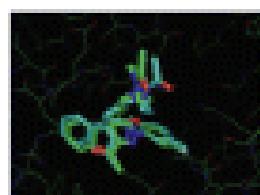




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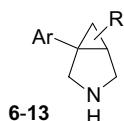
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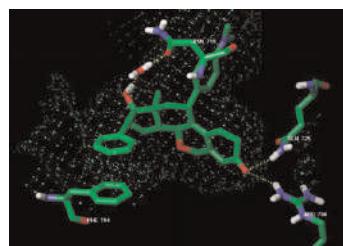
Studies on the structure–activity relationship of bicifadine analogs as monoamine transporter inhibitors

pp 3682–3686

Mingzhu Zhang, Florence Jovic, Troy Vickers, Brian Dyck, Junko Tamiya, Jonathan Grey,
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Fu-An Kang*, Xin Chen, Nareshkumar Jain, George Allan, Pamela Tannenbaum, Scott Lundein, Zhihua Sui

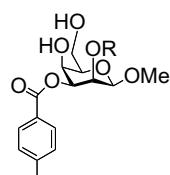


Molecular modeling, in vivo activity, pharmacokinetic and metabolic properties of unnatural 7-oxa-steroids are reported.

Protein subtype-targeting through ligand epimerization: Talose-selectivity of galectin-4 and galectin-8

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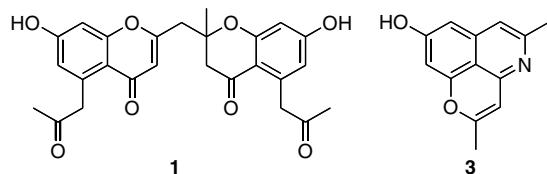
Christopher T. Öberg, Helen Blanchard, Hakon Leffler, Ulf J. Nilsson*

Galectin-4 C-terminal domain and galectin-8 N-terminal domain were found to prefer the β -talopyranose configuration to the natural ligand α -galactopyranose configuration. Methyl β -D-talopyranosides derivatized at O2 and O3 were synthesized and discovered to be selective submillimolar inhibitors of galectin-4C and galectin-8N.

Chrobisiamone A, a new bischromone from *Cassia siamea* and a biomimetic transformation of 5-acetyl-7-hydroxy-2-methylchromone into cassiarin A

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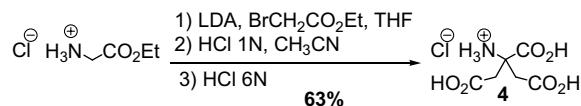


A new bischromone, chrobisiamone A (**1**) with an antiplasmodial activity has been isolated from the leaves of *Cassia siamea* and the structure was elucidated by 2D NMR analysis. Cyclization of 5-acetyl-7-hydroxy-2-methylchromone (**2**) in the presence of ammonium acetate resulted in generation of cassiarin A (**3**) with an unprecedented tricyclic skeleton, supporting a biogenetic pathway proposed for **3**.

2-Aminopropane-1,2,3-tricarboxylic acid: Synthesis and co-crystallization with the class A β -lactamase BS3 of *Bacillus licheniformis*

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Joséphine Beck, Eric Sauvage, Paulette Charlier, Jacqueline Marchand-Brynaert*



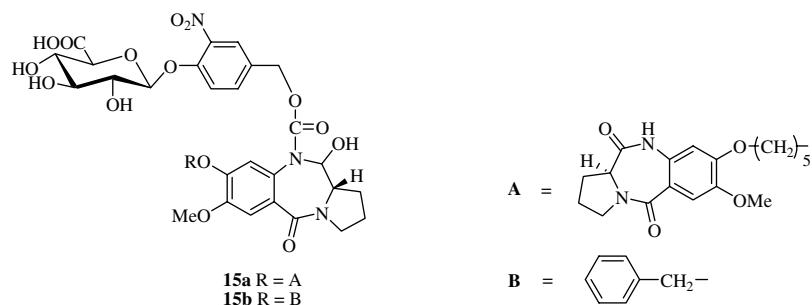
Synthesis and biochemical evaluation against β -lactamases of amino analog of citric acid are presented. The structure of the complex aminocitrate-BS3 has been analyzed by X-ray diffraction and compared to ones with citrate and isocitrate.



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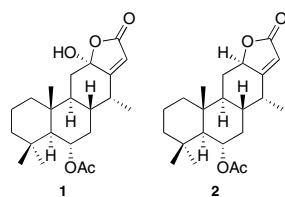
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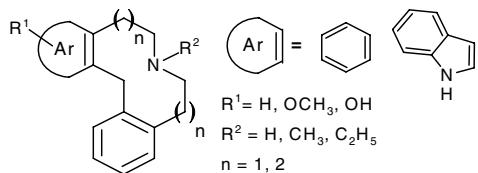
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Two new diterpenes, sucutiniranes A (**1**) and B (**2**), have been isolated from *Bowdichia nitida*. Sucutinirane A (**1**) and 6 α -acetoxyvouacapane (**3**) showed a moderate cytotoxicity and 6 α ,7 β -diacetoxyvouacapane (**4**) showed in vitro antiplasmodial activity against parasite *Plasmodium falciparum* 3D7.

**Dopamine/serotonin receptor ligands. Part 17: A cross-target SAR approach:
Affinities of azecine-styled ligands for 5-HT_{2A} versus D₁ and D₂ receptors**
Christoph Enzensperger, Tilo Görnemann, Heinz H. Pertz, Jochen Lehmann*

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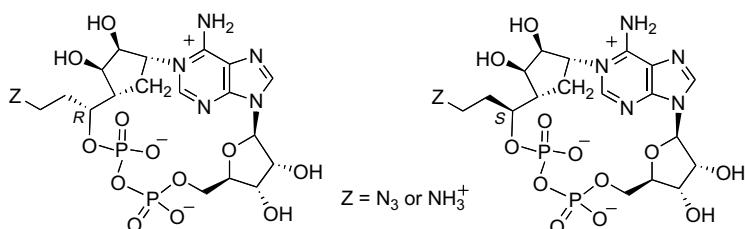
 pK_i for D₁: 6.79 to 9.45; for D₂: 6.41 to 8.45; pA_2 for 5-HT_{2A}: 7.21 to 9.97

A cross-target SAR was conducted with 13 azecine-styled compounds on D₁, D₂ and 5-HT_{2A} receptors. Surprisingly, molecular modifications affect the affinity for the D₁ receptor in the same manner as the 5-HT_{2A} receptor. The protein–ligand interactions were discussed with respect to the different binding pockets.

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Chrobisiamone A, a new bischromone from *Cassia siamea* and a biomimetic transformation of 5-acetonyl-7-hydroxy-2-methylchromone into cassiarin A

Shiori Oshimi^a, Yuichiro Tomizawa^a, Yusuke Hirasawa^a, Toshio Honda^a, Wiwied Ekasari^b, Aty Widyawaruyanti^b, Marcellino Rudyanto^b, Gunawan Indrayanto^b, Noor Cholies Zaini^b, Hiroshi Morita^{a,*}

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ABSTRACT

A new bischromone, chrobisiamone A (**1**) with an antiplasmodial activity has been isolated from the leaves of *Cassia siamea* and the structure was elucidated by 2D NMR analysis. Cyclization of 5-acetonyl-7-hydroxy-2-methylchromone (**2**) in the presence of ammonium acetate resulted in generation of cassiarin A (**3**) with an unprecedented tricyclic skeleton, supporting a biogenetic pathway proposed for **3**.

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Keywords:

Bischromone

Chrobisiamone A

Antiplasmodial activity

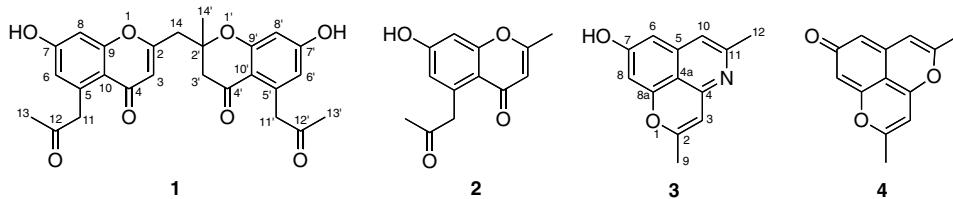
Cassia siamea

Biomimetic transformation

Malaria caused by parasites of the genus *Plasmodium* is one of the leading infectious diseases in many tropical and some temperate regions.¹ The emergence of widespread chloroquine-resistant and multiple-drug-resistant strains of malaria parasites leads to the need for the development of new therapeutic agents against malaria.²

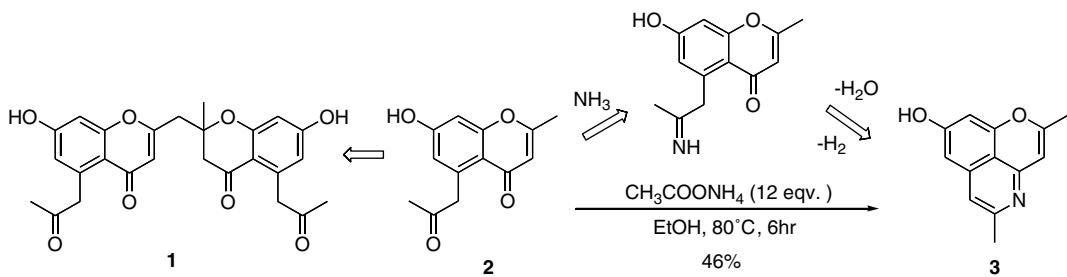
Cassia siamea (Leguminosae), have been widely used in traditional medicine, particularly for the treatment of periodic fever and malaria in Indonesia.³ So far some chromone derivatives, such as anhydrobarakol,⁴ 5-acetonyl-7-hydroxy-2-methylchromone,⁵ 2-methyl-5-propyl-7,12-dihydroxychromone-12-O-β-D-glucopyranoside,⁶ and cassiadinine,⁷ have already been isolated from the bark, leaves, and flowers of *C. siamea*. We have isolated cassiarins

A (**3**) and **B** with an unprecedented tricyclic skeleton and a potent antiplasmodial activity from the leaves of *C. siamea* and proposed a biogenetic path for cassiarin A (**3**) generated from 5-acetonyl-7-hydroxy-2-methylchromone (**2**).⁸ Recently, the first total synthesis of cassiarin A (**3**) was completed via sequential alkynylation of arenes with Sonogashira coupling and 6-*endo*-dig-cyclization of phenolic oxygens to the resulting alkynes in 51% overall yield in seven steps.⁹ On continuing search for chemical constituents of *C. siamea*, we have isolated a new bischromone, chrobisiamone A (**1**) with an antiplasmodial activity. This letter describes the isolation and structural elucidation of **1**, and a biomimetic transformation of 5-acetonyl-7-hydroxy-2-methylchromone (**2**) into cassiarin A (**3**).



* Corresponding author. Tel./fax: +81 354985778.
E-mail address: moritah@hoshi.ac.jp (H. Morita).

The leaves of *C. siamea* (300 g), which were collected at the Purwodadi Botanical Garden, Pasuruan, Indonesia (2005), were



Scheme 1. Biomimetic transformation and plausible biogenetic path for chrobisiamone A (**1**) and cassiarin A (**3**).

Table 1

¹H [δ , Hz] and ¹³C NMR data (δ) of chrobisiamone A (**1**) in CD₃OD at 300 K

	δ _H	δ _C
2		165.5
3	6.00 (1H, s)	113.8
4		180.6
5		139.4
6	6.61 (1H, d, 2.2)	120.2
7		164.1
8	6.68 (1H, d, 2.2)	102.9
9		161.1
10		115.1
11	4.17 (2H, s)	50.5
12		208.2
13	2.31 (3H, s)	30.9
14a	3.06 (1H, d, 14.0)	43.3
14b	2.93 (1H, d, 14.0)	
2'		80.7
3'	2.74 (2H, s)	49.0 ^a
4'		192.6
5'		140.7
6'	6.20 (1H, d, 2.4)	115.9
7'		165.9
8'	6.25 (1H, d, 2.4)	103.8
9'		165.2
10'		112.5
11'a	3.88 (1H, d, 17.1)	50.7
11'b	3.96 (1H, d, 17.1)	
12'		208.7
13'	2.26 (3H, s)	30.8
14'	1.50 (3H, s)	25.7

^a Overlapped with CD₃OD.

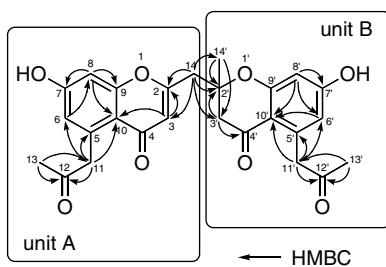


Figure 1. Selected 2D NMR correlations of chrobisiamone A (**1**) in CD₃OD.

extracted with MeOH, and the extract was partitioned with CHCl₃ and H₂O. CHCl₃-soluble materials were subjected to a silica gel column (CHCl₃/MeOH/AcOEt, 1:0:0 → 0:1:0), in which a fraction eluted by CHCl₃/MeOH/AcOEt (1:1:0) was further purified on an LH-20 column with CHCl₃/MeOH (1:1) to afford chrobisiamone A (**1**, 0.00013%) together with 5-acetonyl-7-hydroxy-2-methylchromone (**2**,⁵ cassiarins A (**3**) and B,⁸ anhydrobarakol (**4**),⁴ and 4-(trans)-acetyl-3,6,8-trihydroxy-3-methylidihydro-naphthalenone.¹⁰

Chrobisiamone A (**1**), colorless solid, $[\alpha]_D^{20} -19$ (*c* 0.33, MeOH), showed the molecular formula, C₂₆H₂₄O₈, which was determined by HRESIMS [m/z 465.1549, (M+H)⁺, Δ -0.2 mmu]. IR absorptions implied the presence of OH and/or NH (3100 cm⁻¹) and ketone (1720 and 1670 cm⁻¹) functionalities. UV absorptions at 238 (ε 17,500), 250 (14,400), and 282 (13,500) indicated a conjugated aromatic ring system. ¹H and ¹³C NMR data are presented in Table 1. The ¹³C NMR spectrum revealed 26 carbon signals due to 13 sp² quaternary carbons, one sp³ quaternary carbon, five sp² methines, four methylenes, and three methyls. Among them, six quaternary carbons (δ _C 80.7, 161.1, 164.1, 165.2, 165.5, and 165.9) and four carbonyl carbons (δ _C 180.6, 192.6, 208.2, and 208.7) were ascribed to those bearing an oxygen atom.

The gross structure of **1** was deduced from extensive analyses of the two-dimensional NMR data, including the ¹H-¹H COSY and HMBC spectra in CD₃OD (Fig. 1). The HMBC spectrum revealed connectivities of two chromone structures (C-1 ~ C-10 and C-1' ~ C-10') and two acetonyl groups (C-11 ~ C-13 and C-11' ~ C-13') classed into two units A and B as shown in Figure 1.

The presence of two acetonyl groups at C-5 and C-5' was supported by HMBC correlations for H₂-11 (δ _H 4.17) of C-5 (δ _C 134.1), C-6 (δ _C 120.2), and C-10 (δ _C 115.1), and H₂-11' (δ _H 3.88 and 3.96) of C-5' (δ _C 140.7), C-6' (δ _C 115.9), and C-10' (δ _C 112.5) as shown in Figure 1. HMBC correlations for H₂-14 (δ _H 2.93 and 3.06) of C-2 (δ _C 165.5), C-2' (δ _C 80.7), C-3 (δ _C 113.8), and C-3' (δ _C 49.0), H₃-14' (δ _H 1.50) of C-2' and C-3' gave rise to the connectivity of two partial structures A and B through C-14 and C-2' atoms. Thus, chrobisiamone A (**1**) was concluded to be an unique dimeric ring system consisting of 5-acetonyl-7-hydroxy-2-methylchromone and 5-acetonyl-7-hydroxy-2-methyl-2,3-hydrochromone.¹¹

A plausible biogenetic pathway for chrobisiamone A (**1**) and cassiarin A (**3**) was proposed as shown in Scheme 1. Chrobisiamone A (**1**) might be derived from 5-acetonyl-7-hydroxy-2-methylchromone (**2**) by Michael addition of the chromone carbanion of C-14 to C-2' of a second chromone. Treatment of 5-acetonyl-7-hydroxy-2-methylchromone (**2**) with ammonium acetate as the nitrogen source caused ring cyclization giving cassiarin A (**3**) in 46% yield. This biomimetic transformation might support a biogenetic pathway proposed for cassiarin A (**3**), which might be derived through an imine intermediate **2** followed by cyclization.

Chrobisiamone A (**1**), 5-acetonyl-7-hydroxy-2-methylchromone (**2**), and anhydrobarakol (**4**) showed a moderate in vitro antiplasmodial activity¹² against parasite *Plasmodium falciparum* 3D7 (IC₅₀ **1**: 2.6 µg/ml; **2**: 4.5 µg/ml; **4**: 7.8 µg/ml).

Acknowledgments

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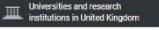
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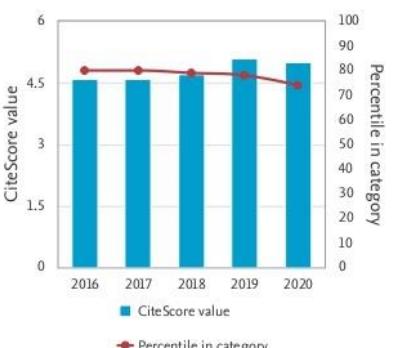
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