텍스트, 도표, 스케치, 기술 도면이(가) 표시된 사진

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Supplementary Figure S1. Basal PDE1A expression and knockdown efficiency in ovarian cancer cell lines.

(A) Basal PDE1A mRNA expression in normal immortalized ovarian epithelial cells and ovarian cancer cell lines. OVCA429 and OVCA433 cells, which show relatively high endogenous PDE1A levels, were selected for functional assays. (B) Validation of PDE1A knockdown by siRNA in OVCA429 and OVCA433 cells assessed by quantitative reverse transcription polymerase chain reaction (qRT-PCR). The number of asterisks indicates statistical significance: \*\* *p* < 0.01, \*\*\* *p* < 0.001. Error bars represent the mean ± SD of three independent experiments, each performed in triplicate.

텍스트, 도표, 스케치, 디자인이(가) 표시된 사진

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Supplementary Figure S2. CTNNB1 mRNA expression after PDE1A knockdown. Relative CTNNB1 mRNA levels in OVCA429 and OVCA433 cells transfected with control or PDE1A siRNAs for 48 h, measured by qRT-PCR. Error bars represent the mean ± SD of values from triplicate experiments.