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Supplemental information

**lncRNA SNHG26 promoted the growth, metastasis,
and cisplatin resistance of tongue squamous cell
carcinoma through PGK1/Akt/mTOR signal pathway**

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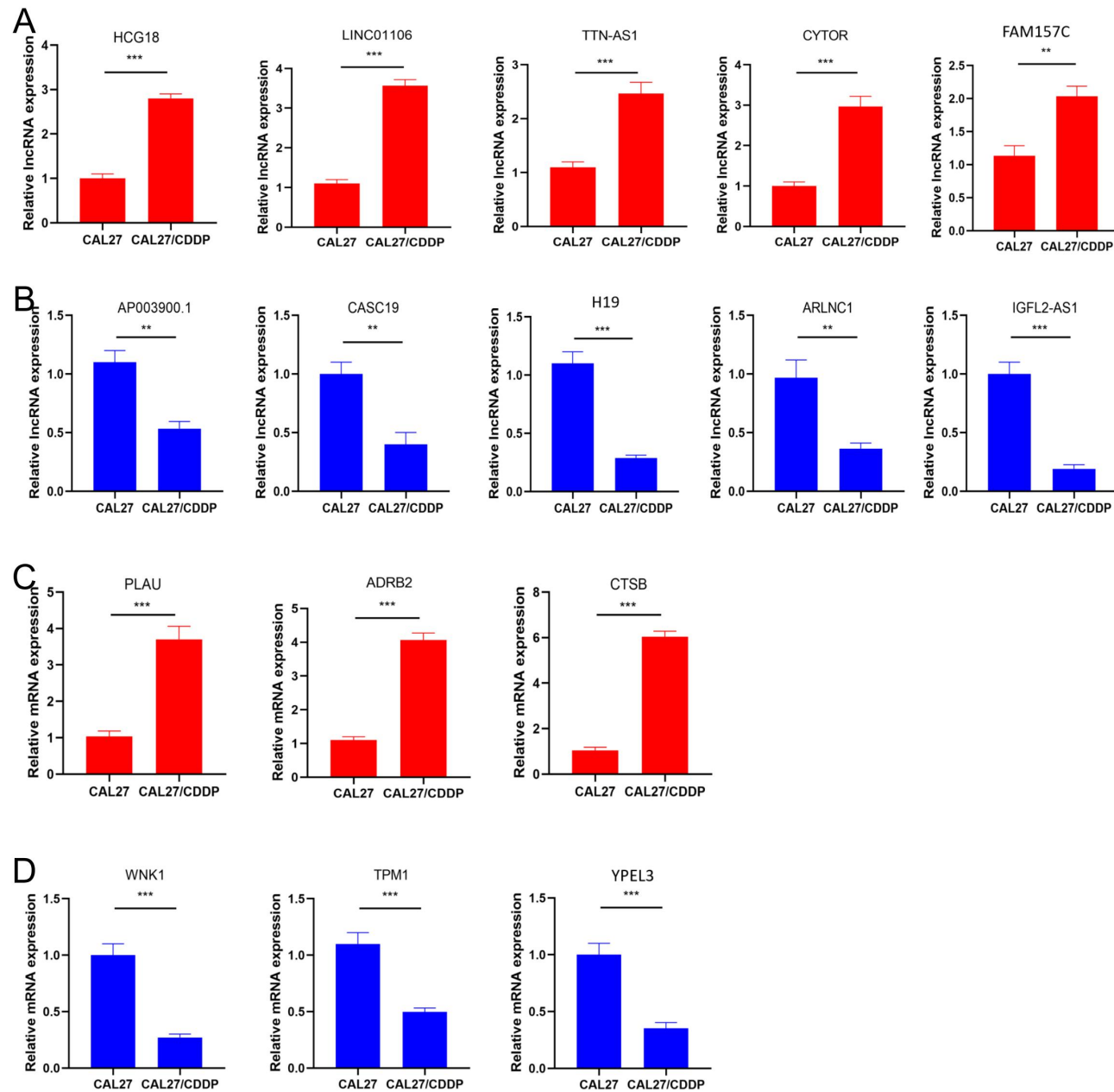
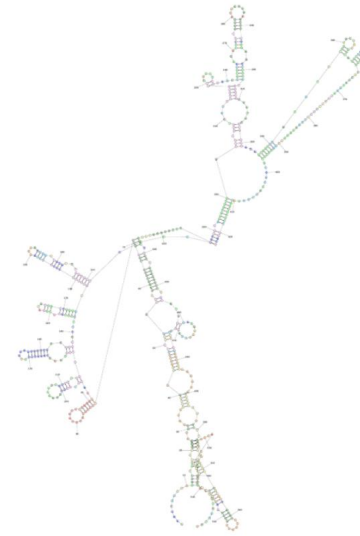


Figure S1. Expression levels of 10 lncRNAs (a, b) and 6 mRNAs (c, d) by qRT-PCR in CAL27 and CAL27/CDDP cells. The mean \pm SD of triplicate experiments were plotted. * $P < 0.05$, ** $P < 0.01$, *** $P < 0.001$.

A

GAAGCAGTGTGAGCTAGCGTCTAAGGATTGTGAAAAGTGAAGTCAAGTGGCC
 AGAACTAGAAAACCTCCAGCCCCCTGAACTCAGCCTCAGGGTAAGCGAGT
 TTAATAGTGCTTCCTTTGATTTTCAAATCTGTGGAAGGTTTGTTCAGTTTTT
 CCCAGACAAATATTGTTGGGATAGTGTTCCTTTCAAAAATGGATAGGATTC
 AATGCTCATCAGTGGGAAGATAGTGACTCCTGACCATTTTCCTCCTAGCTG
 GAAACTACAGAAGACTCCCAAGTATCAGGAAATTTGGAGAATTTGTACAGG
 AATTATATTCAGAGATAGCAATTTATTTACTCATCAAAGAGAGCTATTGGATA
 GCCACCATAAAGACCACTCATTAAAAACAAAATTGCATGATATTCTTAACA
 TTGTCTTTCACTTTTTTAACCAGGCTTGGTGGAGTTTCAAGATGCTGATG
 AACTTAAGGTCTGGCGCTTGAAGAATCAGTGAAGCTTTTCAAGTAGAGG
 AGGTATATTTGAATATGTAACCTTAGTATCTACAGAATAGAGGTTGCTTTTT
 TTTTTTTT



Probability >= 99%
 99% > Probability >= 95%
 95% > Probability >= 90%
 90% > Probability >= 80%
 80% > Probability >= 70%
 70% > Probability >= 60%
 60% > Probability >= 50%
 50% > Probability
 ENERGY = -137.9 SNHG26

B

Protein coding potential

Metric	Raw result	Interpretation
PRIDE reprocessing 2.0	0	non-coding
Lee translation initiation sites	0	non-coding
PhyloCSF score	-280.9938	non-coding
CPAT coding probability	1.22%	non-coding
Bazzini small ORFs	0	non-coding

Figure S2. The full-length sequence of SNHG26 and the secondary mechanism of prediction (a). Prediction of coding ability of SNHG26 (b).