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Supporting Information for

Dicarabrol A, dicarabrone C and dipulchellin A, unique sesquiterpene lactone dimers from Carpesium abrotanoides †

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Experimental Section

X-ray crystallographic analyses.

Table S1. X-ray crystallographic data for dipulchellin A (3)

Figure S1. ¹H NMR spectrum of dicarabrol A (1) in CDCl₃

Figure S2. ¹³C NMR spectrum of dicarabrol A (1) in CDCl₃

Figure S3. ¹H-¹H COSY spectrum of dicarabrol A (1) in CDCl₃

Figure S4. HSQC spectrum of dicarbrol A (1) in CDCl₃

Figure S5. HMBC spectrum of dicarabrol A (1) in CDCl₃

Figure S6. ROESY spectrum of dicarabrol A (1) in CDCl₃

Figure S7. IR spectrum of dicarabrol A (1)

Figure S8. HRESI(+)MS spectrum of dicarabrol A (1)

Figure S9. ¹H NMR spectrum of dicarabrone C (2) in CDCl₃

Figure S10. ¹³C NMR spectrum of dicarabrone C (2) in CDCl₃

Figure S11. ¹H-¹H COSY spectrum of dicarabrone C (2) in CDCl₃

Figure S12. HSQC spectrum of dicarabrone C (2) in CDCl₃

Figure S13. HMBC spectrum of dicarabrone C (2) in CDCl₃

Figure S14. ROESY spectrum of dicarabrone C (2) in CDCl₃

Figure S15. IR spectrum of dicarabrone C (2)

Figure S16. HRESI(+)MS spectrum of dicarabrone C (2)

Figure S17. ¹H NMR spectrum of dipulchellin A (3) in CDCl₃

Figure S18. ¹³C NMR spectrum of dipulchellin A (3) in CDCl₃

Figure S19. ¹H-¹H COSY spectrum of dipulchellin A (3) in CDCl₃

Figure S20. HSQC spectrum of dipulchellin A (3) in CDCl₃

Figure S21. HMBC spectrum of dipulchellin A (3) in CDCl₃

Figure S22. ROESY spectrum of dipulchellin A (3) in CDCl₃

Figure S23. IR spectrum of dipulchellin A (3)

Figure S24. HRESI(+)MS spectrum of dipulchellin A (3)

Experimental Section

X-ray crystallographic analyses. Dipulchellin A (**3**) was crystallized from acetone at room temperature. The X-ray crystallographic data was obtained on a Bruker SMART CCD detector employing graphite monochromated Cu-K α radiation ($\lambda = 1.54178$ Å) (operated in the φ - ω scan mode). The structure was solved by direct method using SHELXS-97 program and refined with full-matrix least-squares calculations on *F*2 using SHELXL-97. Crystallographic data for **3** (key parameters see Table S1) have been deposited at the Cambridge Crystallographic Data Centre (Deposition No.: CCDC 1479445). Copies of these data can be obtained free of charge via the internet at www.ccdc.cam.ac.uk/conts/retrieving.html or on application to CCDC, 12 Union Road, Cambridge CB2 1EZ, UK [Tel: (+44) 1223-336-408; Fax: (+44) 1223-336-033; E-mail: deposit@ccdc.cam.ac.uk].

Crystal data	
$C_{34}H_{44}O_7$	F(000) = 1216
$M_r = 564.69$	$D_{\rm x} = 1.221 {\rm Mg m^{-3}}$
Orthorhombic, $P2_12_12_1$	Cu $K\alpha$ radiation, $\lambda = 1.54178$ Å
a = 10.4583 (1) Å	$\mu = 0.68 \text{ mm}^{-1}$
b = 13.3200 (2) Å	T = 296 K
c = 22.0536 (3) Å	Plate, colourless
V = 3072.17(7) Å ³	$0.15 \times 0.1 \times 0.05 \text{ mm}$
Z = 4	
Data collection	

Table S1. X-ra	y crystallographi	c data for d	ipulchellin A ((3))
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Radiation source: fine-focus sealed tube $R_{int} = 0.034$

graphite	$\theta_{\text{max}} = 67.7^{\circ}, \ \theta_{\text{min}} = 3.9^{\circ}$
40986 measured reflections	$h = -10 \rightarrow 12$
5469 independent reflections	$k = -15 \rightarrow 14$
5246 reflections with $I > 2\sigma(I)$	$l = -26 \rightarrow 25$
Refinement	
Refinement on F^2	Secondary atom site location: difference Fourier map
Least-squares matrix: full	Hydrogen site location: inferred from neighbouring sites
$R[F^2 > 2\sigma (F^2)] = 0.099$	H atoms treated by a mixture of independent and constrained refinement
$wR(F^2) = 0.308$	$w = 1/[\sigma^2(F_o^2) + (0.2P)^2]$ where $P = (F_o^2 + 2F_c^2)/3$
<i>S</i> = 1.54	$(\Delta/\sigma)_{\rm max} = 0.179$
5469 reflections	$\Box \Delta \rangle_{\text{max}} = 0.93 \text{ e} \text{ Å}^{-3}$
377 parameters	$\Box \Delta \rangle_{\min} = -1.14 \text{ e } \text{\AA}^{-3}$
22 restraints	Absolute structure: Flack H D (1983), Acta Cryst. A39, 876-881
Primary atom site location: structure- invariant direct methods	Flack parameter: 0.1 (4)



Figure S1. ¹H NMR spectrum of dicarabrol A (1) in CDCl₃



Figure S2. ¹³C NMR spectrum of dicarabrol A (1) in CDCl₃



Figure S3. ¹H-¹H COSY spectrum of dicarabrol A (1) in CDCl₃



Figure S4. HSQC spectrum of dicarabrol A (1) in CDCl₃



Figure S5. HMBC spectrum of dicarabrol A (1) in CDCl₃

THJ-DG1617 CDC13 ROESY

Sample Name:

Data Collected on: 400MR-vnmrs400 Archive directory:

Sample directory:

FidFile: RDESY

Pulse Sequence: ROESY Solvent: cdc13 Data collected on: May 21 2013

Temp. 20.0 C / 293.1 K Operator: chempack

Relax. delay 1.000 sec Acq. time 0.107 sec Width 5542.0 Hz 2D Width 5542.0 Hz 3D Width 5542.0 Hz 8 repetitions 2 x 280 increments DBSERVE H1, 395.7504842 MHz DATA PROCESSING Gauss apodization 0.045 sec F1 DATA PROCESSING Gauss apodization 0.012 sec FT size 2048 x 2048 Total time 1 hr, 33 min



Figure S6. ROESY spectrum of dicarabrol A (1) in CDCl₃



Figure S7. IR spectrum of dicarabrol A (1)



Figure S8. HRESI(+)MS spectrum of dicarabrol A (1)



Figure S9. ¹H NMR spectrum of dicarabrone C (2) in CDCl₃



Figure S10. ¹³C NMR spectrum of dicarabrone C (2) in CDCl₃



Figure S11. ^{1}H - ^{1}H COSY spectrum of dicarabrone C (2) in CDCl₃



Figure S12. HSQC spectrum of dicarabrone C (2) in CDCl₃



Figure S13. HMBC spectrum of dicarabrone C (2) in CDCl₃



Figure S14. ROESY spectrum of dicarabrone C (2) in CDCl₃



Figure S15. IR spectrum of dicarabrone C (2)

Elemental Composition Report

Tolerance = 10.0 PPM / DBE: min = -1.5, max = 50.0 Isotope cluster parameters: Separation = 1.0 Abundance = 1.0%

Monoisotopic Mass, Odd and Even Electron Ions 36 formula(e) evaluated with 1 results within limits (up to 20 closest results for each mass)



Figure S16. HRESI(+)MS spectrum of dicarabrone C (2)



Figure S17. ¹H NMR spectrum of dipulchellin A (3) in CDCl₃



Figure S18. ¹³C NMR spectrum of dipulchellin A (3) in CDCl₃



Figure S19. ¹H-¹H COSY spectrum of dipulchellin A (3) in CDCl₃



Figure S20. HSQC spectrum of dipulchellin A (3) in CDCl₃



Figure S21. HMBC spectrum of dipulchellin A (3) in CDCl₃

THJ-1602 CDCL3 ROESY

Sample Name:

Data Collected on: 400MR-vmmrs400 Archive directory:

Sample directory:

FidFile: ROESY

Pulse Sequence: RDESY Solvent: cdc13 Data collected on: Sep 5 2013

Operator: chempack

Relax. delay 1.000 sec Aq. time 0.107 sec Width 3542.0 Hz 2D Width 3542.0 Hz 2D Width 3542.0 Hz 8 repetitions 2 x 260 increments 086ERVT NI. 359.7904873 MHz DBTA PROCESSING Geuss apodization 0.043 sec Fi DATA PROCESSING Gauss apodization 0.011 sec Data sec Total time 1 hr. 32 min



Figure S22. ROESY spectrum of dipulchellin A (3) in CDCl₃



Figure S23. IR spectrum of dipulchellin A (3)



Figure S24. HRESI(+)MS spectrum of dipulchellin A (3)