## Fig. S1 Hub gene networks.



The relationship of candidate gene, COX1 and COX3, with other genes

The relationship of candidate gene, DHRS4, with other genes



The relationship of candidate gene, HSD3B, with other genes



The relationship of candidate gene, MSMO1, with other genes



The relationship of candidate gene, NR4A1, with other genes



The relationship of candidate gene, PTX3, with other genes



AVP PTGIR PTGIR VINSL3 CCRHR2 VINSL3

The relationship of candidate gene, RLN2, with other genes

The relationship of candidate gene, SELENOP, with other genes



The relationship of candidate gene, STAR, with other genes



The relationship of candidate gene, CYP11A1, with other genes



The relationship of candidate gene, SCARB1, with other genes



The hub genes for up-regulated DEGs



The hub genes for down-regulated DEGs







The five types AS events were verified by RT-PCR method taking the specific primers outside of AS region (panel A to E), which were detected by 2.5% agarose gel. Diagram of AS types and event locations were arranged on the right. The positions of specific primer were indicated under the exon box in red arrows. Lane 1-6: samples from XS group; Lane 7-12: samples from XL group. A: The A3SS event in exon13 of *LTBP1* were detected with the band of 167 bp in length. B: The band with 194 bp in length showed the A5SS event in exon7 of *SOX5* transcript. C: Fragments with 313/317 bp in length were the MXE event spanned exon3 (151 bp) or exon4 (155 bp) of *CYP19A1*. D: The band in 201 bp denoted the SE event spanned exon27 (126 bp) of *LTBP1*. E: The band (241 bp) showed RI event retained in intron 12 (144 bp) of *HEXB*. F: The expression level of gene transcripts contained AS events were determined by qRT-PCR with one of primer spanned the junction of AS event listed in Table S1.