# Anticancer Mechanism of 7- $\alpha$-Hydroxyfrullanolide on Microtubules and Computational Prediction of its Target Binding in Triple-Negative Breast Cancer Cells 

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Figure S1: Maximum absorbance $\left(\lambda_{\max }\right)$ identification at $1,006.77 \mu \mathrm{M} 7 \mathrm{HF}$ (A-C) The peaks were identified in $1,006.77 \mu \mathrm{M}$ of sample $\mathrm{n} 1, \mathrm{n} 2$, and n 3 , respectively


Figure S2: Protein expressions were evaluated using western blotting. (A) Bub3, cyclin B1, and p-Cdk1 (Tyr15) were detected at $6 \mu \mathrm{M} 7 \mathrm{HF}$ for 0,12 , 24, and 48 h . (B) Bub3, cyclin B1, and p-Cdk1 (Tyr15) were performed at $0,6,12$, and $24 \mu \mathrm{M} 7 \mathrm{HF}$ for 24 h . These proteins were depicted along with their $\beta$-actin. The results were performed in two-three independent experiments (n1-n2/n1-n3). The experiment n1-n3 used for calculation of band intensity (or relative expression) from indicated protein and $\beta$-actin as representing in histogram Figure 2D-E. The full original western blot images in Figure 2D and Figure 2E were showed in page $3-5$ and page 6-8 as follows: Bub3, Cyclin B1, and p-Cdk1 (Tyr15). In full original blot, 0,6 , 12 , and $24 \mu \mathrm{M} ; 7 \mathrm{HF}$ treatment does $(0,6,12$, and $24 \mu \mathrm{M}), 0,12,24$, and $48 \mathrm{~h} ; 7 \mathrm{HF}$ treatment times ( $0,12,24$, and 48 h ), Black outer square; border of full images, Red inner square and black head arrow; indicated protein bands, \#; The blots are shown the same in Figure 2D-E.

Bub3 (40 KDa) and $\beta$-actin (42 KDa)

| Bub3 (n1) <br> $\begin{array}{lllll}0 & 12 & 24 & 48 & \text { (h) }\end{array}$ | Bub3 (n2) <br> $\begin{array}{lllll}0 & 12 & 24 & 48 & \text { (h) }\end{array}$ | \# The result represented in Figure 2D |
| :---: | :---: | :---: |
| $\beta \text {-actin (n1) }$ $\begin{array}{llll} 0 & 12 & 24 & 48 \end{array}$ <br> (h) | $\beta$-actin (n2) <br> $\begin{array}{llll}0 & 12 & 24 & 48\end{array}$ <br> (h) | $\beta$-actin (n3) <br> $\begin{array}{lllll}0 & 12 & 24 & 48 & \text { (h) }\end{array}$ <br> \# The result represented in Figure 2D |

Cyclin B1 (55 KDa) and $\beta$-actin (42 KDa)

| Cyclin B1 (n1) $\begin{array}{lllll} 0 & 12 & 24 & 48 & \text { (h) } \end{array}$ | Cyclin B1 (n2) $\begin{array}{lllll} 0 & 12 & 24 & 48 & \text { (h) } \end{array}$ <br> \# The result represented in Figure 2D | Cyclin B1 (n3) $\begin{array}{lllll} 0 & 12 & 24 & 48 & \text { (h) } \end{array}$ |
| :---: | :---: | :---: |
| $\beta$-actin (n1) $\begin{array}{llll} 0 & 12 & 24 & 48 \end{array}$ <br> (h) | $\beta-\operatorname{actin}(\mathbf{n 2})$ $\begin{array}{ccccc} 0 & 12 & 24 & 48 & \text { (h) } \\ & & & & \\ & & & & \end{array}$ <br> \# The result represented in Figure 2D | $\beta$-actin (n3) <br> $\begin{array}{lllll}0 & 12 & 24 & 48 & \text { (h) }\end{array}$ |

## p-Cdk1 (Tyr15) (34 KDa) and $\beta$-actin (42 KDa)



Bub3 (40 KDa) and $\beta$-actin (42 KDa)

| Bub3 (n1) $\begin{array}{llllll} 0 & 6 & 12 & 24 & (\mu M) \\ & - & - & \end{array}$ | Bub3 (n2) <br> \# The result represented in Figure 2E | Bub3 (n3) |
| :---: | :---: | :---: |
| $\beta$-actin (n1) $\begin{array}{lllll} 0 & 6 & 12 & 24 & (\mu M) \end{array}$ |  | \# The result represented in Figure 2E |

Note: Bub3 at $6 \mu \mathrm{M}$ was used intensity calculation for n1-n2

Cyclin B1 (55 KDa) and $\beta$-actin (42 KDa)

| $$ | Cyclin B1 (n2) <br> \# The result represented in Figure 2E | $\begin{gathered} \text { Cyclin B1 (n3) } \\ \ldots-\infty-\square \end{gathered}$ |
| :---: | :---: | :---: |
| $\beta$-actin (n1) $\begin{array}{cccccc} 0 & 6 & 12 & 24 & (\mu M) \\ & & & & & \end{array}$ | $\beta-\operatorname{actin}(\mathbf{n 2})$ | \# The result represented in Figure 2E |

Note: Cyclin B1 at $6 \mu \mathrm{M}$ was used intensity calculation for n1-n2

## p-Cdk1 (Tyr15) (34 KDa) and $\beta$-actin (42 KDa)

| p-Cdk1 (n1) <br> $\begin{array}{lllll}0 & 6 & 12 & 24 & (\mu M)\end{array}$ <br> \# The result represented in Figure 2E | p-Cdk1 (n2) | p-Cdk1 (n3) |
| :---: | :---: | :---: |
| $\begin{gathered} \beta \text {-actin (n1) } \\ \text { \# The result represented in Figure } 2 \mathrm{E} \end{gathered}$ |  |  |



Figure S3: Cell population was evaluated bromodeoxyuridine plus propidium iodide staining using fluorescence assisted cell sorting analysis in 7HF-treated cells at $0,6,12$, and $24 \mu \mathrm{M}$ for $12 \mathrm{~h}(\mathbf{A})$ and $24 \mathrm{~h}(\mathbf{B})$. Left-lower, middle-top, and right-lower quadrants represent the number of cells in G1, S, and G2 phases, respectively. BrdU; bromodeoxyuridine, n1-n3; the sample 1,2 , and 3 .


Figure S4: Protein expressions were evaluated using western blotting. (A) Rb and $\mathrm{p}-\mathrm{Rb}$ Ser780, (B) Chk1 and p-Chk1 Ser345, (C) Chk2 and p-Chk2 Ser516, and (D) p-H2AX Ser 139 were examined at 6 and $24 \mu \mathrm{M} 7 \mathrm{HF}$ for 24 h . These proteins were depicted along with their $\beta$-actin. The results were performed in two-three independent experiments (n1$\mathrm{n} 2 / \mathrm{n} 1-\mathrm{n} 3$ ). The experiment $\mathrm{n} 1-\mathrm{n} 3$ used for calculation of band intensity (or relative expression) from indicated protein and $\beta$-actin as representing in histogram Figure 3D. The full original western blot images in Figure 3D were showed in page 11-17 as follows: Rb, pRb Ser780, Chk1, p-Chk1 Ser345, Chk2, p-Chk2 Ser516, and p-H2AX Ser139. In full original blot, 0,6 , and $24 \mu \mathrm{M}$; 7 HF treatment does ( 0,6 and $24 \mu \mathrm{M}$ ), Black outer square; border of full images, Red inner square and black head arrow; indicated protein bands, \# and arrow; The blots are shown the same in Figure 3D. Note. Original full fluorescence western blot images were also depicted in page 18-21 as follows: $\mathrm{Rb} / \mathrm{p}-\mathrm{Rb}$, Chk1/p-Chk1, Chk2/pChk2, and p-H2AX. $\beta$-actin was along with the proteins. M; Protein marker, White square and head arrow; indicated protein bands.

Rb (110 KDa) and p-Rb Ser780 (110 KDa)

$\beta$-actin (42 KDa) of Rb and p-Rb


Chk1 (56 KDa) and p-Chk1 Ser345 (56 KDa)

|  |  |
| :---: | :---: |
|  |  |

$\beta$-actin (42 KDa) of Chk1 and p-Chk1

| $\beta$-actin (n1) <br> ( $\mu \mathrm{M}$ ) |  |
| :---: | :---: |

Chk2 (62 KDa) and p-Chk2 Ser516 (62 KDa)

| Chk2 (n1) |  <br> \# The result represented in Figure 3D |
| :---: | :---: |
|  |  |

$\beta$-actin (42 KDa) of Chk2 and p-Chk2


## p-H2AX Ser139 ( 17 KDa)


$\beta$-actin (42 KDa) of p-H2AX

| $\begin{gathered} \beta-\operatorname{actin}(\mathrm{n} 1) \\ \begin{array}{llll} 0 & 6 & 24 & (\mu \mathrm{M}) \end{array} \\ -- \end{gathered}$ |  |
| :---: | :---: |

Rb (110 KDa, green) and p-Rb Ser780 (110 KDa, red)

$\beta$-actin ( 42 KDa , green) of $\mathbf{R b}$ and p -Rb


Chk1 (56 KDa, red) and p-Chk1 Ser345 (56 KDa, green)

$\beta$-actin (42 KDa, green) of Chk1 and p-Chk1


Chk2 (62 KDa, red) and p-Chk2 Ser516 (62 KDa, green)

$\beta$-actin (42 KDa, green) of Chk2 and p-Chk2

p-H2AX (17 KDa, green)

$\beta$-actin (42 KDa, green) of p-H2AX


| 7HF concentration $(\mu \mathrm{M})$ | Absorbance ( 268 nm ) |  |  |
| :---: | :---: | :---: | :---: |
|  | \#1 | \#2 | \#3 |
| 125.85 | 0.09184 | 0.08400 | 0.05441 |
| 251.69 | 0.34277 | 0.38235 | 0.29000 |
| 503.38 | 0.41608 | 0.44129 | 0.49000 |
| 1,006.77 | 0.68008 | 0.64546 | 0.63684 |
| 2,013.53 | 0.70305 | 0.73100 | 0.91060 |
| 4,027.06 | 0.96164 | 0.97003 | 0.83350 |

Table S1: Absorbance of 7HF ranged from 125.85-4,027.06 $\mu \mathrm{M}$ at 268 nm . \#1-\#3, the sample n1-n3

| Time (h) | Absorbance (268 nm) |  |  |  |  |  |
| ---: | ---: | ---: | ---: | ---: | ---: | ---: |
|  | 7HF-cell culture |  |  | 7HF-cell free medium |  |  |
|  | $\# 1$ | $\# 2$ | $\# 3$ | $\# 1$ | $\# 2$ | $\# 3$ |
| 0.00 | 0.000 | 0.000 | 0.000 | 0.000 | 0.000 | 0.000 |
| 0.25 | -0.037 | -0.091 | -0.035 | 0.188 | 0.256 | 0.200 |
| 0.50 | 0.008 | -0.022 | -0.003 | 0.359 | 0.333 | 0.355 |
| 0.75 | 0.205 | 0.239 | 0.141 | 0.249 | 0.226 | 0.306 |
| 1.00 | 0.240 | 0.190 | 0.211 | 0.292 | 0.280 | 0.312 |
| 5.00 | -0.024 | 0.025 | -0.004 | 0.279 | 0.381 | 0.363 |
| 9.00 | -0.181 | -0.092 | -0.095 | 0.387 | 0.332 | 0.358 |
| 13.00 | -0.207 | -0.209 | -0.099 | 0.196 | 0.212 | 0.285 |
| 17.00 | -0.300 | -0.282 | -0.177 | 0.230 | 0.094 | 0.172 |
| 24.00 | -0.269 | -0.295 | -0.248 | -0.104 | -0.063 | -0.150 |

Table S2: $1,006.77 \mu \mathrm{M} 7 \mathrm{HF}$ was measured at wavelength 268 nm for $0.25-24 \mathrm{~h}$ both cell- and cell free condition. \#1-\#3, the sample n1-n3

