Tables:

Table S1: Read-pairs per Library

Table S2: Coverage per Distinguishing SNP by Sex

Table S3: Expressed Genes per Library

Table S4: All Detected Transcripts

Table S5: Male-Derived Transcripts

Table S6: Concordance Within a Cross

Table S7: Concordance Between RAL 304 and RAL 307

Table S8: Male Derived Transcript Abundance in Mean Counts per SNP