









Figure 21.5. Double replacement gene targeting can be used to introduce subtle mutations. Both the methods in Figure 21.4 result in introduction of a substantial amount of exogenous sequence within the endogenous gene. To introduce a subtle mutation without leaving residual exogenous sequence, a double replacement method with positive and negative selection can be used (Melton, 1994). Exons in the endogenous gene are represented as numbered large boxes, and introns as long thin boxes. In order to introduce a subtle mutation, such as a single nucleotide substitution in exon 8, a replacement knock-out vector is used with a marker gene (e.g. the HPRT gene) flanked by homologous sequences from introns 7 and 8, and a second marker, such as the herpes simplex thymidine kinase (TK) gene outside the homologous region. Gene conversion, or double crossover within the flanking intron sequences, can lead to replacement of exon 8 by the HPRT gene, and can be selected for if a mutant HPRT ES cell is used. A positive-negative selection system can be used: selection in the first step is for HPRT+TK cells. Cells containing random vector integrations will contain the TK gene and can be killed with the thymidine analog gancyclovir (see Figure 22.13). The second replacement involves introducing an altered exon 8 with a point mutation (*) to replace the HPRT gene and can be screened by identifying HPRT cells. Note that mice engineered in this way cannot be described as transgenic because of the lack of foreign sequences in the germline.





















