Sorbicillinoid Derivatives with the Radical Scavenging Activities from the Marine-Derived Fungus *Acremonium chrysogenum* C10

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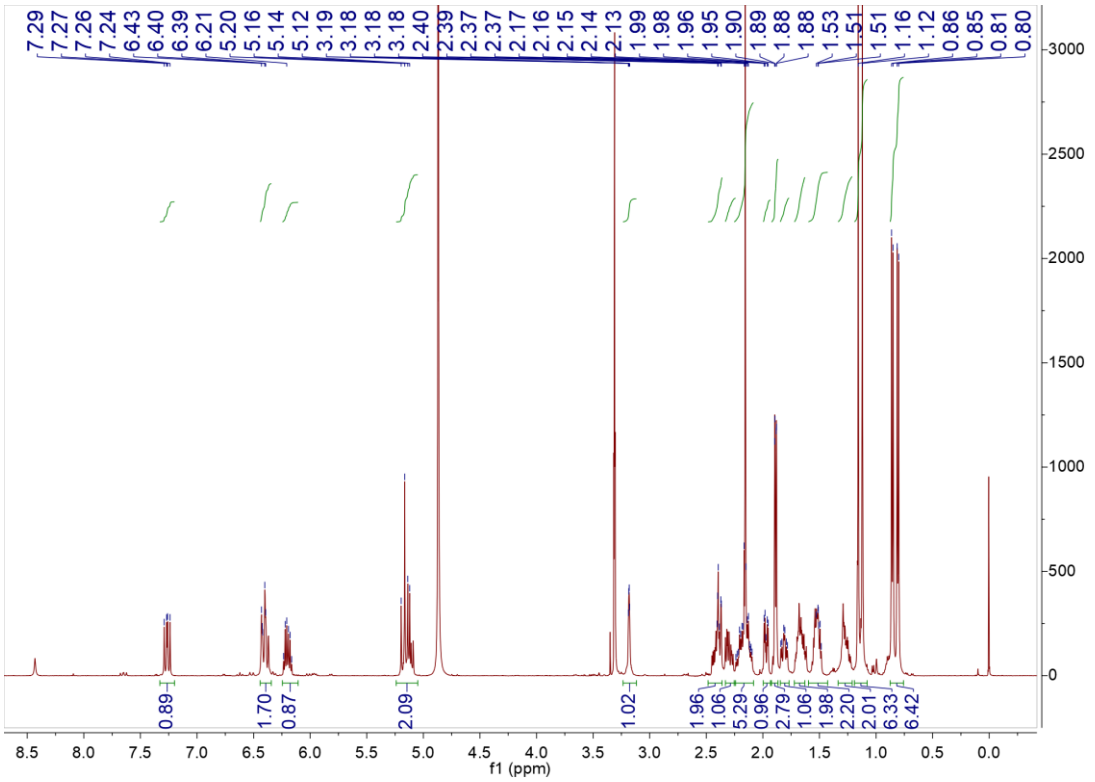


Figure S1. ^{1}H NMR spectrum of acresorbicillinol A (**1**; 500 MHz, CD_3OD)

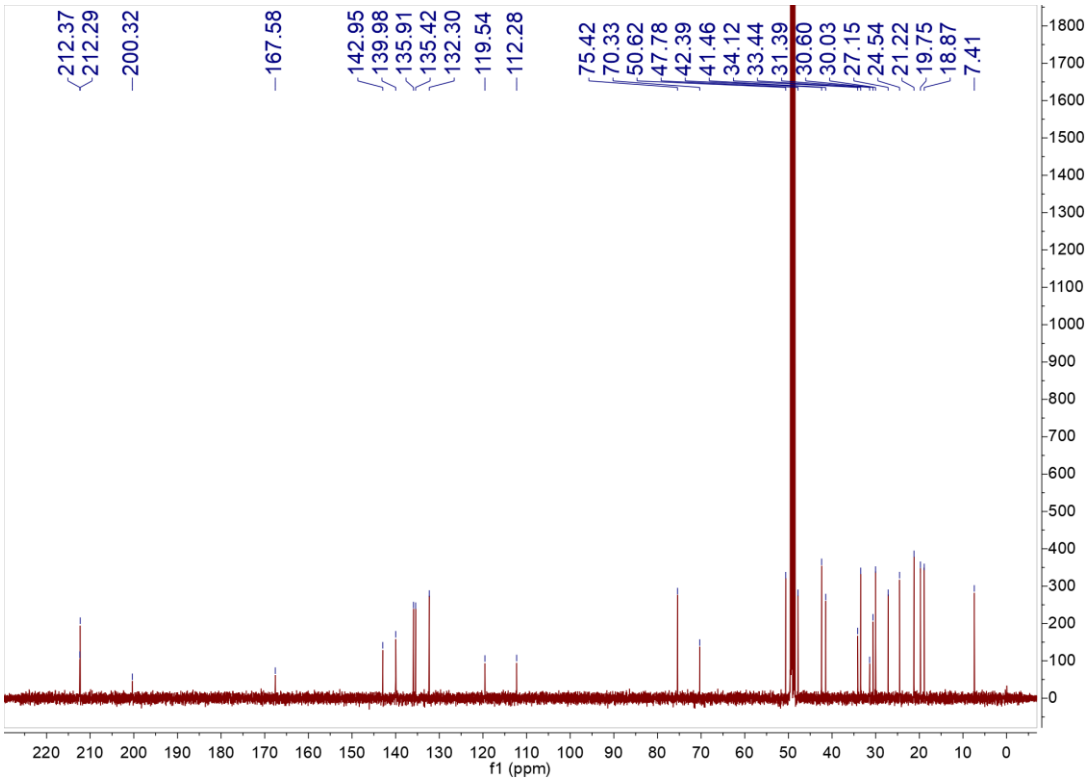


Figure S2. ^{13}C NMR spectrum of acresorbicillinol A (**1**; 125 MHz, CD_3OD)

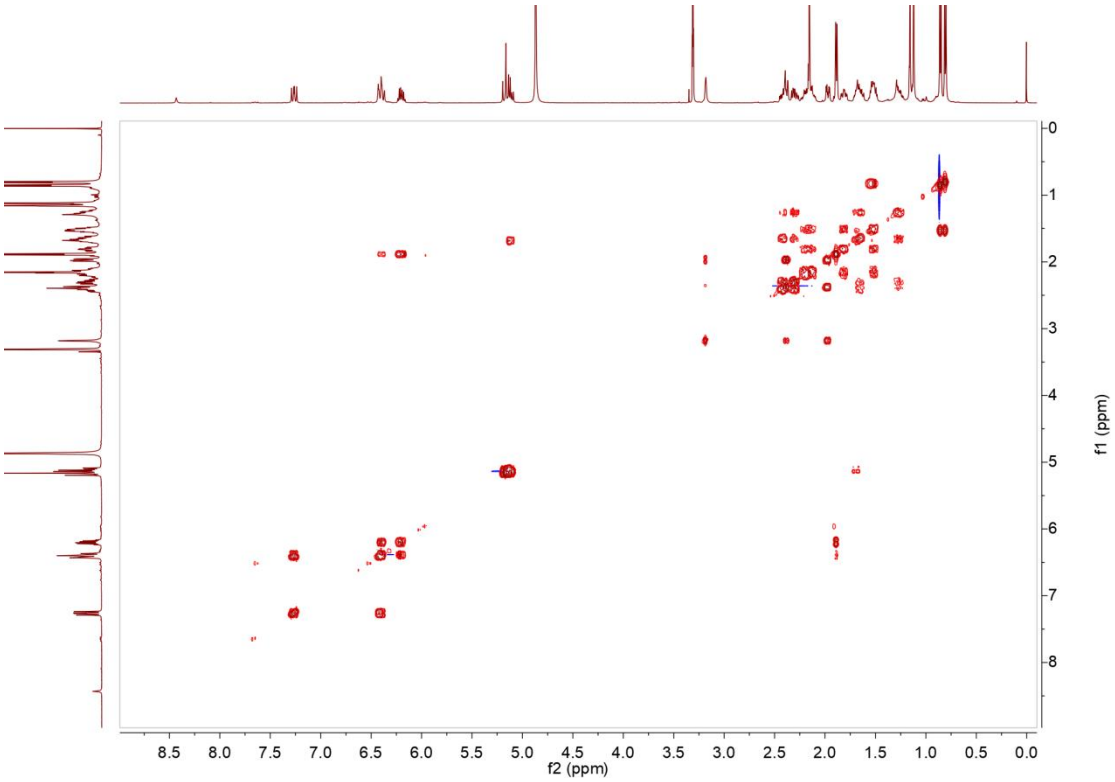


Figure S3. ^{1}H-^{1}H COSY spectrum of acresorbicillinol A (**1**, CD_3OD

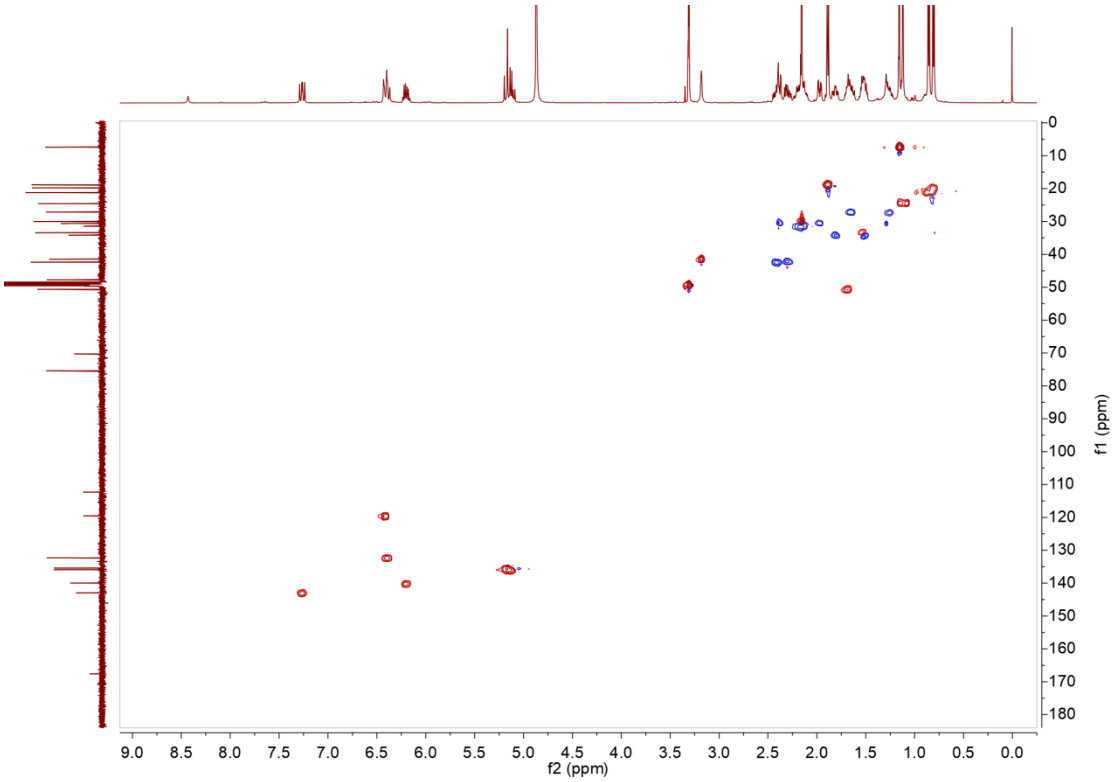


Figure S4. HSQC spectrum of acresorbicillinol A (**1**, CD_3OD)

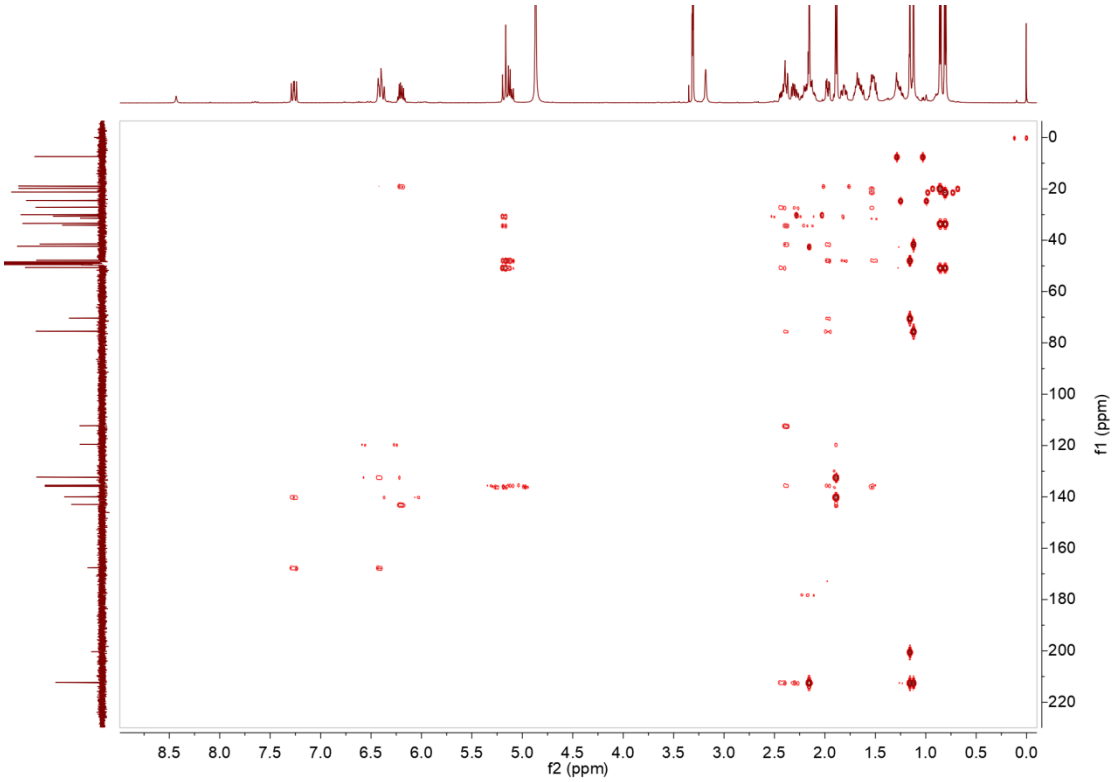


Figure S5. HMBC spectrum of acresorbicillinol A (**1**, CD_3OD)

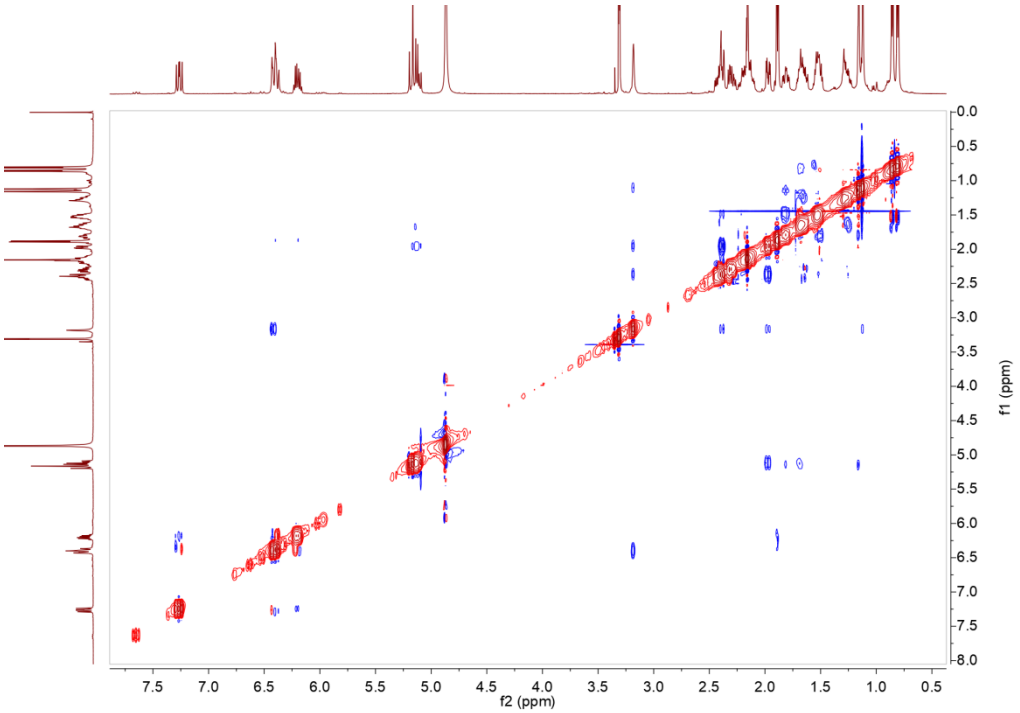


Figure S6. NOESY spectrum of acresorbicillinol A (**1**, CD_3OD)

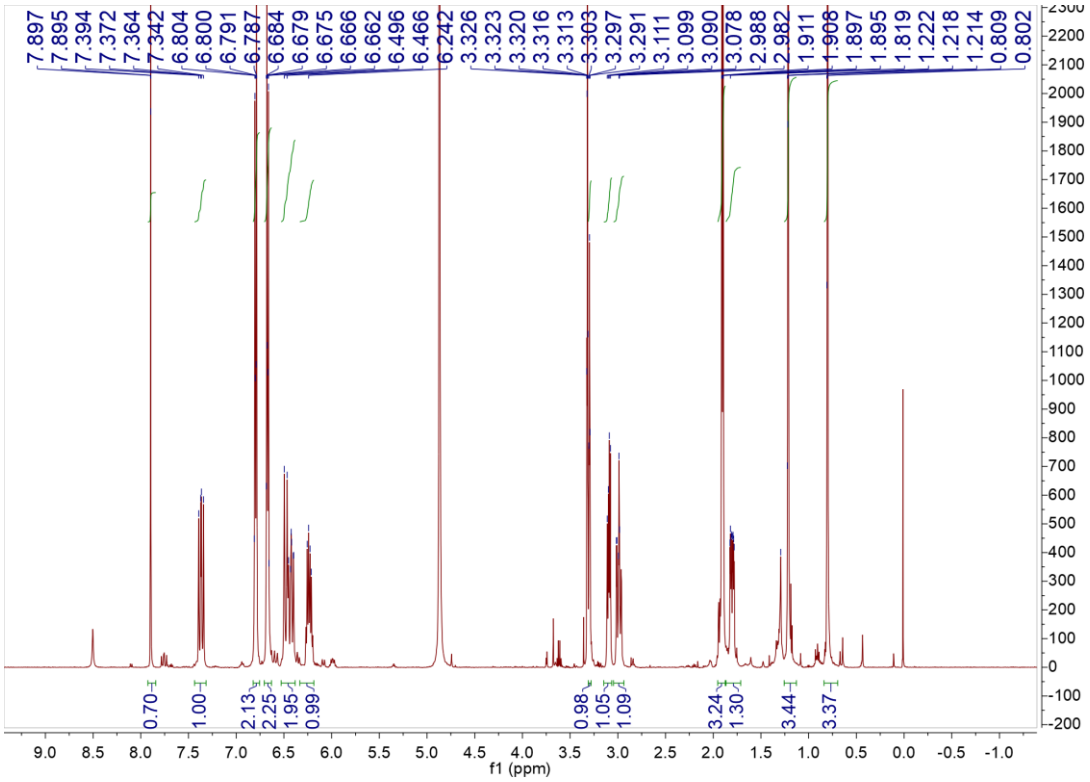


Figure S7. ^{1}H NMR spectrum of acresorbicillinol B (**2**; 500 MHz, CD_3OD)

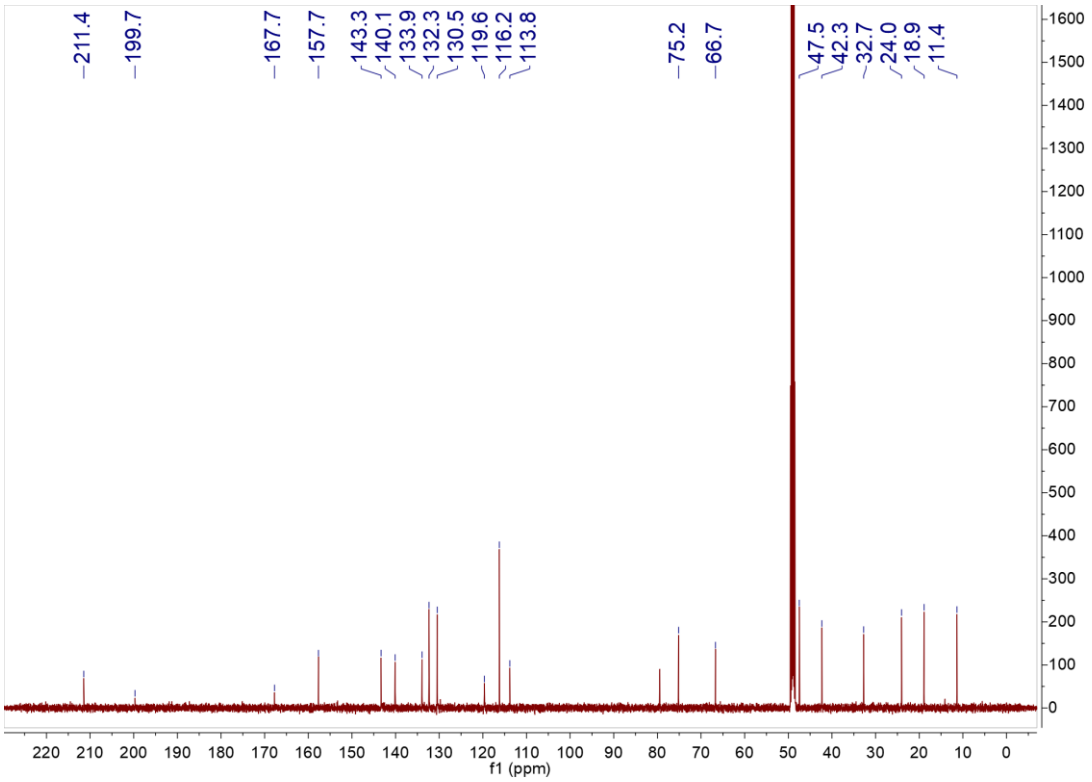


Figure S8. ^{13}C NMR spectrum of acresorbicillinol B (**2**; 125 MHz, CD_3OD)

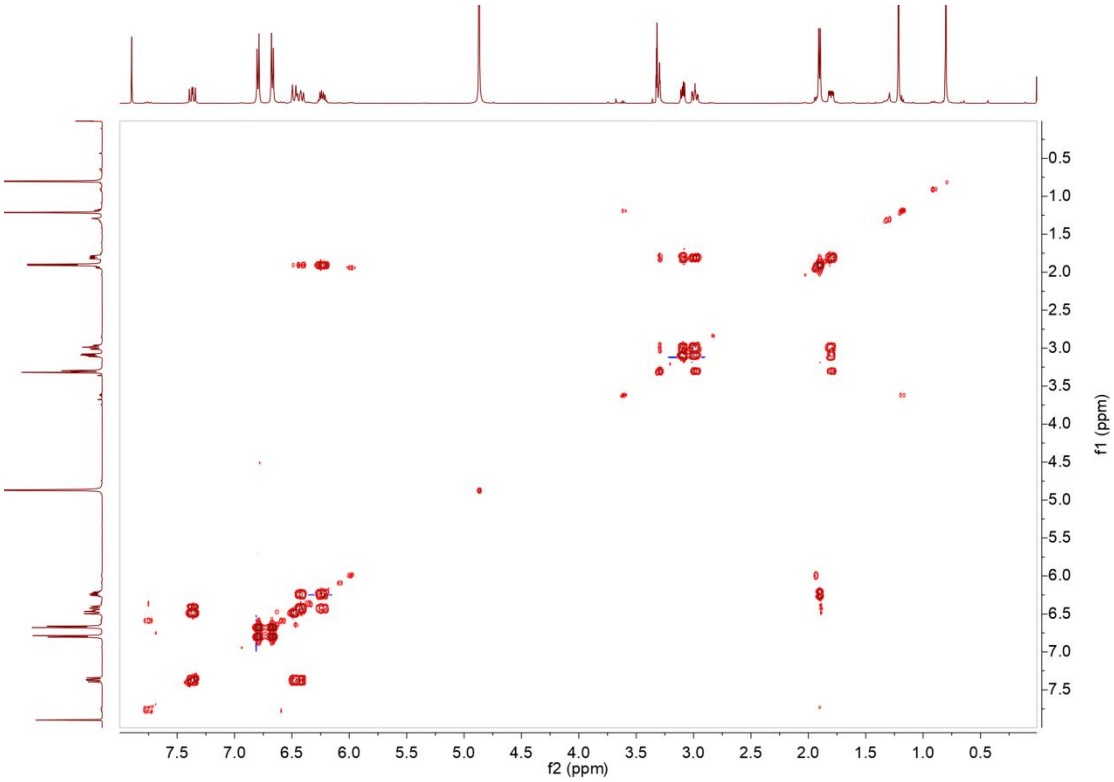


Figure S9. ^{1}H-^{1}H COSY spectrum of acresorbicillinol B (**2**, CD_3OD)

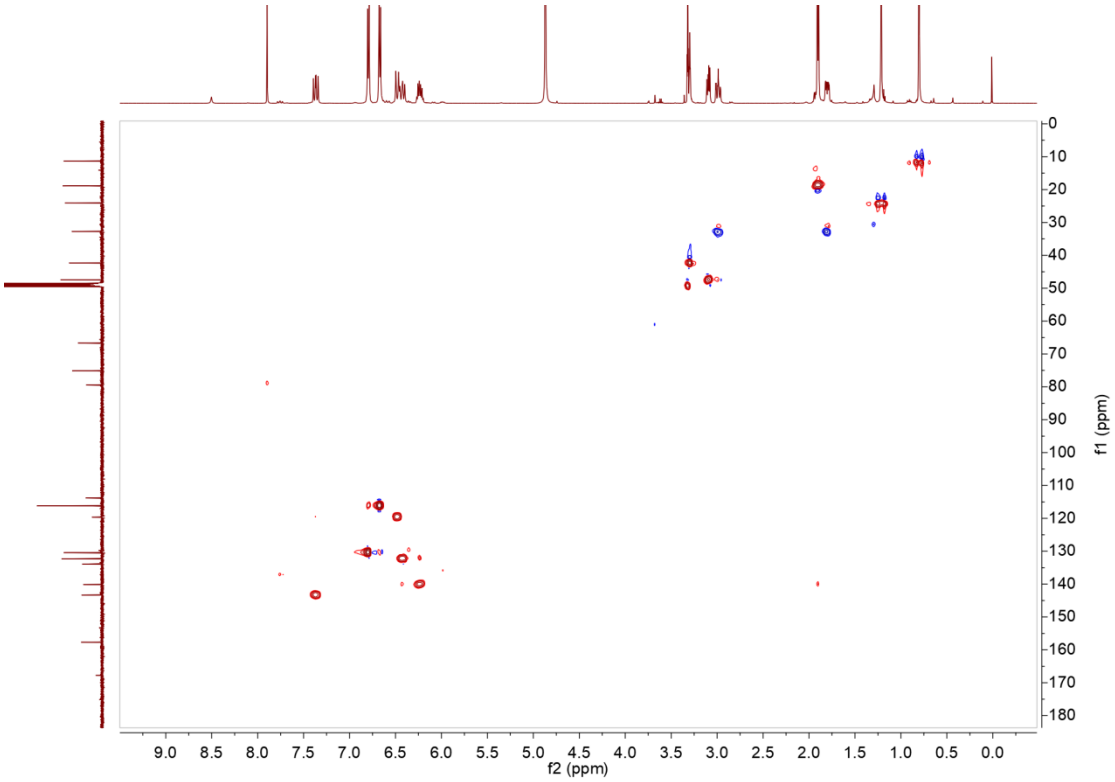


Figure S10. HSQC spectrum of acresorbicillinol B (**2**, CD_3OD)

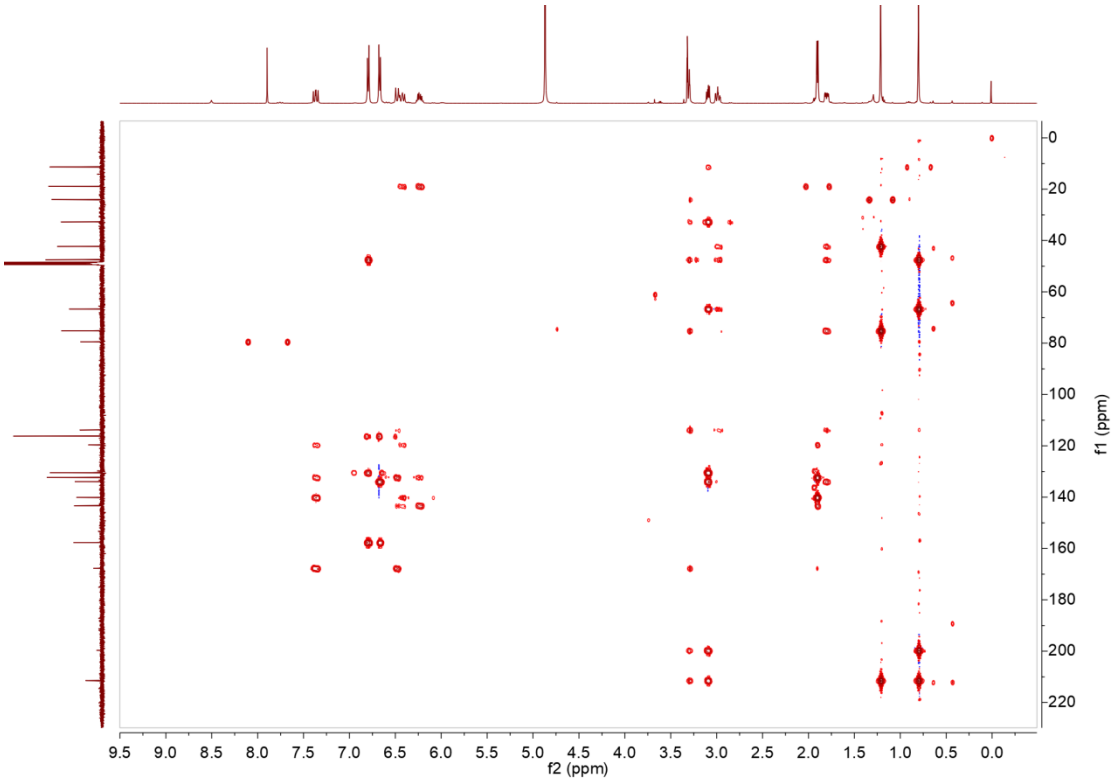


Figure S11. HMBC spectrum of acresorbicillinol B (**2**, CD_3OD)

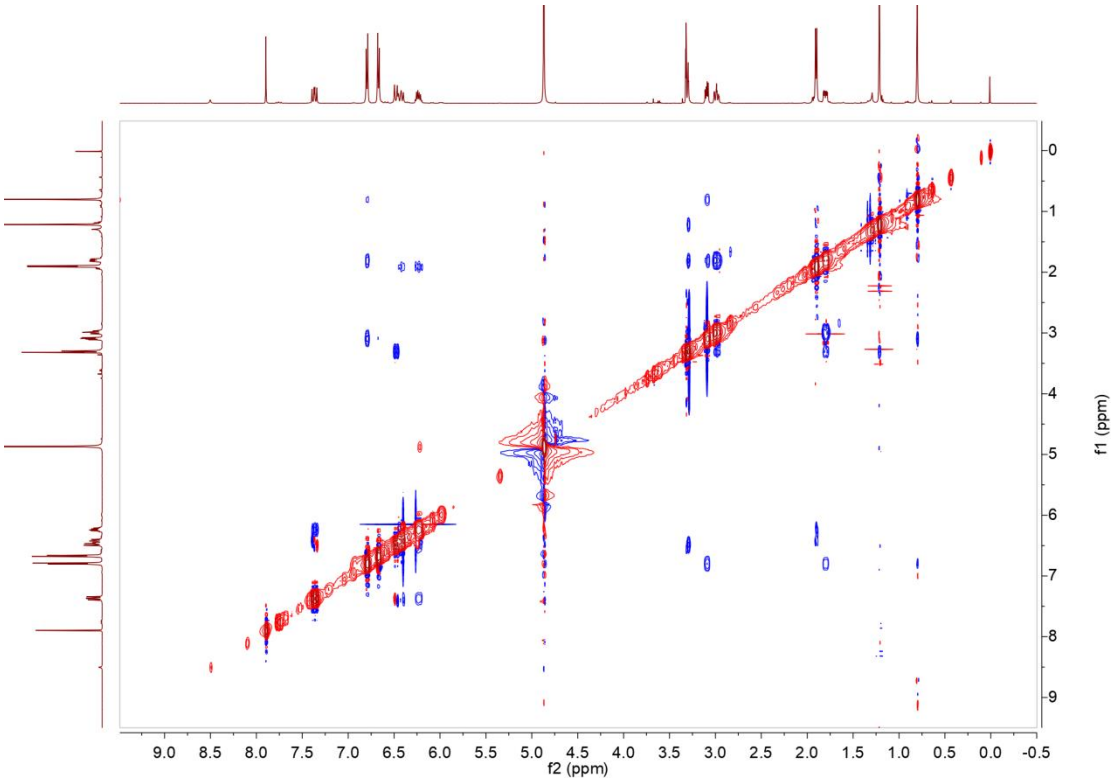


Figure S12. NOESY spectrum of acresorbicillinol B (**2**, CD_3OD)

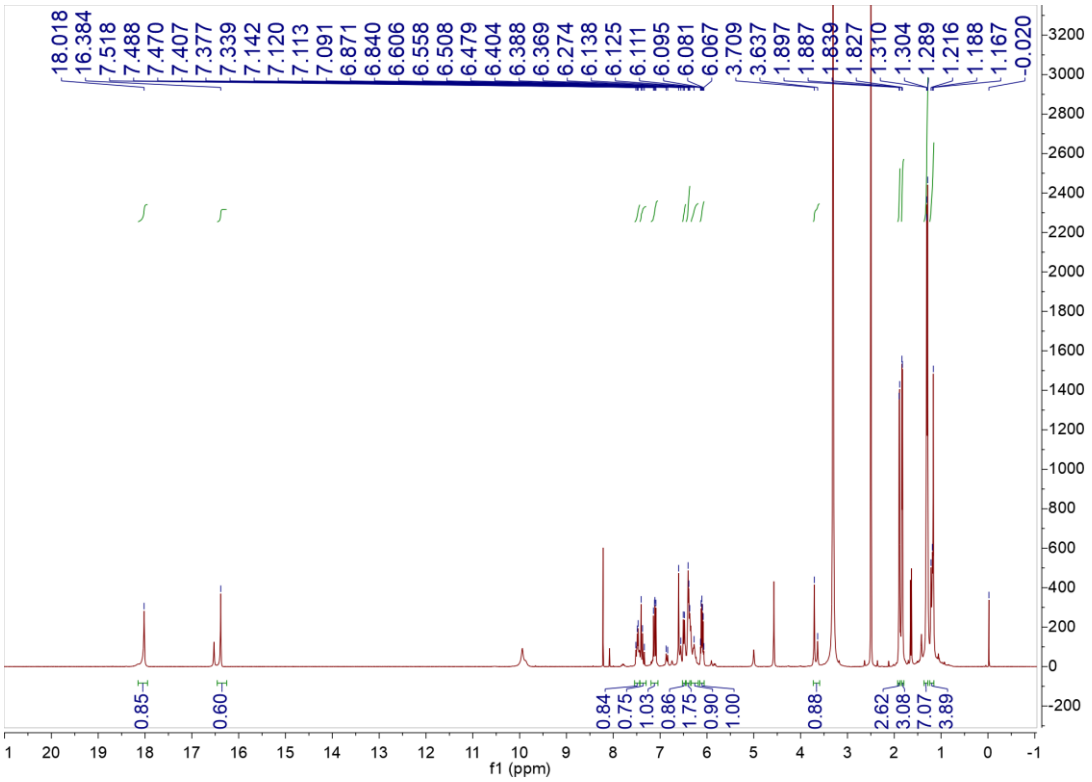


Figure S13. ^{1}H NMR spectrum of acresorbicillinol C (**3**; 500 MHz, DMSO:$CDCl_3$ = 3:1)

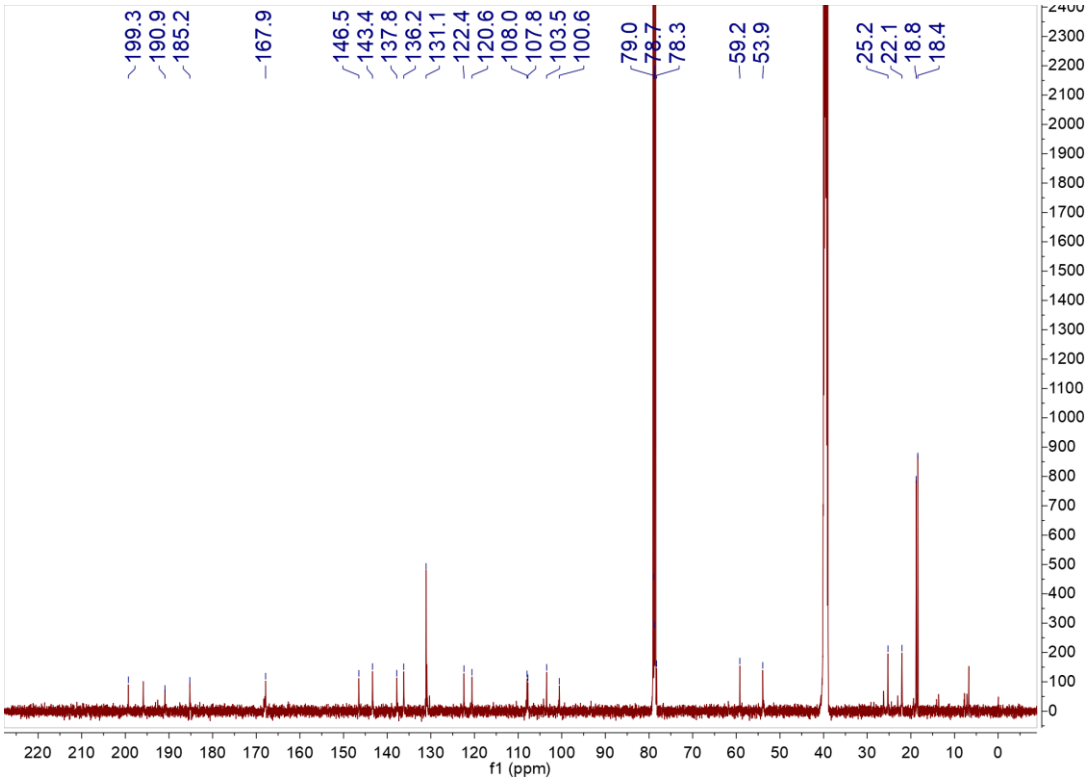


Figure S14. ^{13}C NMR spectrum of acresorbicillinol C (**3**; 125 MHz, DMSO:$CDCl_3$ = 3:1)

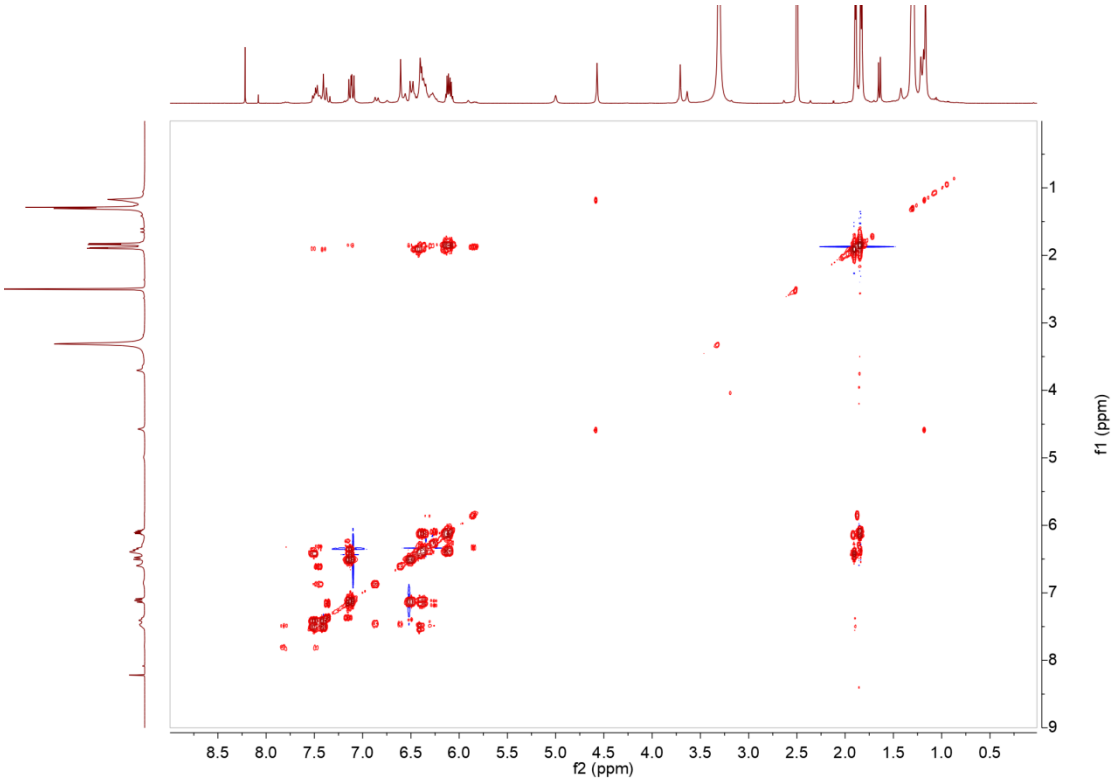


Figure S15. ^{1}H-^{1}H COSY spectrum of acresorbicillinol C (**3**, DMSO:$CDCl_3$ = 3:1)

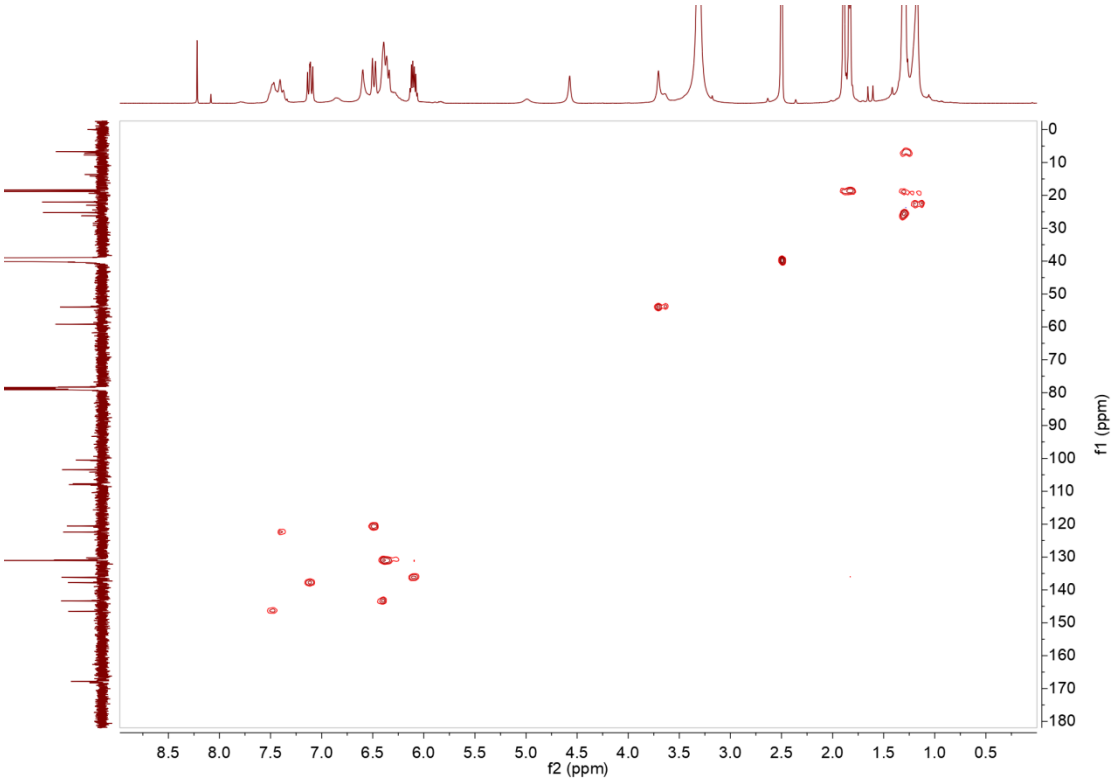


Figure S16. HSQC spectrum of acresorbicillinol C (**3**, DMSO:$CDCl_3$ = 3:1)

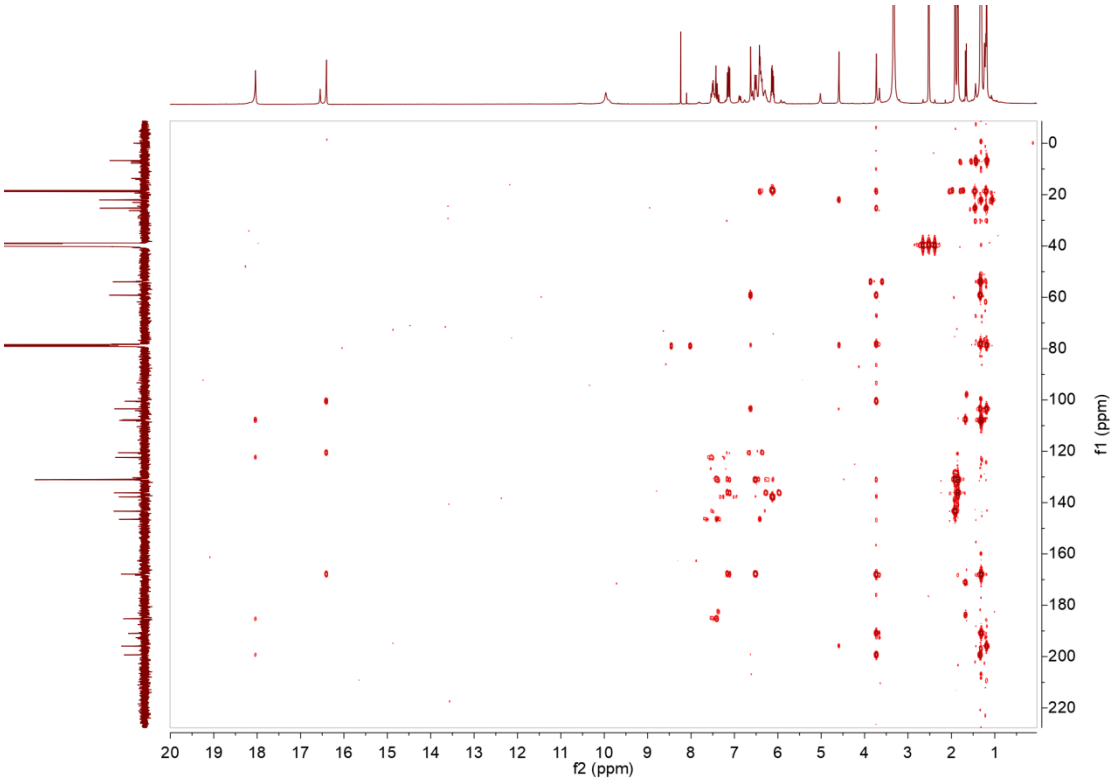


Figure S17. HMBC spectrum of acresorbicillinol C (**3**, DMSO:$CDCl_3$ = 3:1)

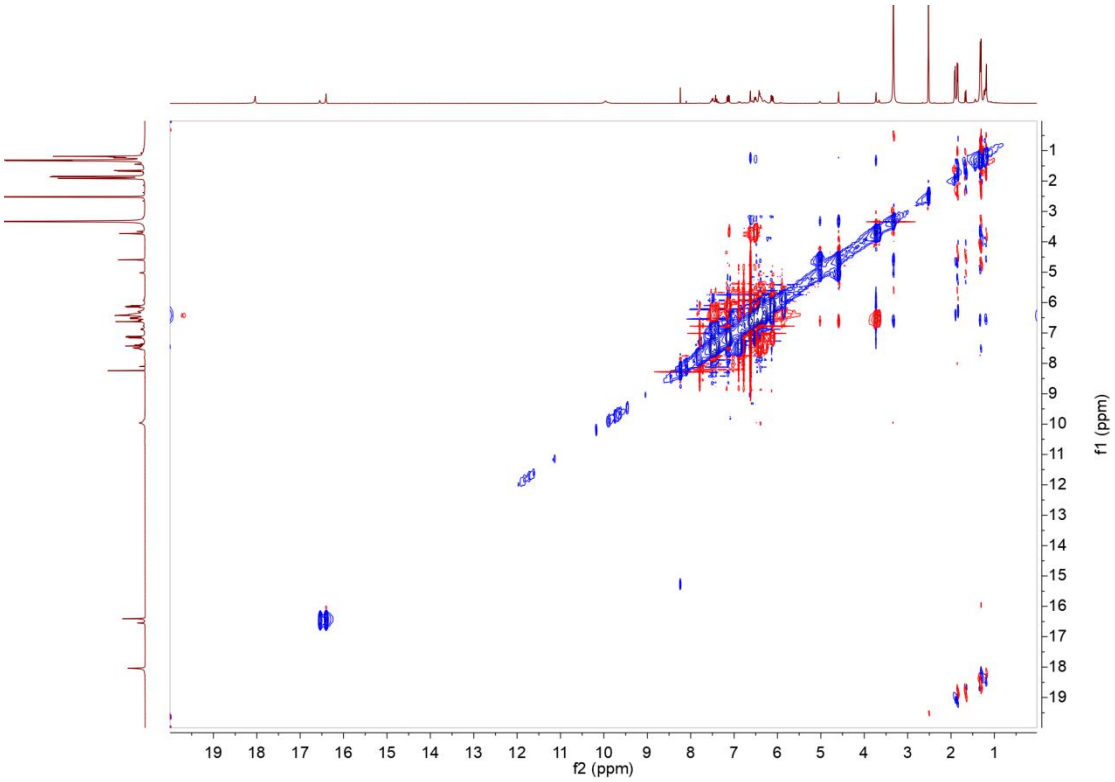
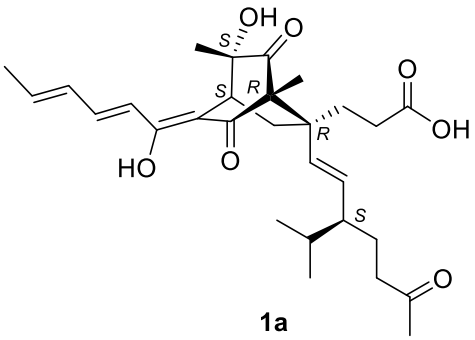
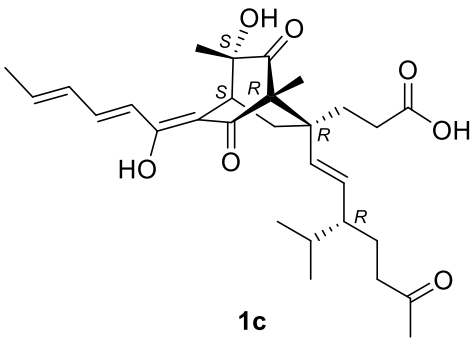


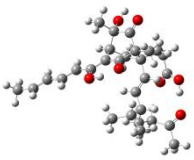
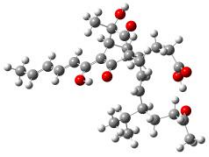
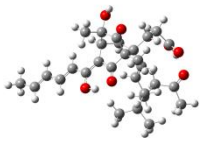
Figure S18. NOESY spectrum of acresorbicillinol C (**3**, DMSO:$CDCl_3$ = 3:1)

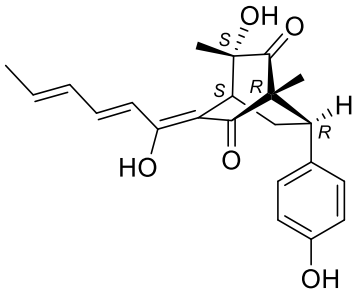


1a

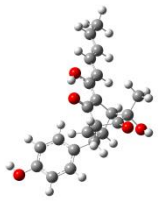
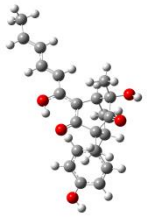


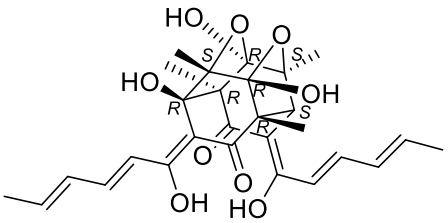
1c

Conformers	Populations	Conformers	Populations
	25.49%		68.26%
	53.89%		31.74%
	20.62%		

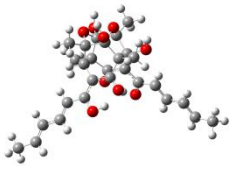
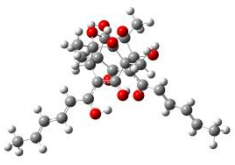


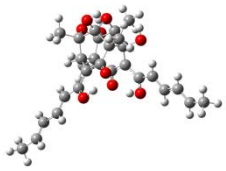
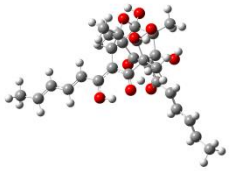
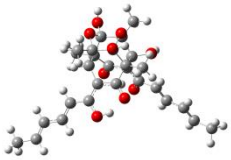
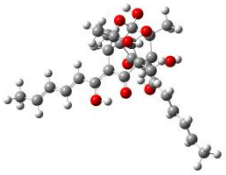
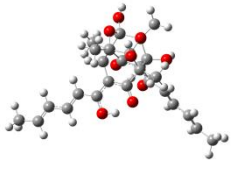
2a

Conformers	Populations
	51.44%
	48.56%



3a

Conformers	Populations
	1.97%
	8.24%

	3.15%
	12.10%
	15.53%
	53.82%
	1.01%

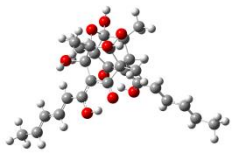
	4.17%

Figure S19. ECD conformers of acresorbicillinols A–C (**1**–**3**)

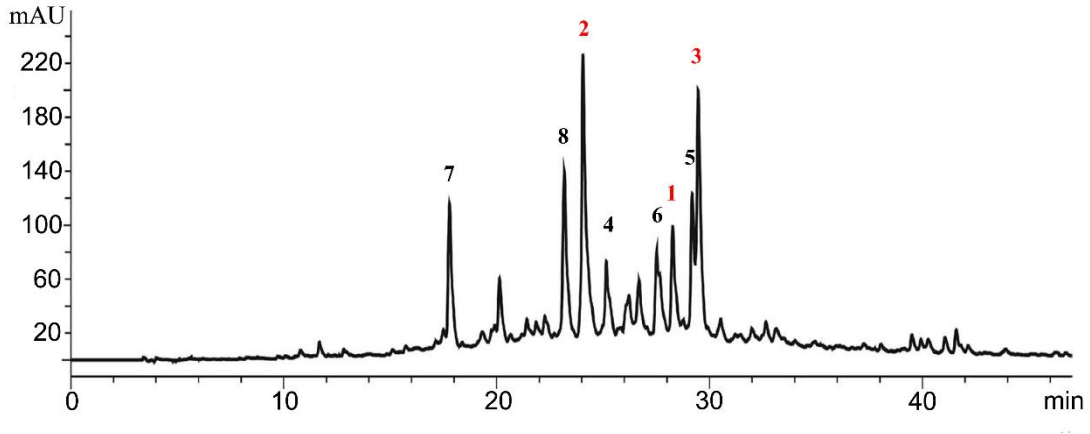


Figure S20. HPLC profiles of the extracts from the rice solid medium of *A. chrysogenum* after 7 days fermentation

Table S1. Primers used in this study

Primer	Sequence
actin-F(RT)	AGTCCAAGCGTGGTATCC
actin-R(RT)	TAGAAGGCAGGGGCGTTG
AcsorA-F(RT)	CGCACTTCTTTGACCTCCG
AcsorA-R(RT)	TTGGCACCGCCCACGAT
AcsorB-F(RT)	GCCATCCTGGGACCACT
AcsorB-R(RT)	TGAACACGCCGAGACAAC
AcsorC-F(RT)	TCGGGGAAGGGTACAAGGC
AcsorC-R(RT)	CTCGACGGACCGCAAGTT
AcsorD-F(RT)	TGATGGCAATTCCGTCTGG
AcsorD-R(RT)	GACTCGTTGAGGGCGTTCC
AcsorE-F(RT)	GGGCTCGTTCCGTCTCGTCT
AcsorE-R(RT)	CCTCGGGCTGGGATTTGAT
AcsorT-F(RT)	GTATTCTCGGCGTCGTCTTC
AcsorT-R(RT)	TTCTTCTTGCCCTGGTTCTT
AcsorR1-F(RT)	CGATGTGGACGACTTGAGG
AcsorR1-R(RT)	TCATACTGCGAGTCCTTGTTAG
AcsorR2-F(RT)	GCTGACCAGTACCGCATCG
AcsorR2-R(RT)	GCCCGCCCTAACGTCTAT
orf1-F(RT)	CGGGATTGTTTCCAAGTCG
orf1-R(RT)	TCCACCAGGGTTGTGCC
orf2-F(RT)	CGGAGATACAGGAAGGC
orf2-R(RT)	CCGCTAAGGCTAGGGTC