“An axonemal dynein at the *Hybrid Sterility 6* locus: implications for *t* haplotype-specific male sterility and the evolution of species barriers” by J. Fossella et al. (*Mammalian Genome*, volume 11, number 1, January 2000, pp 8–15)

Based on recent extensive sequence analysis of *Dnahc8* genomic and cDNA clones, please note following correction to the afore-mentioned article: the authors found that the mRNA Differential Display-PCR clone, *Hst6.7*, originally assumed to be part of the 3' untranslated region of *Dnahc8*, actually derives from what are most likely translated exons from the N-terminal 17% of the *Dnahc8* coding sequence. All previous submissions to GenBank concerning *Hst6.7* have been updated to reflect this new information.