

Supplementary Information for

Switching glycosyltransferase UGT_{BL1} regioselectivity toward polydatin synthesis using semi-rational design

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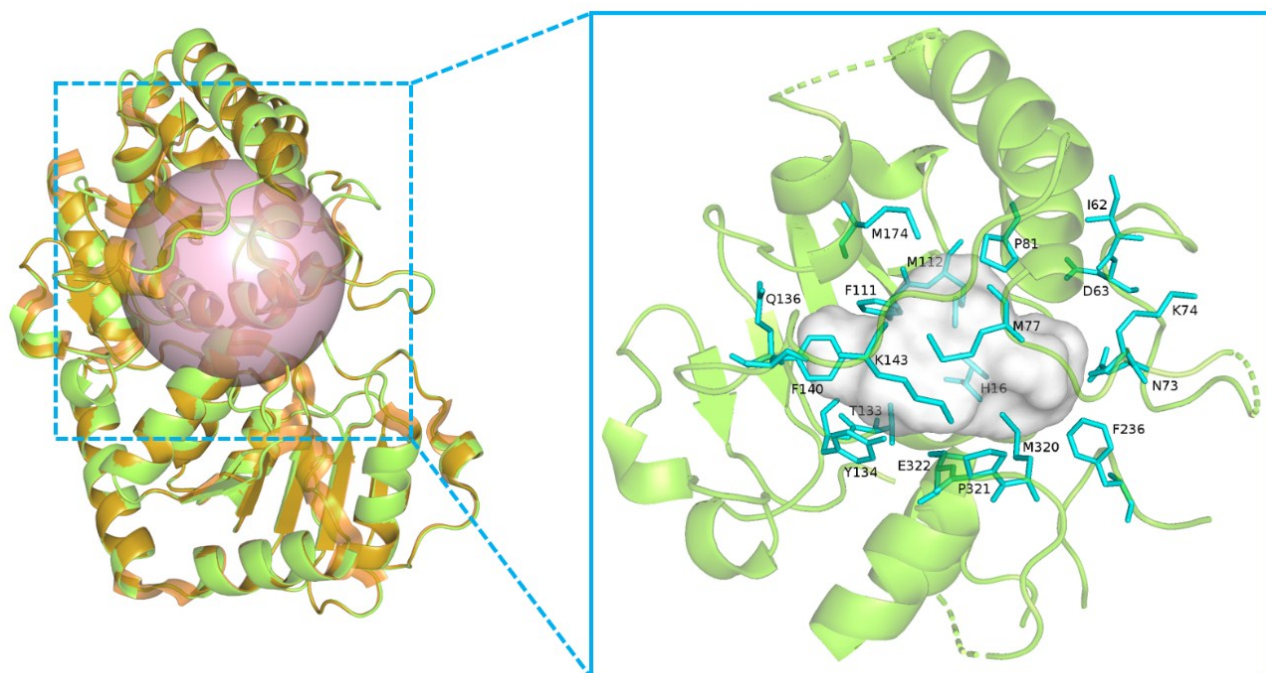


Fig. S1 The superimposition of homology model UGT_{BL1} with template OleI (2IYA), and the residues targeted for mutagenesis.

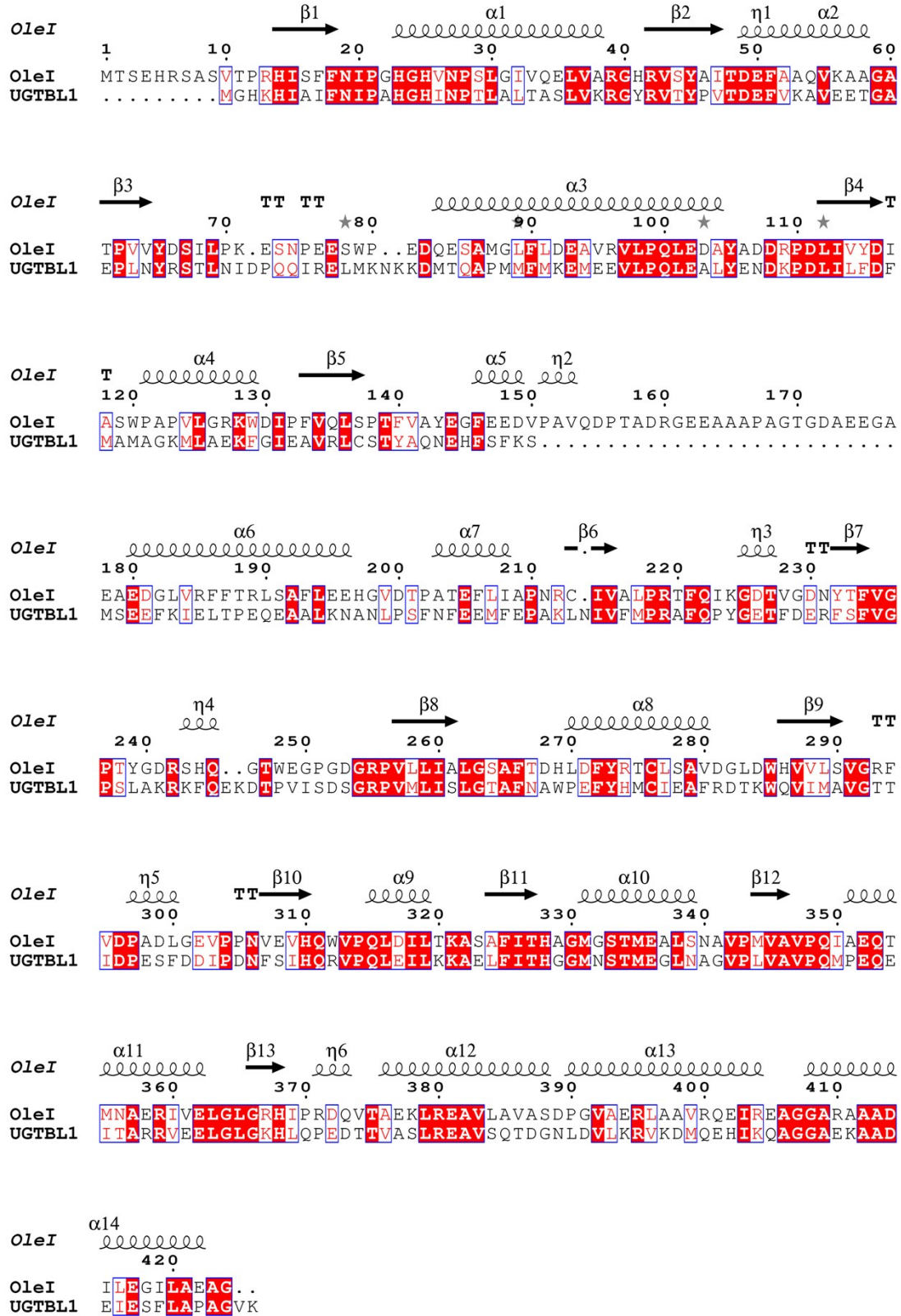


Fig. S2 The alignment of OleI and UGT_{BL}1. Secondary structure of OleI is shown above the OleI sequence.

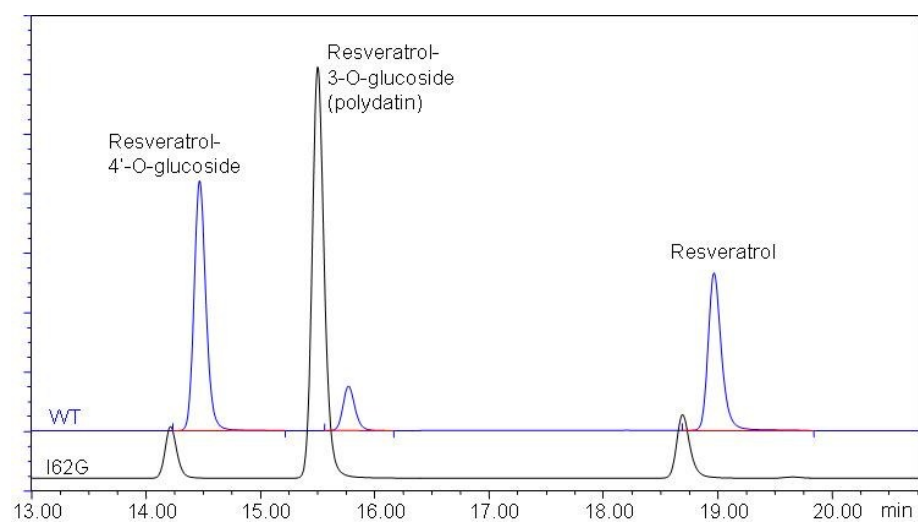


Fig. S3 The HPLC chromatogram of glycosylation of resveratrol catalyzed by UGT_{BL1} WT (blue, up) and UGT_{BL1} I62G (black, down).

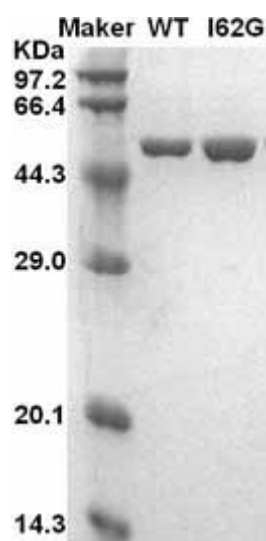


Fig. S4 SDS-PAGE gel of purified UGT_{BL1} WT and I62G.

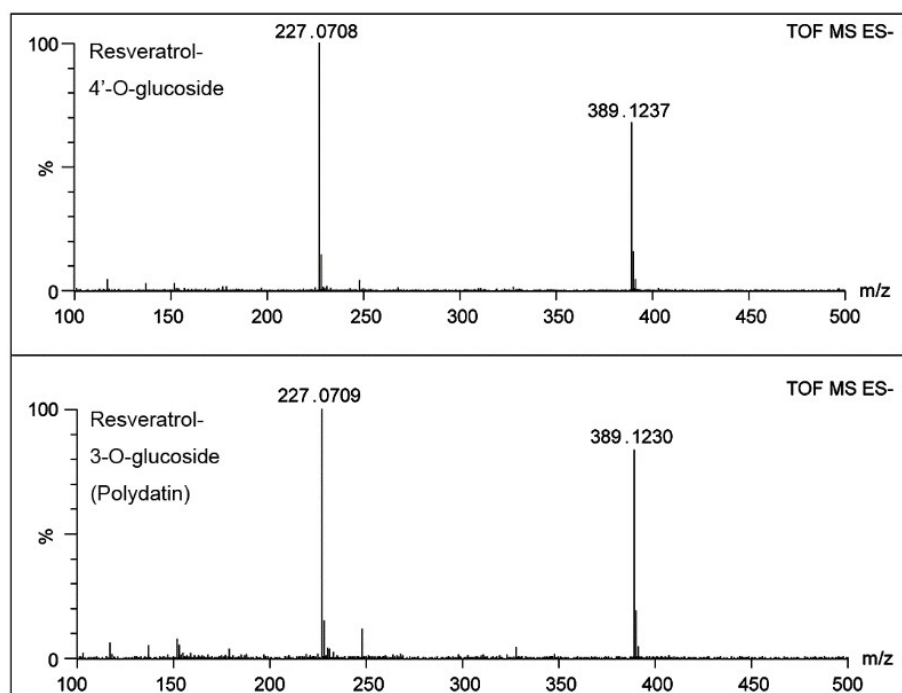


Fig. S5 The HRMS spectrums of resveratrol glucosides. A: product with shorter retention time (speculated to be 4'-O- β -glucoside of resveratrol); B: product with longer retention time (speculated to be 3-O- β -glucoside of resveratrol, polydatin)

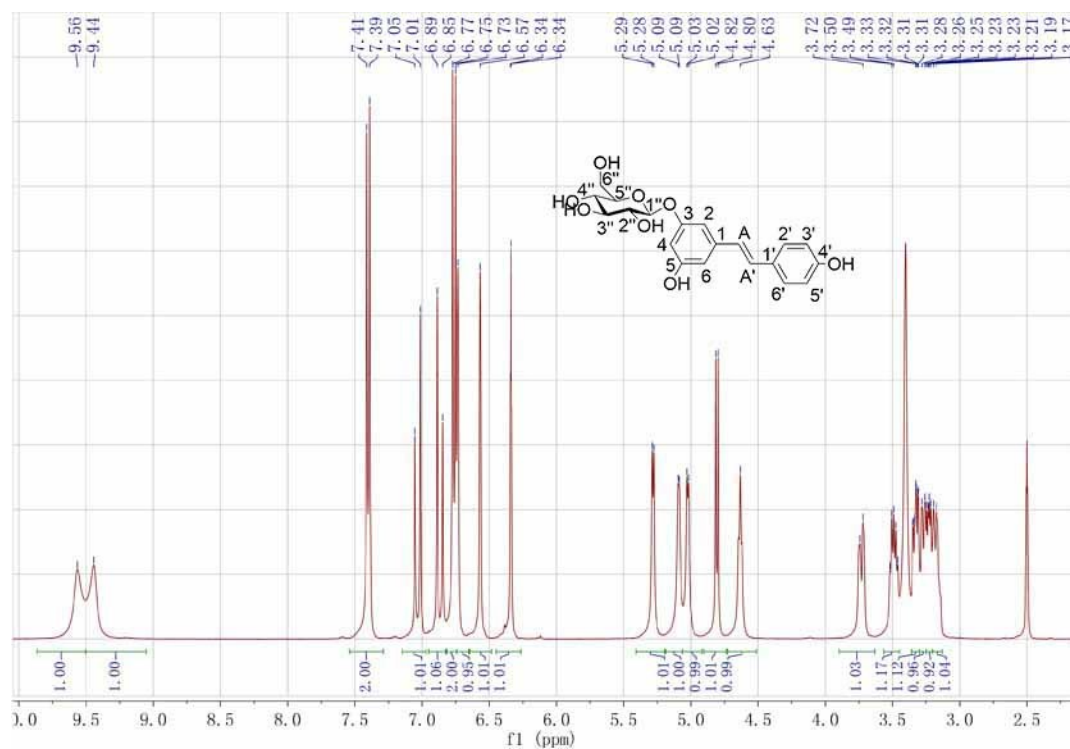


Fig. S6 ^1H NMR of resveratrol 3-O-glucoside (polydatin)

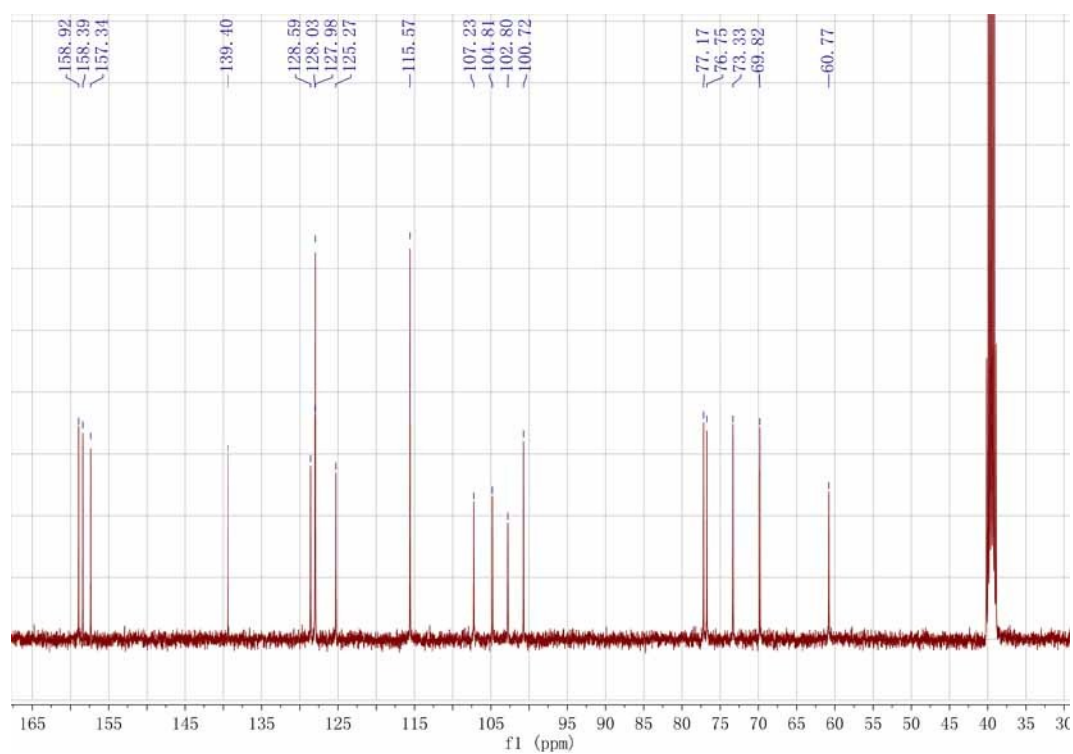


Fig. S7 ^{13}C NMR of resveratrol 3-O-glucoside (polydatin)

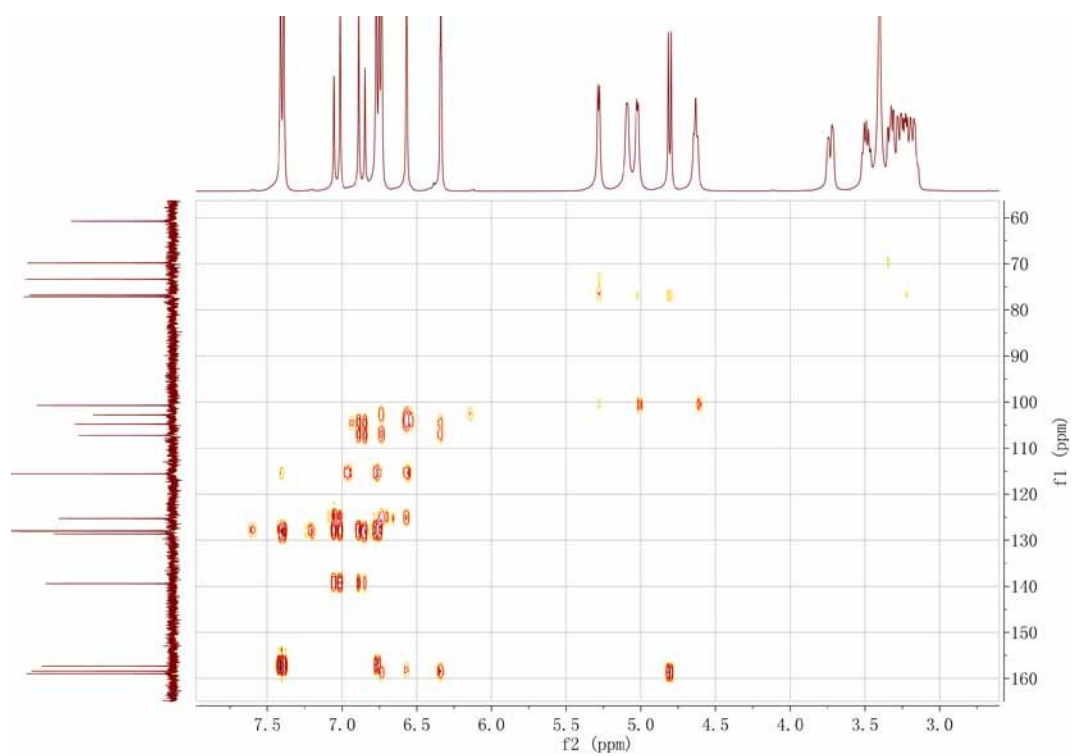


Fig. S8 HMBC NMR of resveratrol 3-O-glucoside (polydatin)

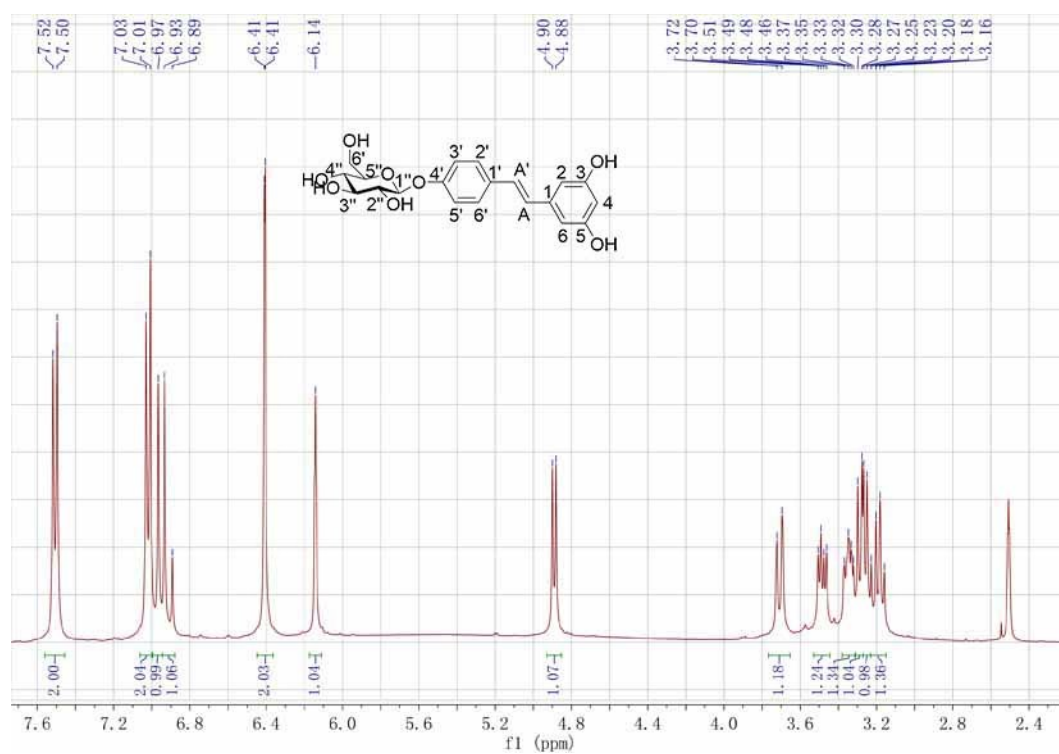


Fig. S9 ¹H NMR of resveratrol 4'-O-glucoside

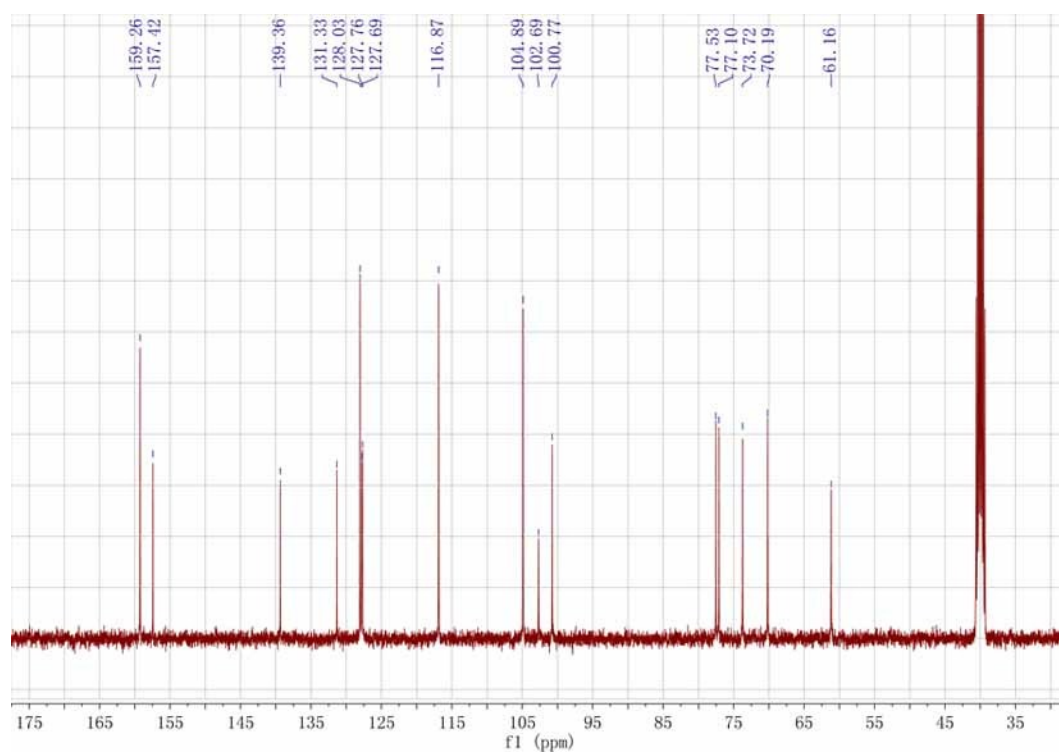


Fig. S10 ¹³C NMR of resveratrol 4'-O-glucoside

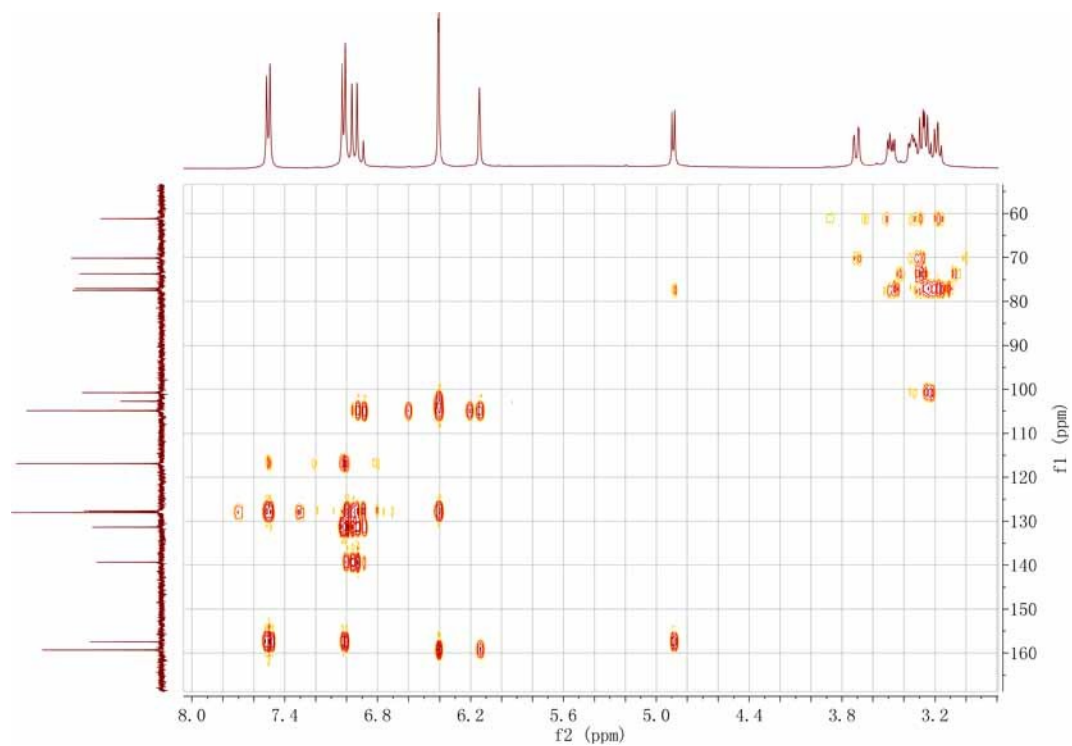


Fig. S11 HMBC NMR of resveratrol 4'-O-glucoside

Table S1: Primers used in mutations

Primer	Sequence (5' to 3')
H16A-F	atattcccgtcacggggcgattaatccgacgttgc
H16A-R	gcaagcgtcggattaatcgccccgtgagcgggaatat
I62A-F	gctcaactaccgctcaactttaaatgccgatccgcagcaa
I62A-R	ttgctgcggatcggcatttaaagttgagcggtagttgagc
N63A-F	ccgctcaactttaaatatcgctccgcagcaaatcgg
N63A-R	ccgaatttgctgcggagcgatatttaaagttgagcgg
N73A-F	gcaaattcgggagctgatgaaagctaaaaaggatatgacacaggct
N73A-R	agcctgtgtcatatccttttagctttcatcagctcccgaattgc
K74A-F	gcaaattcgggagctgatgaaaaatgcaaaggatatgacacaggct
K74A-R	agcctgtgtcatatcctttgcatttttcatcagctcccgaattgc
M77A-F	gggagctgatgaaaaataaaaaggatcgacacaggctccgatg
M77A-R	catcggagcctgtgtcgcacatccttttttttcatcagctccc
P81A-F	aataaaaaggatatgacacaggctgcgatgatgttatgaaagaaa
P81A-R	tttctttcataaacatcatcgcagcctgtgtcatatccttttatt
F111A-F	ctgacctcatcctttttgacgctatggccatggcgggaa
F111A-R	ttcccgccatggccatagcgtcaaaaaggatgaggtcag
M112A-F	ctgacctcatcctttttgactttgcggccatggcggg
M112A-R	cccgcctatggccgcaaaagtcaaaaaggatgaggtcag
T133A-F	cggttcgcctttgttctgcatatgcacagaacgaa
T133A-R	ttcgttctgtgcatatgcagaacaaaggcgaaccg
Y134A-F	agaggcgggtcgcctttgttctacagctgcacagaacgaa
Y134A-R	ttcgttctgtgcagctgtagaacaaggcgaaccgcctct
Q136A-F	gttcgcctttgttctacatatgcagcgaacgaacatttttattcaaatc
Q136A-R	gatttgatgaaaaatgttcgttcgctgcatatgtagaacaaggcgaac
F140A-F	tgttctacatatgcacagaacgaacatgcttcattcaaatcaatgtctgaag
F140A-R	cttcagacattgatttgatgaagcatgttcgttctgtgcatatgtagaaca
K143A-F	tgcacagaacgaacatttttattcgcacatgaatgtctgaagatttaagatc
K143A-R	gatcttaactcttcagacattgatgcgaatgaaaaatgttcgttctgtgca
M174A-F	cttccgtcatttaattttgaagaggcgttcgaaccggcaaaattgaaca
M174A-R	tgttcaattttgccggttcgaacgcctcttcaaaattaaatgacggaag
F236A-F	ccggccaggcattggccgccgtccctaaag
F236A-R	ctttagggacggcggccaatgcctggccgg
M320A-R	gatttcctgttcaggcgttcggggacggcaacaagc
M320A-F	gcttgttgccgtcccgaagcgcctgaacaggaaatc
P321A-R	tgatttcctgttcagccattttgcgggacggc
P321A-F	gccgtcccgcaaatggctgaacaggaaatca
E322A-R	ggcagtgatttcctgtgcaggcattttgcgggac
E322A-F	gtcccgcaaatgcctgcacaggaaatcactgcc
I62F-F	ccgctcaactaccgctcaactttaaatttgatccgcagcaaat
I62F-R	aatttgctgcggatcaaaatttaaagttgagcggtagttgagcgg

I62L-F	gctcaactaccgctcaactttaaatctggatccgcagcaaat
I62L-R	atttgctgcggatccagatttaaagttgagcggtagttgagc
I62V-F	gctcaactaccgctcaactttaaatgtggatccgcagcaaat
I62V-R	atttgctgcggatccacatttaaagttgagcggtagttgagc
I62M-F	tcaactaccgctcaactttaaatatggatccgcagcaaat
I62M-R	atttgctgcggatccatatttaaagttgagcggtagttga
I62P-F	ccgctcaactaccgctcaactttaaatccggatccgcagcaaat
I62P-R	aatttgctgcggatccggatttaaagttgagcggtagttgagcgg
I62T-F	caactaccgctcaactttaataaccgatccgcagca
I62T-R	tgctgcggatcggattttaagttgagcggtagttg
I62Y-F	agccgctcaactaccgctcaactttaattatgatccgcagcaaatc
I62Y-R	gaatttgctgcggatcataatttaaagttgagcggtagttgagcggct
I62H-F	agccgctcaactaccgctcaactttaaatcatgatccgcagcaaatc
I62H-R	gaatttgctgcggatcatgatttaaagttgagcggtagttgagcggct
I62Q-F	ccgctcaactaccgctcaactttaaatcaggatccgcagcaaat
I62Q-R	aatttgctgcggatcctgatttaaagttgagcggtagttgagcgg
I62N-F	caactaccgctcaactttaataacgatccgcagca
I62N-R	tgctgcggatcgtattttaagttgagcggtagttg
I62K-F	cgctcaactaccgctcaactttaataaagatccgcagcaaat
I62K-R	aatttgctgcggatcttatttaaagttgagcggtagttgagcgg
I62D-F	agccgctcaactaccgctcaactttaaatgatgatccgcagcaaatc
I62D-R	gaatttgctgcggatcatcatttaaagttgagcggtagttgagcggct
I62E-F	agccgctcaactaccgctcaactttaaatgaagatccgcagcaaatc
I62E-R	gaatttgctgcggatcttcatttaaagttgagcggtagttgagcggct
I62C-F	gctcaactaccgctcaactttaaattgcgatccgcagcaa
I62C-R	ttgctgcggatcgcaatttaaagttgagcggtagttgagc
I62W-F	ccgctcaactaccgctcaactttaaattgggatccgcagcaaat
I62W-R	aatttgctgcggatcccaatttaaagttgagcggtagttgagcgg
I62R-F	gctcaactaccgctcaactttaaatcgcgatccgcagcaa
I62R-R	ttgctgcggatcgcgatttaaagttgagcggtagttgagc
I62S-F	caactaccgctcaactttaaatagcgatccgcagca
I62S-R	tgctgcggatcgctatttaaagttgagcggtagttg
I62G-F	gctcaactaccgctcaactttaaatggcgatccgcagcaa
I62G-R	ttgctgcggatcgccatttaaagttgagcggtagttgagc

Table S2: ^1H and ^{13}C NMR data of glucosides of resveratrol

Nr	Resveratrol-4'-O- β -glucoside		Nr	Resveratrol-3-O- β -glucoside (Polydatin)	
	δ_{C}	δ_{H}		δ_{C}	δ_{H}
1	139.36		1	139.40	
2/6	104.89	6.41 (d, $J = 1.8$ Hz, 2H)	2	107.23	6.73 (s, 1H)
3/5	159.26		3	158.39	
4	102.69	6.14 (s, 1H)	4	104.81	6.34 (s, 1H)
			5	158.92	
			6	102.80	6.57 (s, 1H)
A	127.69	6.91 (d, $J = 16.3$ Hz, 1H)	A	128.03	6.87 (d, $J = 16.3$ Hz, 1H)
A'	127.76	6.97 (d, $J = 16.3$ Hz, 1H)	A'	128.59	7.03 (d, $J = 16.3$ Hz, 1H)
1'	131.33		1'	125.27	
2'/6'	128.03	7.51 (d, $J = 8.7$ Hz, 1H)	2'/6'	127.98	7.40 (d, $J = 8.6$ Hz, 2H)
3'/5'	116.87	7.02 (d, $J = 8.8$ Hz, 1H)	3'/5'	115.57	6.76 (d, $J = 8.6$ Hz, 2H)
4'	157.42		4'	157.34	
1''	100.77	4.89 (d, $J = 7.4$ Hz, 1H)	1''	100.72	4.81 (d, $J = 7.5$ Hz, 1H)
2''	73.72	3.19 (m, 1H)	2''	73.33	3.18 (m, 1H)
3''	77.10	3.34 (m, 1H)	3''	76.75	3.33 (m, 1H)
4''	70.19	3.26 (m, 1H)	4''	69.82	3.23 (m, 1H)
5''	77.53	3.29 (m, 1H)	5''	77.17	3.27 (m, 1H)
6''	61.16	3.71 (m, 1H) , 3.48 (m, 1H)	6''	60.77	3.73 (m, 1H), 3.51 (m, 1H)