

Superantigen Expression Is Driven by Both Mouse Mammary Tumor Virus Long Terminal Repeat-Associated Promoters in Transgenic Mice

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In addition to the usual retroviral promoter, the mouse mammary tumor virus (MMTV) long terminal repeat carries a second promoter located in the U3 region. Here we show that both of these promoters are independently able to give rise to superantigen activity in transgenic mice. The ability of multiple MMTV promoters to drive superantigen expression underscores its importance in the virus life cycle.

The long terminal repeat (LTR) of mouse mammary tumor virus (MMTV) encodes a superantigen (Sag). After infection of mice with exogenous virus, expression of Sag in antigen-presenting cells, such as B lymphocytes, specifically stimulates the proliferation of whole classes of T cells bearing the cognate V β chain as part of their T-cell receptor (1). This T-cell stimulation results in local cytokine production and, in turn, proliferation of B cells in the vicinity, including those that were infected with MMTV. The virus thus uses the host immune system to establish and amplify a reservoir of infected cells. Later, the virus is passed to mammary epithelial cells by an as-yet-unknown mechanism and causes mammary tumors in susceptible mice (12).

Expression of Sag from endogenous, germ line-transmitted MMTVs leads to deletion and/or anergy of specific reactive V β -bearing classes of T cells (1). The deletion thereby protects mice from subsequent challenge with an MMTV encoding a Sag with the same V β specificity (4).

MMTV is unusual among retroviruses in that it carries, in addition to the classic retroviral promoter (P₁₁₉₆), a second promoter (P₆₉₈) (Fig. 1) (8, 16, 17). Intriguingly, both of these promoters independently give rise to Sag activity in mixed-lymphocyte reactions when coupled 5' to a Sag encoding the 3' LTR (16, 17). To determine which of the two LTR-associated promoters is used *in vivo* for Sag expression, a number of transgenic mouse lines carrying the coding region for the Sag of the Mtv-2 provirus linked to either P₁₁₉₆, P₆₉₈, or both promoters were established. Construction of the plasmids pORFexp, p Δ U3, and p Δ RU5 has been previously described (17). The inserts containing the miniproviruses were excised from each plasmid and used for microinjection into freshly isolated C57BL/B6 mouse oocytes. These were reimplanted after 12 h of cultivation into recipient pseudopregnant NMRI mice as previously described (3). DNA was prepared from tail clips of 6-week-old offspring and analyzed by PCR using the previously described primers located at +702 or +1211 together with primer -1730 (8) or a new primer, -3302 (5'-G

CAACTTCCCCCAATAGCC-3') (Fig. 1). Since all mice carry endogenous MMTV copies, signals specific for the deleted miniproviruses were indicative of transgenicity. Mice transgenic for p Δ RU5 gave an indicative 0.5-kb fragment with the primers +702 and -1730 as the result of the deletion of the R and U5 sequences. Both the pORFexp and the p Δ U3 mice gave an indicative 2.1-kb fragment with the primers +1211 and -3302 as the result of the deletion of the *pol* and *env* sequences (data not shown).

Expression of the Mtv-2 Sag results in the deletion or anergy of V β 14-bearing T cells (8). After 100 μ l of blood from the retro bulbar plexus of anesthetized transgenic mice was sampled using a glass capillary, the T cells were stained with R-phycoerythrin-labeled anti-CD3 monoclonal antibody and a fluorescein-conjugated anti-V β 14 monoclonal antibody and analyzed by flow cytometry (fluorescence-activated cell sorting) (Elite Coulter Inc.) to determine the percentage of V β 14⁺ T cells. Table 1 shows that these T cells were deleted (between <1 and 1.4%) in mice regardless of the promoter carried by the construct but that this deletion did not occur in nontransgenic mice (8.3%). Thus, both LTR-associated promoters have the potential to give rise to Sag activity *in vivo* in the temporal window during ontogeny, allowing deletion of V β 14-bearing T cells. Additionally, these results show that both promoters are active in the antigen-presenting cells involved in the establishment of Sag-mediated tolerance. These data are consistent with previous data obtained with a mixed-lymphocyte reaction *in vitro* (17) in that the p Δ RU5 construct, which carries only the new P₆₉₈ promoter, shows the best Sag activity. Although we cannot rule out the possibility that the sequences deleted in our constructs normally silence one promoter or the other, this seems unlikely, since transcripts from both promoters can be detected in B cells (and mammary tumor GR cells) carrying the Mtv-2 locus (8).

Evidence has been presented that a promoter located within the *env* coding sequences (P₇₂₄₆) (10, 15, 18), in association with an enhancer element located in the *pol* region (E_{pol}), directs between 98 and 99.7% of Sag gene expression in B cells (11). However, these elements are not included in the constructs described here (Fig. 1). There is an additional promoter (P₈₄₉₈) in our construct; however, only low-level expression (0.3% maximum) has been attributed to this promoter (11).

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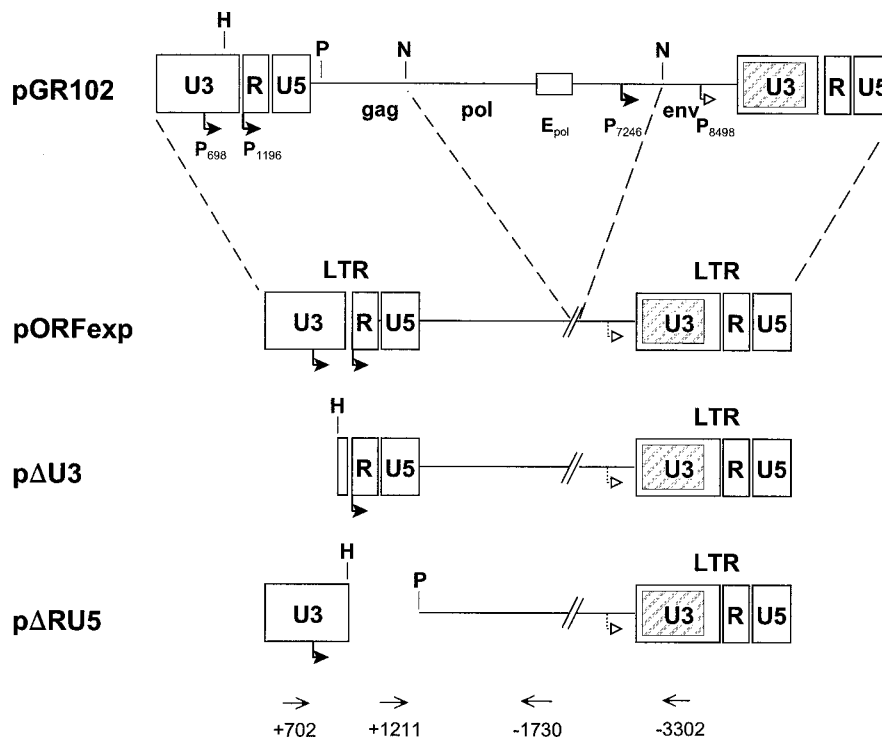


FIG. 1. Schematic representation of the MMTV constructs used. The plasmid pGR102 carries a hybrid provirus in which the 5' half is from Mtv-8 and the 3' half is from Mtv-2 (13). The Sag, encoded by the open reading frame located in the 3' LTR (shaded box), is thus derived from the Mtv-2 provirus and specifically interacts with Vβ14-bearing T cells (2). The plasmid pORFexp was derived from pGR102 by deletion of the internal *Nco*I fragment. Also shown are the locations of the classic promoter (P₁₁₉₆) and other promoters present in the LTR (P₆₉₈) or in the *env* gene (P₇₂₄₆ and P₈₄₉₈) (nomenclature as given in reference 11). The enhancer (E_{pol}) shown to be important for Sag activity from P₇₂₄₆ is indicated as an open box, as are the *Hpa*I (H), *Pvu*II (P), and *Nco*I (N) restriction sites used in the construction of the plasmids pORFexp, pΔU3, and pΔRU5. At the bottom of the figure, the positions of the primers used for the PCR analysis are shown.

Thus, taken together, our data suggest that, in the absence of the P₇₂₄₆ promoter, either of the LTR promoters can also drive expression of a functional Sag in vivo, during ontogeny. We have previously shown the presence of spliced messages from these promoters to the Sag open reading frame (8, 9).

The presence of multiple promoters in MMTV may act as a fail-safe feature, ensuring Sag expression. Such Sag expression from endogenous MMTVs protects mice against infection with exogenous virus since the cognate T-cell classes are deleted (4), whereas the use of redundant promoters in exogenous viruses may ensure establishment of infection. Further, transcripts from the P₆₉₈ promoter in particular may be important for the generation of new MMTV variants, in order to overcome infection restrictions imposed by the Sag-mediated T-cell deletion (14). Such MMTV variants arising from recombinations between pre-existing endogenous and exogenous viruses have been recently detected (5–7).

TABLE 1. Levels of Vβ14⁺ T cells in normal and transgenic mice

Construct	LTR promoter(s) present	Mean % Vβ14 ⁺ T-cells ^a
Nontransgenic	None	8.3
pORFexp	P ₆₉₈ and P ₁₁₉₆	1.3
pΔU3	P ₁₁₉₆	1.4
pΔRU5	P ₆₉₈	<1.0

^a Data are the percentage measured in nontransgenic mice and the averages of the measurements for at least five transgenic mice from each line carrying a construct.

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