

Figure S1. Expression of mmu_circ_0000712, mmu_circ_0000898, novel_circ_0001857 and novel_circ_0001778 in kidney tissue of *db/db* mice. Reverse transcription-quantitative PCR estimated the expression levels of (A) mmu_circ_0000712, (B) mmu_circ_0000898, (C) novel_circ_0001857 and (D) novel_circ_0001778 in diabetic nephropathy mice. ** $P < 0.01$ *db/db* group vs. *Db/dm* group. circRNA, circular RNA.

