

Ginsenoside Rh2 epigenetically regulates cell-mediated immune pathway to inhibit proliferation of MCF-7 breast cancer cells

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Supplementary Information

Table S1. Sequences of primers employed for real time polymerase chain reaction

Gene Symbol	Primer sequence (5'→3')
<i>SAMM50</i>	F : TCAAGGTGATGACGCACTTC
	R : AAGCTTGAGGCCAAGTACCA
<i>CASP1</i>	F : AAATTTTCCGCAAGGTTCG
	R : ACTTCCTGCCACAGACATT
<i>INSL5</i>	F : ATCTCAGAAGTGCGGAGCAA
	R : CAGCTTGCTGAGCTTGAGG
<i>TCF4</i>	F : AAAAGGGACAGCGAGTTTGA
	R : GATGGAGCATAGACCGCTTT
<i>OR52A1</i>	F : AAT CTG AGC GCA GTC TCC AT
	R : CAC ATT TGA AGC AAG CAG GA
<i>CLINT1</i>	F : CAGCCAGTCAACAGGAGGAT
	R : TGTTGCTGTTACTTGGGAAGG
<i>ST3GAL4</i>	F : ATCTCCCGGGAAGACAGTTT
	R : CGGGAGTAGTTGCCAAAGAG
<i>C1orf198</i>	F : TCCTGGGAAACAAAGAGTCAG
	R : GGAAGCTTTGGACAGTGGTC
<i>RNLS</i>	F : AGGACTCAGGGGGAAGAATG
	R : TGGCATAATGAGGAGTGCAG
<i>GAPDH</i>	F : ACATCGCTCAGACACCATG
	R : TGTAGTTGAGGTCAATGAAGGG

Fig. S1. Effect of ginsenoside Rh2 on the *Alu* methylation level in MCF-7 cells. The methylation levels of the three CpGs on the *Alu* from the MCF-7 cells were determined by pyrosequencing after treatment with Rh2. (A) The *Alu* sequence adopted in this study. The analyzed three CpGs are indicated in red and numbered. (B) Bar graphs representing the methylation levels of CpGs of *Alu* in the MCF-7. Five independent experiments were carried out for individual CpG and average levels are given with the standard errors.

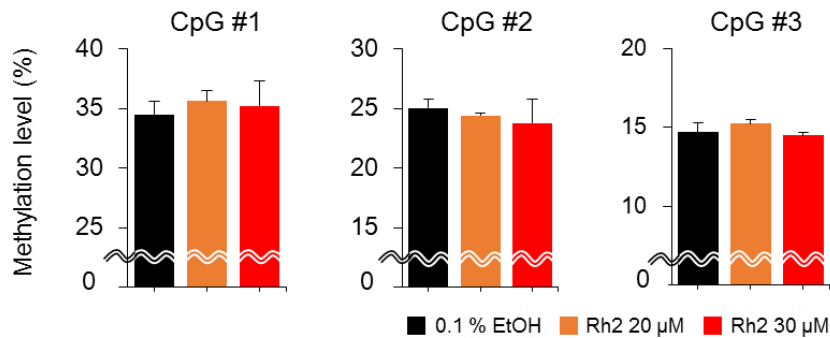
A

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1  ggccgggcgga ggccggggcgc ggtggctcac gcctgtaate ccagcacttt gggaggaaga
61  tcacctgagg tcaggagtgc gagaccagcc tggccaacat ggtgaaaccc cgtctctact
121  aaaaatacaa aaattagccg ggcgtggtgg cgccgcgcctg taatcccagc tactcgggag
                                     1 2 3
181  gctgaggcag gagaatcgc tgaacccggg aggcggaggt tgcagtgagc cgagatcgcg
231  ccactgcact ccagcctggg cgacagagcg agactccgtc tcaaaaaaaaa

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B



EtOH, ethanol.