

Figure S1. The flowchart of constructing the web tool OSacc.

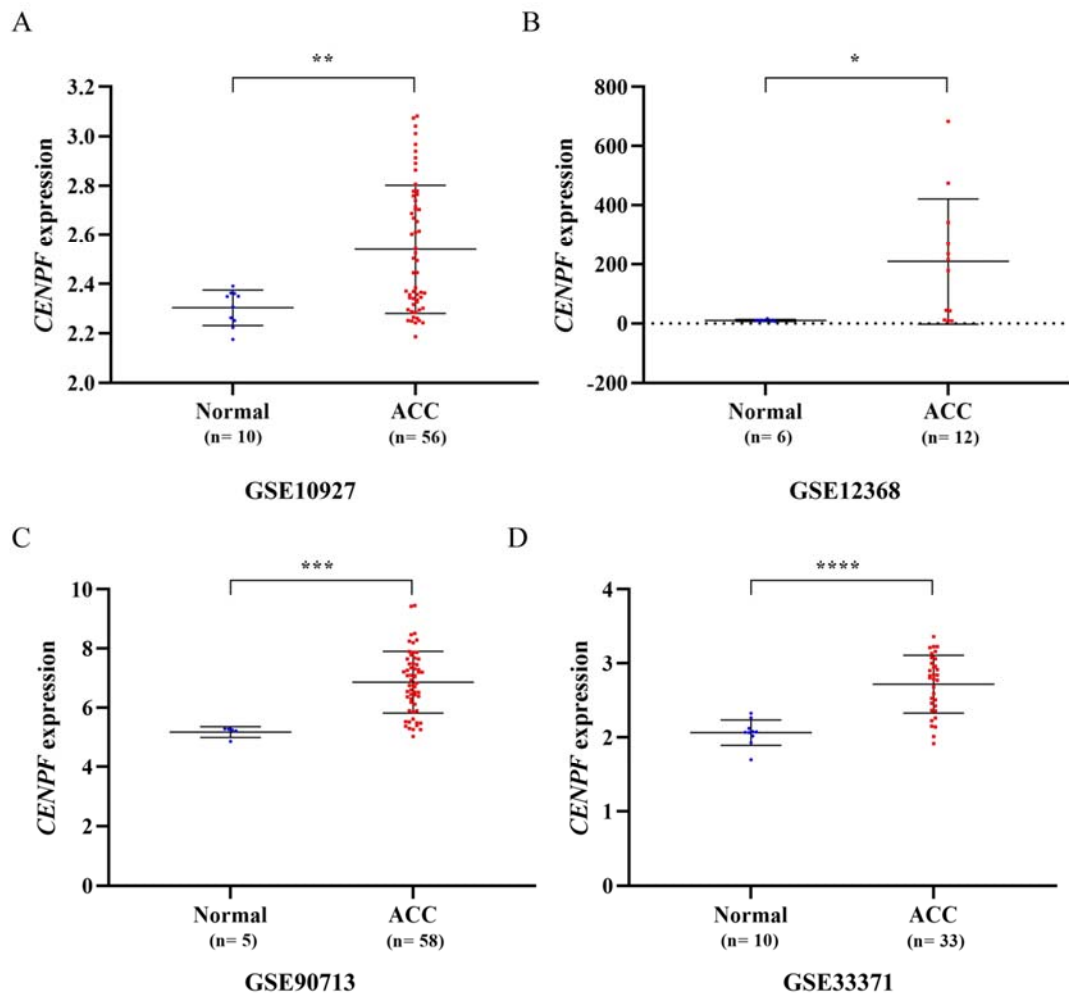


Figure S2. Gene expression of *CENPF* in 4 GEO datasets (GSE10927, GSE12368, GSE90713 and GSE33371). The four datasets all show *CENPF* highly expressed in ACC tissues compared with normal tissues. \* represents  $p < 0.05$ ; \*\* represents  $p < 0.01$ ; \*\*\* represents  $p < 0.001$ ; \*\*\*\* represents  $p < 0.0001$ .