Figure S1. The expression of *SCGB2A1* mRNA in cancer cell lines from the Broad Institute Cancer Cell Line Encyclopedia database. The x axis represents different cancer cell lines, and the y axis represents *SCGB2A1* mRNA expression values. *SCGB2A1*, secretoglobin family 2A member 1.

mRNA expression (RNAseq): SCGB2A1

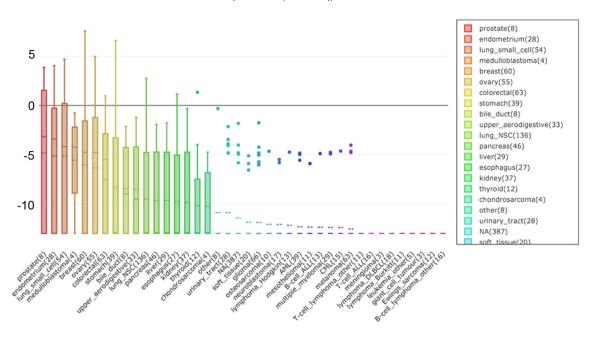


Figure S2. *SCGB2A1* expression in two patients with endometrial cancer detected by immunohistochemistry from the The Human Protein Atlas database. The expression of *SCGB2A1* in endometrial cancer was strong positive in brown color. *SCGB2A1*, secretoglobin family 2A member 1.

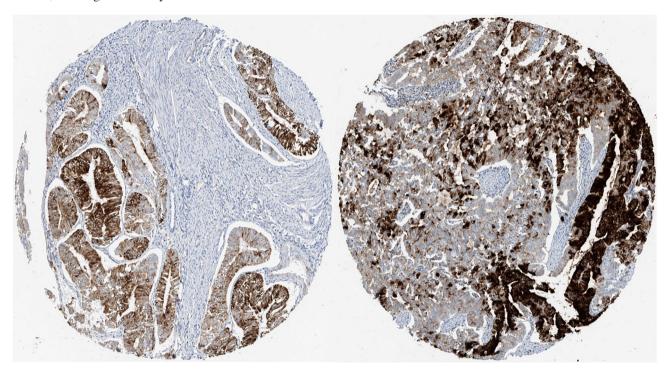


Figure S3. DNA methylation analysis of SCGB2A1 in UCEC using the Mexpress tool. The solid lines represent six SCGB2A1 methylation sites and r-value represents the correlation between SCGB2A1 DNA methylation and SCGB2A1 gene expression regulation. Probe 1 =cg01919620, r=-0.017, P>0.05; probe 2 =cg05277881, r=-0.339, P<0.001; probe 3 =cg16986846, r=-0.210, P<0.001; probe 4 =cg06334737, r=-0.295, P<0.001; probe 5=cg23206745, r=-0.378, P<0.001 and probe 6=cg14265033, r=-0.198, P<0.001. SCGB2A1, secretoglobin family 2A member 1.

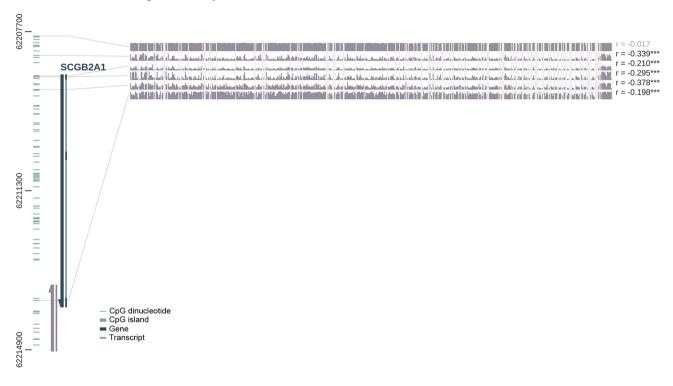


Table SI. Clinical characteristics of 47 patients with endometrial cancer from the Fudan University Shanghai Cancer Center.

2 (42-69) 0 (0) 0 (0) 7 (100)
0) (0)
0) (0)
7 (100)
2 (25.53)
5 (74.47)
7 (100)
3(17.02)
9 (82.98)
0) (0)
8 (80.85)
9 (19.15)
0) (0)
3 (91.49%)
4 (8.51%)
2 (4.26)
5 (95.74)
7 (100)
0) (0)
7 (100)
(0)
3 (91.49)
1 (8.51)
,
7 (100)
(100)
(-)
9 (82.98)
4 (8.51)
1 (8.51)
) (0)