SUPPLEMENTAL MATERIAL - FIGURE LEGENDS


Figure S1. Allele combination counts of SNPs in the M. fascicularis genome draft

Figure S2. Tissue specific gene expression in M. fascicularis

Genes with tissue specific expression and robust expressions level (EST-counts) were selected from NCBI's Unigene EST database (http://www.ncbi.nlm.nih.gov/UniGene/). The expression levels (log2) of M. fascicularis orthologs of 6 liver (F7, TTR, AGT, APOB, CRP, CYP3A5) and 7 testes (KIF2B, TNP2, SPATA20, SPATA21, SPZ1, TSGA13, TSKS) specific genes is shown in liver, testes, bladder, spleen and kidney samples from three animals. The expression levels of three housekeeping genes (GAPDH, POLR2B and POLR2C) are shown as reference and the detection limit of the microarray platform is depicted by a hatched line.