



Fig. S2. Evolutionary conservation of the *Cyp24a1* gene. Analysis of the *Cyp24a1* gene including 5'-500 bp flanking transcriptional start and 3'-1,000 bp were compared between human and mouse. The analysed SNPs are located as follows: rs2248359 (3,972 bp) and rs2296241 (9,271 bp). Pink areas indicate evolutionary conserved regions (>75% homology). Sequence displayed in 5'-3'-format.