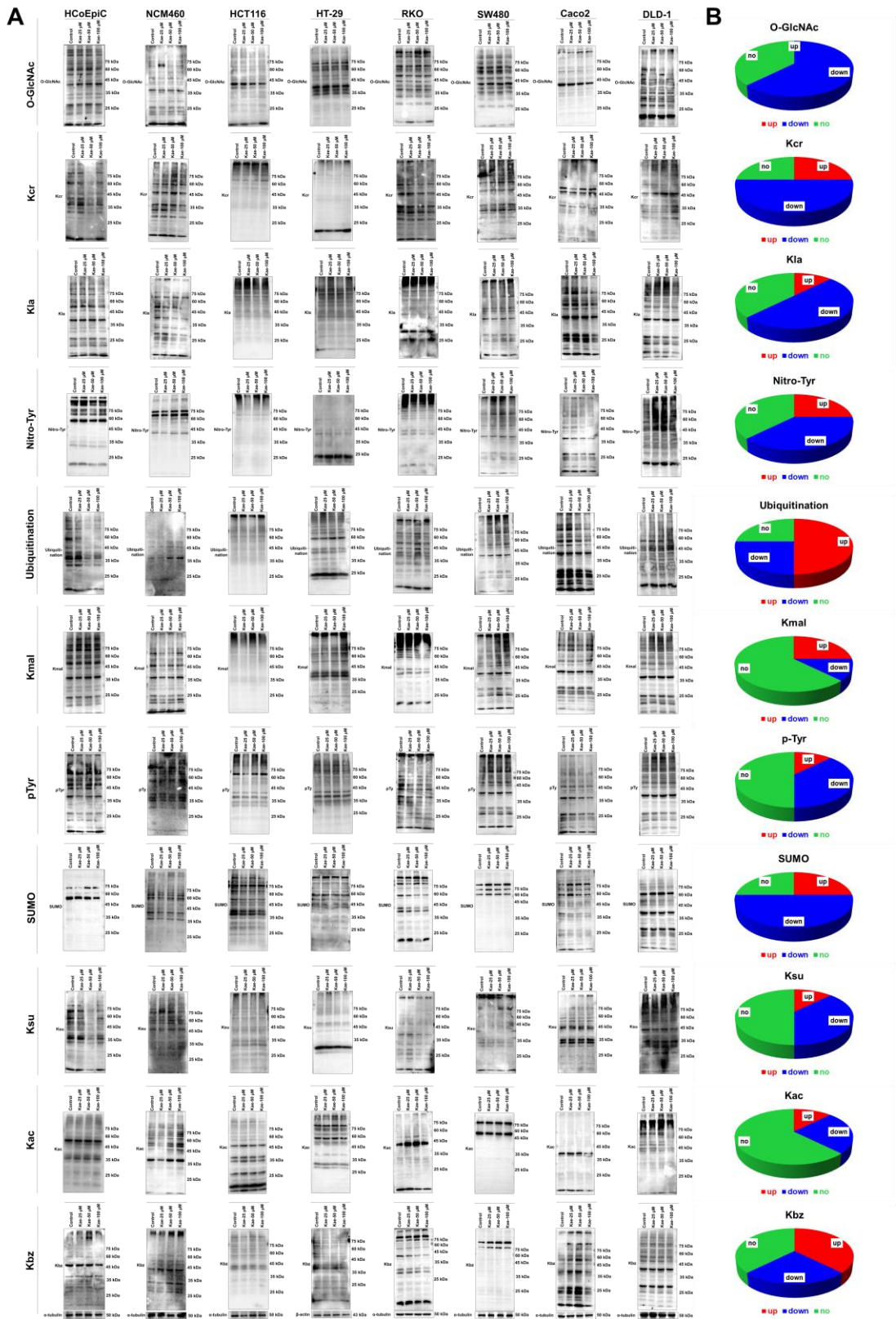
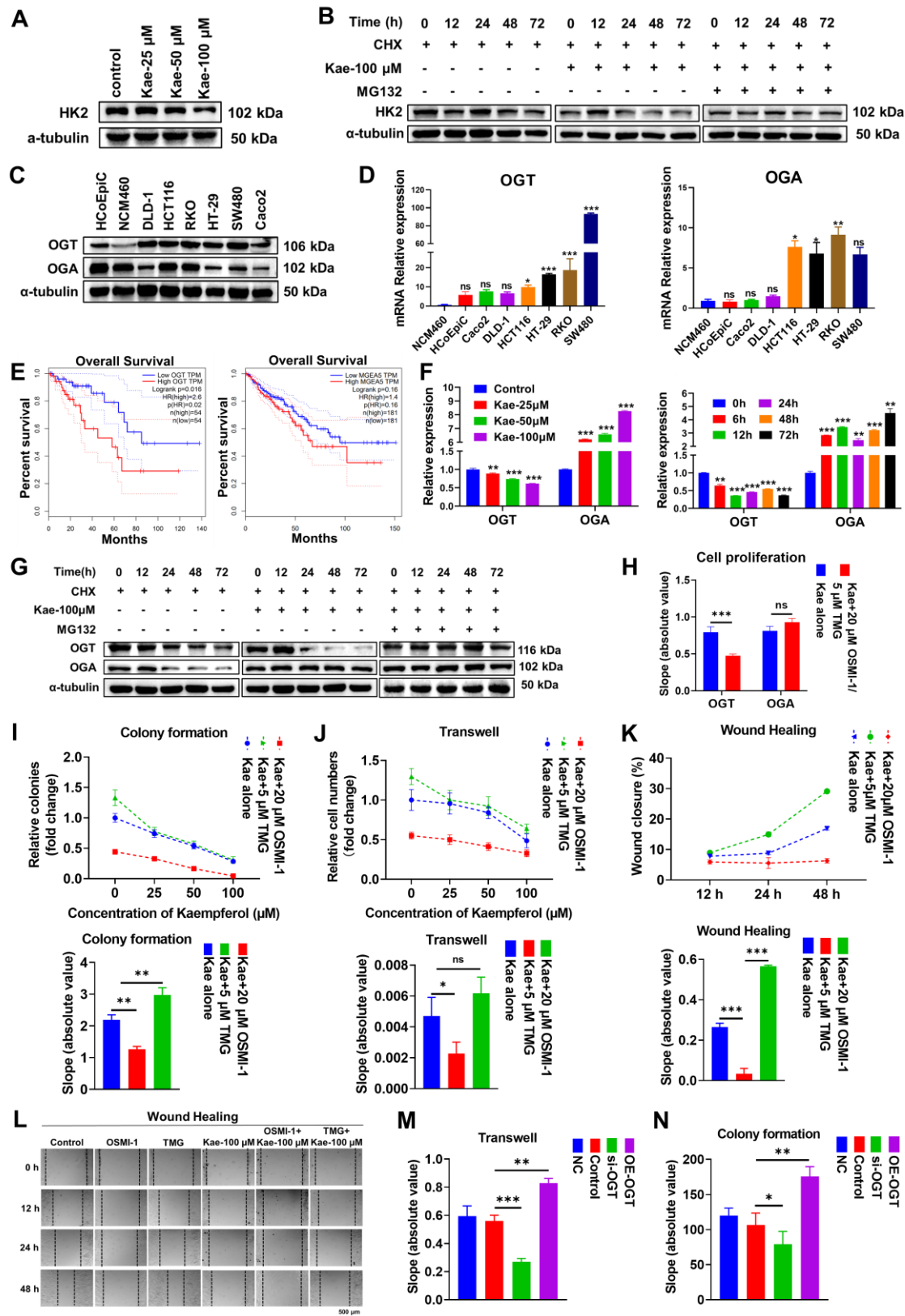


**Supplemental Figure 1.** Comprehensive antibody detection of 11 posttranslational modifications across 8 cell lines at various concentrations of kaempferol.



**Supplementary Figure 2. Expression and role of OGT and OGA in CRC.**



**A**, Effect of kaempferol on HK2 protein expression. **B**, Effect of kaempferol on HK2

protein stability. **C**, Differences in OGT and OGA protein expression between normal colonic epithelial cells and CRC cell lines. **D**, Differences in OGT and OGA mRNA expression between normal colonic epithelial cells and CRC cell lines. **E**, Survival analysis on the basis of OGT and OGA expression. **F**, Quantitative analysis of the effects of different concentrations of kaempferol on OGT and OGA protein expression at different time points. **G**, Effect of kaempferol on the half-life of OGT and OGA proteins. **H**, Quantitative analysis of the effects of OGT and OGA inhibition on the ability of kaempferol to suppress HCT116 cell viability. **I**, Quantitative analysis of the effects of OGT and OGA inhibition on the ability of kaempferol to suppress HCT116 cell stemness. **J**, Quantitative analysis of the effects of OGT and OGA inhibition on the ability of kaempferol to suppress HCT116 cell migration. **K**, Quantitative analysis of the effects of OGT and OGA inhibition on kaempferol-mediated suppression of HCT116 cell wound healing. **L**, Effects of OGT and OGA inhibition on HCT116 cell wound healing. **M**, Quantitative analysis of the effects of OGT knockdown or overexpression on the ability of kaempferol to suppress HCT116 cell migration. **N**, Quantitative analysis of the effects of OGT knockdown or overexpression on the ability of kaempferol to suppress HCT116 cell stemness. The data are presented as the means  $\pm$  SDs, \*  $P < 0.05$ , \*\*  $P < 0.01$ , \*\*\*  $P < 0.001$ , ns indicates no statistically significant difference.

**A**

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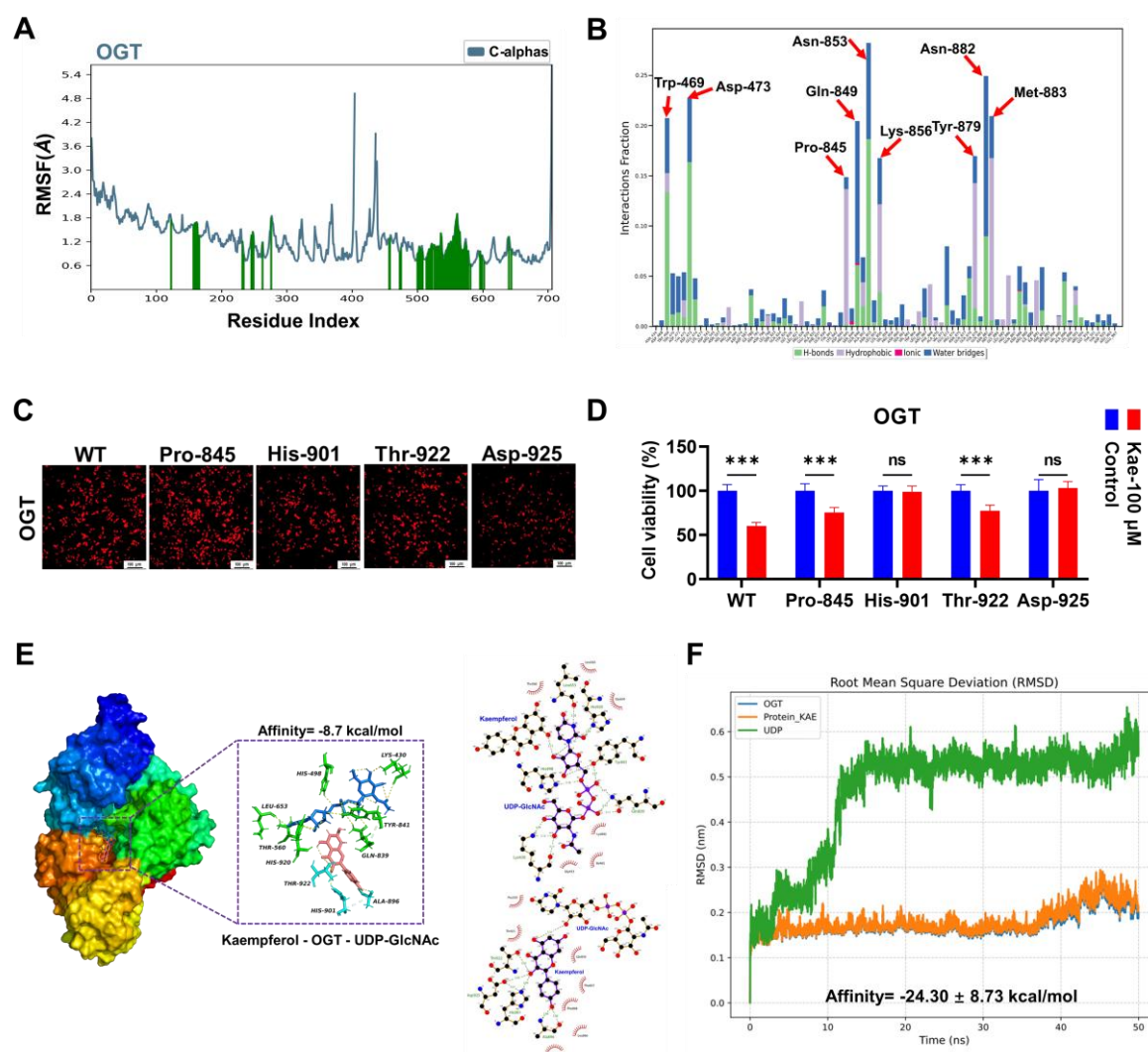
**K**

**A**, Western blotting analysis of the effect of combining kaempferol with the OGA inhibitor TMG on protein O-GlcNAcylation. **B**, The effect of OGA knockdown on the

inhibition of cell proliferation by kaempferol. **C**, The effect of OGA knockdown on the inhibition of cell stemness by kaempferol. **D**, The effect of OGA knockdown on the inhibition of cell migration by kaempferol. **E**, Fluorescence colocalization of kaempferol with OGA. **F**, SIP assay verifying the binding interaction between kaempferol and OGA. **G**, CETSA assay verifying the binding interaction between kaempferol and OGA. **H**, Root mean square deviation (RMSD) changes in OGA during molecular dynamics simulation with kaempferol. **I**, RMSF changes in the OGA during the molecular dynamics simulation. **J**, Changes in various interaction bonds between kaempferol and OGA during molecular dynamics simulation. **K**, Molecular docking analysis of kaempferol with OGA. The data are presented as the means  $\pm$  SDs, \*  $P < 0.05$ , \*\*  $P < 0.01$ , \*\*\*  $P < 0.001$ , ns indicates no statistically significant difference.

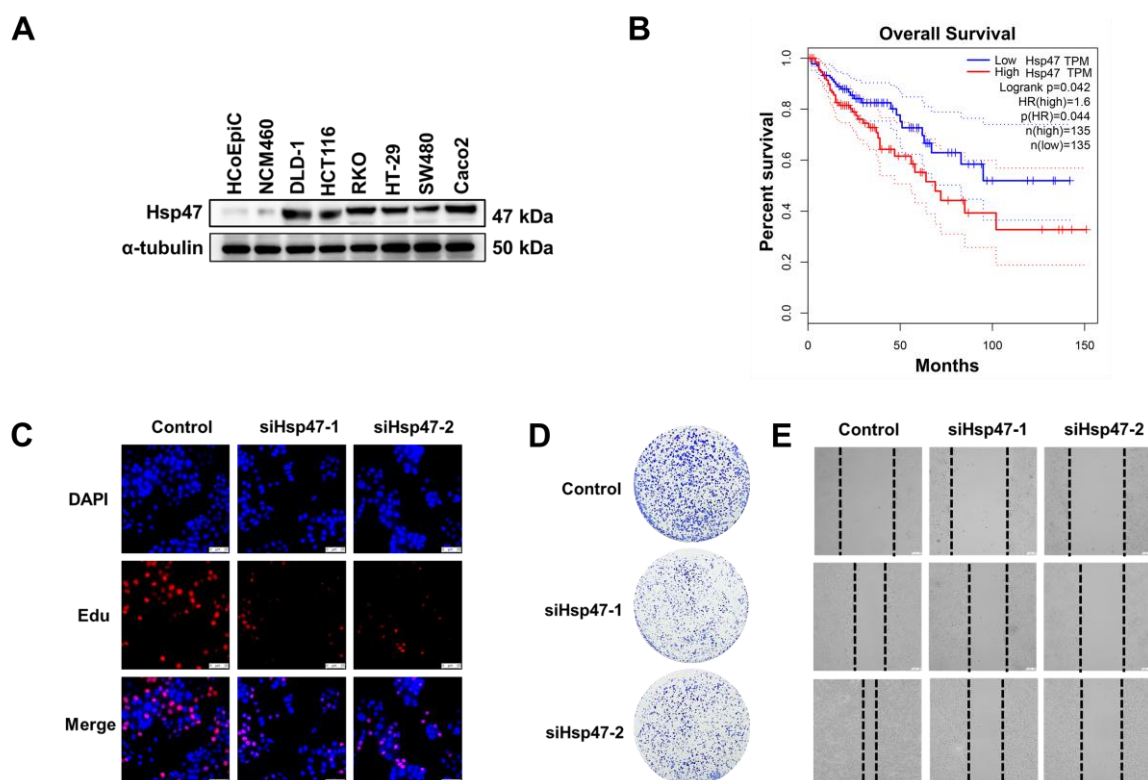


**Supplementary Figure 4.** Binding analysis of kaempferol/UDP-GlcNAc with OGT.



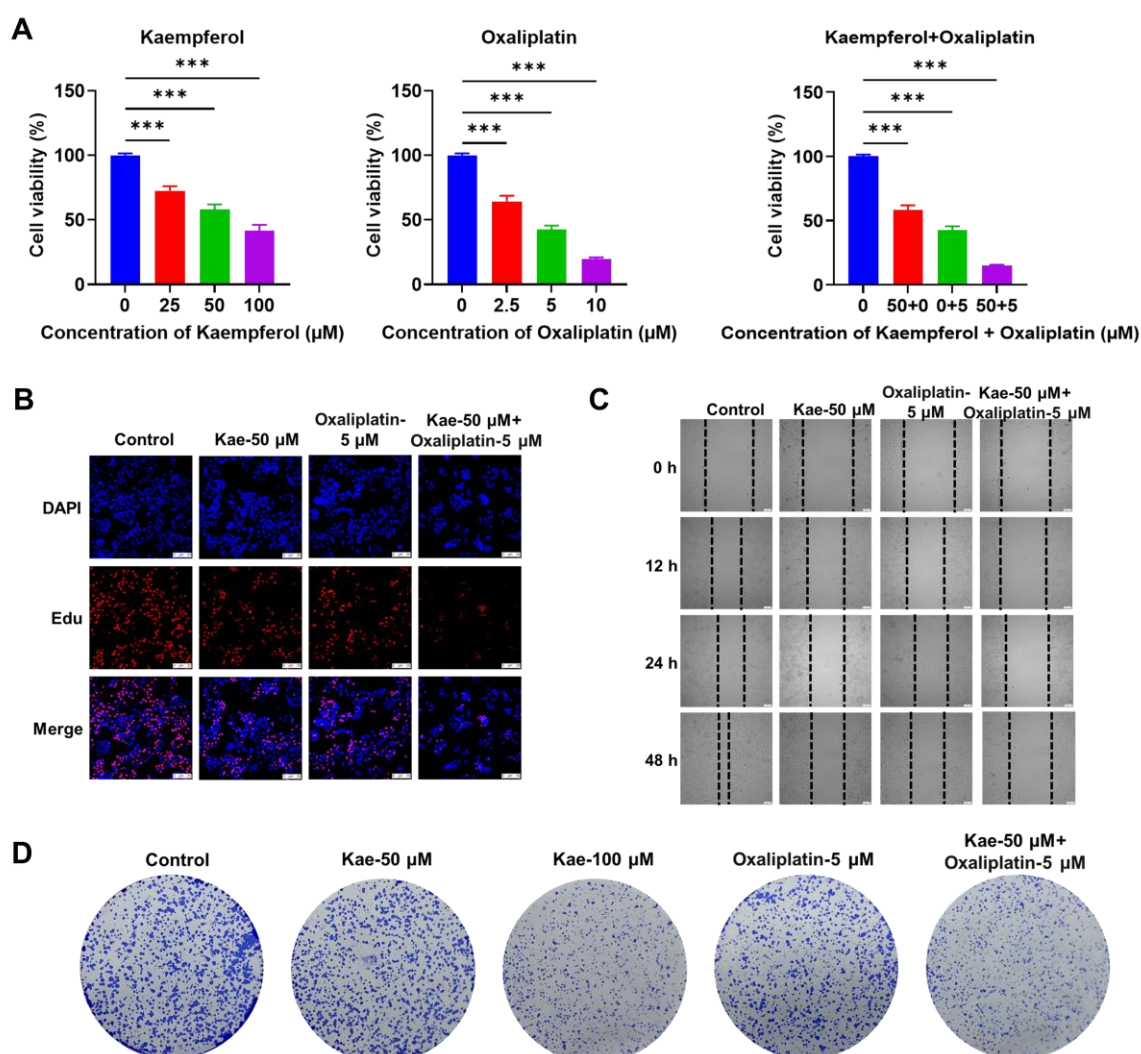
**A**, Root mean square fluctuation (RMSF) changes in OGT during molecular dynamics simulation. **B**, Analysis of interactions between OGT and kaempferol throughout the simulation, categorized into hydrogen bonds, hydrophobic interactions, ionic bonds, and water bridges. **C**, Fluorescence detection of OGT expression after key amino acid site mutation. **D**, Effects of mutations at the OGT binding sites of kaempferol on the inhibition of cell viability. **E**, Molecular docking of UDP-GlcNAc with kaempferol and OGT complex. **F**, Molecular dynamics simulation of UDP-GlcNAc with kaempferol and OGT complex. The data are presented as the means  $\pm$  SDs, \*  $P < 0.05$ , \*\*  $P < 0.01$ , \*\*\*  $P < 0.001$ , ns indicates no statistically significant difference.

**Supplementary Figure 5.** Analysis of the correlation between Hsp47 and colorectal cancer.



**A**, Protein expression of Hsp47 in normal intestinal epithelial cells and CRC cells. **B**, Survival analysis of Hsp47 in CRC patients. **C**, Edu detection of the inhibitory effect of Hsp47 knockdown on CRC cell proliferation. **D**, Plate colony formation assay to detect the inhibition of CRC cell stemness by Hsp47 knockdown. **E**, The cell scratch assay detects the inhibition of CRC cell migration by Hsp47 knockdown.

**Supplementary Figure 6.** The inhibitory effect of combining kaempferol with the chemotherapy drug oxaliplatin on CRC.



**A**, The SRB assay was used to evaluate the effect of kaempferol combined with oxaliplatin on CRC cell proliferation. **B**, Edu experiment to evaluate the effect of kaempferol combined with oxaliplatin on CRC cell proliferation. **C**, Wound healing assay to evaluate the effect of kaempferol combined with oxaliplatin on CRC cell migration. **D**, Colony formation assay to evaluate the effect of kaempferol combined with oxaliplatin on the stemness of CRC cells. The data are presented as the means  $\pm$  SDs, \*  $P < 0.05$ , \*\*  $P < 0.01$ , \*\*\*  $P < 0.001$ , ns indicates no statistically significant difference.