Figure S1

A Active ingredient Ö ŌН OH OH. ŌΗ .OH HO¹ H Bufalin Ö Rhein Salicylic acid Change metabolism Prodrug Enhance activity Clinical drug OH Ħ ŌΗ O Diacerein 0Ac-Bufalin Aspirin

Figure S2

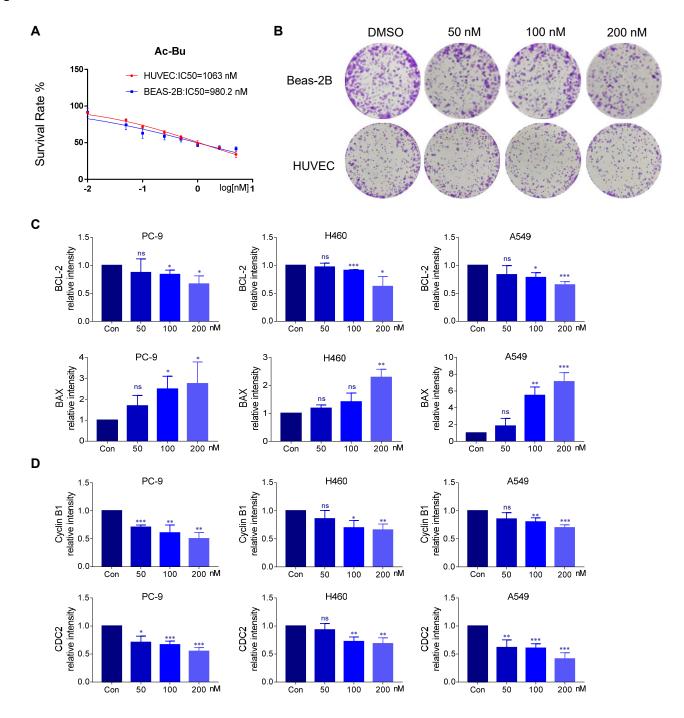


Figure S3

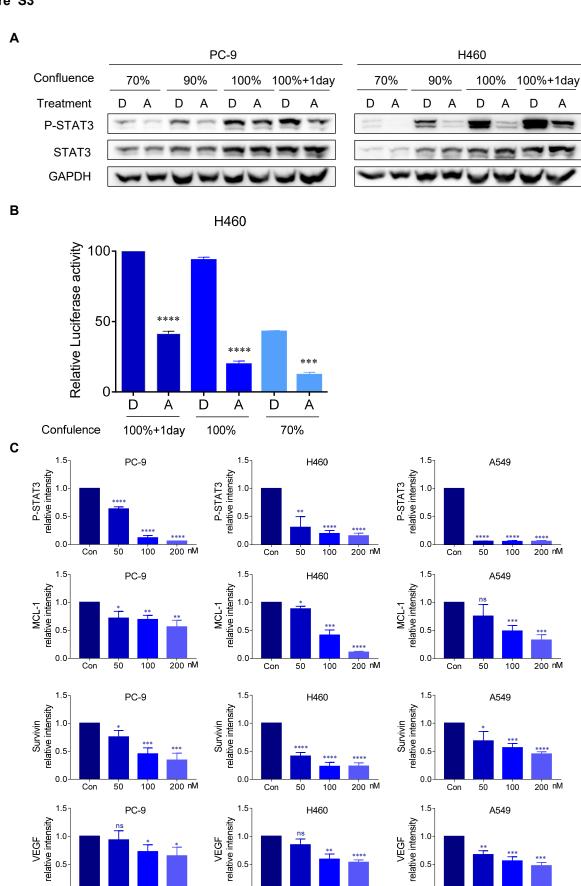
0.5

50

Con

100

200 nM



0.5

50

100

200 nM

Con

0.5

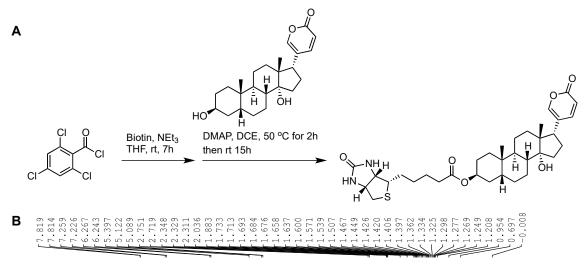
50

100

200 nM

Con

Figure S4



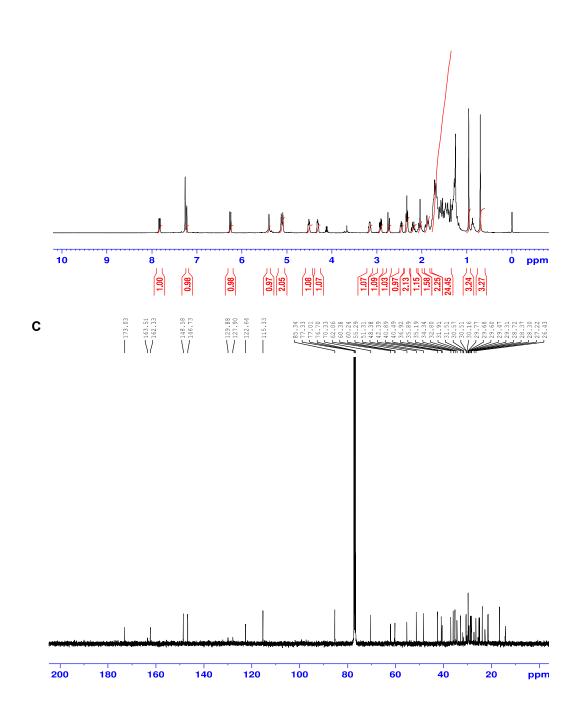


Figure S5

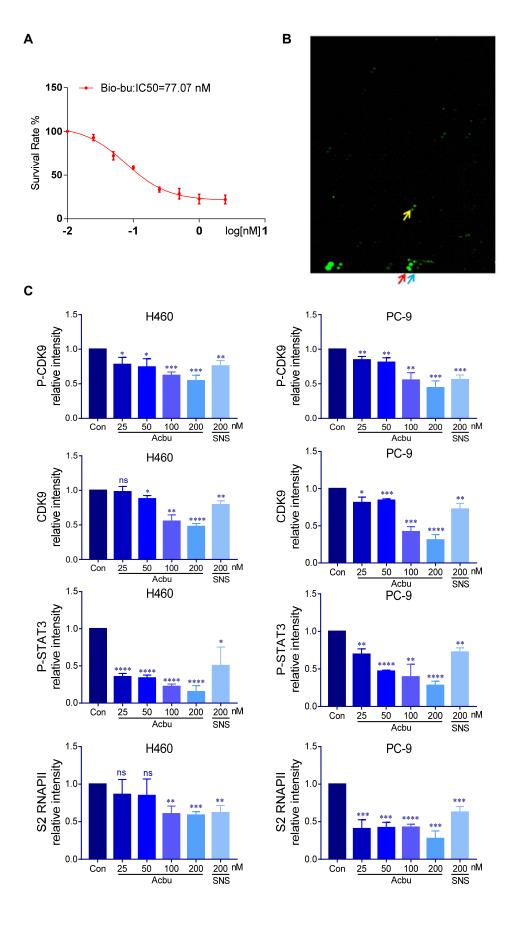


Figure S6

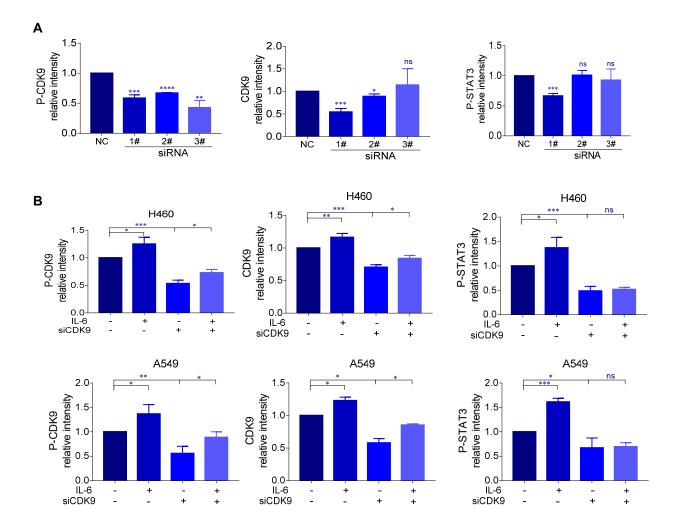
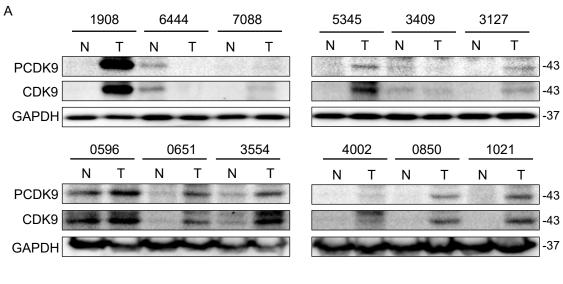


Figure S7



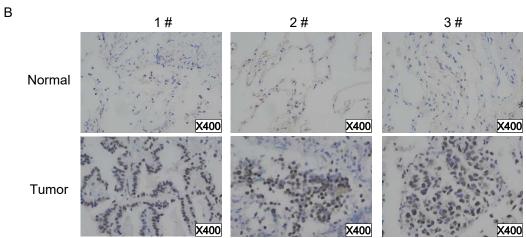


Fig. S1 Designed and synthesized acetyl-bufalin

- 2 A. Designed and synthesized acetyl-bufalin (Ac-bufalin) base on aspirin (acetylsalicylic acid) and
- 3 diacerein (diacetoxy-rhein). **B.** Chemistry experimental section of Ac-bufalin: To a mixture of
- 4 bufalin (15 mg, 0.0388 mmol) and NEt₃ (0.39 mmol) in DCE (dry, 1 mL) was added AcCl (0.39
- 5 mmol), the reaction mixture was stirred at rt for 19 h. The mixture was then purified by a silica gel
- 6 flash chromatography (DCM/MeOH = 50:1 to 20:1) to afford target compound (17 mg, 100% yield)
- 7 as a pale yellow solid. ¹H NMR (400 MHz, CDCl₃) δ 7.82 (dd, J = 9.6, 4.0 Hz, 1H), 7.22 (s, 1H),
- 8 6.26 (d, J = 9.6 Hz, 1H), 5.07 (s, 1H), 2.49-2.40 (m, 1H), 2.26-2.15 (m, 1H), 2.09-1.99 (m, 4H),
- 9 1.94-1.81 (m, 2H), 1.80-1.35 (m, 18H), 1.27 (s, 3H), 0.69 (s, 3H); 13 C NMR (100 MHz, CDCl₃) δ
- 10 170.7, 162.3, 148.6, 146.7, 122.6, 115.3, 85.4, 70.4, 51.3, 48.4, 42.4, 40.9, 36.8, 35.9, 35.2, 32.8,
- 11 30.5, 30.5, 28.7, 26.4, 25.1, 23.7, 21.5, 21.4, 21.3, 16.5; ESI-HRMS: calcd. for $C_{26}H_{37}O_5^+(M+H)^+$
- 12 429.2636, found 429.2637.

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Fig. S2 Antitumor activity of acetyl-bufalin in vitro

- 15 A. HUVEC and BEAS-2B cells were incubated with increasing doses of acetyl-bufalin for 48 h.
- 16 Cell viability was determined by the MTT assay. B. HUVEC and BEAS-2B cells were incubated
- with acetyl-bufalin for 24 h and allowed to form colonies for one week. Colonies were then fixed,
- stained with crystal violet and photographed. C. Cells were treated with acetyl-bufalin at different
- concentrations as indicated for 24 h, the cell lysates were processed for Western blot analysis for
- protein expression of BCL-2 and BAX, and the relative intensity was calculated. **D.** Western blot
- analysis was used to detect the levels of cyclin B1 and CDC2 after acetyl-bufalin treatment and the
- 22 relative intensity was calculated. *P<0.05, **P<0.01, ***P<0.001.

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Fig. S3 Acetyl-bufalin inhibited the STAT3 signaling pathway in human NSCLC cells

- 25 A. PC-9 and H460 cells were grown to 70, 90 or 100% confluence and up to 1 day postconfluence
- as indicated, in the presence of 50 nM acetyl-bufalin or the DMSO diluent. Cell lysates were used
- to detect P-STAT3, STAT3 and GAPDH as a control. **B.** H460 cells were grown to 70, or 100%
- confluence and up to 1 day postconfluence, then transfected with luciferase reporter gene plasmid
- and treated with 50 nM acetyl-bufalin or the DMSO diluent. The results were normalized to the

- Renilla luciferase activity. C. H460, A549 and PC-9 cells treated with concentration gradients of
- 31 acetyl-bufalin. STAT3 and its downstream target genes, including MCL-1, Survivin and VEGF,
- were detected via western blot analysis. The relative intensity was calculated. *P<0.05, **P<0.01,
- 33 ***P<0.001, ****P<0.0001.

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Fig.S4 Chemistry experimental section of Bio-bufalin

- To a mixture of Biotin (0.3 mmol) and NEt₃ (0.45 mmol) in THF (dry, 3 mL) was added 2,4,6-
- 37 trichlorobenzoyl chloride (0.3 mmol), the reaction mixture was stirred at rt for 7 h. The bufalin (20
- 38 mg, 0.052 mmol), DMAP (0.4 mmol) and DCE (dry, 4 mL) were added. The suspension was stirred
- 39 at 100 °C for 2 h and then rt for 15 h. The mixture was then washed with 1 N HCl (aq, x 3) and
- brine, purified by a silica gel flash chromatography (DCM/MeOH = 30:1 to 10:1) to afford target
- 41 compound (18.0 mg, 57% yield) as a white solid.
- ¹H NMR (400 MHz, CDCl₃) δ 7.83 (dd, J = 9.6, 4.0 Hz, 1H), 7.23 (s, 1H), 6.26 (d, J = 9.6 Hz, 1H),
- 43 5.40 (s, 1H), 5.12 (s, 1H), 5.09 (s, 1H), 4.52-4.49 (m, 1H), 4.33-4.30 (m, 1H), 3.16-3.15 (m, 1H),
- 44 2.92 (dd, J = 12.8, 4.8 Hz, 1H), 2.74 (d, J = 12.8 Hz, 1H), 2.45-2.44 (m, 1H), 2.33 (t, J = 7.6 Hz,
- 45 2H), 2.20-2.17 (m, 1H), 2.09-1.94 (m, 1H), 1.96-1.81 (m, 2H), 1.80-1.35 (m, 24H), 1.21 (s, 3H),
- 46 0.70 (s, 3H); ¹³C NMR (100 MHz, CDCl₃) δ 173.0, 163.5, 162.3, 148.6, 146.7, 122.6, 115.3, 85.3,
- 47 70.3, 62.1, 60.2, 55.3, 51.3, 48.4, 42.4, 40.9, 40.5, 36.9, 35.9, 35.2, 34.3, 32.8, 30.6, 30.5, 29.7, 29.5,
- 48 28.7, 26.4, 25.5, 25.1, 23.8, 21.4, 21.3, 16.5; ESI-HRMS: calcd. for C₃₄H₄₉N₂O₆S⁺ (M+H)⁺
- 49 613.3306, found 613.3307.

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Fig.S5 CDK9 is a direct target of acetyl-bufalin

- A. Viability of PC-9 cells exposed to biotin-labeled bufalin as determined by MTT assay. **B**.
- Representative image of an experimental microarray (blue = negative control, red = positive control,
- 54 yellow= positive spot). C. The expression of proteins involved in the CDK9/STAT3 pathway in
- 55 H460 and PC-9 cells was examined by Western blotting after acetyl-bufalin (Ac-bu) and SNS-032
- SNS) treatment, and the relative intensity was calculated. *p < 0.05, **p < 0.01, ***p < 0.00,
- 57 ****P<0.0001.

58 59 Fig.S6 Acetyl-bufalin inhibited the CDK9/STAT3 signaling pathway 60 A. CDK9 and STAT3 expression levels by western blotting after transduction with CDK9 siRNA 61 in H460 cells, the relative intensity was calculated. B. Western blot analysis of CDK9 and STAT3 62 expression levels in H460 and PC-9 cells transfected with CDK9 siRNA1 and treated for 30 min 63 with or without IL-6 (25 ng/ml). The relative intensity was calculated. *p < 0.05, **p < 0.01, ***p < 0.01, *** 64 <0.00, ****P<0.0001. 65 66 Fig.S7 High CDK9 expression in human NSCLC tissues 67 A. Protein expression of CDK9 in each paired human NSCLC tissue sample (T) and adjacent normal 68 lung tissues (N) from the same patient was detected by western blotting. B. Immunohistochemical 69 analysis for P-CDK9 from human NSCLC tissue samples and normal tissue samples. 70 71 Supplementary Table 1: IC50 of the main active components of cinobufagin in different NSCLC 72 cells. 73 **Supplementary Table 2:** The clinicopathologic characteristics of samples 74

Supplementary Table 1

 IC_{50} of main active components of cinobufagin in different NSCLC cells

| Drug | | IC ₅₀ μM | |
|------------------------|--------|---------------------|---------|
| | PC-9 | A549 | H460 |
| Arenobufagin | 0.46 | 0.01274 | |
| Bufalin | 0.1409 | 0.08323 | 0.06472 |
| Bufotalin | 0.8133 | | |
| Pseudobufarenogin | > 100 | 4.268 | |
| Cinobufagin | 0.4144 | | |
| Cinobufotalin | 0.5868 | 0.8186 | |
| Desacetylcinobufagin | 10.06 | 1.818 | |
| Desacetylcinobufotalin | 2.888 | 0.8867 | |
| Gamabufotalin | 0.281 | 0.7302 | |
| Resibufogenin | 4.64 | 1.129 | 1.061 |
| Telocinobufagin | 0.5741 | 0.1457 | |
| Marinobufagin | 15.00 | 1.547 | 1.301 |
| Resibufagin | 16.02 | | |

Supplementary Table 2

The clinicopathologic characteristics of samples

| Number | Age | Sex | Pathological diagnosis | TNM |
|--------|-----|--------|-------------------------|---------|
| 1 | 63 | Male | Adenocarcinoma | T1bN0M0 |
| 2 | 59 | Male | Squamous cell carcinoma | T3N0M0 |
| 3 | 52 | Female | Adenocarcinoma | T1bN0MX |
| 4 | 64 | Female | Adenocarcinoma | T1cN1MX |
| 5 | 71 | Male | Adenocarcinoma | T2aN2M0 |
| 6 | 67 | Female | Adenocarcinoma | T1bN0MX |
| 7 | 74 | Female | Squamous cell carcinoma | T4NXM0 |
| 8 | 66 | Female | Adenocarcinoma | T1bN0M0 |
| 9 | 79 | Female | Adenocarcinoma | T1cN0M0 |
| 10 | 68 | Male | Adenocarcinoma | T1bN0M0 |
| 11 | 60 | Male | Adenocarcinoma | T1bN0MX |
| 12 | 54 | Male | Adenocarcinoma | T1bN0M0 |