

**GLSP and it-derived triterpenes attenuate atherosclerosis and aortic calcification by improving ABCA1/G1-mediated cholesterol efflux and inactivating RUNX2-mediated osteogenesis**

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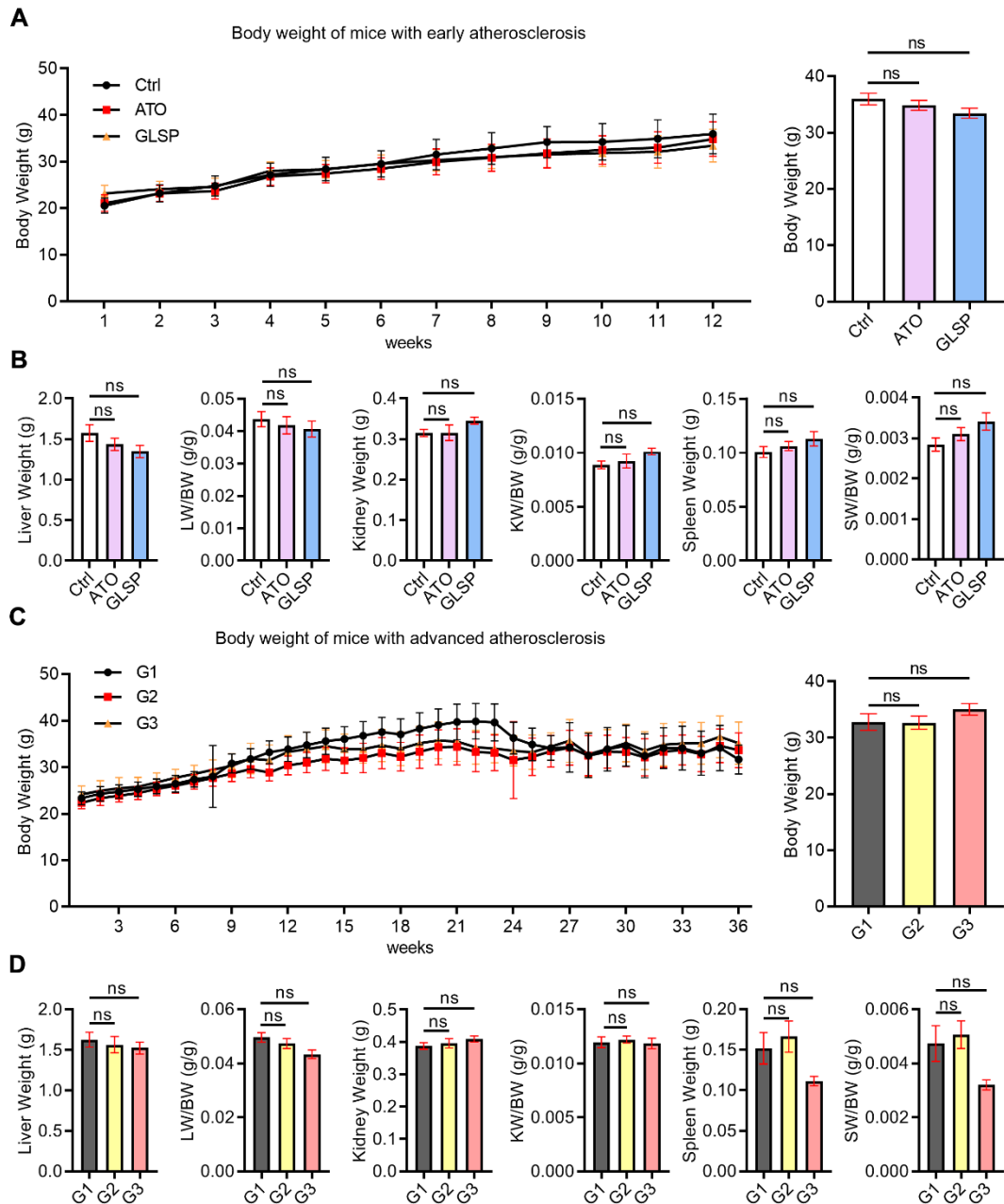
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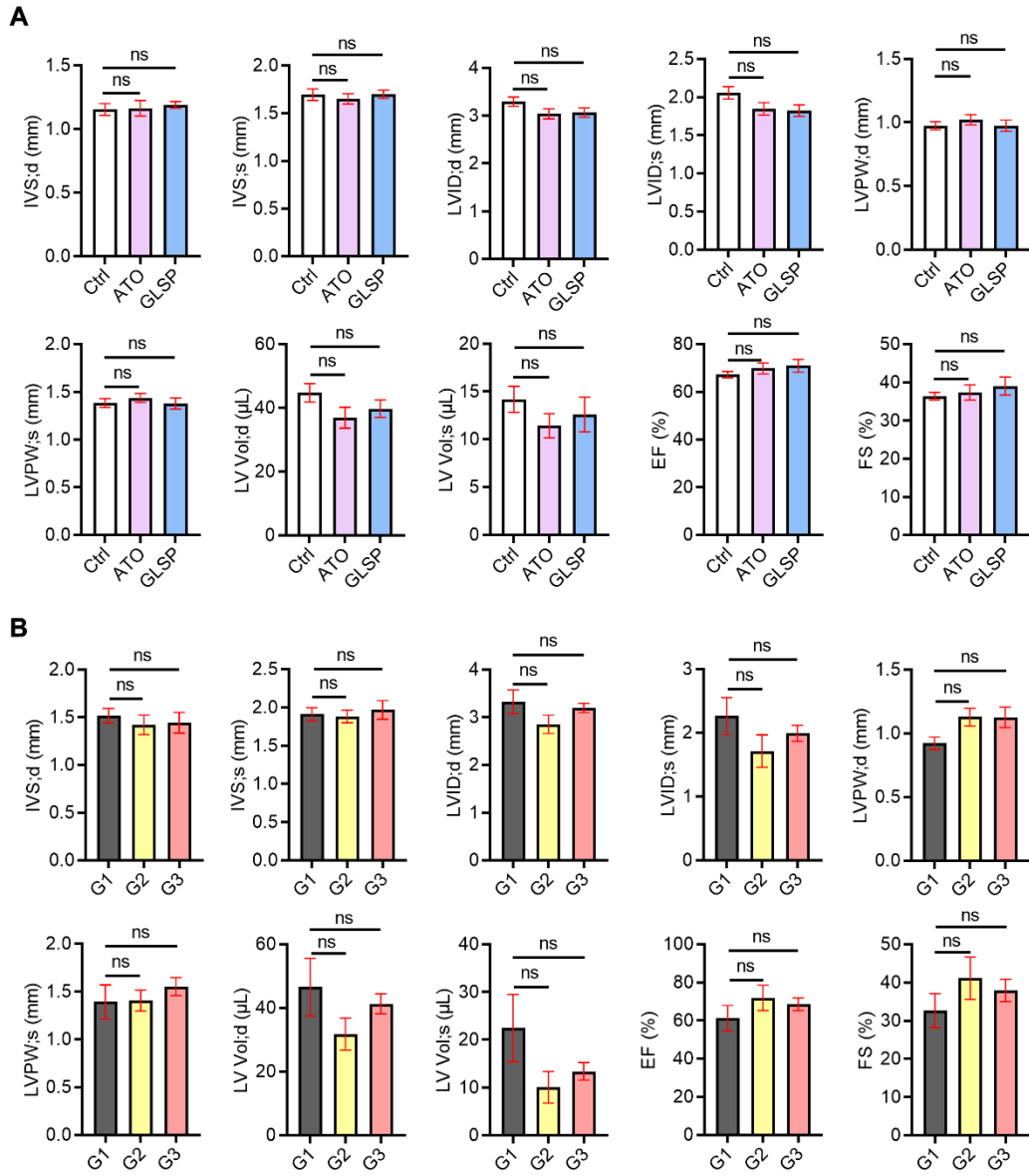
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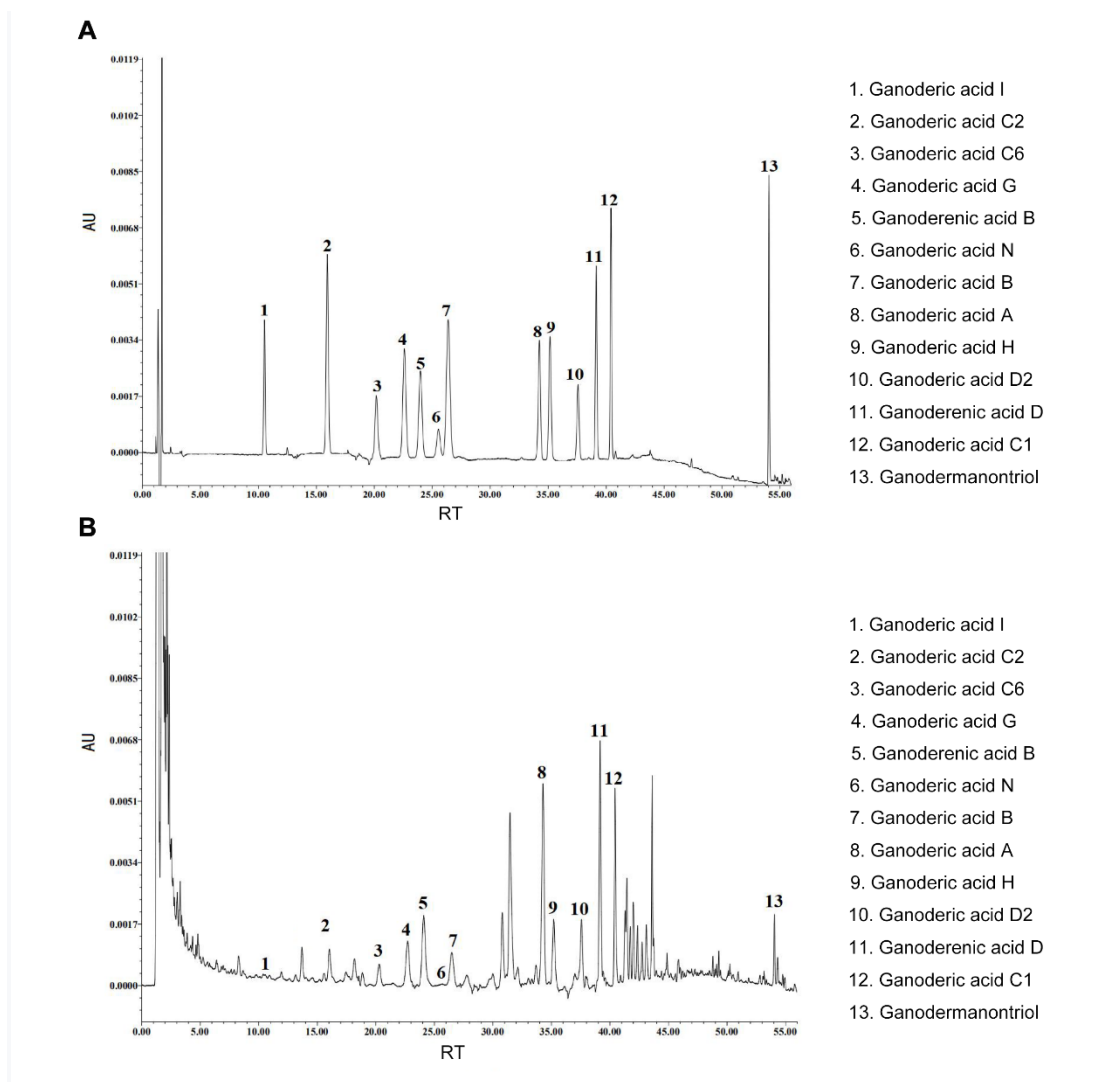
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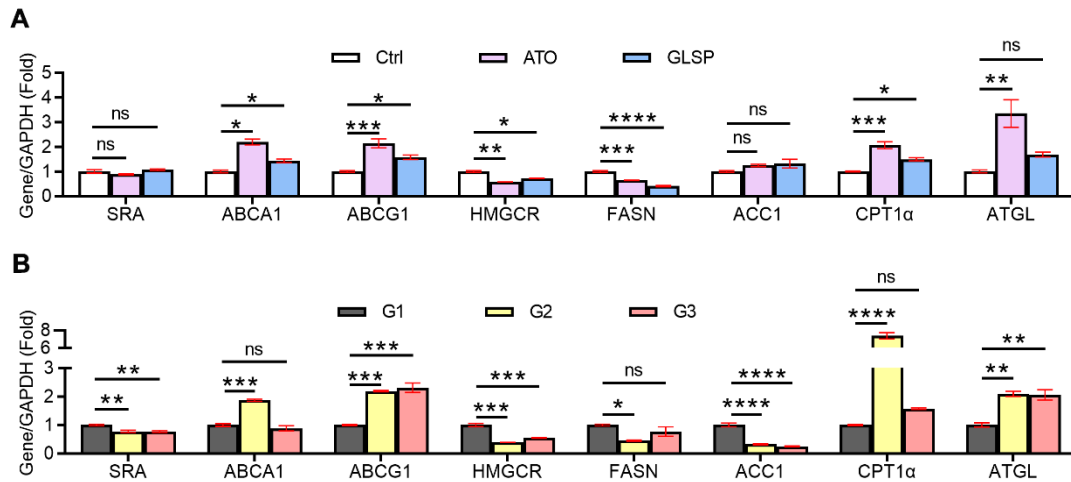
**Figure S1. Body and organ weights of early and advanced  $LDLR^{-/-}$  mice.** (A) Early  $LDLR^{-/-}$  mice weight change curve, endpoint weight and endpoint change difference. (B) Liver, kidney and spleen weights, and liver-to-body, kidney-to-body ratios and spleen-to-body ratios in early  $LDLR^{-/-}$  mice. (C) Advanced  $LDLR^{-/-}$  mice weight change curve, endpoint weight and endpoint change difference. (D) Liver, kidney and spleen weights, and liver-to-body, kidney-to-body ratios and spleen-to-body ratios in advanced  $LDLR^{-/-}$  mice. \* $P < 0.05$ , \*\* $P < 0.01$ , \*\*\* $P < 0.001$ , \*\*\*\* $P < 0.0001$  compared with the Ctrl or the G1 group.



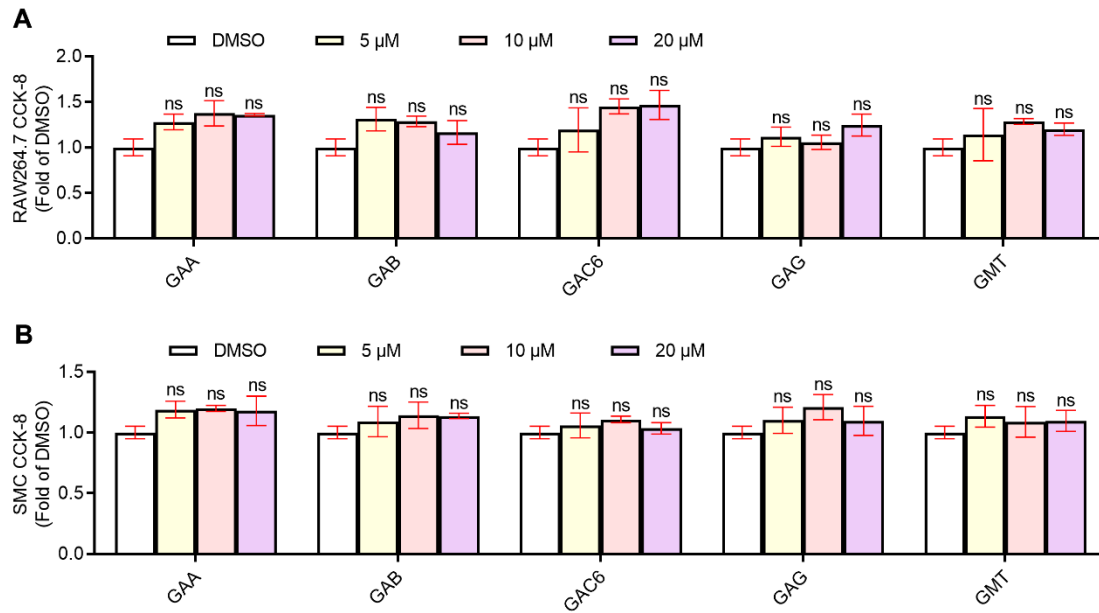
**Figure S2. Cardiac ultrasound data in early and advanced LDLR<sup>-/-</sup> mice. (A) Cardiac ultrasound data in early LDLR<sup>-/-</sup> mice. (B) Cardiac ultrasound data in advanced LDLR<sup>-/-</sup> mice.**



**Figure S3. Chromatogram of thirteen triterpene components in GLSP by HPLC. (A) Chromatogram of standard. (B) Chromatogram of sample.**



**Figure S4. Effects of GLSP on the expression of genes related to lipid metabolism in LDLR<sup>-/-</sup> mice peritoneal macrophages. (A) Effects of GLSP on the expression of genes related to lipid metabolism in early LDLR<sup>-/-</sup> mice peritoneal macrophages. (B) Effects of GLSP on the expression of genes related to lipid metabolism in advanced LDLR<sup>-/-</sup> mice peritoneal macrophages. \*P<0.05, \*\*P<0.01, \*\*\*P<0.001, \*\*\*\*P<0.0001 compared with the Ctrl or G1 group.**



**Figure S5. Cell viability of RAW264.7 and HASMC was determined by CCK-8 assay.** (A)  $4 \times 10^3$  RAW264.7 cells were seeded in 96-well plates, GAA, GAB, GAC6, GAG and GMT (5, 10, 20  $\mu$ M) were administered and cell viability was measured after 16 h. (B)  $4 \times 10^3$  HASMCs were seeded in 96-well plates, GAA, GAB, GAC6, GAG and GMT (5, 10, 20  $\mu$ M) were administered and cell viability was measured after 16 h. ns: no significance.

**Table S1. Identification and content determination of thirteen triterpene components in *Ganoderma lucidum* spore powder**

Number	Compound name	Content (%)
1	Ganoderic acid I	0.004
2	Ganoderic acid C2	0.026
3	Ganoderic acid C6	0.026
4	Ganoderic acid G	0.058
5	Ganoderenic acid B	0.036
6	Ganoderic acid N	0.002
7	Ganoderic acid B	0.044
8	Ganoderic acid A	0.338
9	Ganoderic acid H	0.100
10	Ganoderic acid D2	0.094
11	Ganoderenic acid D	0.070
12	Ganoderic acid C1	0.112
13	Ganodermanontriol	0.022
Gross weight		0.932

**Table S2. The sequences of primers for qRT-PCR analysis**

Gene	Forward	Backward
<i>Mus GAPDH</i>	AGGTCGGTGTGAACGGATTG	TGTAGACCATGTAGTTGAGGTCA
<i>Mus eNOS</i>	GGCTGGGTTTAGGGCTGTG	CTGAGGGTGTCGTAGGTGATG
<i>Mus IL-1<math>\beta</math></i>	GCAACTGTTCTGAACTCAACT	ATCTTTTGGGGTCCGTCAACT
<i>Mus IL-10</i>	GCTCTACTGACTGGCATGAG	CGCAGCTCTAGGAGCATGTG
<i>Mus Arg1</i>	CTCCAAGCCAAAGTCCTTAGAG	AGGAGCTGTCATTAGGGACATC
<i>Mus TGF<math>\beta</math></i>	CTCCCGTGGCTTCTAGTGC	GCCTTAGTTTGGACAGGATCTG
<i>Mus SRA</i>	GCACAATCTGTGATGATCGCT	CCCAGCATCTTCTGAATGTGAA
<i>Mus LXR<math>\alpha</math></i>	ATGTCTTCCCCACAAGTTCT	GACCACGATGTAGGCAGAGC
<i>Mus ABCA1</i>	AAAACCGCAGACATCCTTCAG	CATACCGAAACTCGTTCACCC
<i>Mus ABCG1</i>	CTTTCCTACTCTGTACCCGAGG	CGGGGCATTCCATTGATAAGG
<i>Mus HMGCR</i>	TGTTACCCGGCAACAACAAGA	CCGCGTTATCGTCAGGATGA
<i>Mus FASN</i>	GGCTCTATGGATTACCCAAGC	CCAGTGTTTCGTTCTCCGGA
<i>Mus ACC1</i>	CTCCCATTGATAATTGGGTCTG	TCGACCTTGTTTTACTAGGTGC
<i>Mus CPT1<math>\alpha</math></i>	AGATCAATCGGACCCTAGACAC	CAGCGAGTAGCGCATAGTCA
<i>Mus ATGL</i>	CTGAGAATCACCATTCCCACATC	CACAGCATGTAAGGGGGAGA
<i>Mus SREBP1</i>	GATGTGCGAACTGGACACAG	CATAGGGGGCGTCAAACAG
<i>Mus SCD1</i>	TTCTTGCGATACTCTGGTGC	CGGGATTGAATGTTCTTGTCGT
<i>Mus HSL</i>	TGGCACACCATTTGACCTG	TTGCGGTTAGAAGCCACATAG
<i>Mus PPAR<math>\gamma</math></i>	GGAAGACCACTCGCATTCTT	GTAATCAGCAACCATTGGGTCA