Supplementary Figure 4
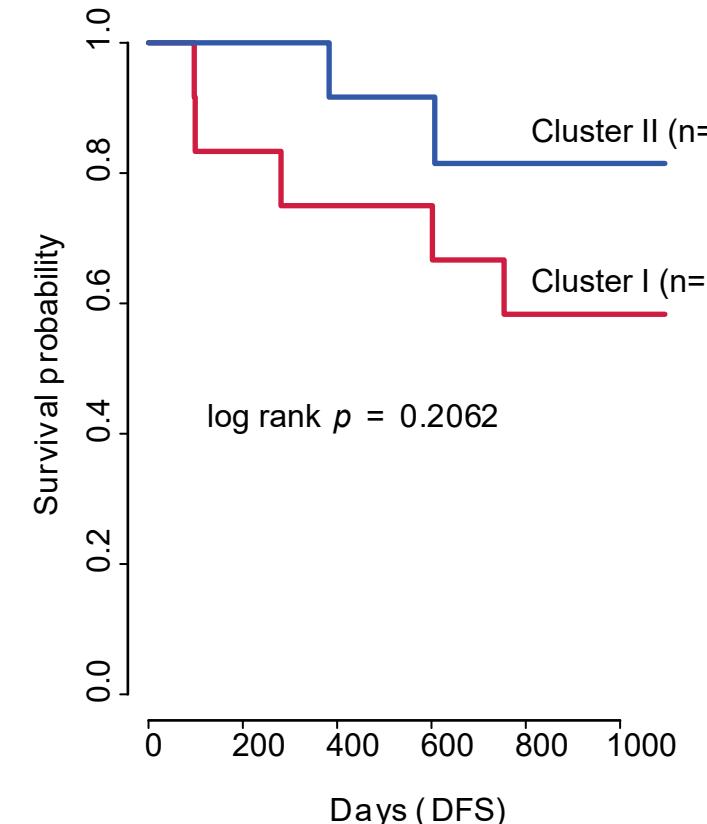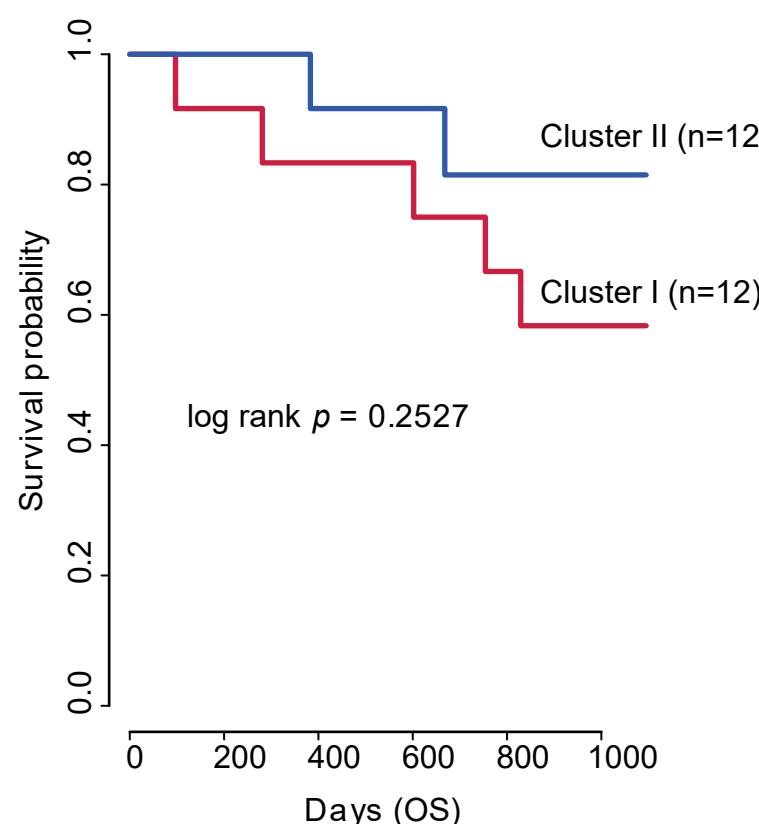
A

B

PSAT1

C.

Enrichment plot:
KEGG_UBIQUITIN_MEDIATED_PROTEOLYSIS

NES=1.42
FDR=0.18
p-value=0.03

D

LUSC with good prognosis

LUSC with poor prognosis

SAE1^low (n=12)

SAE1^high (n=58)

log-rank *p* =0.0095

Supplementary Figure 4. Analysis of two clusters.
A. The prognosis of two clusters within 3-year follow-up.
B. PSAT1 expression between Cluster I and Cluster II.
C. Interaction of proteins that were enriched in the Ubiquitin mediated proteolysis pathway. The minimum required interaction score is 0.7 which means high confidence.
D. SAE1 expression and survival analysis in tissue microarray (TMA) assay.