Supporting Information for:

Assessing the Regioselectivity OleD-Catalyzed Glycosylation with a Diverse Set of Acceptors

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Daidzein 7-O-β-D-glucoside (4): ¹H NMR (DMSO-d6, 500 MHz): 9.55 (s, 1H), 8.40 (s, 1H), 8.04 (d, J = 9.0 Hz, 1H), 7.40 (d, J = 8.5 Hz, 2H), 7.23 (d, J = 2.5 Hz, 1H), 7.14 (dd, J = 9.0, 2.0 Hz, 1H), 6.81 (d, J = 8.5 Hz, 2H), 5.43 (d, J = 5.0 Hz, 1H), 5.14 (d, J = 4.5 Hz, 1H), 5.10 (d, J = 7.0 Hz, 1H), 5.08(d, J = 5.0 Hz, 1H), 4.61 (t, J = 5.5 Hz, 1H), 3.71 (m, 1H), 3.45 (m, 1H), 3.4-3.1 (m, 4H); ¹³C NMR (DMSO-d6, 125 MHz): 180.1, 166.8, 162.6, 162.4, 158.7, 135.5(2C), 132.4, 129.1, 127.7, 123.9, 121.0, 120.4 (2C), 108.8, 105.4, 82.7, 82.0, 78.6, 75.1, 66.2; HRESIMS *m/z* 439.1003 [M+Na]⁺ (cacld for C₂₁H₂₀O₉Na, 439.1000).

Daidzein 4'-O-β-D-glucoside (5): ¹H NMR (DMSO-d6, 500 MHz): 10.8 (s, 1H), 8.35 (s, 1H), 7.97 (d, J = 9.0 Hz, 1H), 7.50 (d, J = 8.5 Hz, 2H), 7.08 (d, J = 8.5 Hz, 2H), 6.94 (dd, J = 9.0, 2.5 Hz, 1H), 6.88 (d, J = 2.5 Hz, 1H), 5.32 (d, J = 5.0 Hz, 1H), 5.08 (d, J = 4.5 Hz, 1H), 5.01 (d, J = 5.5 Hz, 1H), 4.90 (d, J = 7.0 Hz, 1H), 4.56 (t, J = 5.0 Hz, 1H), 3.70 (m, 1H), 3.47 (m, 1H), 3.4-3.1 (m, 4H); ¹³C NMR (DMSO-d6, 125 MHz): 179.9, 168.0, 162.8, 162.5, 158.7, 135.4(2C), 132.7, 130.9, 128.5, 122.0, 121.4(2C), 120.6, 107.6, 105.8, 82.5, 82.1, 78.7, 75.2, 66.2; HRESIMS *m/z* 439.1003 [M+Na]⁺ (cacld for C₂₁H₂₀O₉Na, 439.1000).

Daidzein 7,4'-di-*O*-**β**-**D**-glucoside (6): ¹H NMR (DMSO-d6, 500 MHz): 8.46 (s, 1H), 8.05 (d, J = 9.0Hz, 1H), 7.52 (d, J = 8.5, 2H), 7.25 (d, J = 2.0, 1H), 7.15 (dd, J = 9.0, 2.0 Hz, 1H), 7.09 (d, J = 8.5, 2H), 5.11 (d, J = 7.0, 1H), 4.91 (d, J = 7.5, 1H), 3.80-3.10 (m, 10H); ¹³C NMR (DMSO-d6, 125 MHz): 174.6, 161.5, 157.2, 157.0, 153.8, 130.0 (2C), 126.9, 125.3, 123.3, 116.0(2C), 118.4, 115.6, 103.4, 100.3, 100.0, 77.2, 77.0, 76.5, 76.4, 73.2, 73.1, 69.7, 69.6, 60.65, 60.60; HRESIMS *m*/*z* 601.1526 [M+Na]⁺ (cacld for C₂₇H₃₀O₁₄Na, 601.1528).

Resveratrol 4'-O-β-D-glucoside (8): ¹H NMR (CD₃OD, 500 MHz): 7.47 (d, J = 9.0, 2H), 7.10 (d, J = 9.0, 2H), 7.02 (d, J = 16.5, 1H), 6.90 (d, J = 16.5, 1H), 6.488 (s, 1H), 6.484 (s, 1H), 6.2 (t, J = 2.0, 1H), 4.94 (d, J = 7.5, 1H), 3.92 (dd, J = 12.0, 2.0, 1H), 3.72 (dd, J = 12.0, 6.0, 1H), 3.52-3.36 (m, 4H); ¹³C NMR (CD₃OD, 125 MHz): 159.9 (2C), 158.8, 141.1, 133.4, 129.0, 128.73 (2C), 128.69, 118.1 (2C), 106.1(2C), 103.1, 102.4, 78.3, 78.1, 75.1, 71.5, 62.7; HRESIMS *m/z* 413.1189 [M+Na]⁺ (cacld for $C_{20}H_{22}O_8Na$, 413.1207).

Resveratrol 3-O-β-D-glucoside (9): ¹H NMR (CD₃OD, 500 MHz): 7.38 (d, J = 8.5, 2H), 7.03 (d, J = 16.0, 1H), 6.86 (d, J = 16.5, 1H), 6.80 (s, 1H), 6.77 (d, J = 8.5, 2H), 6.62 (s, 1H), 6.46 (t, J = 2.0, 1H), 4.90 (d, J = 7.5, 1H), 3.94 (dd, J = 12.0, 2.0, 1H), 3.72 (dd, J = 12.0, 6.0, 1H), 3.52-3.34 (m, 4H); ¹³C NMR (CD₃OD, 125 MHz): 160.6, 159.7, 158.6, 141.6, 130.5, 130.1, 129.1(2C), 126.8, 116.6 (2C), 108.5, 107.2, 104.6, 102.6, 78.4, 78.2, 75.1, 71.6, 62.7; HRESIMS *m*/*z* 413.1204 [M+Na]⁺ (cacld for C₂₀H₂₂O₈Na, 413.1207).

Resveratrol 3,4'-di-O-β-D-glucoside (10): ¹H NMR (CD₃OD, 500 MHz): 7.47 (d, J = 8.5, 2H), 7.09 (d, J = 9.0, 2H), 7.06 (d, J = 16.0, 1H), 6.94 (d, J = 16.0, 1H), 6.82 (s, 1H), 6.64 (s, 1H), 6.48 (t, J = 2.5, 1H), 4.93 (d, J = 7.5, 1H), 4.90 (d, J = 7.0, 1H), 3.94 (dd, J = 12.0, 2.0, 1H), 3.92 (dd, J = 12.0, 2.0, 1H), 3.72 (dd, J = 12.0, 5.0, 2H), 3.52-3.30 (m, 8H); ¹³C NMR (CD₃OD, 125 MHz): 160.6, 159.8, 158.9, 141.3, 133.2, 129.6, 128.8(2C), 128.3, 118.1(2C), 108.7, 107.3, 104.5, 102.5, 102.4, 78.39, 78.32, 78.19, 78.13, 75.09, 75.06, 71.6, 71.5,62.73, 62.65; HRESIMS *m/z* 575.1730 [M+Na]⁺ (cacld for C₂₆H₃₂O₁₃Na, 575.1736).

Resveratrol 3,5-di-O-β-D-glucoside (11): ¹H NMR (CD₃OD, 500 MHz): 7.40 (d, J = 9.0, 2H), 7.10 (d, J = 16.0, 1H), 6.977 (s, 1H), 6.973 (s, 1H), 6.92 (d, J = 16.0, 1H), 6.79 (d, J = 8.5, 2H), 6.78 (t, J = 2.0, 1H), 4.97 (d, J = 7.0, 2H), 3.97 (dd, J = 12.0, 2.0, 2H), 3.72 (dd, J = 12.0, 6.0, 2H), 3.60-3.36 (m, 8H); ¹³C NMR (CD₃OD, 125 MHz): 160.3(2C), 158.7, 141.6, 130.7, 130.4, 129.2(2C), 126.4, 116.6(2C), 109.9, 107.3, 105.0, 102.4(2C), 78.4 (2C), 78.2(2C), 75.1(2C), 71.6(2C), 62.9(2C); HRESIMS *m*/*z* 575.1729 [M+Na]⁺ (cacld forC₂₆H₃₂O₁₃Na, 575.1736).

10-hydroxycamptothecin 10-*O*-β**-D-glucoside (15**): ¹H NMR (DMSO, 500 MHz): 8.55 (s, 1H), 8.11 (d, J = 9.5 Hz, 1H), 7.63 (s, 1H), 7.59 (s, 1H), 7.30 (d, J = 9.5 Hz, 1H), 6.54 (s, 1H), 5.42 (m, 2H), 5.28 (m, 2H), 5.12 (d, J = 6.0 Hz, 1H), 4.64 (m, 1H), 3.77-3.16 (m, 6H), 1.88 (dd, J = 14.0, 7.0 Hz, 1H), 1.83 (dd, J = 14.0, 8.0 Hz, 1H), 0.87 (dd, J = 8.0, 7.0 Hz, 3H); ¹³C NMR (DMSO, 125 MHz): 172.4, 156.7, 155.9, 150.6, 149.9, 145.6, 144.2, 130.33, 130.30, 130.1, 129.0, 123.0, 118.5, 110.4, 100.3, 96.2, 77.1, 76.6, 73.2, 72.4, 69.6, 65.2, 60.6, 50.3, 30.3, 7.8; HRESIMS *m*/*z* 549.1484 [M+Na]⁺ (cacld for C₂₆H₂₆N₂O₁₀Na, 549.1480).



Figure S1. Key ¹H and ¹³C NMR data and HMBC correlations of 13, 15, 17, 18, 19 and 20.



Figure S2. Plots of the MD-simulated internuclear distances and RMSD for atomic positions of the ligand *versus* the simulation time for OleD binding with compounds **3** (panel a) and **12** (panel b). Top: Trace D1 represents the internuclear distance between the oxygen of the hydroxyl group at position C4' and the NE2 atom of the His19 side chain. Trace D2 represents the internuclear distance between the oxygen atom of Asp179. Bottom: Trace D1 represents the internuclear distance between the oxygen of the hydroxyl group at position C7 and the side chain of Asp179. Bottom: Trace D1 represents the internuclear distance between the oxygen of the hydroxyl group at position C3' and the NE2 atom of His19 side chain. Trace D2 represents the internuclear distance between the oxygen of the hydroxyl group at position C3' and the NE2 atom of His19 side chain. Trace D2 represents the internuclear distance between the oxygen of the hydroxyl group at position C3' and the NE2 atom of His19 side chain. Trace D2 represents the internuclear distance between the oxygen of the hydroxyl group at position C3' and the NE2 atom of His19 side chain. Trace D2 represents the internuclear distance between the oxygen atom of the carbonyl group and the hydroxyl group of Tyr114 side chain.



Figure S3. Ribbon view of the binding modes of compounds 3 (a, b), 4 (c), 5 (d) and 12 (e, f) with OleD.



Figure S4. Ribbon view of the two binding modes of compound 6 with OleD.



Figure S5. Ribbon view of the binding modes of compounds 7 (a, b), 8 (c), 9 (d, e), and 10 (f) with OleD.



Figure S6. Ribbon view of the binding modes of compounds 14 (a) and 15 (b) with OleD.



Figure S7. Ribbon view of the binding modes of compounds 16 (a, b), 17 (c), and 18 (d) with OleD.

























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