Supporting Information

Glycosylation of luteolin in hydrophilic organic solvents and

structure-antioxidant relationships of luteolin glycosides

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Figure S1. The ¹H NMR spectrum of luteolin in DMSO- d_6 (500 MHz).



Figure S2. The ¹H NMR spectrum of luteolin-3'-O-glucoside in DMSO-*d*₆ (500 MHz).



Figure S3. The ¹H NMR spectrum of luteolin-4'-O-glucoside in DMSO-*d*₆ (500 MHz).



Figure S4. The ¹H NMR spectrum of luteolin-7-O-glucoside in DMSO-*d*₆ (500 MHz).



Figure S5. The ¹H NMR spectrum of luteolin-7,3'-di-O-glucoside in DMSO-d₆ (500 MHz).



Figure S6. The ¹H NMR spectrum of luteolin-7,4'-di-O-glucoside in DMSO-*d*₆ (500 MHz).



Figure S7. The HR-MS spectrum of quercetin glucosides.



Figure S8. The HR-MS spectrum of apigenin glucosides.



Figure S9. The HR-MS spectrum of naringenin glucosides.



Figure S10. The HR-MS spectrum of baicalein glucoside.



Figure S11. The HR-MS spectrum of genistein glucoside.



Figure S12. The HR-MS spectrum of daidzein glucoside.



Figure S13. The HR-MS spectrum of emodin.



Figure S14. The HR-MS spectrum of emodin-6-O-glucoside.



Figure S15. The HR-MS spectrum of catenarin.



Figure S16. The HR-MS spectrum of catenarin-6-O-glycoside.



Figure S17. The HR-MS spectrum of skyrin.



Figure S18. The HR-MS spectrum of skyrin-6-O-glycoside.



Figure S19. The HR-MS spectrum of skyrin-6,6'-di-O-glycoside.







Figure S21. The HR-MS spectrum of emodin bianthrone (rac).







Figure S23. The HR-MS spectrum of emodin bianthrone (rac)-6-O-glycoside.



Figure S24. The HR-MS spectrum of emodin bianthrone (*meso*)-6,6'-di-O-glycoside.



Figure S25. The HR-MS spectrum of emodin bianthrone (rac)- 6,6'-di-O-glycoside.



Figure S26. The HR-MS spectrum of emodin-catenarin bianthrone.



Figure S27. The HR-MS spectrum of emodin-catenarin bianthrone-O-glucoside.

16S rDNA sequence of Bacillus cereus A46

GAGCGTAGCGTGCGGTCGAGCGAATGAGATTAAGAGCTTGCTGTTATGAAGTTAGCGGCGGAC GGGTGAGTAACACGTGGGTAACCTGCCCATAAGACTGGGATAACTCCGGGAAACCGGGGCTAA TACCGGATAACATTTTGAACCGCATGGTTCGAAATTGAAAGGCGGCTTCGGCTGTCACTTATGG ATGGACCCGCGTCGCATTAGCTAGTTGGTGAGGTAACGGCTCACCAAGGCAACGATGCGTAGC CGACCTGAGAGGGTGATCGGCCACACTGGGACTGAGACACGGCCCAGACTCCTACGGGAGGC GGCTTTCGGGTCGTAAAACTCTGTTGTTAGGGAAGAACAAGTGCTAGTTGAATAAGCTGGCACC TTGACGGTACCTAACCAGAAAGCCACGGCTAACTACGTGCCAGCAGCCGCGGTAATACGTAGG GAAAGCCCACGGCTCAACCGTGGAGGGTCATTGGAAACTGGGAGACTTGAGTGCAGAAGAGG AAAGTGGAATTCCATGTGTAGCGGTGAAATGCGTAGAGATATGGAGGAACACCAGTGGCGAAG GCGACTTTCTGGTCTGTAACTGACACTGAGGCGCGAAAGCGTGGGGGGGCAAACAGGATTAGAT ACCCTGGTAGTCCACGCCGTAAACGATGAGTGCTAAGTGTTAGAGGGTTTCCGCCCTTTAGTGC TGAAGTTAACGCATTAAGCACTCCGCCTGGGGAGTACGGCCGCAAGGCTGAAACTCAAAGGAA TTGACGGGGGCCCGCACAAGCGGTGGAGCATGTGGTTTAATTCGAAGCAACGCGAAGAACCTT ACCAGGTCTTGACATCCTCTGAAAACCCTAGAGATAGGGCTTCTCCTTCGGGAGCAGAGTGACA GGTGGTGCATGGTTGTCGTCAGCTCGTGTCGTGAGATGTTGGGTTAAGTCCCGCAACGAGCGCA ACCCTTGATCTTAGTTGCCATCATTAAGTTGGGCACTCTAAGGTGACTGCCGGTGACAAACCGG AGGAAGGTGGGGATGACGTCAAATCATCATGCCCCTTATGACCTGGGCTACACACGTGCTACA ATGGACGGTACAAAGAGCTGCAAGACCGCGAGGTGGAGCTAATCTCATAAAACCGTTCTCAGT TCGGATTGTAGGCTGCAACTCGCCTACATGAAGCTGGAATCGCTAGTAATCGCGGATCAGCATG CCGCGGTGAATACGTTCCCGGGCCTTGTACACACCCCCGTCACACCACGAGAGTTTGTAACAC CCGAAGTCGCTGGGGTCACCTCGTAGCAACAGCCGCCTAGAGTC