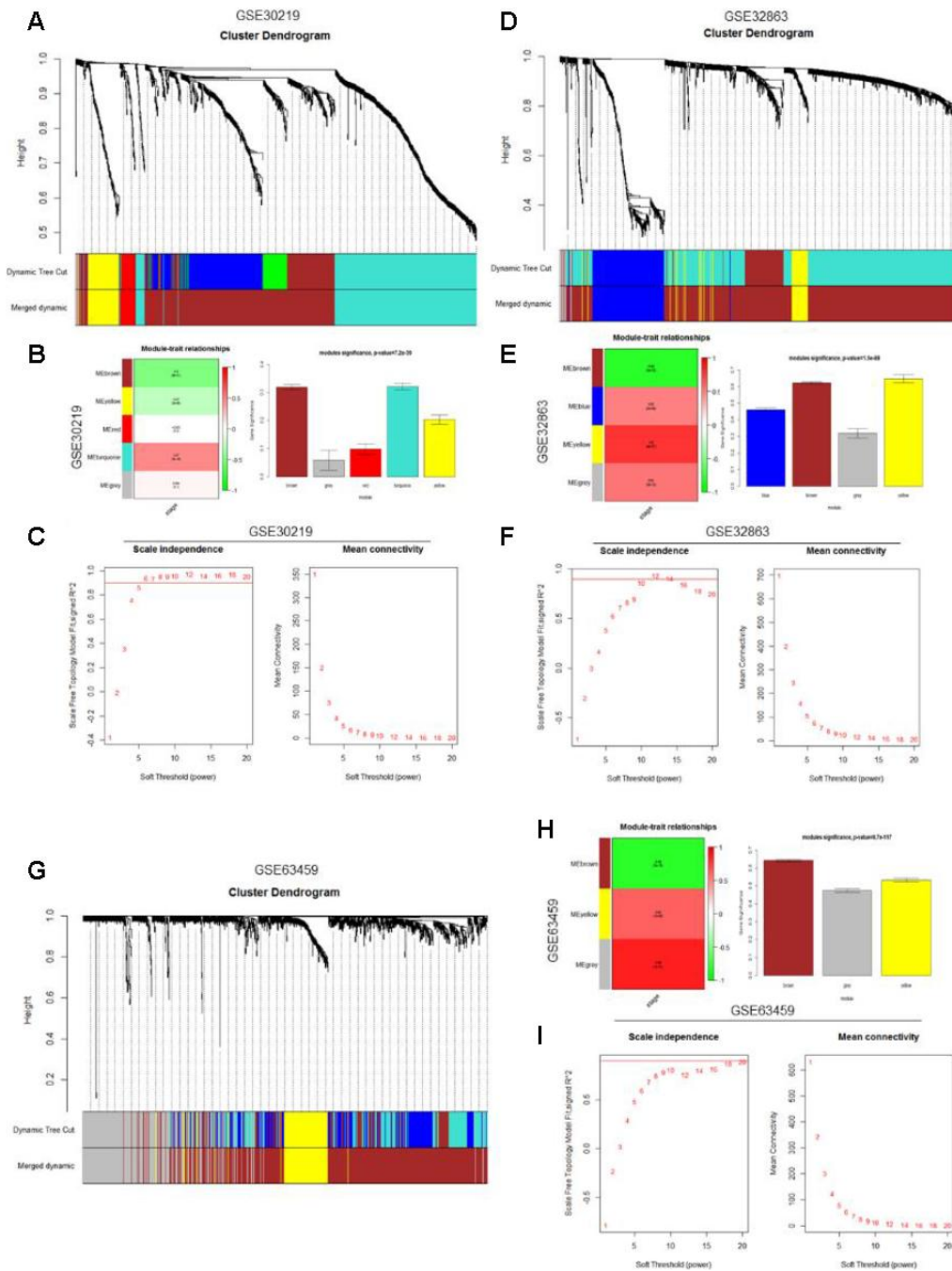
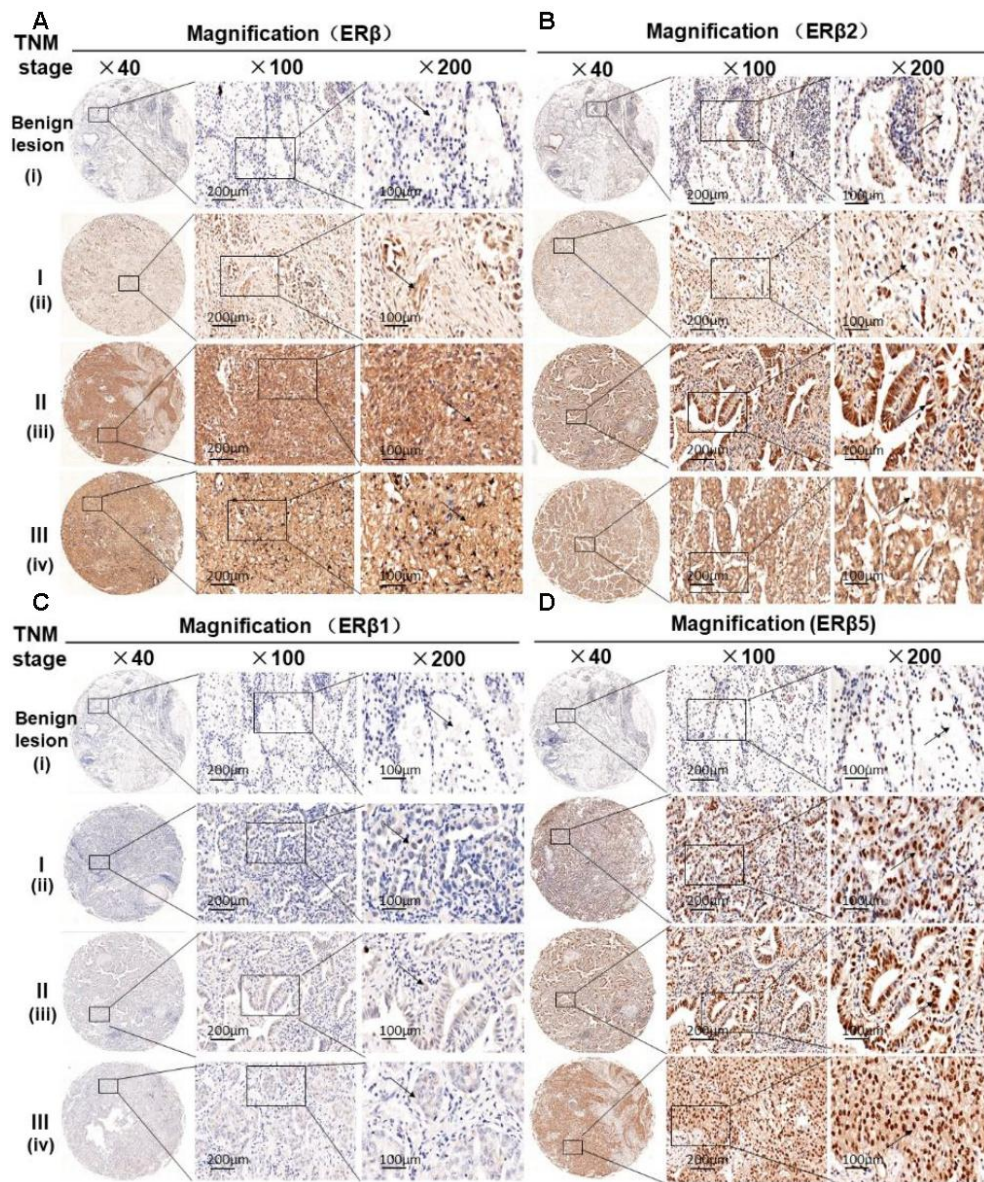


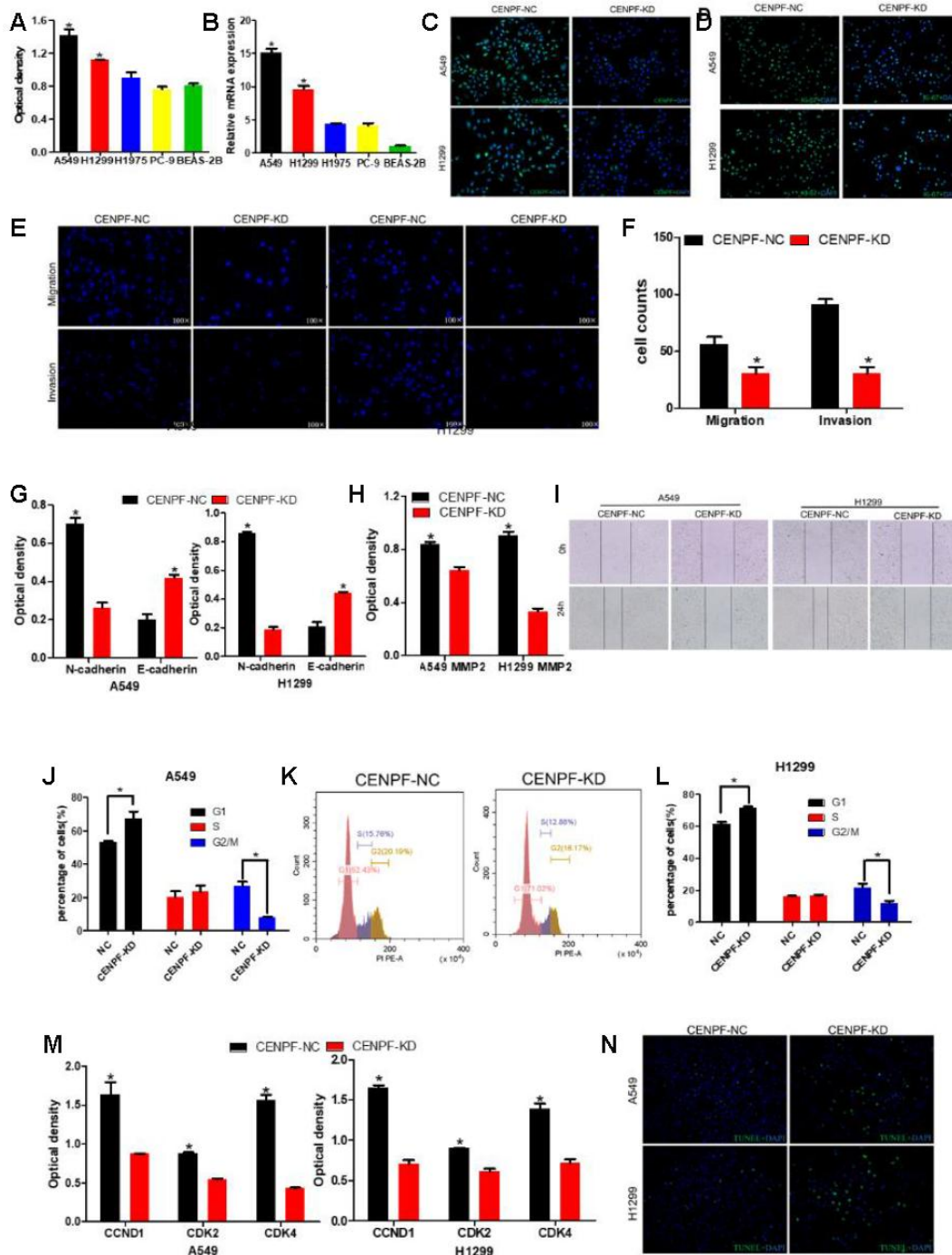
SUPPLEMENTARY FIGURES



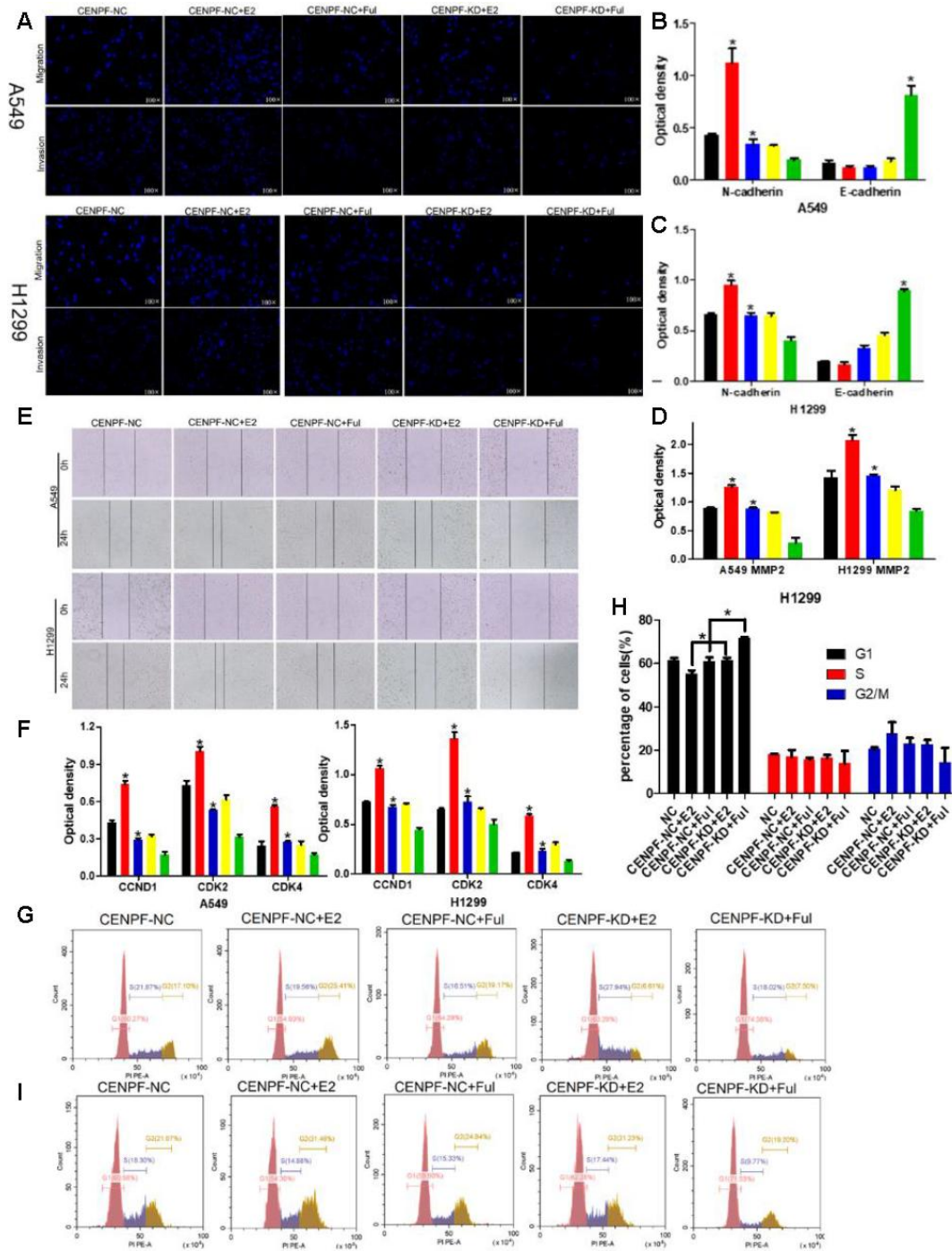
Supplementary Figure 1. WGCNA analysis and determination of the CENPF gene. (A, D, G) Dendrogram of all differentially expressed genes clustered based on a dissimilarity measure (1-TOM) (GSE30219, GSE32863, GSE63459). (B, E, H) Heat maps and distribution of differential genes for different modules related to NSCLC staging (GSE30219) and LUAD staging (GSE32863, GSE63459). (C, F, I) Analysis of the scale-free fit index for various soft-thresholding power (β) and analysis of the mean connectivity for various soft-thresholding power (GSE30219, GSE32863, GSE63459).



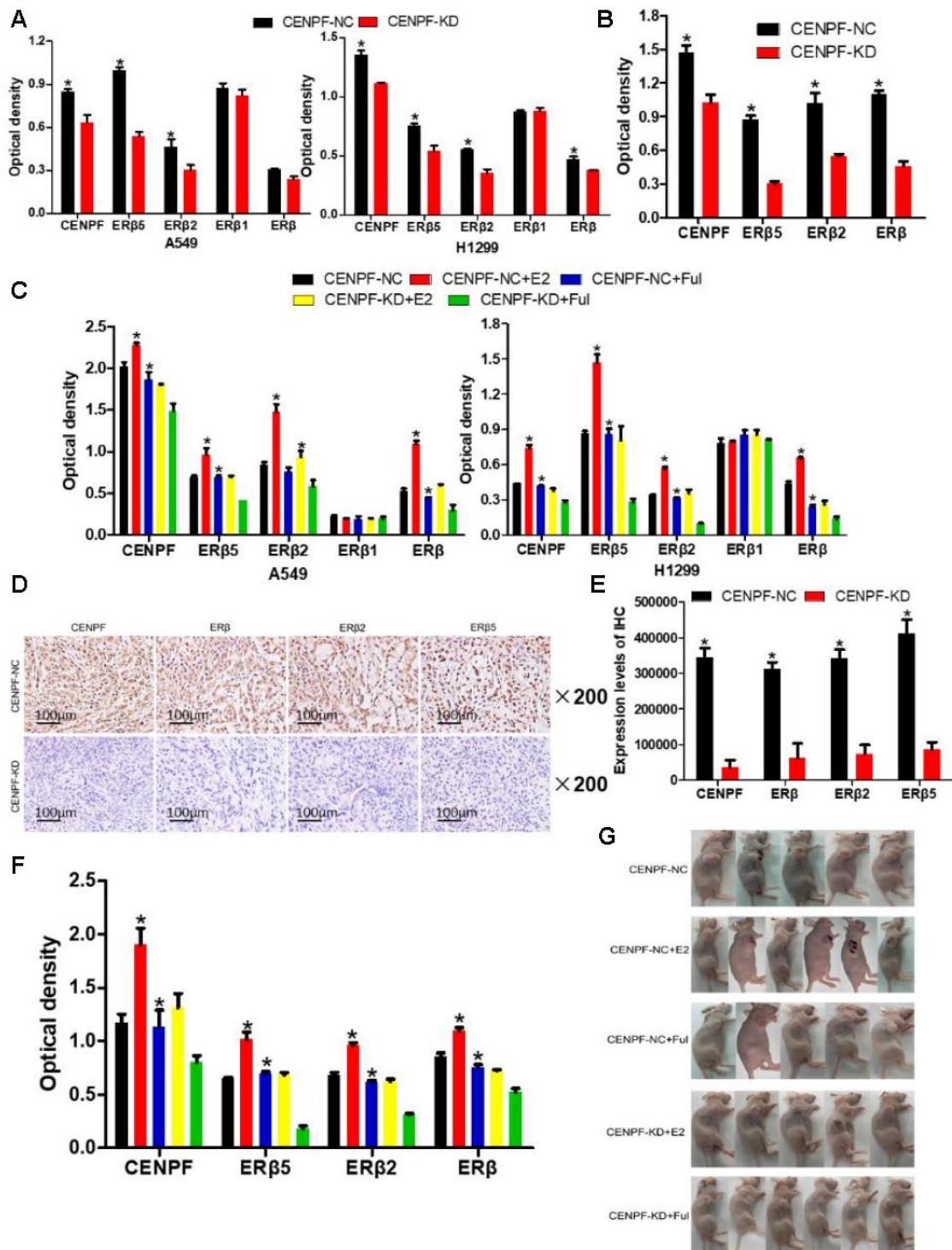
Supplementary Figure 2. (A–D) Expression of ERβ, ERβ1, ERβ2 and ERβ5 in benign lung lesions (i) and different TNM staging of LUAD (ii=I stage, iii=II stage, iv=III stage). The magnification of each slice is 40×, 100×, 200× in order.



Supplementary Figure 3. Knockdown of CENPF inhibits the biological effects of LUAD cells. (A) The protein level of CENPF in A549 and H1299 cell lines were higher than in normal cell lines BEAS-2B and other LUAD cells. GAPDH served as the internal control. * $P < 0.05$ vs other cells. (B) mRNA expression of CENPF in different cell lines. * $P < 0.05$ vs other cells. (C) Representative cellular immunofluorescence images after transfection of CENPF (200 \times). Green stands for CENPF and blue stands for DAPI. (D) Representative Ki67 staining (green) shows cell proliferation of LUAD cells after CENPF-NC or KD treatment (200 \times). The nuclei were counterstained with DAPI (blue). (E, F) Migration assays and invasion assays revealed that CENPF-KD decreased cell migration and invasion abilities of A549 and H1299. (G, H) Corresponding gray value analysis showed that the related protein E-cadherin was significantly increased ($P=0.009$, Figure 4F), and N-cadherin and MMP2 were significantly decreased when compared with NC group ($P=0.004$, 0.012) of N-cadherin, E-cadherin and MMP2 in A549 and H1299 cells. (I) Representative scratched pictures of A549 and H1299 cells. (J–L) Percentage of CENPF-KD cells H1299 at different stages of the cell cycle (G1, S and G2/M) and corresponding quantified histograms of A549 and H1299. (M) Corresponding gray value analysis showed that the expression of CCND1, CDK2 and CDK4 was significantly lowered in CENPF-KD group ($P=0.022$, 0.001, 0.002). (N) Representative TUNEL staining (green) shows (200 \times). P values were calculated with two-tailed unpaired Student's t-test, or one-way analysis of variance.



Supplementary Figure 4. Knockdown of CENPF inhibits proliferation, invasion and migration of LUAD cells via the ER β 2/5 pathway. (A) Migration and invasion pictures of A549 and H1299 cells. (B–D) Corresponding quantified histograms of MMP2, N-cadherin and E-cadherin in A549 and H1299 cells. (E) Representative scratched images of A549 and H1299 cells. (F) Corresponding quantified histograms of CCND1, CDK2 and CDK4 in A549 and H1299. *P < 0.05. (G–I) Percentage of the A549 cells and H1299 cells at different stages of the cell cycle (G1, S and G2/M) and corresponding quantified histograms. P values were calculated with two-tailed unpaired Student's t-test, or one-way analysis of variance.



Supplementary Figure 5. Knockdown of CENPF can inhibit the expression of ERβ2/5 *in vitro* and *in vivo*. (A, B) Corresponding gray value analysis showed that knockdown of CENPF inhibited the expression of ERβ2/5 *in vitro* and *in vivo* experiment. (C) Corresponding gray value analysis of CENPF, ERβ, ERβ1, ERβ2 and ERβ5 *in vitro* experiment after treated with E2 and Ful treatment. (D, E) Immunohistochemical analysis of CENPF, ERβ, ERβ2 and ERβ5 expression in nude mice tumor tissues and corresponding quantitative histograms. *P < 0.05. (F) Corresponding gray value analysis of CENPF, ERβ, ERβ1, ERβ2 and ERβ5 *in vivo* experiment after treated with E2 and Ful treatment. (G) Pictures of nude mice sacrificed at 45 days. *P < 0.05. P values were calculated with two-tailed unpaired Student's t-test, or one-way analysis of variance.