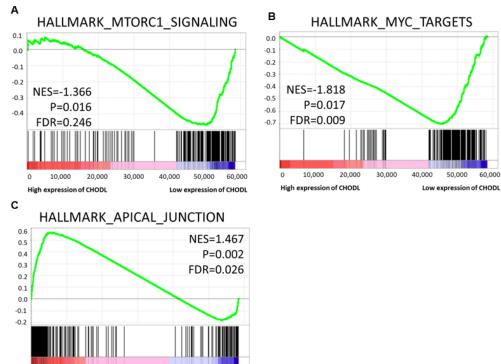
## **Supplementary Materials**

Table S1

Part of differentially expressed genes in CHODL over-expressed cells

Gene	Fold-change	Gene	Fold-change
Up-regulated	(p<0.05)	Down-regulated	(p<0.05)
LOC100129203	23	TRIM17	11
RAET1E-AS1	17	AKR1C1	10
HOXD-AS2	15	PLSCR3	10
CHRNA1	13	CDH11	9
RPRML	13	TULP2	8
SERTAD4	13	CXCR2	8
PRR34	13	PTGS2	7
ANGPTL6	13		
LOC100506747	13		
PCDH8	13		

Figure S1



60,000

Figure S1 Classified the differentially expressed genes by GSEA.

50.000

High methylation of CHODL

60.000

40.000

Figure S2

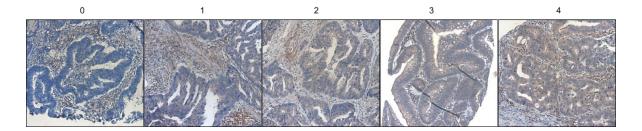
0

10,000

Low methylation of CHODL

20,000

30,000



**Figure S2** Scores indicated CHODL levels in representative tumor tissues as determined by IHC staining. Scores were calculated by intensity and percentage of stained cells. Scale bars, 100 um.