



**S6 Fig.** Gene ontology (GO) analysis of the differentially expressed genes in uterine sarcomas (US). The GO categories include biological processes (BP), cellular components (CC), and molecular functions (MF). (A, B) GO results of the up- (A) and down-regulated (B) genes in endometrial stromal sarcoma (ESS) vs. uterine leiomyosarcoma (uLMS). (C, D) GO results of the up- (C) and down-regulated (D) genes in high-grade endometrial stromal sarcoma (HGESS) vs. low-grade endometrial stromal sarcoma (LGEES).