental transfer may result in T-cell tolerance and explain the greater than 90% chronicity rates of infants born to HBV-infected, HBeAg-positive mothers (31). Because the ability of the HBeAg to cross the murine placenta has been questioned (40), we performed an experiment to determine whether early postnatal exposure to the HBeAg may also be tolerogenic. For this purpose, B10.S/HBe(hi)-Tg mice were used because the developmentally regulated major urinary protein promoter is not active until after birth and
B10.S/HBe(hi)-Tg mice are HBeAg negative at birth (18; data not shown) but serum HBeAg becomes detectable by postnatal day 7 and reaches a stable plateau by 4 weeks of age (18). Therefore, young (3 to 4 weeks of age) B10.S/ + or B10.S/HBe(hi)-Tg mice, which were HBeAg negative at birth, were immunized with rHBeAg (10 μg) (incomplete Freund adjuvant). Ten days later, spleen cells were harvested and cultured with various concentrations of the dominant HBC/HBeAg-specific T-cell site in B10.S mice, p120-131. After 2 days of spleen culture, supernatants were collected and analyzed for IL-2 by ELISA. Comparative IL-2 levels were expressed as OD492 readings of undiluted supernatants in the IL-2-and analyzed for IL-2 by ELISA. Comparative IL-2 levels were expressed as OD492 readings of undiluted supernatants in the IL-2-specific ELISA. The symbols represent individual mice. This experiment was performed on two occasions, and the results are representative.

(iii) Immunomodulatory role for HBeAg. Because 7/16–5 × HBe(lo) or 7/16–5 × HBe(lo) double-Tg mice do not appear to be tolerant to the HBe/HBeAgs (Fig. 1), these mice provide a model of T cells of intermediate avidity coexisting with at least low-level expression of their specific antigen. These models provide an opportunity to determine the consequences of activating these apparently quiescent T cells in vivo by injecting the TCR-specific p129-140 peptide. Injection of the TCR-specific peptide activates HBc/HBeAg-specific T cells but not B cells. Anti-HBC/HBeAg-specific antibody production is dependent on B-cell recognition of the endogenous transgene-encoded HBc/HBeAg. A single dose of p129-140 activates TCR T cells in 7/16–5 × HBe(lo) double-Tg mice and results in episodes of sporadic chronic liver injury (sALT elevations) of relatively low magnitude (Fig. 8). Single 7/16–5 TCR-Tg or single HBc(lo)-Tg mice injected with p129-140 do not demonstrate liver injury (data not shown). Note that anti-HBc antibody production correlates with periods of liver injury most likely due to the release of intracellular HBcAg by injured hepatocytes. A similar injection of p129-140 into 7/16–5 × HBe(lo) double-Tg mice results in anti-HBe seroconversion (Fig. 9) but no or only limited liver injury (Fig. 10). Also note that the anti-HBe antibody elicited by p129-140 injection in 7/16–5 × HBe(lo) double-Tg mice is transient, whereas the anti-HBe antibody elicited by p129-140 injection in 7/16–5 × HBe(lo) double-Tg mice is long lived (Fig. 9), suggesting an immunoregulatory function for the HBeAg.

Interestingly, p129-140 injection in 7/16–5 × HBe(lo) × HBe(lo) triple-Tg mice elicited anti-HBe antibodies but not anti-HBc antibodies even though anti-HBc was produced in 7/16–5 × HBe(lo) double-Tg mice (Fig. 9). This suggests that the presence of HBeAg in the serum can down-regulate antibody production to another viral protein, the HBcAg. Because the HBcAg and HBcAgs are cross-reactive at the level of T-cell recognition, the regulation by serum HBeAg is most likely mediated at the T-cell level. This was confirmed by examining liver injury in double- and triple-Tg mice (Fig. 10). Injection of 2 doses of p129-140 resulted in a relatively high frequency of ALT elevations in a group of six 7/16–5 × HBe(lo) double-Tg mice and a low frequency of liver injury in a group of six 7/16–5 × HBe(lo) double-Tg mice. Because of the sporadic and unpredictable nature of liver injury and the fact that each mouse could not be bled each week, 2 to 3 mice per group were bled on alternate weeks and the data are presented as the frequency of sALT elevations (ALT ≥ 100 U/liter) observed in the combined bleeds in each group during months 1, 2, and 4. As shown in Fig. 10, p129-140 injection in 7/16–5 × HBe(lo) × HBe(lo) triple-Tg mice resulted in a relatively low frequency of sALT elevations, indicating that p129-140-induced liver injury was inhibited by the presence of HBeAg in the serum. These data suggest that the HBeAg may serve an immunomodulatory role in natural HBV infection and that secreted HBeAg can down-modulate the immune response to the HBeAg. Because the HBcAgspecific T-cell
response is predominantly Th1-like (34), inhibition of this response may cause predisposition to chronicity.

(iv) T-cell tolerance split between HBeAg and HBcAg. A number of experiments described herein suggest that secreted HBeAg is a superior T-cell tolerogen compared to the intracellular HBcAg, even though the two forms of the antigen are cross-reactive at the T-cell level. T-cell tolerance split between the HBc/HBeAgs is best exemplified in studies of 7/16–5/H11003 HBc or HBe double-Tg mice expressing high and low levels of the HBc/HBeAgs. Figure 11 represents a summary of studies comparing 7/16–5/H11003 HBc and 7/16–5/H11003 HBe double-Tg mice in terms of spontaneous anti-HBc/HBe seroconversion, induced anti-HBc/HBe antibody production in vivo, and the ability of naive 7/16–5 TCR-Tg T cells to elicit antibody production after adoptive transfer into HBcAg- or HBeAg-Tg recipients. As shown in Fig. 2A, 7/16–5/HBe(hi) double-Tg mice spontaneously seroconvert to anti-HBc positivity between 4 and 6 weeks of age and 7/16–5 TCR+ double-Tg mice expressing a lower level of HBeAg do not. Therefore, the low HBcAg level appears limiting for immunogenicity, and the high level of HBeAg is immunogenic rather than tolerogenic. In contrast, 7/16–5 TCR+ double-Tg mice expressing either high or low levels of the secreted HBeAg do not spontaneously produce anti-HBe (Fig. 11, top panel). Induced anti-HBe/HBe antibody production in TCR/Ag double-Tg mice was examined by injecting the 7/16–5 TCR-specific peptide p129-140. High- and low-HBcAg-expressing double-Tg mice could be induced to produce anti-HBc and high HBcAg expression correlated with higher titer. In contrast, only low expressers of the HBeAg could be induced to produce anti-HBe antibody, consistent with the concentration dependence of T-cell tolerance to the secreted HBeAg (Fig. 11, middle panel). The ability to induce anti-HBe antibodies by p129-140 injection in 7/16–5/H11003 HBe(lo) double-Tg mice indicates that even this low concentration of serum HBeAg (10 to 20 ng/ml) is not limiting as an immunogen and further indicates that the lack of anti-HBe production at the higher level of HBeAg is due to tolerance mechanisms as opposed to limiting amounts of antigen. Lastly, adoptive transfer of naive 7/16–5 TCR-Tg spleen cells into HBcAg-Tg recipients elicits efficient anti-HBc antibody production in high-HBcAg-expressing recipients and lower levels of anti-HBe antibody in low-HBcAg-expressing recipients. In contrast, adoptive transfer of the same number of 7/16–5 TCR-Tg spleen cells into high-HBeAg-expressing Tg recipients elicited no anti-HBe antibody production and only very low anti-HBe antibody production in HBeAg(lo) expresser Tg recipients (Fig. 11, bottom panel). This indicates that serum HBeAg need not be present at birth or even perinatally to function as a tolerogen. Rather, exposure of adult, naive 7/16–5 TCR-Tg T cells to circulating HBeAg for a relatively brief period of time in the HBeAg-Tg recipients is sufficient to render the T cells tolerant or at least incapable of mediating
production and proliferative T-cell responses observed in 7/16-5 × HBe(hi) double-Tg mice, the failure to produce either spontaneous or induced anti-HBe antibodies indicates that serum concentrations of HBeAg of 4 to 10 μg/ml are highly tolerogenic for 7/16-5 TCR-Tg T cells. No similar tolerance induction in the same 7/16-5 TCR-Tg T cells was observed in response to the HBcAg regardless of the level of antigen expression. This apparent split tolerance between the HBeAg and HBcAg is likely due to the efficient secretion of monomeric HBeAg as opposed to the intracellular location of the particulate HBcAg. Secreted HBeAg has access to the thymus and the secondary lymphoid system, whereas only very small amounts of the HBcAg exit the liver. Monomeric HBeAg is significantly less immunogenic than particulate HBcAg at the T- and B-cell levels even when formulated in adjuvants (29). Furthermore, monomeric secreted antigens tend to be more tolerogenic than immunogenic in general (41, 52).

**DISCUSSION**

The purpose of this study was to gain insight into the condition of the HBc/HBeAg-specific CD4+ T-cell repertoire during long-term exposure to the secreted and intracellular forms of the HBV nucleocapsid antigens, as occurs in chronically infected patients. As a model of chronic HBc/HBeAg exposure, we have used TCR × HBc or HBeAg double-Tg mice. The function of the secreted HBeAg has been the subject of much speculation, since it is not required for viral assembly, infection, or replication. We have previously suggested that the HBeAg acts as an immunomodulatory protein via the induction of tolerance and Th1/Th2 cross-regulation (35, 36). In natural HBV infection, a dichotomy exists between the apparent tolerogenic and immunogenic functions of the HBeAg (28). In this present study, several mechanisms are described which may help to explain this dichotomy.

Serum HBeAg was shown to be tolerogenic for HBc/ HBeAg-specific TCR-Tg T cells, and three distinct phenotypes of tolerance were observed in double-Tg mice. One phenotype, illustrated by the 8/12-2 TCR-Tg lineage, was profound tolerance induction by the HBeAg even at very low serum concentrations (10 ng/ml), most likely mediated by clonal deletion and FAS-mediated apoptosis in the periphery. A second phenotype, represented by the 11/4-12 TCR-Tg lineage, was profound tolerance induction in the same 7/16-5 TCR-Tg cells was observed. The symbols in this figure represent individual mice; however, the results are representative of experiments performed on multiple occasions with a minimum of 6 mice per group. wk, week.

anti-HBe antibody production. The absence of serum anti-HBe antibodies was not the result of masking by an excess of serum HBeAg because no IgG anti-HBe antibody production occurred in cultures of the HBeAg-Tg recipient spleen (data not shown). Although relative comparisons of the concentration of serum HBeAg versus cellular HBcAg are not possible, the positive correlation between in vivo HBcAg concentration and immunogenicity and conversely the inverse correlation between HBeAg serum concentration and immunogenicity is noteworthy. Taken together with the low levels of cytokine

![Image](https://via.placeholder.com/150)
terms of sensitivity to tolerance induction. What explains multiple HBeAg-specific T-cell tolerance phenotypes? The immunogenicity and tolerogenicity of a T-cell epitope are likely to be proportional at least over a large range of TCR-peptide-major histocompatibility protein (TCR-pMHC) avidities. For example, a high-avidity TCR-pMHC interaction is likely to result in strong immunogenicity for a given T-cell epitope unless that epitope is a self-epitope, in which case T-cell tolerance is the likely outcome. An important contributor to overall TCR-pMHC avidity is the affinity of the peptide-MHC interaction.

In previous studies, we have demonstrated that the immunodominant HBeAg-specific 120–131/I-As-specific 7/16–5 TCR-Tg lineage are highly sensitive to HBeAg-induced T-cell tolerance and the 129–140/I-Ab-specific 7/16–5 and 114–12 TCR-Tg lineages are less sensitive and demonstrate different tolerance phenotypes. We have functionally classified the 8/12–2, 7/16–5, and 114–12 TCRs as high, intermediate, and low avidity, respectively, based on parameters of T-cell activation, and this hierarchy also appears to correlate well with sensitivity to T-cell tolerance.

In the context of chronic HBV (CHB) infection, a multiplicity of HBeAg-specific T-cell clones are likely to coexist even within the same patient, and the balance between activation or tolerance status among the heterogeneous HBeAg-specific T-cell repertoire may influence degrees of liver injury and viral clearance. The question of whether the HBeAg acts as a tolerogen or an immunogen during HBV infection must be addressed at the clonal level. In the so-called tolerance phase of CHB infection, the majority of HBeAg-specific T-cell clones would be expected to be tolerant, although the mechanism of tolerance may vary (i.e., deletion, anergy, or ignorance). The status quo of the various clonal tolerance phenotypes would likely be maintained as long as the HBeAg concentration and/or the noninflammatory hepatic environment remains unchanged. However, a nonspecific increase in hepatic inflammation or a decrease in HBeAg serum concentration, perhaps due to the emergence of core promoter region (4) or precore region (37) mutants, may allow activation of low-avidity ignorant HBeAg-specific T cells and/or reverse the anergic state of others, respectively. Such a shift from HBeAg-specific T-cell tolerance to T-cell activation may precipitate the so-called clearance phase of CHB infection. As previously discussed, high-avidity HBeAg-specific T-cell clones are likely to be physically or functionally deleted and not available to participate in antiviral clearance mechanisms after a chronic infection has become established. Intermediate- or low-avidity HBeAg-specific T-cell clones that are not physically deleted at least have the potential to be activated either through the reversal of adaptive tolerance (anergy) or the primary activation of ignorant HBeAg-specific T cells. Therefore, treatment modalities for chronic HBV infection should be directed at activating the relatively low-avidity HBeAg-specific T cells. Given the efficacy of the present antiviral drugs in terms of reducing HBV and antigen load, reversing HBeAg-specific adaptive tolerance by reducing or eliminating circulating levels of the HBeAg may be an achievable goal and may in fact explain viral clearance in a percentage of chronic patients on long-term antiviral therapy. As demonstrated by 7/16–5 TCR-Tg T cells, the total elimination of the tolerogen (HBeAg) may not be required, since maintenance of HBeAg-specific T-cell tolerance for these T cells requires relatively high concentrations of HBeAg. Furthermore, if circulating HBeAg can be eliminated for a sufficient period of time, even deleted high-avidity HBeAg-specific T-cell clones may repopulate the peripheral T-cell repertoire, possibly depending on the age of the patient. Because antiviral therapy reduces the immunogenic stimulus as well as the tolerogenic stimulus, a dual antiviral/vaccine strategy may be necessary in the treatment of chronic HBV. Such a treatment strategy has shown some efficacy in a woodchuck model of chronic infection (27).

The clonal heterogeneity of HBeAg-specific T-cell tolerance may also explain how a primarily tolerogenic protein can exert pressure on the immune response to select an HBeAg-negative mutant. For example, high-avidity HBeAg-specific T-cell clones may be tolerized and simultaneously lower-avidity T-cell clones may be activated and involved in selecting HBeAg-negative mutants in the same patient.

Why is the reversal of T-cell tolerance to the secreted, nonstructural HBeAg an important goal in terms of treating chronic HBV infection? The HBeAg and HBcAg are cross-reactive at the level of T-cell recognition, and T-cell tolerance to the HBeAg would be expected to pertain to the HBcAg as well. However, the present studies demonstrate a split in T-cell tolerance between the HBeAg and the HBcAg. The secreted HBeAg appears significantly more efficient at eliciting T-cell tolerance than the HBcAg. If T-cell tolerance towards the nucleocapsid of HBV is an important determinant of chronicity, a secreted tolerogenic form of the antigen may be necessary to insur virological persistence and may actually be the primary function of the HBeAg in the virus life cycle. During a chronic infection with wild-type HBV, both immunogenic (HBcAg) and tolerogenic (HBeAg) forms of the same antigen coexist. As shown herein, the presence of serum HBeAg can inhibit anti-HBc antibody production in B10.S/HBeAg-Tg mice immunized with rHBcAg (Fig. 6) and in 7/16–5 × HBcAg-lo triple-Tg mice (Fig. 9). Furthermore, the presence of serum HBeAg reduced the frequency of HBcAg/HBeAg-specific liver injury in triple-Tg mice (Fig. 10) as well as down-regulated anti-HBc antibody production. We suggest that the HBeAg functions as an immunoregulatory protein by virtue of eliciting tolerance (i.e., deletion and/or anergy) in CD4+ T cells specific for both the Hbc/HBeAg in a manner which is not likely to occur to the HBcAg alone. The effects of HBeAg-specific T-cell tolerance may be apparent in utero (31), during the perinatal period (Fig. 7), or during adult immunization and/or infection, as demonstrated by the absence of anti-HBe antibody production in HBeAg(hi)-Tg recipients of 7/16–5 TCR-Tg T cells (Fig. 11, bottom panel).

A number of clinical observations suggest that the HBeAg plays a role in HBV chronicity (19, 28). For example, neonatal
infection with an HBeAg-negative mutant virus often results in acute fulminant rather than chronic HBV infection (7, 49). Similarly, infection of young woodchucks with a WHAg-negative mutant of woodchuck hepatitis virus results in a low rate of chronic infection in contrast to the high rate of chronicity observed in young woodchucks infected with wild-type woodchuck hepatitis virus (12). Furthermore, adult infection with an HBeAg-negative mutant is often associated with a fulminant course of infection rather than the relatively benign acute course which characterizes most adult-onset infections with wild-type HBV (5, 13, 25, 39, 48). In addition, emergence of an HBeAg-negative mutant during a chronic active HBV infection, especially in the presence of a high viral load, can correlate with an exacerbation of liver injury and a worse prognosis (3, 11). A number of relatively common mutations in the precore and core promoter regions inhibit HBeAg production to correlate between precore and core promoter mutations and core and core promoter regions inhibit HBeAg production to (3, 11). A number of relatively common mutations in the precore and core promoter regions inhibit HBeAg production to correlate between precore and core promoter mutations and core and core promoter regions inhibit HBeAg production to (3, 11). A number of relatively common mutations in the precore and core promoter regions inhibit HBeAg production to correlate between precore and core promoter mutations and core and core promoter regions inhibit HBeAg production to (3, 11). A number of relatively common mutations in the precore and core promoter regions inhibit HBeAg production to correlate between precore and core promoter mutations and core and core promoter regions inhibit HBeAg production to (3, 11). A number of relatively common mutations in the precore and core promoter regions inhibit HBeAg production to.