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1 ***Alicyclobacillus acidocaldarius* squalene-hopene cyclase: the critical role of steric**
2 **bulk at Ala306 and the first enzymatic synthesis of epoxydammarane from 2,3-**
3 **oxidosqualene**

4

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17

1 Abstract

2 The acyclic molecule squalene (**1**) is cyclized into a 6,6,6,6,5-fused pentacyclic hopene
3 (**2**) and hopanol (**3**) (ca. 5:1) by *Alicyclobacillus acidocaldarius* squalene-hopene
4 cyclase (*AaSHC*). The polycyclization reaction proceeds with regio- and stereochemical
5 specificity under precise enzymatic control. This pentacyclic hopane skeleton is
6 generated by folding **1** into an all-chair conformation. The Ala306 in *AaSHC* is
7 conserved in known SHCs; however, increases in steric bulk (A306T and A306V) led to
8 the accumulation of 6,6,6,5-fused tetracyclic scaffolds possessing 20*R* stereochemistry
9 in high yield (94% for A306V). The production of the 20*R* configuration indicated that
10 **1** had been folded in a chair-chair-chair-boat conformation; in contrast, the normal
11 chair-chair-chair-chair conformation affords the tetracycle with 20*S* stereochemistry, but
12 the yield produced by the A306V mutant was very low (6%). Consequently, bulk at
13 position 306 significantly affects the stereochemical fate during the polycyclization
14 reaction. SHC also accepts (3*R*) and (3*S*)-2,3-oxidosqualenes (OXSQs) to generate 3 α ,
15 β -hydroxyhopenes and 3 α , β -hydroxyhopanols through a polycyclization reaction
16 initiated at the epoxide ring. However, the Val and Thr mutants generated
17 epoxydammarane scaffolds from (3*R*)-OXSQ, which indicated that the polycyclization
18 cascade started at the terminal double bond position. This work is the first to report the
19 polycyclization of oxidosqualene starting from the terminal double bond.

21 Introduction

22 The cyclization mechanisms by which acyclic squalene (**1**) and (3*S*)-2,3-
23 oxidosqualene form polycyclic triterpenes have attracted much attention from chemists
24 and biochemists for over a half a century.^[1,2] *Alicyclobacillus acidocaldarius* squalene-
25 hopene cyclase (*AaSHC*) catalyzes the conversion of **1** into hopene (**2**) and hopanol (**3**)
26 (ca 5:1), pentacyclic triterpenes consisting of 6,6,6,6,5-fused A/B/C/D/E-ring systems,
27 as shown in Scheme 1A.^[2a,b] The X-ray crystallographic structure of the SHC from *A.*
28 *acidocaldarius* (*AaSHC*) was first reported in 1997^[3] and further refined later.^[4]
29 Compound **1** is folded into an all-prechair conformation inside the reaction cavity and is
30 cyclized in a regio- and stereospecific fashion via a series of carbocationic
31 intermediates, leading to the formation of 5 new C-C bonds and 9 stereocenters. This
32 reaction consists of 8 reaction steps^[2b] (Scheme 1B): (1) the acyclic molecule **1** is folded
33 into a chair conformation to give the intermediary monocyclic cation **4**; (2) a second
34 ring formation reaction occurs to afford cation **5**, consisting of a 6,6-fused bicyclic
35 cation; (3) a third ring formation reaction produces the 6,6,5-fused tricyclic cation **6**; (4)
36 cation **6** undergoes a ring expansion to generate the secondary cation **7** with a 6,6,6-

1 fused A/B/C-ring system; (5) a further ring closure gives the tertiary cation **8** with a
2 6,6,6,5-fused tetracycle (17-*epi*-dammarenyl cation); (6) cation **8** undergoes a second
3 ring expansion reaction to provide the secondary cation **9** (prohopanyl cation); (7) the
4 final ring formation reaction furnishes the 6,6,6,6,5-fused pentacyclic cation **10**
5 (hopanyl cation); and (8) proton elimination from Me-29 introduces a double bond to
6 produce **2**, and a water molecule attacks the C-22 cation of **10** to give **3**. Contrary to this
7 step-wise mechanism, there are reports that the polycyclization reactions proceed in a
8 concerted manner.^[5a, b] (3*S*)-2,3-Oxidosqualene (**11**) is also accepted as a substrate by
9 SHC, yielding 3β-hydroxyhopene (**12**) and 3β-hydroxyhopanol (**13**) through an all-chair
10 conformation in a similar cyclization pathway to that of **1** (Scheme 1C). (3*R*)-2,3-
11 Oxidosqualene (**14**) is also converted into 3α-hydroxyhopene (**15**) and 3α-
12 hydroxyhopanol (**16**) through a boat-chair-chair-chair-chair conformation.^[6a-d]
13

14 <Scheme 1>

15 To identify the active site residues of *Aa*SHC and their functions, site-specific
16 mutagenesis experiments were conducted. The central Asp residue (Asp376) in the
17 D³⁷⁴XD³⁷⁶D³⁷⁷ motif functions as the proton donor to initiate the polycyclization
18 reaction.^[7a,b] His451 hydrogen bonds with Asp376 to enhance the motif acidity.^[2a,6a]
19 Asp377 likely stabilizes cation **4**.^[7a] Trp312 likely binds with the terminal methyl group
20 via a hydrophobic interaction (possibly CH-π interaction) to place **1** in close proximity
21 to the DXDD motif, thus facilitating the initiation of the polycyclization reaction.^[8]
22 Phe365 stabilizes cation **5**.^[9] Phe601 likely stabilizes cations **6** and **7**,^[10] and Phe605
23 stabilizes the intermediary cations **9** and **10**.^[11] All the Phe residues mentioned above
24 function to stabilize transient cations via cation/π interactions.^[12] The main function of
25 Y420 is to stabilize the bicyclic cation **5** by pointing towards the active site cavity and
26 providing additional negative electrostatic potential.^[2a,4,13] The steric bulk at the Leu607
27 position is critical for the correct chair-folding around the B-ring formation site.^[13] The
28 steric bulk of another aliphatic residue, Ile261, is also crucial for enforcing the correct
29 folding around the D/E-ring formation sites.^[14] The I261A variant generates cations **6**
30 and **7** with 13α-H and 17α-H configurations, respectively, which are opposite to the
31 configurations of the genuine intermediary cations **6** (13β-H) and **7** (17β-H); thus, the
32 steric bulk at this position directs the stereochemical fate. There have been many recent
33 reports on the use of SHC mutants as biocatalysts, by which many known and unknown
34 compounds have been successfully created.^[15]

35 Herein, we report the function of the Ala306 residue in *Aa*SHC, which is widely
36 conserved in known SHCs (Fig. S1, Supporting Information). The X-ray crystal

1 structure of *Aa*SHC complexed with a modeled hopene molecule indicates that Ala306
2 is located near the squalene molecule (Fig. S2).^[3,4] To investigate the function of Ala306
3 of *Aa*SHC in hopene biosynthesis, we constructed site-directed mutants in which the
4 steric bulk at this position is altered, incubated **1**, **11**, and **14** with these variants, and
5 determined the structures of the resulting enzymatic products. We provide clear
6 evidence that the Ala306 residue is located in close proximity to the D-ring formation
7 site and that the steric bulk at this position is a key determinant of the stereochemical
8 fate during the polyolefin cyclization reaction, similar to the Ile216 residue.
9 Furthermore, we report the first case in which the polycyclization reaction of
10 oxidosqualene was initiated from the terminal double bond instead of the terminal
11 epoxide ring, resulting in the production of a 17-*epi*-epoxydammarane scaffold.
12

13 Results

14 Isolation of products generated from the incubation of **1** with A306X mutants

15 Fig. 1A shows the GC profile obtained by incubating **1** with the cell-free
16 homogenates of the A306V variant. Squalene **1** (200 mg) was incubated (pH 6.0, 45°C,
17 16 h) with the cell-free extracts (300 mL) obtained from a 6 L culture of the A306V
18 variant. The incubation mixture was freeze-dried, and the lipophilic materials, including
19 the enzymatic products, were extracted with hexane. These were then subjected to SiO₂
20 column chromatography by eluting with 100% hexane to remove the Triton X-100
21 present in the incubation mixture and to partially separate the products into several
22 fractions. Only a small amount of **1** was recovered, indicating that the reaction by the
23 variant was almost complete. The purity of the partially separated fractions was
24 monitored by GC-MS. The fraction containing a mixture of **17**, **19**, and **20** was obtained
25 by partial fractionation with 100% hexane as the eluent. Complete separation of the
26 products was attained by column chromatography with activated SiO₂ (heated in an
27 oven at 180°C for 2 h), yielding 2.2 mg of **17**, 1.0 mg of **19**, and 1.5 mg of **20** in a pure
28 state. However, separation of the other products was more difficult. The fraction
29 containing products **18**, **21**, **22**, **23**, and **24** was chromatographed on a 5% AgNO₃-
30 impregnated SiO₂ column, and pure **23** was obtained in a yield of 1.4 mg by eluting
31 with hexane/EtOAc (100:0.2), but the separation of **18** and **21** failed. This separation
32 was achieved by C₁₈ reverse-phase HPLC (THF/H₂O=60:40), which yielded 0.8 mg of
33 **18** and 1.3 mg of **21** in a pure state. The fraction obtained by eluting with hexane/EtOAc
34 (100:1) contained a mixture of **21**, **22**, **23**, and **24**. Pure **24** (2.1 mg) was obtained by
35 reverse-phase HPLC (THF/H₂O=70:30), but the separation of **21**, **22**, and **23** was not
36 successful. The fractions containing **18**, **21**, **22**, and **23**, which were not separable by 5%

1 argentation chromatography, were combined and subjected to 10%-AgNO₃ SiO₂ column
2 chromatography, and eluted with hexane/EtOAc (100:0.1) to yield a mixture of **18**, **21**,
3 and **22**, and then C₁₈-HPLC (THF/H₂O=60:40) successfully gave pure **22** in a yield of
4 0.6 mg. The isolation yields of **18**, **21**, **22**, and **23** were very low, despite the large peaks
5 detected by GC (Fig. 1A). The low yields could be ascribed to both the separation
6 difficulty and the use of an AgNO₃-SiO₂ column (as the recovery from AgNO₃
7 chromatography is generally low). In addition, there were many remaining fractions that
8 have yet to be purified. Fig. S3.1 depicts the GC trace from the incubation of **1** with the
9 A306T variant. Products **25–27**, which were not detected as products of the Val mutant,
10 were newly revealed in the GC traces, and **28** was identified as a new product of the
11 A306F mutant (Fig. S3.2). Products **25–28** were all identified by comparing their EIMS
12 fragment patterns and retention times to those of previously isolated authenticated
13 compounds.^[2b]

14
15 <Figure 1>

16 **Isolation of products generated from the incubation of **11** and **14** with A306X**

17 **mutants**

18 Fig. 1B shows the GC chromatogram of the reaction mixture obtained by
19 incubating a mixture of **11** and **14** with the A306V mutant. To isolate the enzymatic
20 products **29–33**, a racemic mixture of **11** and **14** (100 mg) was incubated (pH 6.0, 45°C,
21 20 h) with the cell-free extract (300 mL) from a 6 L culture of the A306V mutant. After
22 lyophilization of the reaction mixture, the reaction products were extracted with hexane,
23 which was passed through a short SiO₂ column by elution with hexane/EtOAc (100:20)
24 to remove Triton X-100. The residues obtained after removing Triton X-100 were
25 subjected to SiO₂ column chromatography in a stepwise gradient fashion
26 (hexane/EtOAc=100:1 to 100:2). Product **33** was isolated in a pure state (6.0 mg), but
27 the other products could not be separated. The inseparable fractions were divided into
28 the following 2 fractions: a mixture (22.4 mg) of **29**, **31**, and **33**, and a mixture (14.7
29 mg) of **30** and **32**. Products **29** and **31** were successfully separated by normal-phase
30 HPLC (hexane/2-propanol=100:0.1) to yield **29** (2.3 mg) and **31** (1.0 mg) in a pure
31 state. Products **30** and **32** were separated by normal-phase HPLC
32 (hexane/THF=100:1.0) with yields of 2.0 mg and 2.3 mg, respectively. The A306F
33 variant generated products **34**, **35**, and **36** from a mixture of **11** and **14**, as shown in Fig.
34 S4. Fifty mg of a mixture of **11** and **14** was incubated with the A306F mutant. SiO₂
35 column chromatography (hexane/EtOAc=100:1 to 100:5) afforded pure **36** (14.5 mg)
36 and a mixture (6.0 mg) of **34** and **35**, which were then acetylated with Ac₂O/Py (1:1).

1 Normal-phase HPLC (hexane:2-PrOH=100:0.02) successfully isolated the pure acetates
2 of **34** and **35** in yields of 0.6 mg and 0.8 mg, respectively. The isolation yields are not
3 consistent with the production yields because some fractions still contained inseparable
4 products.

6 **Structural determination of products 17–36.**

7 Fig. 2 lists the structures of enzymatic products that were identified. Products **17–**
8 **28**, with the exception of **23**, were previously isolated from the site-specific variants of
9 *AaSHC*.^[2b] Detailed NMR analyses of **23**, a new product generated from *AaSHC*-
10 mutants, are described below. The EIMS and NMR spectra of **17–22** and **24** are shown
11 in Figs. S5–S10, and brief descriptions for proposing the structures are provided in the
12 Supporting Information. We isolated **17** from the L607F,^[13] Y420A,^[13] and Y609F
13 variants.^[8] Compound **18** is identical to the product isolated from variants I261A^[14] and
14 F605A.^[11] Product **19** was isolated from mutants Y609A,^[8] Y612A^[8], and F365A.^[9]
15 Product **20** is identical to the compound generated by the F601A mutant.^[10] Compound
16 **21** was isolated from the W489 variant.^[2b, 16, 17] Product **22** was previously isolated from
17 the F605A mutant.^[11] Product **24** was isolated from the W169F, W169H, and W489F
18 variants.^[17] The EIMS spectra of minor products **25–28** are shown in Fig. S13. The
19 spectra and GC retention times of **25–28** were the same as those of samples we isolated
20 previously: **25** and **27** from F605A,^[11] **26** from P263G,^[19] and **28** from F365A.^[9]
21 Accordingly, these structures are illustrated in Fig. 2.

22 The detailed NMR analyses of product **23**, including 2D NMR spectra (600 MHz,
23 C₆D₆, Fig. S11.9), revealed that **23** had the same tetracyclic skeleton as **22**. Furthermore,
24 **22** and **23** had identical EIMS spectra (Figs. S10.1 and S11.1), but their GCMS
25 retention times were different (Fig. 1A), indicating that the 2 compounds are
26 stereoisomers. The relative configurations of **23**, which is present in the A/B/C/D-fused
27 tetracyclic scaffolds, were determined to be identical to those of **22** by detailed analyses
28 of the NOESY spectrum (Fig. S11.6). Thus, the C-20 stereochemistry of **23** is opposite
29 to that of **22** and can be assigned as the 20*S* configuration for **23**. Consequently, **23** is
30 20*S*-tirucalla-7(8),24-diene, as shown in Fig. 2. The δ_{H} of Me-21 of 20*S*-tirucalla-
31 7(8),24-diene in CDCl₃ was reported to be δ_{H} 0.895 (d, *J*=6.8 Hz),^[18] which was close
32 to that of **23** (δ_{H} 0.885, *J*=6.8 Hz) in CDCl₃ solution. Thus, the 20*S* stereochemistry of
33 **23** is credibly assigned.

34 Next, the structural analyses of the enzymatic products **29–36**, which were
35 generated by incubating a mixture of **11** and **14**, are described. We have not reported the
36 enzymatic reactions of **11** and **14** by the *AaSHC* variants; all the products are new SHC-

1 metabolites, and the NMR analyses are described below.

2 The ^1H -NMR spectrum of **29** (400 MHz, C_6D_6 , Fig. S14.2) showed the presence of
3 an isopropylidene residue: δ_{H} 1.83 (s, 3H, Me-26) and 1.75 (s, 3H, Me-27). The ^{13}C
4 NMR spectrum (100 MHz, C_6D_6 , Fig. S14.4) revealed the presence of a tetrasubstituted
5 double bond (δ_{C} 139.7, s; 134.9, s). Detailed HMBC analyses (Fig. S14.10) indicated
6 that **29** consists of a 6,6,6,5-fused tetracyclic system and that the double bonds of **29**
7 exist at C-13–C-17 and C-24–C-25. The multiplicity of H-3 (δ_{H} 3.29) showed a broad
8 singlet, indicating that the hydroxyl group is α -oriented. The doublet Me was located at
9 C-20 because Me-21 (δ_{H} 1.14, d, $J=7.2$ Hz, 3H) showed a strong HMBC cross peak for
10 C-17. The chemical shift of Me-21 (δ_{H} 0.918 d, $J=6.8$ Hz) in the CDCl_3 solution (see
11 Fig. S14.3) was very close to the reported value (δ_{H} 0.910 d, $J=6.9$ Hz) for 20*R*-
12 dammara-13(17),24-diene.^[18] Thus, **29** was determined to be 3 α -hydroxy-20*R*-
13 dammara-13(17)-24-diene, as shown in Fig. 2.

14 The EIMS spectrum of **30** (Fig. S15.1) is almost identical to that of **29** (Fig. S14.1),
15 indicating that **30** is a stereoisomer of **29**. Detailed NMR analyses (Fig. S15.9) showed
16 that dammara-13(17),24-diene can be assigned to **30** as described in the structural
17 determination of **29**. The C-20 stereochemistry of **30** was the same as that of **29** because
18 the Me-21 of **30** showed a δ_{H} 0.919 (d, $J=6.8$ Hz, 3H) identical to that of **29** in CDCl_3
19 solution (cf. in the C_6D_6 solution, δ_{H} 1.14 for Me-21 of **29**; δ_{H} 1.15 for Me-21 for **30**;
20 both compounds had the same chemical shifts in C_6D_6). The OH was determined to be
21 β -oriented because H-3 (δ_{H} 3.16) gave a dd splitting pattern (dd, $J=10.2$, 6.1 Hz). The
22 entire structure of **30** can be deemed 3 β -hydroxy-20*R*-dammara-13(17),24-diene, as
23 shown in Fig. 2.

24 The ^1H -NMR (400 MHz, C_6D_6) and ^{13}C -NMR (100 MHz, C_6D_6) spectra of **31** (Fig.
25 S16.3 and S16.4) revealed the presence of 2 double bonds: δ_{H} 5.41 (2H: m, 1H, H-12;
26 m, 1H, H-24) and δ_{C} 125.5 (d, C-24), 130.9 (s, C-25), 115.9 (d, C-12) and 149.2 (s, C-
27 13). Two Me groups (δ_{H} 1.88, s, 3H, Me-26; 1.73, s, 3H, Me-27) had clear HMBC cross
28 peaks with C-25 and C-24, indicating that the isopropylidene moiety remained without
29 participation of the cyclization reaction, and thus, **31** is a tetracyclic product. Me-30 (δ_{H}
30 1.13, s, 3H) showed a distinct HMBC correlation with C-13; Me-21 (δ_{H} 1.05, d, $J=6.8$
31 Hz, 3H) had a clear HMBC contour with C-17 (δ_{C} 48.94, d); and the multiplicity of C-
32 17 (d) indicated the presence of a proton at C-17. Thus, another double bond is
33 positioned at C-12–C-13. Indeed, the chemical shifts of H-11 [δ_{H} 1.92 (m) and 2.16 (m)]
34 clearly indicated that H-11 is located at the allylic position. The OH group was α -
35 oriented because the multiplicity of H-3 (δ_{H} 3.29) showed a broad singlet. Detailed
36 NMR analyses (Fig. S16.10) indicated that **31** possesses 3 α -hydroxy-17-*epi*-dammara-

12(13),24-diene with *R* stereochemistry at C-20 because the chemical shift of Me-21 in CDCl₃ solution was δ_{H} 0.789 (d, $J=6.4$ Hz, see Fig. S16.2.2), which was very close to the reported value (δ_{H} 0.784, d, $J=6.8$ Hz),^[18] and to that of Me-21 (δ_{H} 0.793, d, $J=6.8$ Hz, measured in CDCl₃, see Fig. S9.9) in product **21**.

Detailed NMR analyses (Fig. S17.9) indicated that product **32** also has a 17-*epi*-dammarane-12(13),24-diene scaffold, which was further confirmed by the fact that the EIMS spectra of **31** (Fig. S16.1) and **32** (Fig. S17.1) were identical. The OH function was oriented at the β -position, as the multiplicity of H-3 (δ_{H} 3.14) was a double doublet ($J=11.4, 4.7$ Hz). Furthermore, the chemical shift of Me-21 (δ_{H} 1.06, d, $J=6.7$ Hz, Fig. S17.2.2) of **32** in C₆D₆ solution was the same as that (δ_{H} 1.05, d, $J=6.8$ Hz, Fig. S16.3.2) of **31** (cf. δ_{H} 0.784 for **32** and δ_{H} 0.789 for **31**, in CDCl₃ solution, Fig. S17.2.1). Thus, **32** was proposed to be 3 β -hydroxy-17-*epi*-20*R*-dammarane-12(13),24-diene.

No olefinic protons and no double bonds were found in the ¹H-NMR (600 MHz, acetone *d*₆, Fig. S18.2) and ¹³C-NMR spectra (150 MHz, acetone *d*₆, Fig. S18.3) of product **33**, indicating that the full cyclization reactions occurred. Three carbons connected to the oxygen atom were found: δ_{C} 86.06 (s, C-20), 88.23 (d, C-24) and 70.43 (s, C-25). Three Me groups had distinct HMBC correlations with these 3 carbons: Me-21 (δ_{H} 1.19, s, 3H)/C-20 and both Me-26 (δ_{H} 1.13, s, 3H) and Me-27 (δ_{H} 1.05, s, 3H)/C-24/C-25, where the chemical shift assignments of Me-26 and Me-27 are exchangeable. Me-21 exhibited further HMBC contours with C-22 (δ_{C} 38.70, t) and C-17 (δ_{C} 49.30, d), and H-24 (δ_{H} 3.71, dd, $J=10.6, 5.2$ Hz) was spin-coupled with H-23 (δ_{H} 1.75, m, 1H; 1.85, m, 1H). Thus, the moiety 2-(tetrahydrofuran-2-yl)propan-2-ol is connected to C-17. This THF moiety was further confirmed by EIMS, in which the fragment ion *m/z* 143 was observed as the base peak (Fig. 18.1). More detailed HMBC analyses revealed the presence of a 6,6,6,5-fused tetracyclic ring, in which H-17 is positioned in a β -orientation because H-17 had a clear NOE with H-13 (δ_{H} 2.09, m, 1H). Further analyses, including 2D NMR (Fig. S18.9), indicated that product **33** is 20,24-epoxydammarane-25-ol with 17 β -H (17-*epi*-epoxydammarane skeleton). No NOE was found between Me-21 and H-24, indicating that either (20*R*, 24*R*) or (20*S*, 24*S*) can be assigned for the THF ring. We have reported that epoxydammarane scaffolds are created by the SHC-mediated reactions of non-natural (3*R*)- and (3*S*)-2,3-squalene diols and 2,3:22,23-dioxidosqualenes,^[20] but this work is the first to report the production of the epoxydammarane from oxidosqualene. The ¹H- and ¹³C-NMR data of **33** (Fig. S18.9) were indistinguishable from those of (20*R*, 24*R*)-17-*epi*-epoxydammarane (Fig. S18.10), which was obtained from the reaction of (3*R*)-2,3-squalene diol with native SHC.^[20] (20*S*, 24*S*)-17-*epi*-Epoxydammarane and (20*R*, 24*R*)-17-*epi*-epoxydammarane

1 are diastereomers, thus the NMR spectra of the 2 diastereomers must be different, which
2 allowed us to propose (20*R*, 24*R*)-stereochemistry for **33**.

3 Product **34**-acetate showed peaks for 5 vinylic Me groups: δ_{H} 1.69 (s, 3H, Me-29);
4 δ_{H} 1.81 (s, 3H, Me-30); δ_{H} 1.72 (s, 6H, Me-26 and Me-28); and δ_{H} 1.78 (s, 3H, Me-27).
5 One tetrasubstituted double bond (δ_{C} 126.2, s, C-8 and δ_{C} 140.4, s, C-9) was found.
6 Furthermore, 3 aliphatic Me signals were observed: δ_{H} 1.05 (s, 3H, Me-25), δ_{H} 0.892 (s,
7 3H, Me-24), and δ_{H} 1.05 (s, 3H, Me-23). These findings are suggestive of a bicyclic
8 ring system. Me-25 and Me-26 had distinct HMBC cross peaks with C-9 and C-8,
9 respectively, thus the tetrasubstituted double bond is arranged at C-8 and C-9. The H-3
10 (δ_{H} 5.03) showed a broad singlet, indicating α -oriented OH. Consequently, the structure
11 of **35** is polypoda-8(9),13,17,21-tetraen-3 α -ol, as shown in Fig. 2.

12 Detailed NMR analyses of **35**-acetate (Fig. S20.9) and **36** (Fig. S21.9) revealed that
13 the structures of **35** and **36** are polypoda-7(8),13,17,21-tetraen-3 α -ol and polypoda-
14 7(8),13,17,21-tetraen-3 β -ol, respectively.

15
16 <Figure 2>

17 Discussion

18 Mechanistic insights into the production of 17-36 from substrates 1, 11, and 14.

19 Scheme 2 shows the formation mechanisms of 6,6-fused bicyclic products.
20 Substrates **1** and **11** were folded into a chair-chair conformation to give 6,6-fused
21 bicyclic cations **5** and **37**, respectively. Substrate **14** was folded into a boat-chair
22 conformation to afford the same bicyclic skeleton **38**, but the OH-orientation (α -
23 arrangement) from **14** was opposite to that (β -orientation) from **11**. This finding is
24 consistent with the cyclization mechanisms of **1**, **11**, and **14** (Scheme 1). Proton
25 elimination from Me-26 gave **17** (path *a*). The deprotonation of 7 α -H produced **19**, **35**,
26 and **36** (path *b*). The elimination of H-9 generated **34** (path *d*), which was obtained only
27 from substrate **14**. Product **28** was produced by the following successive antiparallel
28 1,2-shifts: H-9 \rightarrow C-8 cation, Me-25 \rightarrow C-9, H-5 \rightarrow C-10, and the deprotonation of 6 β -H to
29 give the double bond at C-5 and C-6 (path *c*).

30 Scheme 3A shows the formation mechanism of tricyclic **20**. This compound was
31 produced only from **1**. Substrate **1** was folded into a chair-chair-boat conformation (**39**)
32 to furnish the 6,6,5-fused A/B/C-ring system with 13 α -H (**40**). On the other hand, the
33 chair-chair-chair conformation (**41**) gives the 6,6,5-fused tricycle with 13 β -H (**42**),
34 which is identical to cation **7**. Deprotonation from Me-27 of cation **40** afforded product
35 **20** (path *a*).

36 Scheme 3B depicts the cyclization mechanisms leading to the pentacyclic products

1 **25** and **26**, the skeletons of which were produced only from **1** and not from **11** and **14**.
2 Squalene **1** was folded into an all-chair conformation to afford the final hopanyl cation
3 **10**. The following successive rearrangement reactions of **10** with the C-22 cation
4 afforded **25** (path *b*): H-21 to C-22 cation, H-17 to C-21, Me-28 to C-17, followed by
5 deprotonation of H-13. H-21 was eliminated to give **26** (path *c*).

6 Scheme 3C illustrates the formation mechanism of product **33**, which is produced
7 by the A306V and the A306T mutants (Fig. 1B and Fig. S4). The (20*R*, 24*R*)
8 stereochemistry of **33** indicates that this compound was produced from (3*R*)-
9 oxidosqualene **14**, but not from **11**. Substrate **14** was folded into an all-chair
10 conformation. Proton attack on the terminal double bond, but not on the epoxide ring,
11 triggered successive ring formation reactions to yield the 17-*epi*-6,6,6,5-fused
12 dammarenyl cation **43**. The epoxide ring of **43** attacked the C-20 cation (*Re*-face
13 attack), and then the nucleophilic attack of a water molecule on the C-25 cation afforded
14 **33**. Intriguingly, the cyclization reaction started from the terminal double bond position
15 and not from the epoxide ring. This study is the first to report a polycyclization reaction
16 that initiated from the terminal double bond, despite the epoxide ring being present in
17 the substrate. As shown in Scheme 1B, the SHC-mediated cyclization reaction starts
18 from the epoxide ring side but not from the terminal double bond side. This
19 phenomenon has been clearly explained in terms of the higher nucleophilicity of
20 epoxide than that of double bond π -electrons towards the DXDD motif, a proton donor
21 that can initiate the polycyclization reaction.^[7] However, why epoxydammarane
22 compounds such as **33** are never produced from **11** and **14** by the native SHC, even in
23 small amounts, is unclear. An increase in the steric bulk at position 306 (A306V and
24 A306T variants) may have altered the geometry of the substrate inside the reaction
25 cavity, and the terminal double bond may have been positioned nearer to the DXDD
26 motif than the epoxide ring. In the case of the wild type SHC, the epoxide ring may be
27 in closer contact with the DXDD motif, and the more highly nucleophilic nature of the
28 epoxide ring may further facilitate the polycyclization reaction. We have reported that
29 native SHC can accept both (3*R*)- and (3*S*)-2,3-squalene diols to afford 24*R*- and 24*S*-
30 epoxydammarane, respectively.^[20] Furthermore, the four different 2,3;22,23-
31 dioxidosqualenes (3*R,S* and 22*R,S* isomers) are also converted into four 3-hydroxy-
32 epoxydammarane scaffolds with four different configurations (3*R*, 24*R*; 3*R*, 24*S*; 3*S*,
33 24*R*; 3*S*, 24*S*);^[20] that is, native SHC can accept both 22*R*- and 22*S*-isomers as
34 substrates, which correspond to (3*R*)- and (3*S*)-isomers in the case of the
35 polycyclization reactions of 2,3-oxidosqualene, respectively, resulting in the production
36 of both 24*R*- and 24*S*-epoxydammaranes. This fact that native SHC can accept both

1 (3*R*)-**14** and (3*S*)-**11** raises the question why (3*S*)-**11** was not cyclized by the A306V and
2 A306T variants to produce 24*S*-epoxydammarane, while both (3*R*)-**14** and (3*S*)-**11**
3 underwent cyclization reactions from the epoxide ring to generate 3 α - and 3 β -hydroxy-
4 tetracycle **29–32** and bicycle **34–36**. It is also unclear why the mutated SHCs cannot
5 accept the monocyclic and bicyclic products of the premature cyclization step to yield
6 further cyclized products through reentry of the mon- and/or bicyclic product into the
7 reaction cavity after exiting. The substrate uptake and the product release occur through
8 the same channel,^[3,4] which may be very compact; thus, the truncated cyclization
9 products cannot reenter the reaction cavity. In contrast, tetraprenyl- β -curcumene cyclase
10 from *Bacillus megaterium* (BmeTC) can accept 3-deoxyachilleol A (monocyclic
11 triterpene) to yield (+)-ambrein as a result of the re-entry of this monocyclic product
12 into the reaction cavity.^[21] Further studies are necessary to provide detailed insight into
13 the cyclization mechanism underlying how **33** is produced from **14**.
14

15 <Scheme 2>

16 Scheme 4 shows the formation mechanisms of tetracyclic products from **1**, **11**, and
17 **14**. Two types of formation mechanisms are proposed: (A) the chair-chair-chair-boat
18 folding conformation **44** from **1**, leading to the tetracycle **45** with 17 α -H; (B) the chair-
19 chair-chair-chair folding conformation **48** from **1**, resulting in the tetracycle **49** with
20 17 β -H. Substrates **11** and **14** were also cyclized according to the folding conformation
21 of Type A, yielding the 3 β -hydroxytetracyclic cation **46** and 3 α -hydroxytetracyclic
22 cation **47**, respectively. By contrast, Type B is limited to the cyclization reactions of **1**.
23 The Newman projection of cations **45**, **46**, and **47** is shown in Scheme 4A. A minimal
24 motion (60° rotation) in the reaction cavity leads to 20*R* stereochemistry, but a large
25 rotation (120°) gives the 20*S* configuration. A minimal motion is preferable to a large
26 motion in the reaction cavity and is thus allowed to afford 20*R* stereochemistry. This
27 theory was proposed by Abe and Rohmer.^[18] By contrast, the minimal motion (60°
28 rotation) of cation **49** with 17 β -H affords a 20*S* configuration (allowed, Scheme 4B),
29 but the large motion (120° rotation) results in the generation of 20*R* products (not
30 allowed, Scheme 4B). Thus, cation **49** led to products with 20*S* stereochemistry. Cation
31 **45** underwent further structural modifications. A hydride shift of H-17 to the C-20
32 cation, followed by the deprotonation of H-13, furnished **18**, **29**, and **30** (path *a*). 1,2-
33 Shifts of H-17 to the C-20 cation and H-13 to C-17, followed by the removal of H-12 α ,
34 gave **21**, **31**, and **32** (path *b*). The following consecutive 1,2-rearrangement reactions
35 occurred: H-17→C-20 cation, H-13→C-17, Me-30→C-13, Me-18→C14, and the
36 subsequent deprotonation of H-7 α generated product **22** (path *c*). Proton elimination of

1 Me-21 of cation **49** provided product **24** (path *d*). Cation **49** underwent sequential 1,2-
2 shift reactions (path *c*): H-17→C-20, H-13→C-17, Me-30→C-13, Me-18→C14. The
3 subsequent deprotonation reaction of H-7 α furnished **23** (path *c*). The nucleophilic
4 attack of a water molecule on the C-20 cation gave **27** (path *e*).
5

6 <Scheme 3>

7 **Distribution ratio of products generated by altering the steric bulk at position 306**

8 Table 1 shows the product distribution ratio obtained by the reactions of squalene **1**.
9 To address the steric bulk, we employed the van der Waals volumes (nm³) reported by
10 Lin et al.: Gly, 0.00279; Ala, 0.05702; Thr, 0.19341; Val, 0.25674; Leu, 0.37876; and
11 Phe, 0.55298.^[22] These values indicate that the steric bulk increases as follows:
12 Gly<Ala<Thr<Val<Leu<Phe. The Gly mutant, with decreased steric bulk, afforded the
13 normal cyclization products **2** and **3** in large quantities and the aberrant cyclization
14 products **25** and **26** (pentacycle) in small amounts, but did not produce any bi-, tri-, or
15 tetracyclic products. The Thr variant, with increased bulk, afforded decreased amounts
16 of **2** and **3** (25%), while the tetracyclic products were generated in increased quantities
17 (57%). The generation of the tetracyclic products suggests that the Ala306 residue is
18 located near the D-ring formation site. A further increase in steric bulk (A306V variant)
19 afforded small amounts of **2** (2%) and of bicyclic (4%) and tricyclic products (2%), but
20 large quantities of tetracyclic products were produced (90%). Furthermore, almost all
21 the tetracyclic products possessed a 20*R* configuration (94%) derived from cation **45**,
22 while only 6% were 20*S* tetracycles derived from cation **49**. In the case of the A306T
23 variant, 63% of the products consisted of **18**, **21**, and **22**, derived from cation **45**
24 (Scheme 4A), and 37% consisted of **23**, **24**, and **27**, derived from cation **49** (Scheme
25 4B). Therefore, an increase in the steric bulk facilitates the formation of cation **45**.
26 Cation **8**, an intermediary tetracyclic cation (Scheme 1A), is the same as cation **49**
27 (Scheme 4B), indicating that 20*S* tetracyclic products should be generated from the
28 intermediary cation **8** or **49** during the normal cyclization process. However, increased
29 steric bulk at the 306 position drastically increased the production of 20*R* tetracycles,
30 strongly indicating that the size of the residue at 306 is critical to the normal
31 polycyclization pathway. The Leu mutant furnished bicyclic products with a
32 significantly high yield (43%), and an almost equal amount of tetracyclic products
33 (39%), as shown in Table 1 and Fig. S3.3. The Leu and Val residues have the same
34 hydrophobic properties, but possess no polar OH and π -electrons; the difference
35 between them is only the steric size (Leu>Val). Replacing Phe with bulk larger than Val
36 and Leu generated only bicyclic products and no tetracyclic products. This finding

1 indicates that the introduction of significantly larger substituents at the 306 position
2 altered the orientation of substrate **1** inside the reaction cavity. Consequently, the steric
3 bulk at position 306 significantly affects the stereochemical fate and the truncation steps
4 during the polycyclization cascade.

5 Table 2 shows the product distribution ratios obtained by the incubation of **11** and
6 **14**. The wild type produced **12** and **13** from **11**, and **15** and **16** from **14**, but no other
7 product was produced, as shown in Scheme 1B. The A306T variant produced tetracyclic
8 products in larger amounts (82%), in addition to **12** and **15** (8%). This result is
9 consistent with the outcomes from the incubation of **1** (Table 1), but a small amount
10 (8%) of epoxydammarane **33** was found by GC (Fig. S4). The A306V variant also
11 produced tetracyclic products with a 20*R* configuration (47%), and no products with
12 20*S* stereochemistry were present above the limit of GC detection. Interestingly, a larger
13 amount of **33** was produced (20%) than that by the A306T variant. The A306F mutant
14 produced only bicyclic products **34–36**, which was consistent with the results of
15 squalene incubation with the A306F variant. Notably, products **29–32** were all generated
16 from cations **46** and **47** (equivalent to **45**), which resulted from a chair-chair-chair-boat
17 folding conformation (Scheme 4A). In addition, **33** was produced only from **14** but not
18 from **11**. The higher production of **33** by the Val mutant than by the Thr mutant indicates
19 that the production of **33** is also dependent on the steric bulk at position 306. In fact, the
20 larger Phe variant did not generate any **33** (Table 2). Thus, an appropriate steric size is
21 required for the formation of **33**.

22
23 This study clearly demonstrates that the appropriate steric bulk at position 306 of
24 AaSHC is critical for directing the normal polycyclization cascade. We describe the
25 enzymatic reactions of various substrate analogs with SHC, e.g., norsqualenes lacking
26 Me-group^[6a, 23] and homosqualenes bearing ethyl groups on the squalene backbone,^[24]
27 and we provide insight into how the polycyclization reaction is affected by the specific
28 modifications. The steric bulk at the Me-positions of squalene substrate significantly
29 altered the polycyclization pathway, e.g., alteration of the folding conformation leading
30 to different stereochemistry and a truncation of the ring-forming cascade, which in some
31 cases resulted in the generation of novel triterpene scaffolds.^[23b] The results of the site-
32 specific mutagenesis experiments provide further evidence that the appropriate steric
33 size of the active site residues is critical to correct folding and the full cyclization
34 pathway.

35 36 Conclusions

1 The following conclusions can be drawn from our study: (1) the appropriate steric
2 bulk of the Ala306 residue directs the normal polycyclization reaction without
3 premature termination; (2) the Ala side chain is in close proximity to the D-ring
4 formation site, as the A306V mutant produced large amounts of tetracyclic products
5 with the 20*R* configuration; and (3) this is the first report describing the initiation of the
6 polycyclization reaction from the terminal double bond rather than from the epoxide
7 ring. As discussed above, the production of **33** raises the question why the native SHC
8 does not produce epoxydammarane structure(s) from **11** or **14**, specifically why the
9 proton attack occurs only at the epoxide ring, but not the terminal double bond, despite
10 both the terminal double bond and the epoxide ring entering the reaction cavity through
11 the substrate channel. The A306V and A306T variants are able to accept both heads of
12 oxidosqualene **14** because products **29–32** are generated by a proton attack on the
13 epoxide ring, while product **33** is produced by a proton attack on the terminal double
14 bond.

17 **Experimental**

19 **General analytical methods:** NMR spectra were recorded in C₆D₆ on a Bruker
20 DMX 600 or DPX 400 spectrometer, the chemical shifts (δ) given in ppm relative to the
21 residual solvent peak δ_{H} 7.280 and δ_{C} 128.0 as the internal reference for ¹H- and ¹³C-
22 NMR spectra, respectively. In the case of CDCl₃ solutions, the chemical shifts are given
23 in ppm relative to the solvent peak (δ_{H} 7.26; δ_{C} 77.0). The coupling constants *J* are given
24 in Hz. GC analyses were performed on a Shimadzu GC-8A chromatograph equipped
25 with a flame ionization detector (DB-1 capillary column, 0.53 mm × 30 m). GC-MS
26 spectra were performed on a JEOL SX 100 spectrometer under electronic impact at 70
27 eV with a DB-1 capillary column (0.32 mm × 30 m). HR-EIMS was performed by
28 direct inlet system. Specific rotation values were measured at 25°C with a Horiba
29 SEPA-300 polarimeter.

31 **Site-directed mutagenesis**

32 The *shc* genes were mutated via oligonucleotide-directed *in vitro* mutagenesis using
33 the Pharmacia Unique Site Elimination Mutagenesis kit (Pfizer, New York, NY, USA).
34 Plasmid pKS, a pUC119 derivative containing a 1 kb *KpnI-SacI* fragment, was used as a
35 template for all mutagenesis reactions. The synthetic oligonucleotides for the mutations
36 (with changes to the wild type sequence shown in bold type) are as follows:

1 A306G: 5'-GGATGTTTCAGGGGTCGATATCGCCGGTGTG-3'
2 A306T: 5'-GGATGTTTCAGACTTCGATATCGCCGGTGTG-3'
3 A306V: 5'-GGATGTTTCAGGTTTCGATATCGCCGGTGTG-3'
4 A306L: 5'-GGATGTTTCAGTTGTCGATATCGCCGGTGTG-3'
5 A306F: 5'-GGATGTTTCAGTTTTTCGATATCGCCGGTGTG-3'

6 Silent mutations were also used to screen the desired mutants by restriction fragment
7 analysis (underlined sequences). To ascertain that the desired mutation had been
8 achieved, the entire region of the inserted DNA was sequenced.

9

10 **Incubation conditions and GC analysis**

11 The SHC described here is from *A. acidocaldarius*.^[7-14, 16-17, 19,20] Standard culture of
12 the cloned *Escherichia coli* and incubation conditions were performed according to our
13 published protocols.^[7] The cell-free extract was prepared as follows. One liter of *E. coli*
14 culture encoding the native and mutated SHCs (pET3a) was harvested by centrifugation
15 50 mL of citrate buffer solution (pH 6.0, 50 mM) was added to the pellets, and
16 ultrasonication was then performed to disrupt the cells. The supernatant was used for
17 incubations after removal of all cell debris by centrifugation. One mL of the supernatant
18 contained *ca.* 200 μ g of the pure SHCs. One mg of the substrates, **1** and a mixture of **11**
19 and **14**, and 20 mg of Triton X-100 were emulsified with 1.0 mL dH₂O and 3.0 mL Na-
20 citrate buffer solution (pH 6.0, 0.5 M). To this solution, 2.0 mL of the cell-free extract
21 was added, followed by incubation at 45°C for 12 h. To terminate the enzyme reaction,
22 6 mL of 15% KOH/MeOH was added and heated at 80°C for 30 min. The lipophilic
23 enzyme products and the substrate analogs, which remained unreacted, were extracted 4
24 times with hexane (5 mL) from the incubation mixtures, and the quantities of the
25 products and the starting materials were estimated by GC analyses with a DB-1
26 capillary column (30 m in length, J&W Scientific, Folsom, CA, USA). GC-EIMS
27 product profiles and substrate distributions, prepared after the saponification treatment
28 (quenching reaction), were identical to those obtained prior to the saponification
29 reaction, and hence there was no change in product distribution and structural
30 modification, or the substrates produced by the quenching conditions. The epoxide ring
31 was also unmodified by this quenching treatment. GC analyses shown in Fig. 1A were
32 performed under the following conditions: injection temperature, 290°C; column
33 temperature, 290°C; N₂ carrier pressure, 0.75-1.5 kg/cm². GC analyses shown in Fig.
34 1B were performed on the same GC instrument and GC column as (A), but the N₂
35 carrier pressure was 0.5 kg/cm².

36 **Spectroscopic data of enzymatic products.**

1 Almost the enzymatic products obtained from squalene incubation were reported in our
2 previous papers, as described in Text.^[2b] To validate the identification, we isolated
3 products **17–24** and determined by the detailed NMR analyses: see the Figs. S5–S12 in
4 Supporting Information. Production of **23** by the mutated SHCs has not been reported
5 before. Production yields of **25–28** were significantly small, thus identifications were
6 made by GC-EIMS, as described in Fig. S13. NMR and EIMS data of products **29–36**,
7 which have not been reported before. Therefore, the spectroscopic data of **23** and **29–36**
8 are given here.

9 **Product 23:** $[\alpha]_{\text{D}}^{25} = -0.565$ ($c=0.018$, EtOH); ^1H NMR (600 MHz, C_6D_6): $\delta=0.964$ (s,
10 3H, Me-19), 0.967 (s, 3H, Me-28), 1.03 (s, 3H, Me-29), 1.04 (m, 1H, H-1), 1.05 (s, 3H,
11 Me-18), 1.12 (d, $J=6.4$ Hz, Me-21), 1.18 (s, 3H, Me-30), 1.27 (m, 2H, H-22 and H-3),
12 1.42 (m, 1H, H-16), 1.48 (dd, $J=12.1, 5.4$ Hz, 1H, H-5), 1.52 (m, 2H, H-2 and H-3),
13 1.58 (m, 1H, H-20), 1.60 (m, 1H, H-11), 1.62 (m, 1H, H-15), 1.63 (m, 1H, H-17), 1.66
14 (m, 1H, H-11), 1.68 (m, 1H, H-2), 1.72 (m, 1H, H-22), 1.74 (m, 1H, H-1), 1.75 (s, 3H,
15 Me-27), 1.77 (m, 2H, H-12 and H-15), 1.83 (s, 3H, Me-26), 1.92 (m, 1H, H-12), 2.00
16 (m, 1H, H-6), 2.09 (m, 1H, H-16), 2.14 (m, 1H, H-23), 2.25 (m, 1H, H-6), 2.33 (m, 1H,
17 H-23), 2.42 (m, 1H, H-9), 5.42 (bt, $J=6.2$ Hz, H-24), 5.47 (bd, $J=2.6$ Hz, 1H, H-7); ^{13}C
18 NMR (150 MHz, C_6D_6): $\delta=13.36$ (q, C-19), 17.72 (q, C-27), 18.48 (t, C-11), 18.62 (q,
19 C-21), 19.44 (t, C-2), 21.50 (q, C-29), 22.25 (q, C-18), 24.70 (t, C-6), 25.53 (t, C-23),
20 25.88 (q, C-26), 27.56 (q, C-30), 28.60 (t, C-16), 33.16 (q, C-28), 33.29 (s, C-4), 34.29
21 (t, C-12), 34.46 (t, C-15), 35.39 (s, C-10), 36.37 (d, C-20), 36.64 (t, C-22), 39.26 (t, C-
22 1), 42.69 (t, C-3), 43.82 (s, C-13), 49.50 (2C, d, C-9 and s, C-14), 51.66 (d, C-5), 53.42
23 (d, C-17), 118.5 (d, C-7), 125.7 (d, C-24), 130.7 (s, C-25), 146.1 (s, C-8). Me-21
24 showed $\delta_{\text{H}} 0.885$ (d, $J=6.8$ Hz) in the CDCl_3 solution. The assignments of C-13 and C-
25 14 may be exchangeable. MS(EI): m/z : 69 (30%), 109 (28%), 395 (100%), 410 (27%);
26 $[\text{M}^+]$; HRMS (EI): m/z : calcd. for $\text{C}_{30}\text{H}_{50}$: 410.39125; found: 410.39187.

27 **Product 29:** $[\alpha]_{\text{D}}^{25} = -17.15$ ($c=0.206$, EtOH); ^1H NMR (400 MHz, C_6D_6): $\delta=0.872$ (s,
28 3H, Me-29), 0.949 (s, 3H, Me-19), 1.045 (s, 3H, Me-28), 1.105 (s, 3H, Me-18), 1.141
29 (d, $J=7.2$ Hz, 3H, Me-21), 1.253 (s, 3H, Me-30), 1.39 (m, 1H, H-11), 1.40 (m, 1H, H-
30 15), 1.43 (m, 1H, H-6), 1.47 (m, 2H, H-1), 1.48 (m, 1H, H-7), 1.53 (2H, m, 1H, H-22;
31 m, 1H, H-5), 1.54 (m, 1H, H-6), 1.55 (m, 1H, H-2), 1.62 (m, 1H, H-22), 1.67 (m, 1H,
32 H-11), 1.70 (m, 1H, H-7), 1.71 (m, 1H, H-9), 1.752 (s, 3H, Me-27), 1.828 (s, 3H, Me-
33 26), 1.93 (m, 1H, H-2), 2.02 (m, 1H, H-12), 2.07 (m, 1H, H-15), 2.10 (m, 1H, H-23),
34 2.25 (m, 1H, H-23), 2.28 (m, 1H, H-16), 2.46 (m, 1H, H-16), 2.59 (dd, $J=13.6, 3.6$ Hz,
35 1H, H-12), 2.72 (m, 1H, H-20), 3.29 (br s, 1H, H-3), 5.42 (t, $J=6.4$ Hz, H-24); ^{13}C NMR
36 (100 MHz, C_6D_6): $\delta=16.50$ (q, C-19), 17.53 (q, C-18), 17.79 (q, C-27), 18.53 (t, C-6),

1 20.18 (q, C-21), 22.09 (t, C-11), 22.39 (q, C-29), 23.23 (q, C-30), 23.30 (t, C-12), 25.86
2 (q, C-26), 25.94 (t, C-2), 27.18 (t, C-23), 28.59 (q, C-28), 29.47 (t, C-16), 31.05 (t, C-
3 15), 32.24 (d, C-20), 33.90 (t, C-1), 35.72 (t, C-7), 36.02 (t, C-22), 37.74 (s, C-10),
4 37.79 (s, C-4), 41.60 (s, C-8), 49.71 (d, C-5), 51.82 (d, C-9), 56.94 (s, C-14), 75.80 (d,
5 C-3), 125.5 (d, C-24), 130.9 (s, C-25), 134.9 (s, C-17), 139.7 (s, C-13). The assignments
6 of C-4 and C-10 may be exchangeable. MS (EI) see Fig. S14.1. HRMS (EI): m/z : calcd.
7 for $C_{30}H_{50}O$: 426.38616; found: 410.38896.

8 **Product 30**: $[\alpha]_D^{25} = -10.9$ ($c=0.278$, EtOH); 1H NMR (600 MHz, C_6D_6): $\delta=0.796$ (dd,
9 $J=11.8$, 1.4 Hz, 1H, H-5), 0.909 (s, 3H, Me-19), 0.912 (s, 3H, Me-29), 0.92 (m, 1H, H-
10 1), 1.070 (s, 3H, Me-18), 1.149 (s, 3H, Me-28), 1.152 (d, $J=6.7$ Hz, 3H, Me-21), 1.275
11 (s, 3H, Me-30), 1.40 (2H, m, 1H, H-11; m, 1H, H-15), 1.44 (m, 1H, H-6), 1.46 (m, 1H,
12 H-7), 1.47 (m, 1H, H-9), 1.53 (m, 1H, H-22), 1.56 (m, 2H, H-2), 1.62 (2H: m, 1H, H-
13 22; m, 1H, H-7), 1.63 (m, 1H, H-11), 1.64 (m, 1H, H-6), 1.69 (m, 1H, H-1), 1.753 (s,
14 3H, Me-27), 1.831 (s, 3H, Me-26), 2.02 (m, 1H, H-12), 2.07 (m, 1H, H-15), 2.09 (m,
15 1H, H-23), 2.24 (m, 1H, H-23), 2.28 (m, 1H, H-16), 2.47 (m, 1H, H-16), 2.59 (ddd,
16 $J=12.5$, 4.7, 1.8 Hz, 1H, H-12), 2.72 (m, 1H, H-20), 3.16 (dd, $J=10.2$, 6.1 Hz, 1H, H-3),
17 5.41 (t, $J=7.0$ Hz, H-24); ^{13}C NMR (150 MHz, C_6D_6): $\delta=15.83$ (q, C-29), 16.61 (q, C-
18 19), 17.52 (q, C-18), 17.79 (q, C-27), 18.59 (t, C-6), 20.18 (q, C-21), 22.17 (t, C-11),
19 23.11 (q, C-30), 23.31 (t, C-12), 25.86 (q, C-26), 27.17 (t, C-23), 27.89 (t, C-2), 28.35
20 (q, C-28), 29.47 (t, C-16), 31.06 (t, C-15), 32.26 (d, C-20), 35.84 (t, C-7), 36.01 (t, C-
21 22), 37.62 (s, C-10), 39.15 (s, C-4), 39.26 (t, C-1), 41.34 (s, C-8), 52.02 (d, C-9), 56.29
22 (d, C-5), 56.83 (s, C-14), 78.46 (d, C-3), 125.4 (d, C-24), 130.9 (s, C-25), 135.1 (s, C-
23 17), 139.5 (s, C-13). MS (EI) see Fig. S15.1. HRMS (EI): m/z : calcd. for $C_{30}H_{50}O$:
24 426.38616; found: 410.38648.

25 **Product 31**: $[\alpha]_D^{25} = +6.3$ ($c=0.074$, EtOH); 1H NMR (400 MHz, C_6D_6): $\delta=0.900$ (s,
26 3H, Me-29), 1.011 (s, 3H, Me-18), 1.030 (s, 3H, Me-19), 1.047 (d, $J=6.8$ Hz, 3H, Me-
27 21), 1.065 (s, 3H, Me-28), 1.131 (s, 3H, Me-30), 1.32 (2H: m, 1H, H-1; m, 1H, H-15),
28 1.44 (m, 1H, H-22), 1.46 (m, 1H, H-7), 1.50 (m, 1H, H-6), 1.51 (m, 1H, H-2), 1.52 (m,
29 1H, H-1), 1.57 (2H: m, 1H, H-5; m, 1H, H-6), 1.62 (m, 1H, H-22), 1.63 (m, 2H, H-16),
30 1.730 (s, 3H, Me-27), 1.77 (m, 1H, H-15), 1.80 (m, 1H, H-7), 1.85 (m, 1H, H-2), 1.884
31 (s, 3H, Me-26), 1.92 (2H: m, 1H, H-11; m, 1H, H-9), 1.98 (m, 1H, H-20), 2.16 (m, 1H,
32 H-11), 2.23 (m, 2H, H-23), 2.59 (m, 1H, H-17), 3.29 (bs, 1H, H-3), 5.41 (2H: m, 1H, H-
33 12; m, 1H, H-24); ^{13}C NMR (100 MHz, C_6D_6): $\delta=15.48$ (q, C-21), 15.71 (q, C-19),
34 17.05 (q, C-18), 17.74 (q, C-27), 18.69 (t, C-6), 22.44 (q, C-29), 22.85 (q, C-30), 23.86
35 (t, C-11), 24.49 (t, C-16), 25.76 (t, C-2), 25.89 (q, C-26), 26.53 (t, C-23), 28.59 (q, C-
36 28), 32.07 (t, C-15), 33.46 (t, C-1), 34.71 (d, C-20), 35.06 (t, C-7), 36.35 (t, C-22),

1 37.60 (s, C-10), 37.61 (s, C-4), 38.52 (s, C-8), 48.36 (d, C-9), 48.94 (d, C-17), 49.75 (d,
2 C-5), 50.57 (s, C-14), 75.77 (d, C-3), 115.9 (d, C-12), 125.5 (d, C-24), 130.9 (s, C-25),
3 149.2 (s, C-13). The assignments of C-4 and C-10 may be exchangeable. MS (EI) see
4 Fig. S16.1. HRMS (EI): m/z : calcd. for $C_{30}H_{50}O$: 426.38616; found: 410.38432.

5 **Product 32:** $[\alpha]_D^{25} = +8.0$ ($c=0.213$, EtOH); 1H NMR (600 MHz, C_6D_6): $\delta=0.865$ (dd,
6 $J=12.0, 10.3$ Hz, 1H, H-5), 0.92 (m, 1H, H-1), 0.947 (s, 3H, Me-29), 0.973 (s, 3H, Me-
7 18), 0.991 (s, 3H, Me-19), 1.061 (d, $J=6.7$ Hz, 3H, Me-21), 1.148 (s, 3H, Me-30), 1.157
8 (s, 3H, Me-28), 1.30 (dd, $J=6.3, 11.8$ Hz, 1H, H-15), 1.42 (m, 1H, H-7), 1.47 (m, 1H, H-
9 22), 1.50 (m, 1H, H-6), 1.52 (m, 1H, H-2), 1.56 (2H: m, 1H, H-1; m, 1H, H-2), 1.62 (m,
10 2H, H-16), 1.63 (m, 1H, H-22), 1.64 (m, 1H, H-6), 1.68 (m, 1H, H-7), 1.72 (m, 1H, H-9),
11 1.732 (s, 3H, Me-27), 1.78 (m, 1H, H-15), 1.832 (s, 3H, Me-26), 1.89 (m, 1H, H-11), 1.99
12 (m, 1H, Me-20), 2.08 (m, 1H, H-11), 2.24 (m, 2H, H-23), 2.59 (m, 1H, H-17), 3.14 (dd,
13 $J=11.4, 4.7$ Hz, 1H, H-3), 5.39 (bs, 1H, H-12), 5.47 (t, $J=7.4$ Hz, H-24); ^{13}C NMR (150
14 MHz, C_6D_6): $\delta= 15.48$ (q, C-21), 15.88 (2C; q, C-19; q, C-29), 16.97 (q, C-18), 17.73 (q,
15 C-27), 18.72 (t, C-6), 22.83 (q, C-30), 23.91 (t, C-11), 24.50 (t, C-16), 25.88 (q, C-26),
16 26.53 (t, C-23), 27.71 (t, C-2), 28.32 (q, C-28), 32.08 (t, C-15), 34.72 (d, C-20), 35.08 (t,
17 C-7), 36.34 (t, C-22), 37.48 (s, C-10), 38.30 (s, C-8), 38.96 (t, C-1), 39.06 (s, C-4), 48.57
18 (d, C-9), 48.94 (d, C-17), 50.47 (s, C-14), 56.32 (d, C-5), 78.50 (d, C-3), 115.9 (d, C-12),
19 125.5 (d, C-24), 130.9 (s, C-25), 149.3 (s, C-13). MS (EI) see Fig. S17.1. HRMS (EI):
20 m/z : calcd. for $C_{30}H_{50}O$: 426.38616; found: 410.38648.

21 **Product 33:** $[\alpha]_D^{25} = -20.8$ ($c=0.239$, EtOH); 1H NMR (600 MHz, acetone d_6): $\delta=0.817$
22 (s, 3H, Me-29), 0.82 (m, 1H, H-5), 0.849 (s, 3H, Me-28), 0.86 (m, 1H, H-1), 0.861 (s,
23 3H, Me-19), 0.969 (s, 3H, Me-18), 1.025 (s, 3H, Me-30), 1.050 (s, 3H, Me-27), 1.06 (m,
24 1H, H-15), 1.135 (s, 3H, Me-26), 1.14 (m, 1H, H-3), 1.188 (s, 3H, Me-21), 1.21 (m, 1H,
25 H-11), 1.26 (m, 1H, H-7), 1.32 (m, 1H, H-3), 1.38 (m, 1H, H-2), 1.39 (m, 1H, H-6),
26 1.43 (2H: m, 1H, H-15; m, 1H, H-9), 1.49 (m, 1H, H-12), 1.52 (m, 1H, H-6), 1.53 (m,
27 1H, H-16), 1.58 (m, 1H, H-11), 1.59 (m, 1H, H-22), 1.61 (2H: m, 1H, H-2; m, 1H, H-7),
28 1.65 (m, 1H, H-1), 1.75 (m, 1H, H-23), 1.76 (m, 1H, H-16), 1.85 (m, 1H, H-23), 1.92
29 (2H: m, 1H, H-12; m, 1H, H-22), 2.09 (m, 1H, H-13), 2.24 (m, 1H, H-17), 3.71 (dd,
30 $J=10.6, 5.2$ Hz, 1H, H-24); ^{13}C NMR (150 MHz, acetone d_6): $\delta= 16.33$ (q, C-18), 16.64
31 (q, C-19), 16.94 (q, C-30), 19.37 (t, C-6), 19.47 (t, C-2), 21.86 (q, C-29), 23.21 (t, C-
32 11), 25.83 (q, C-27), 26.40 (t, C-23), 26.84 (t, C-12), 27.37 (q, C-26), 28.29 (t, C-16),
33 28.91 (q, C-21), 33.12 (t, C-15), 33.76 (q, C-28), 33.95 (s, C-4), 36.09 (t, C-7), 38.19 (s,
34 C-10), 38.70 (t, C-22), 41.35 (t, C-1), 41.68 (s, C-8), 42.89 (t, C-3), 44.60 (d, C-13),
35 49.36 (d, C-17), 49.85 (s, C-14), 51.71 (d, C-9), 57.79 (d, C-5), 70.43 (s, C-25), 86.06
36 (s, C-20), 88.23 (d, C-24). MS (EI) see Fig. S18.1. HRMS (EI): m/z 429 ($M^+ - Me$):

1 calcd. for $C_{29}H_{49}O_2$: 426.37325; found: 429.37229.

2 **Product 34-acetate**: $[\alpha]_D^{25} = -9.5$ ($c=0.014$, EtOH); 1H NMR (600 MHz, C_6D_6): $\delta = 0.892$
3 (s, 3H, Me-24), 1.051 (s, 3H, Me-25), 1.052 (s, 3H, Me-23), 1.50 (m, 1H, H-6), 1.62 (m,
4 1H, H-6), 1.68 (m, 1H, H-1), 1.691 (s, 3H, Me-29), 1.724 (s, 3H, Me-26), 1.724 (s, 3H,
5 Me-28), 1.779 (s, 3H, Me-27), 1.806 (s, 3H, Me-30), 1.82 (s, 3H, Me-32, acetyl Me),
6 1.85 (m, 1H, H-1), 1.88 (2H, m, H-2), 1.92 (m, 1H, H-5), 2.01 (dd, $J=6.0$, 18.2 Hz, 1H,
7 H-7), 2.14 (m, 1H, H-12), 2.16 (m, 1H, H-7), 2.24 (4H: m, H-15 and H-19), 2.32 (m,
8 1H, H-12), 2.33 (m, 4H, H-16 and H-20), 2.38 (m, 2H, H-11), 5.03 (bs, 1H, H-3), 5.37
9 (t, $J=6.8$ Hz, 1H, H-21), 5.42 (t, $J=6.8$ Hz, 1H, H-13), 5.43 (t, $J=6.8$ Hz, 1H, H-17); ^{13}C
10 NMR (150 MHz, C_6D_6): $\delta = 16.12$ (q, C-28), 16.21 (q, C-27), 169.7 (s, C-31, acetyl
11 \underline{CO}), 17.72 (q, C-29), 18.87 (t, C-6), 19.70 (q, C-26), 20.02 (q, C-25), 20.76 (q, C-32,
12 acetyl $\underline{CH_3}$), 21.84 (q, C-24), 23.71 (t, C-2), 25.83 (q, C-30), 27.20 (2C: t, C-16 and t,
13 C-20), 27.88 (q, C-23), 28.74 (t, C-12), 29.65 (t, C-11), 30.75 (t, C-1), 33.66 (t, C-7),
14 36.99 (s, C-4), 38.98 (s, C-10), 40.19 (t, C-15), 40.19 (t, C-19), 46.52 (d, C-5), 77.48 (d,
15 C-3), 124.7 (d, C-17), 124.9 (d, C-21), 125.3 (d, C-13), 126.2 (s, C-8), 131.1 (s, C-22),
16 134.8 (s, C-18), 135.1 (s, C-14), 140.4 (s, C-9). The assignments of C-14 and C-18 may
17 be exchangeable. MS (EI) see Fig. S19.1. HRMS (EI): m/z 408 ($M^+ - CH_3CO$): calcd. for
18 $C_{30}H_{48}$: 408.37560; found: 408.37590.

19 **Product 35-acetate**: $[\alpha]_D^{25} = -20.0$ ($c=0.025$, EtOH); 1H NMR (600 MHz, C_6D_6):
20 $\delta = 0.867$ (s, 3H, Me-25), 0.893 (s, 3H, Me-24), 0.957 (s, 3H, Me-23), 1.11 (m, 1H, H-1),
21 1.44 (m, 1H, H-11), 1.56 (m, 1H, H-1), 1.61 (m, 1H, H-11), 1.691 (s, 3H, Me-29), 1.740
22 (s, 6H, Me-27 and Me-28), 1.806 (s, 3H, Me-30), 1.83 (3H: m, 2H, H-2; m, 1H, H-5),
23 1.88 (m, 1H, H-9), 1.889 (s, 3H, Me-26), 1.890 (s, 3H, Me-32, acetyl Me), 1.96 (very
24 broad, 2H, H-6), 2.15 (m, 1H, H-12), 2.24 (m, 4H, H-15 and H-19), 2.33 (m, 4H, H-16
25 and H-20), 2.34 (m, 1H, H-12), 4.99 (t, $J=2.4$ Hz, H-3), 5.37 (t, $J=6.8$ Hz, H-21), 5.42
26 (t, $J=6.8$ Hz, H-13), 5.43 (t, $J=6.8$ Hz, H-17), 5.45 (very br s, 1H, H-7); ^{13}C NMR (150
27 MHz, C_6D_6): $\delta = 13.57$ (q, C-25), 16.10 (q, C-28), 16.28 (q, C-27), 169.6 (s, acetyl \underline{CO}),
28 17.72 (q, C-29), 20.77 (q, acetyl $\underline{CH_3}$), 21.90 (q, C-24), 22.36 (q, C-26), 23.08 (t, C-2),
29 23.59 (t, C-6), 25.83 (q, C-30), 27.20 (2C: t, C-16 and C-20), 27.67 (2C: q, C-23; t, C-
30 11), 30.61 (t, C-12), 32.29 (t, C-1), 36.54 (s, C-4), 36.58 (s, C-10), 40.19 (t, C-19),
31 40.22 (t, C-15), 44.99 (d, C-5), 54.12 (d, C-9), 77.90 (d, C-3), 122.4 (d, C-7), 124.7 (d,
32 C-17), 124.9 (d, C-21), 125.3 (d, C-13), 131.1 (s, C-22), 135.1 (s, C-18), 135.2 (s, C-14),
33 135.6 (s, C-8). The assignments may be exchangeable between C-4 and C-10, between
34 C-14 and C-18, between C-27 and C-28 and C-15 and C-19. MS (EI) see Fig. S20.1.
35 HRMS (EI): m/z 408 ($M^+ - CH_3CO$): calcd. for $C_{30}H_{48}$: 408.37560; found: 408.37759.

36 **Product 36**: $[\alpha]_D^{25} = -34.4$ ($c=1.697$, EtOH); 1H NMR (400 MHz, C_6D_6): $\delta = 0.878$ (s,

1 3H, Me-25), 0.975 (s, 3H, Me-24), 1.084 (s, 3H, Me-23), 1.11 (m, 1H, H-1), 1.22 (t,
2 $J=8.4$ Hz, H-5), 1.40 (m, 1H, H-11), 1.41 (m, 1H, H-2), 1.55 (m, 1H, H-2), 1.60 (m, 1H,
3 H-11), 1.69 (m, 1H, H-9), 1.694 (s, 3H, Me-29), 1.750 (s, 3H, Me-28), 1.768 (s, 3H,
4 Me-27), 1.807 (s, 3H, Me-30), 1.835 (s, 3H, Me-26), 1.87 (m, 1H, H-1), 2.02 (very br s,
5 2H, H-6), 2.16 (m, 1H, H-12), 2.27 (m, 4H, H-15 and H-19), 2.27-2.38 (m, 4H, H-16
6 and H-20), 2.38 (m, 1H, H-12), 3.15 (dd, $J=10.8, 4.8$ Hz, H-3), 5.37 (t, $J=6.8$ Hz, 1H,
7 H-21), 5.42 (t, $J=6.8$ Hz, 1H, H-13), 5.44 (t, $J=6.8$ Hz, 1H, H-17), 5.53 (very br s, 1H,
8 H-7); ^{13}C NMR (100 MHz, C_6D_6): $\delta=13.71$ (q, C-25), 131.1 (s, C-22), 16.12 (q, C-28),
9 16.26 (q, C-27), 17.73 (q, C-29), 22.27 (q, C-26), 23.81 (t, C-6), 25.30 (q, C-24), 25.84
10 (q, C-30), 27.13 (t, C-20), 27.23 (t, C-16), 27.70 (t, C-11), 27.93 (t, C-2), 28.12 (q, C-
11 23), 30.66 (t, C-12), 36.78 (s, C-10), 37.54 (t, C-1), 38.83 (s, C-4), 40.22 (2C, t, C-15
12 and C-19), 49.84 (d, C-5), 54.37 (d, C-9), 78.76 (d, C-3), 122.5 (d, C-7), 124.7 (d, C-
13 17), 124.9 (d, C-21), 125.3 (d, C-13), 135.1 (2C, s, C-14 and C-18), 135.2 (s, C-8). The
14 assignments may be exchangeable between C-14 and C-18, between C-16 and C-20 and
15 between C-27 and C-28. MS (EI) see Fig. S21.1. HRMS (EI): m/z : calcd. for $\text{C}_{30}\text{H}_{50}\text{O}$:
16 426.38616; found: 410.38354.

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22

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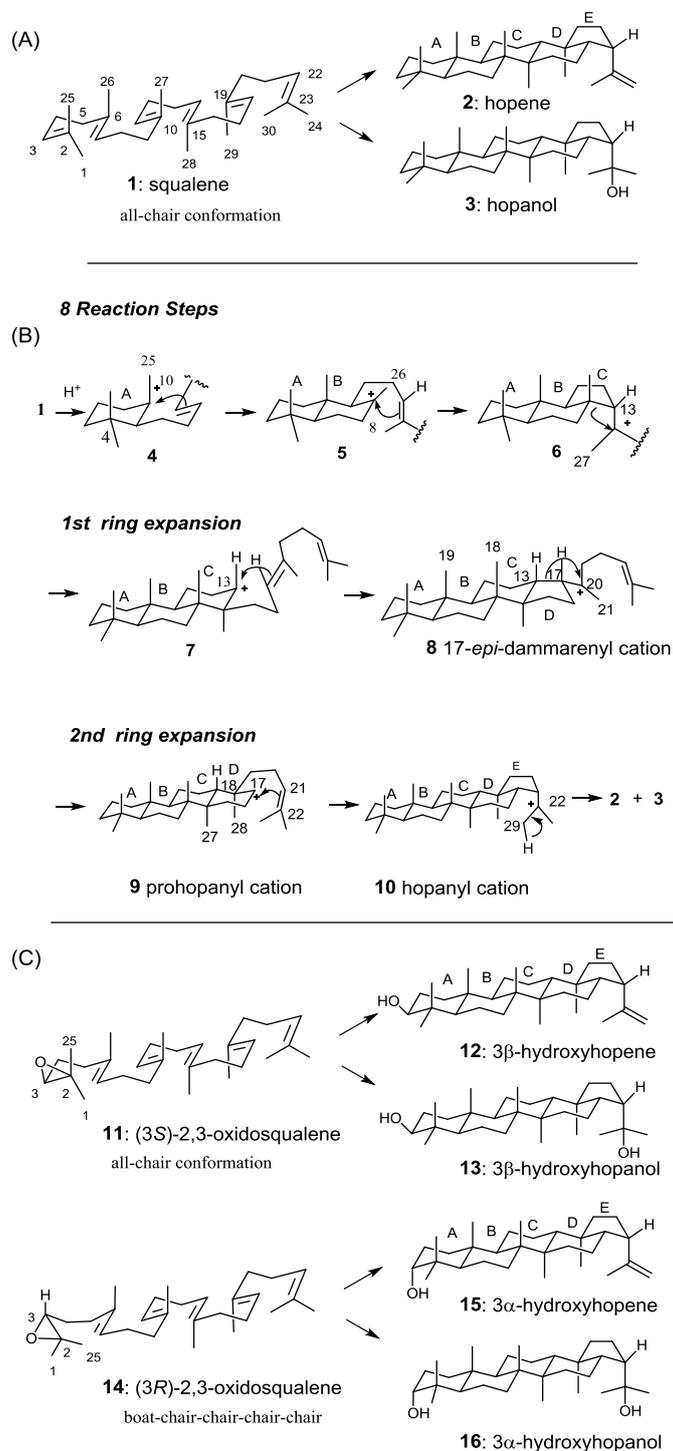
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Figure, Scheme, and Table Legends



Scheme 1. (A) Polycyclization pathways of squalene (**1**) into hopene (**2**) and hopanol (**3**) by squalene-hopene cyclase. Squalene **1** folds in an all-prechair conformation in the

1 reaction cavity. (B) The cyclization consists of 8 reaction steps. (C) Polycyclization
2 reactions of (3*S*)-2,3-oxidosqualene (**11**) and (3*R*)-2,3-oxidosqualene (**14**). Here, **11**
3 folds into an all-prechair structure similar to that in the polycyclization reaction of **1**,
4 leading to 3 β -hydroxyhopene (**12**) and 3 β -hydroxyhopanol (**13**). However, **14** folds into
5 a boat-chair-chair-chair-chair conformation, resulting in the production of 3 α -
6 hydroxyhopene (**15**) and 3 α -hydroxyhopanol (**16**).
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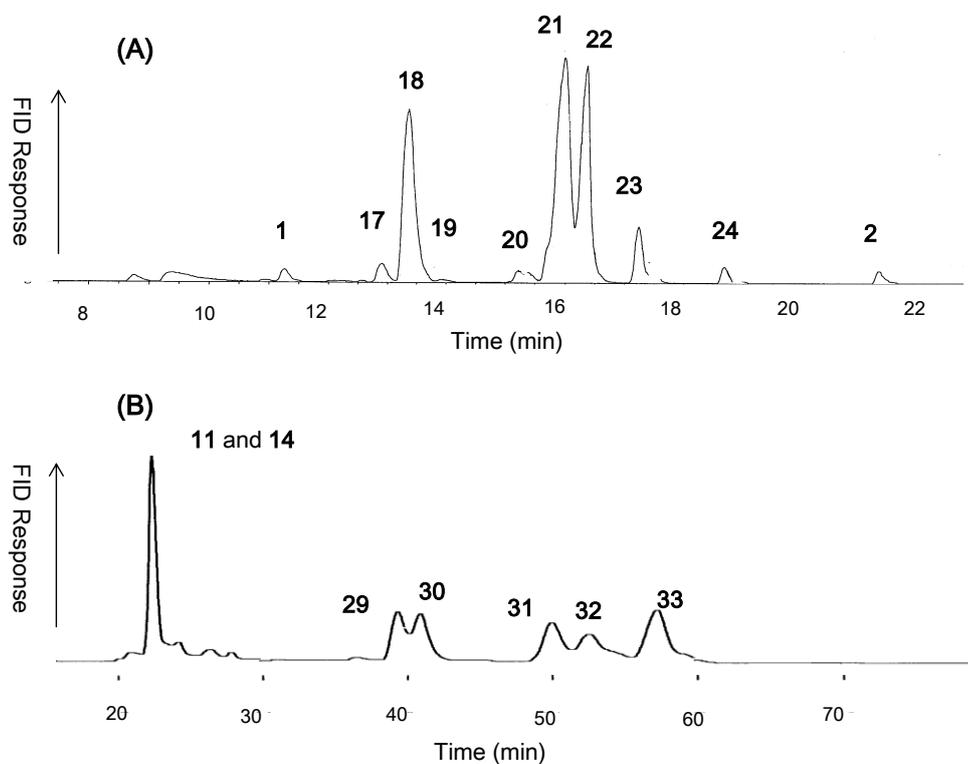


Fig. 1. (A) Incubation mixture of squalene (**1**) with the A306V variant. (B) Incubation mixture of 2,3-oxidosqualene (a mixture of **11** and **14**) with the A306V variant. The cell-free extracts (5 mL) were prepared from a 100 mL culture of the A306V variant. Incubation of 1.0 mg of **1** and that of a mixture of **11** and **14** (1.0 mg) with 2.0 mL of cell-free homogenates was carried out at 45°C for 24 h. The lipophilic materials were extracted with hexane. Triton X-100 present in the hexane extract was removed by passage through a short SiO₂ column using hexane/EtOAc (100:20) as eluent. The lipophilic materials thus collected were subjected to GC measurements. The GC conditions are described in the Experimental section.

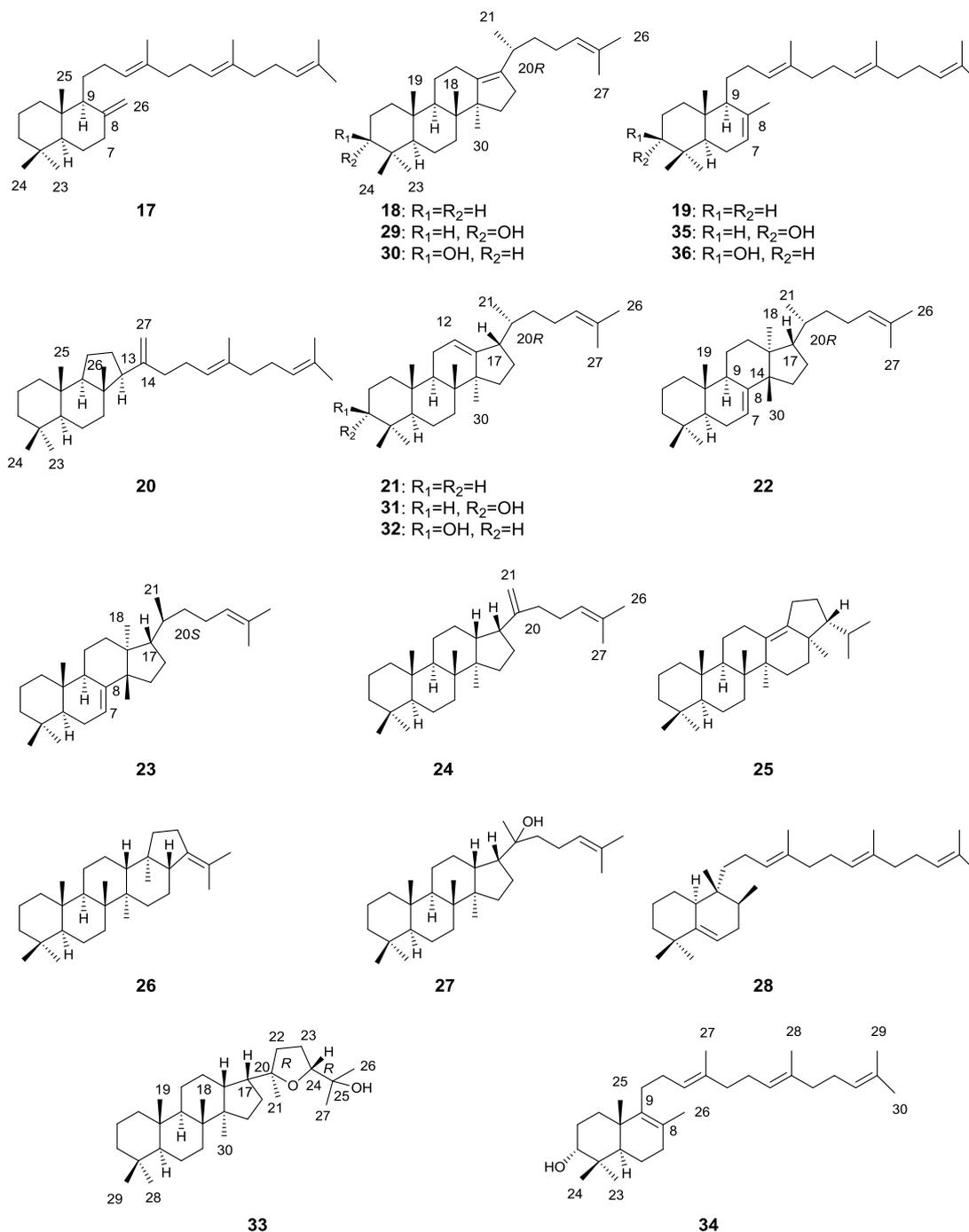
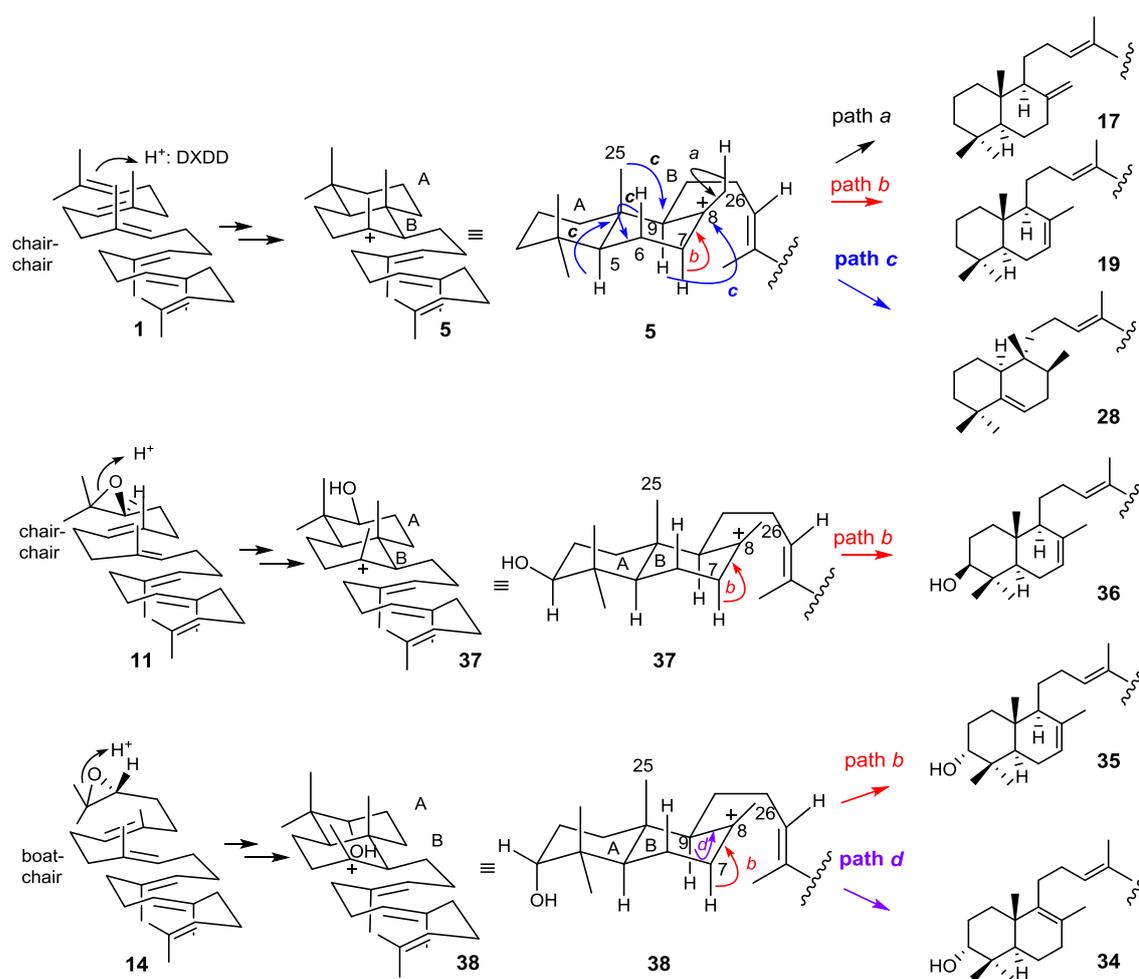


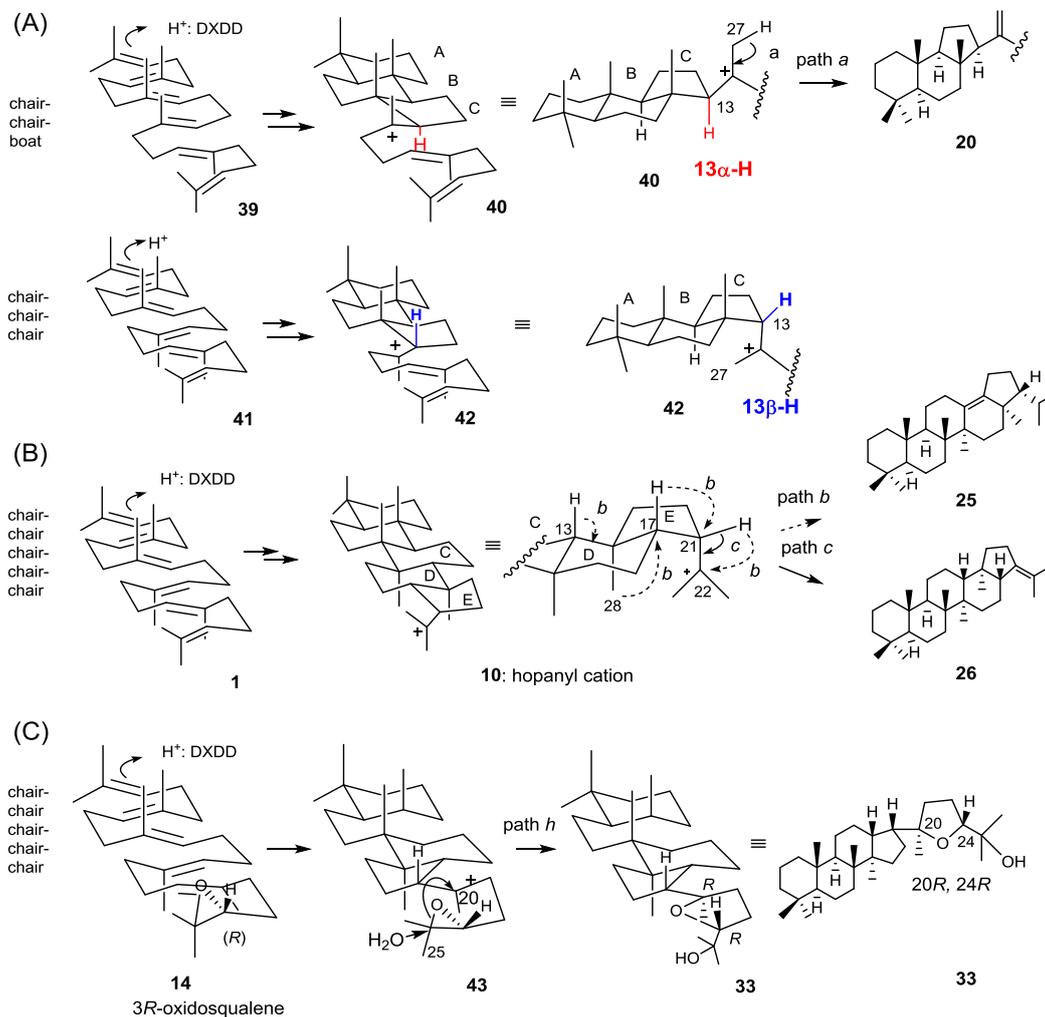
Fig. 2. Structures of the enzymatic products produced by A306X variants. Products **17-24** were produced from the incubation of **1** with the A306V mutant. Products **18, 21-27**

1 were generated by the incubation of **1** with the A306T variant (Fig. S3.1). Products **17**,
2 **19**, and **28** were produced by the reaction of **1** with the A306F mutant (Fig. S.1.2). Fig.
3 S4 shows the GC profiles of the incubation mixtures of a racemic mixture of **11** and **14**
4 with the A360T and A306F variants. Products **29–33** were produced by the A306T
5 mutant. Products **34–36** were produced by the A306F mutant.

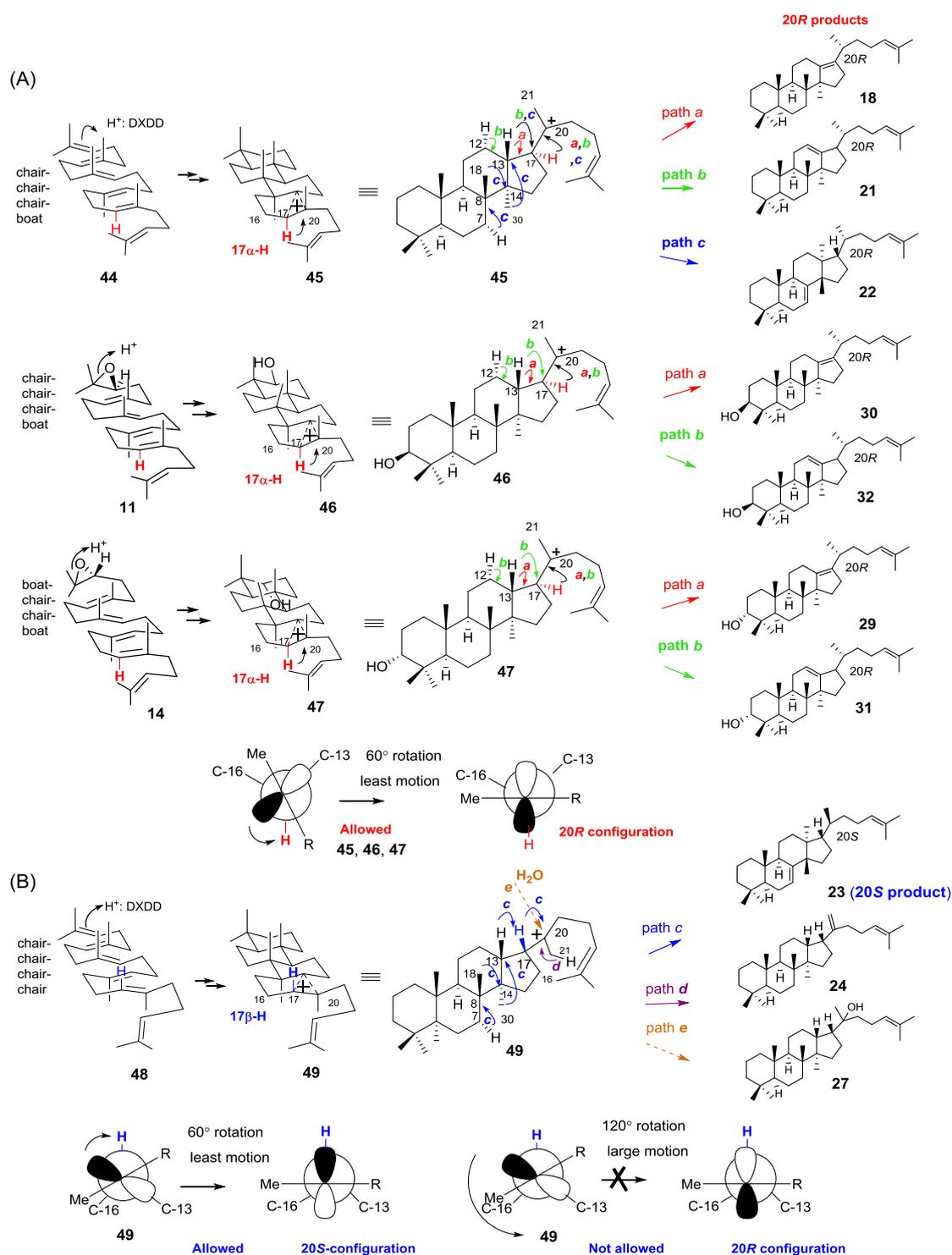
6 Compound names. **17**: α -polypodateraene, **18**: dammara-13(17)-24-diene, **19**: γ -
7 polypodateraene, **20**: (17*E*)-(13 α H)-malabarica-14(27),17,21-triene, **21**: 20*R*-dammara-
8 12(13), 24-diene, **22**: 20*R*-eupha-7(8),24-diene, **23**: 20*S*-tirucalla-7(8),24-diene (a new
9 SHC-mediated product), **24**: 17-*epi*-dammara-20(21),24-diene, **25**: neohop-13(18)-ene,
10 **26**: hop-21(22)-ene, **27**: 20-hydroxy-17-*epi*-dammaraene, **28**: neopolypoda-
11 5(6),13,17,21-teraene (rearranged skeleton of bicyclic polypodatetraene), **29**: 20*R*-
12 dammara-13(17),24-dien-3 α -ol, **30**: 20*R*-dammara-13(17),24-dien-3 β -ol, **31**: 20*R*-
13 dammara-12(13),24-dien-3 α -ol, **32**: 20*R*-dammara-12(13),24-dien-3 β -ol, **33**: (20*R*,
14 24*R*)-17-*epi*-epoxydammarane, **34**: polypoda-8(9),13,17,21-tetraen-3 α -ol, **35**:
15 polypoda-7(8),13,17,21-tetraen-3 α -ol (3 α -hydroxy- γ -polypodatetraene), **37**: polypoda-
16 7(8),13,17,21-tetraen-3 β -ol (3 β -hydroxy- γ -polypodatetraene).
17



Scheme 2. Cyclization mechanisms of **1**, **11**, and **14** leading to bicyclic products **17**, **19**, **28**, and **34–36**. Substrates **1** and **11** fold in a chair-chair conformation, but substrate **14** folds in a boat-chair structure.



Scheme 3. Cyclization mechanisms of **1** and **14** leading to tricyclic product **20** and pentacyclic products **25**, **26**, and **33**. (A) A chair-chair-boat conformation yielded the tricyclic **20**. (B) The all-chair conformation furnished the neohopane scaffold **25** and hopane skeleton **26**. (C) ($3R$)-oxidosqualene **14** was folded into an all-chair conformation to yield $20R, 24R$ -17-*epi*-epoxydammarane **33**. Notably, cyclization started from the isopropylidene moiety rather than from the epoxide ring.



Scheme 4. Formation mechanisms of 6,6,6,5-fused tetracyclic products with 20R and 20S configurations. Cation **8** is formed by the folding of **1** into a chair-chair-chair-chair

1 conformation (Scheme 1). This indicates that the native SHC furnishes the tetracyclic
2 product(s) with the 20S configuration, in contrast to the A306V mutant.

3

4 **Table 1.** Product distribution ratio (%) obtained from the reactions of squalene (**1**) with
5 A306X variants. The reported van der Waals volumes (nm³) are as follows: Gly,
6 0.00279; Ala, 0.05702; Thr, 0.19341; Val, 0.25674; Leu, 0.37876; Phe, 0.55298,^[22] thus,
7 the steric bulk increases as follows: Gly<Ala<Thr<Val<Leu<Phe.

8

Compounds		Normal products		Bicycle			Tricycle	Tetracycle						Pentacycle		
		1	2	3	17	19	28	20	18	21	22	23	24	27	25	26
Wild type SHC	–	84	16	–	–	–	–	–	–	–	–	–	–	–	–	–
A306G	43	45	7	–	–	–	–	–	–	–	–	–	–	4	2	
A306T	2	23	2	–	–	–	–	11	13	12	8	9	4	1	15	
A306V	2	2	–	3	1	–	2	22	31	27	8	2	–	–	–	
A306L	20	1	–	6	31	–	2	21	13	3	–	3	–	–	–	
A306F	56	–	–	1	42	1	–	–	–	–	–	–	–	–	–	

9 –: little or no production, below the limit of GC detection. There might be other
10 products that have not been characterized, but the production amounts are a little, if any
11 present.

12

13

14 **Table 2.** Product distribution ratios (%) generated by incubating oxidosqualene (**11** and
15 **14**) with A306X mutants. The steric bulk increases as follows: Ala<Thr<Val<Phe.^[22]

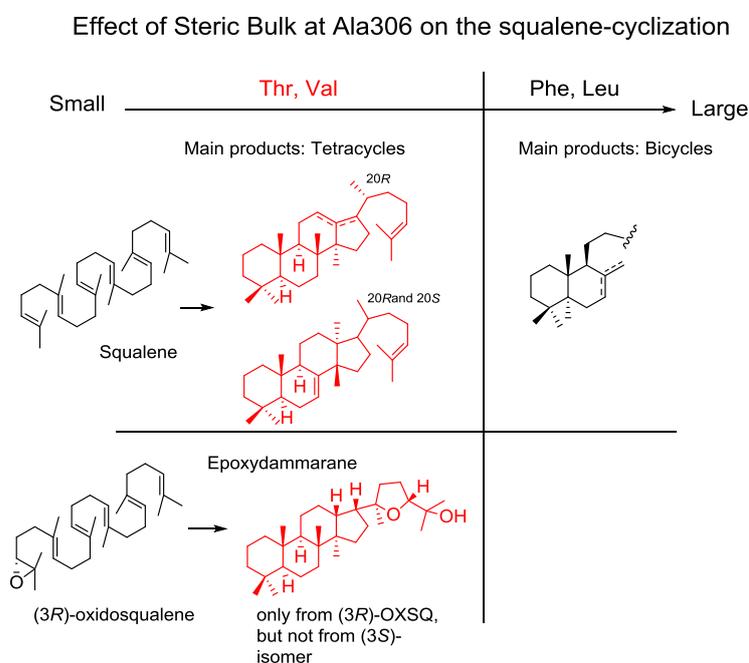
16

Compounds		Normal products				Bicycle			Tetracycle				Epoxy-dammarane
		11 & 14	12	13	15	16	34	35	36	29	30	31	32
Wild type SHC	–	54.2	9.5	31.3	4.9	–	–	–	–	–	–	–	–
A306T	2	4	–	4	–	–	–	–	24	26	18	14	8
A306V	33	–	–	–	–	–	–	–	15	13	11	8	20
A306F	42	–	–	–	–	8	12	38	–	–	–	–	–

17

18 –: negligible or trace amount

TOC graphic



The Ala306 in *A. acidocaldarius* SHC is widely conserved in known SHCs. Increases in steric bulk (A306T and A306V) led to the accumulation of 6,6,6,5-fused tetracyclic scaffolds possessing 20*R* stereochemistry in high yield (94% for A306V). The production of the 20*R* configuration indicates that squalene had been folded in a chair-chair-boat conformation, indicating that the bulk at position 306 significantly affects the stereochemical fate during the polycyclization reaction. Intriguingly, epoxydammarane was generated from (3*R*)-oxidosqualene.