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Computational methods and points for attention in absolute configuration determination

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With the rapid development of high performance computers and computational methods, including software, an increasing number of experimental chemists have tried to use computational methods such as optical rotation (OR, including the matrix model), optical rotatory dispersion (ORD), electronic circular dichroism (ECD or CD), vibrational circular dichroism (VCD), and magnetic shielding constants—nuclear magnetic resonance (NMR)—to explain and/or assign absolute configuration (AC) for various compounds. Such reports in the field of natural products have increased dramatically. However, every method has its range of application. This leads, in some cases, to incorrect conclusions by researchers who are not familiar with these methods. In this review, we provide experimental chemists and researchers with more computational details and suitable suggestions, and especially hope that this experience may help readers avoid computational pitfalls. Finally, we discuss the use of simplified models to replace original complex structures with a long side chain. The fundamental basis for using models to represent complex chiral compounds, such as in OR calculations, is the existence of conformation pairs with near canceling conformer contributions that justify the use of models rather than the original compounds. Using examples, we here introduce the transition state (TS) calculation, which may benefit readers in this area for use and mastery for their AC study. This review will summarize the general concepts involved in the study of AC determinations.

KEYWORDS

optical rotation, matrix model, electronic circular dichroism, vibrational circular dichroism, carbon NMR, conformer pair, simplified model, transition state

1 Introduction

With the rapid development of high performance computers and computational methods, including software, an increasing number of experimental chemists have tried to use computational methods such as optical rotation (OR, including optical rotatory dispersion (ORD)), electronic circular dichroism (ECD)—also known as CD before vibrational circular dichroism (VCD) was widely used—and magnetic shielding constants—nuclear magnetic resonance (NMR)—to explain and/or assign absolute or relative configurations for various compounds and polymers (fibers) (Kurouki et al., 2014; Marty et al., 2014; Li et al., 2016a; Fernández et al., 2019), including metallic complex (Merten et al., 2014; Li et al., 2019a). Reports of such work in bioactive compounds and natural products chemistry has dramatically increased (Sherer et al., 2014a; Sherer et al., 2014b; Yang et al., 2018), while traditional methods, such as Mosher ester (Hoye et al., 2007; Zheng et al., 2019) and X-rays (Wang et al., 2022)—in which Cu-K α (1.5418 Å) radiation is preferred to Mo-K α (0.7107 Å) radiation, due to its stronger anomalous scattering effect (Bijvoet et al., 1951)—are also widely used in absolute

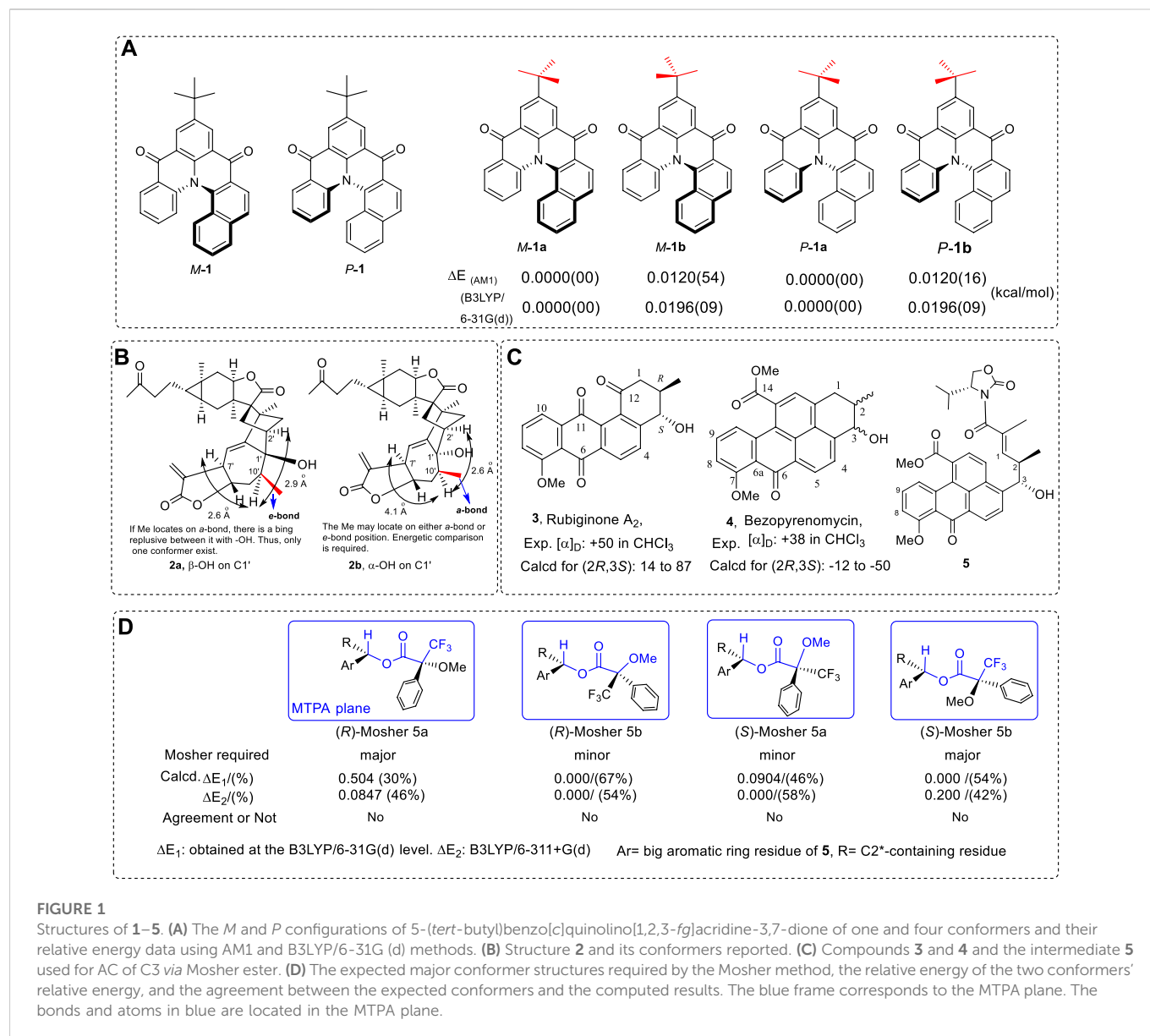


FIGURE 1

Structures of **1–5**. (A) The *M* and *P* configurations of 5-(*tert*-butyl)benzo[*c*]quinolino[1,2,3-*fg*]acridine-3,7-dione of one and four conformers and their relative energy data using AM1 and B3LYP/6-31G (d) methods. (B) Structure **2** and its conformers reported. (C) Compounds **3** and **4** and the intermediate **5** used for AC of C3 via Mosher ester. (D) The expected major conformer structures required by the Mosher method, the relative energy of the two conformers' relative energy, and the agreement between the expected conformers and the computed results. The blue frame corresponds to the MTPA plane. The bonds and atoms in blue are located in the MTPA plane.

configuration (AC) studies. Natural products have also been called the “engines of its development and ... links to the domain of biology” in organic chemistry (E. J. Corey, cited in Barton et al., 1999). However, every method has its range of use (María-Magdalena and Jorge, 2015). This can lead to incorrect conclusions by some researchers who are not familiar with such methods. On the other hand, TS calculations are often used in organic reaction mechanism studies. They have been applied in AC studies in order to understand the AC of final reaction product results. This review also discusses the use of simplified models to replace original complex structures in calculations. The fundamental basis for using models to represent complex chiral compounds, such as in OR calculations, is the existence of conformation pairs that can be investigated rather than the original compounds. It is important and necessary to provide experimental chemists and researchers with more computational details and suitable suggestions, especially to avoid pitfalls in their computations.

2 Importance of conformational search

The first, most important, step is to find the lowest energy conformation for a specific chiral molecule. For example, a 2-butanol should have four stable conformations at different energy that can be found manually. The most stable conformation has a zigzag carbon chain. However, the maximum conformation number is proportional to the bond numbers (*n*) that can rotate freely (3^{n-2} , $n = 3, 4, \dots$). Thus, with the increase in free rotation bonds, the possible conformations increase quickly. Therefore, it is necessary to have software which uses various force fields to investigate all possible conformations and list these geometries according to their relative energy.

Much commercially available software is frequently used, such as ComputeVOA (ComputeVOA, 2022), GaussView16 (GaussView6, 2022; Schrodinger LLC, 2022; RDKit, 2022), and Merck molecular force field (MMFF94) and its developed MMFF94S (Halgren, 1996; Halgren, 1999). Other force fields or methods can also be used.

Each piece of software has its advantages and limitations. Thus, it is suggested that researchers use at least two software packages in conformational searches. This can ensure that the lowest energy conformations can be found. If one or two important conformations are missed in a procedure, the ^{13}C NMR, OR, ORD, ECD, or VCD simulations may be affected. If the missed conformation is too different from the other conformation's physical data, this missed conformation may lead to significant error in the simulated ^{13}C NMR, OR, ORD, ECD, or VCD. Comparing this simulated data with the experimental results may lead to an incorrect conclusion.

Although everyone knows the importance of computations of an OR, ECD, and VCD in AC study, the errors can also appear in different situations. To avoid such mistakes, it is important to list reasons for the errors and the methods that can prevent them. As a very typical example, 5-(*tert*-butyl)benzo[*c*]quinolino[1,2,3-*fg*]acridine-3,7-dione (Figure 1A) was used as one example of conformational analysis.

It has been reported that the relative energy of *M*-1 is more stable than that of *P*-1 by about 0.02 kcal/mol (Pandith et al., 2011). This is an incredible result since the structures are enantiomers. Theoretically, they should have the same energy. The reason for reaching this result must be a mistake in the conformational analysis.

In this example, the key substituent is the *t*-butyl group in calculations. The tertiary butyl group should have two stable conformations in the *M* and *P* structures, as illustrated below (Figure 1A). For *M*-1, the geometry *M*-1a had lower energy of 0.0196 kcal/mol than that of *M*-1b at the B3LYP/6-31G(d) level in the gas phase (Zhu, 2015). Once the total mirror-image geometries *M*-1a and *P*-1a were used in calculations, they both exhibited almost the same energy, the difference between the two geometries being less than 10^{-6} Kcal/mol in calculation. To conclude that *M*-1 has lower energy of 0.012 Kcal/mol than *P*-1 must have been due to incorrect use of the geometries of *M*-1a and *P*-1b (different conformers) from the energy calculations.

In OR, ECD, or VCD calculations, such a small energy difference conformer may be ignored in simulations. However, if it is said that this pair of enantiomers has different energies, then this is an absolutely wrong conclusion. It must strongly indicate that the origin of the chirality does not require any other force, such as chiral catalyst, or other power to form a small excessive enantiomer (%ee) in the prebiotic environment (pool). This is because the 0.012 kcal/mol energy difference is a definite large effect which may bring the %ee value up to of 0.49%ee for the enantiomers. Indeed, the two enantiomers must have the same energy in nature. Professional software is helpful for complex molecular conformational study; however, it is not sufficient to completely find all conformations with relatively low energy.

Another similar example is helicene structures, which consist of non-planar polycyclic aromatic hydrocarbons that form an *ortho*-condensed aromatic ring. Many helicenes exhibit interesting characteristics, such as their use as organic light-emitting diodes (Riobé et al., 2015; Joly et al., 2016). They also exhibit circularly polarized luminescence (CPL) activities (Goto et al., 2012; Hellou et al., 2017). Thus, this is an important kind of intermediate or chiral material.

The following Example 2 contains a seven-membered ring which may have different geometries (Figure 1B) (Xu et al., 2016a). To assign the configuration of C1', NOE signals between the H7' and H10' via H10' with H2' were used. As illustrated in Figure 1B, two conformers

were reported. Since the distance of H7' to H10' in conformer 2a (β -OH) is shorter than that of H10' to H2' in structure 2b (α -OH), the AC of C1' was assigned as β -OH. This is a good example of using geometries to support an experimental conclusion. This compound has a seven-membered ring. The Me on C10' may orientate on an *e*-bond or *a*-bond. If the energetics of these conformers were investigated and reported, the conclusion should be clearer.

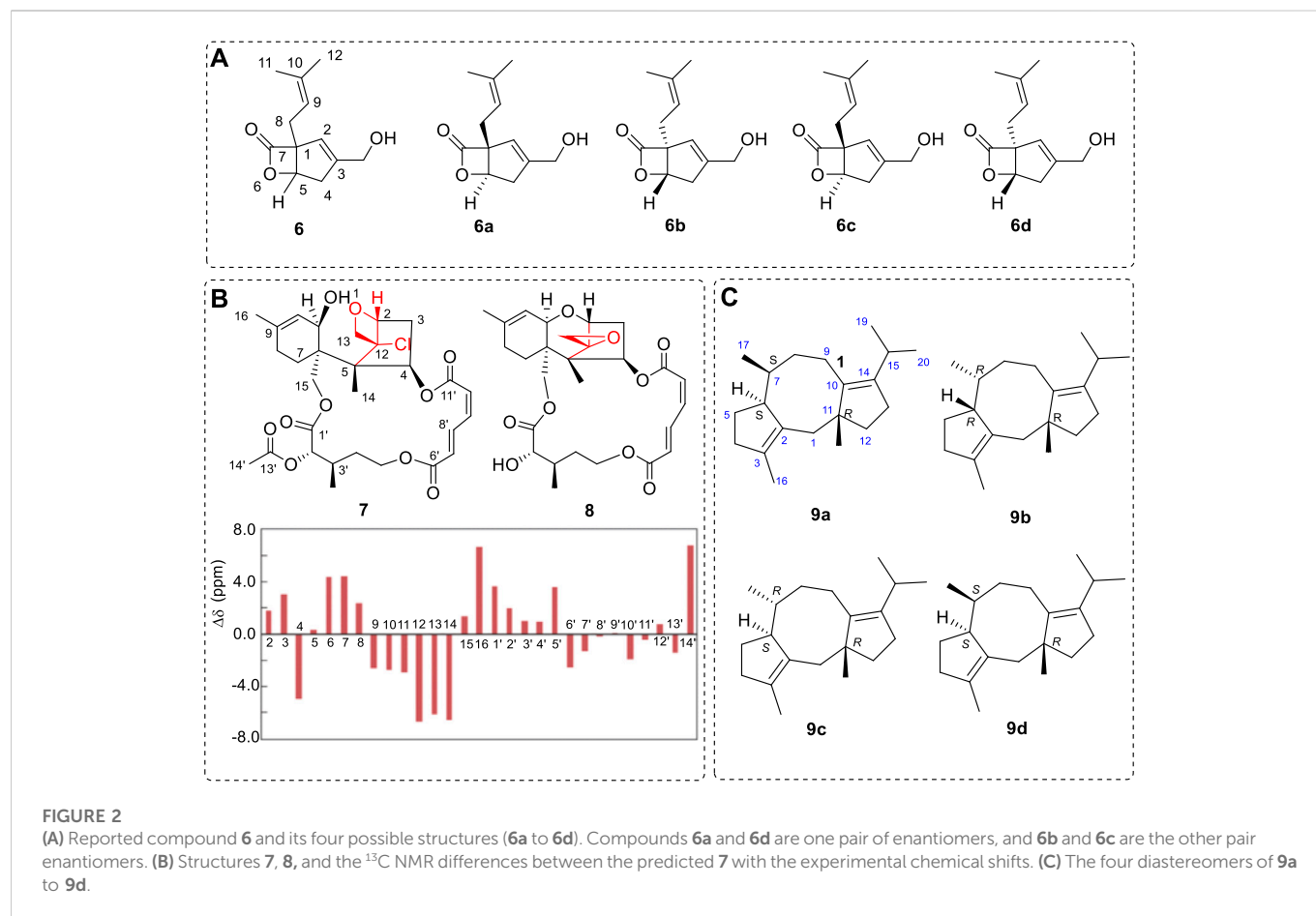
Historically, both rubiginone A₂ (3) and benzopyrenomycin (4) have very close OR values, and it was suggested that both have the same AC on C2 and C3 (Figure 1C) (Pezzuto et al., 1976). Benzopyrenomycin (4) was recently synthesized (Oka et al., 1990; Huang et al., 2008). By determining the AC of C3 in intermediate 5 using Mosher methods (Ancheeva et al., 2017), researchers thought that 4 should have (2*R*,3*S*) configurations. It seems that the AC of (2*R*,3*S*)-4 should be correct. However, it is incorrect, based on the computed OR value and sign. Why does the Mosher method used here give the wrong conclusion?

The reason involves which conformer had the major population in solution. As an empirical method, the Mosher tool is normally valid, and most compounds have the correct conformation distribution in solution. For this specific case 5 (Figure 1C), its conformational searches were performed using three pieces of software and two more MM methods. Thousands of geometries were found for the Mosher ester of 5 using MMFF94 and AMBER force fields and followed by DFT methods (Figure 1D) in order to find the lowest and the second-lowest conformers. The expected major conformers were (*R*)-Mosher 5a and (*S*)-Mosher 5b according to the Mosher methods required. However, the major conformers were (*R*)-Mosher 5b and (*S*)-Mosher 5a by referring to the MTPA plane. The specific energy data and distributions are listed in Figure 1D. The major conformers that Mosher methods require are not those predicted by computational results. This caused the errors in the use of Mosher methods.

In the example, it is an additionally surprising result that the Mosher method requires that the substituent near the stereogenic center not be a large group, such as the substituent of IMI. Indeed, if the substituent is extremely rigid and close to a linear structure, such as two or two more $-\text{C}\equiv\text{C}-$ connected directly, one would need to be careful to use the Mosher method to assign their ACs since this group may change the conformer distribution of the expected major or minor conformers that Mosher methods require in solution.

The three examples are given at the beginning of this review to show the importance of the conformational search not only in AC study but also in other research areas. Indeed, it is very important for chemists in structural study (Perez-Mellor et al., 2019; Wang and Zhu, 2021; Zhu, 2022a), and it is also hard to find some conformers of rigid chiral compounds using software. The reason why some software cannot find the corresponding conformers is very complex. In some cases, some compounds have slightly different conformations exhibiting computed spectra that are almost mirror image (Bringmann et al., 2009; Zajac et al., 2015). Thus, obtaining the conformers with low energy and the correct geometries are very important. We recommend using multiple software to perform the same conformational search for a molecule in order to reduce the possibility of a wrong conclusion. Developing powerful conformational search software is still a big challenge for computational chemists.

Finally, the concept of "most stable conformation" must be mentioned here. There are many stable conformers in a solution.



However, only one most-stable conformer exists in solution instead of multiple most-stable conformers—followed by a second most-stable and third most-stable conformation. However, some authors mention “the most stable conformations” during OR, ECD, or VCD calculations. This is in no way acceptable. If it is a racemic chiral compound in solution, then there are only two most-stable enantiomers. As an example, a general statement is listed here: 38 conformations were found with low energy from 0–3.5 kcal/mol using a MMFF94 force field. These geometries were then used in further optimizations at the B3LYP/6–31+G(d) level (in the gas phase), obtaining one most-stable conformer. To posit “the most stable conformations” is wrong; however, this statement is frequently found in reports.

3 ^{13}C NMR method

NMR calculations involve the position of all atoms in a space and the chemical environment. Many atoms' chemical shifts can be computed, such as ^1H (Wang et al., 2003), ^3He (Ramalho and Buehl, 2005), and ^{15}N (ComputeVOA, 2022). The study of ^{13}C NMR is more attractive for organic chemists—the number methods developed for ^{13}C NMR computations is more than the number of NMR methods for other atoms. The gauge including the atomic orbitals (GIAO) model has been used widely.

Firstly, it must be clear that ^{13}C NMR can be used to assign the relative configuration of a chiral molecule having two or two more

stereogenic centers. If one chiral compound has a known stereogenic center, it is possible to use ^{13}C NMR to assign its AC by comparing the other stereogenic center(s) with the known center. These computed ^{13}C NMR can be compared with the experimental data to assign the configurations (Lauro and Bifulco, 2020). The use of ^{13}C NMR to study the configuration of chiral organic compounds is effective in stereochemistry study (Bifulco et al., 2007; Lodewyk et al., 2012; Gussem et al., 2014; Lauro and Bifulco, 2020).

The maximum principle must be applied to the examination of structures. If the coefficients between two structures are small enough, they cannot be used as evidence for judging structural evaluation. In some cases, more evidence is needed for further confirmation of structures.

The originally computed ^{13}C NMR data may require treatment for further uses (Forsyth and Sebag, 1997; Barone et al., 2002; Liu et al., 2006; Yang et al., 2011; Cao et al., 2020a; Li et al., 2022). For example, after linear corrections (Forsyth and Sebag, 1997), there are two methods for judging whether the structure is correct. The first is to use to maximum of $\Delta\delta$ values. If the maximum of $\Delta\delta$ is over 8.0 ppm, the structure is not reliable; if it is less than 8.0 ppm, then the structure is located in the reliable structure range. However, more evidence may be required for further confirmation of its configuration, such as comparing coefficients of two structures: the bigger the coefficient, the more reliable the structure will be (Yang et al., 2011).

For example, chiral compound **6** was originally obtained and assigned as either **6a** or **6b** since no interaction signal was obtained between the protons H5 and H8 in ROESY spectra (Figure 2A) (Liu

et al., 2006). The ^{13}C NMR was investigated. After conformational searches were performed using the MMFF94 force field, all six conformations were used for the Boltzmann sum.

After conformational searches were performed using a MMFF94 force field and optimizations for the geometries using DFT methods, six conformations were used in ^{13}C NMR calculations. After linear corrections (Forsyth and Sebag, 1997), the coefficients were 0.9947 and 0.9940 (Liu et al., 2006). Other methods could be used to assign the AC (Cao et al., 2020a; Li et al., 2022). It appears that **6a** should be the preferred structure. However, the maximum of the chemical shift errors between C2, C8, and the experiments were -9.1 -10.2 , and -11.2 -11.4 ppm in **6a** and **6b**, respectively. Thus, both **6a** and **6b** are not the correct structures. The other two *cis*-structures (**6c** and **6d**) have bigger coefficients (0.99932 and 0.99935). All the shift errors were smaller than 8.0 ppm. In this case, if it is regarded as **6d** (coefficient was 0.99935), it is still not correct. The OR was then computed. The experimental OR was -135 , and the predicted OR for **6c** was -127° at the B3LYP/6-311++G(2d,p) level; compound **6d** was $+127^\circ$. Therefore, the correct structure is **6c** after its enantiomer **6b** was excluded at the beginning. The structure **6c** was synthesized and the synthetic **6c** had the OR value of -129° (Zhou and Snider, 2008a; Zhou and Snider, 2008b).

Other methods are used in the assignment of the chiral compounds. For example, the mean absolute error (MAE), corrected mean absolute error (CMAE), and the root-mean square deviation (RMSD) (Smith and Goodman, 2010; Grimblat et al., 2015) were applied for the study, including DP4 and DP4+. The concepts used here, for the *i*th nuclei, has a chemical shift error of $\Delta\delta_i = \delta_{i\text{ calcd}} - \delta_{i\text{ exp}}$. $\text{MAE} = \sum(\Delta\delta_i)/n$. $\text{CMAE} = \sum(\Delta\delta_{i\text{ scaled}})/n$. Here, *n* is the total number of nuclei, and $\delta_{i\text{ scaled}}$ can be computed using a linear fit of the calculated ($\delta_{i\text{ calcd}}$) versus experimental ($\delta_{i\text{ exp}}$) chemical shifts for calculating the related $\Delta\delta_{i\text{ scaled}}$ values. For more details, see Baassou et al. (1983), Li et al. (2022), and Cao et al. (2020a), who also compared the experimental shifts with the predicted values in the study. Consequently, comparing the computed ^{13}C NMR is quite valid for assigning chiral compounds' relative configurations, including some AC assignments (Lauro and Bifulco, 2020).

If one compound's configuration is known, it can be used to determine the AC of another moiety's structures for a compound. This is more conveniently used for organic synthetic compound study. For example, compound **7** is derived from natural product **8** by reaction with 2-chloroacetyl chloride (Jia et al., 2018). This three-membered oxirane structure disappeared and a new four-membered oxetane moiety formed while the six-membered ring in **8** decomposed in the reaction. Thus, it had a big difference in ^{13}C NMR from those of **8**. This example is a strong hint that, during the modification of a natural product, even the reaction condition is mild and the reagent may be frequently used, such as where example **8** was reacted with 2-Cl-acetyl chloride, leading the three-membered-ring opened (red moiety) to **7**. Use of the ^{13}C NMR can easily identify the structure of **7**. In this case, the whole aliphatic carbon chemical shift difference is less than 8.0 ppm, within a reasonable error range (Figure 2B).

After their development (Goodman Group, 2017), of DP4 methods have been applied to the AC determination of many chiral compounds by computing the possibility of assigning their structures, such as fusicoca-2,10(14)diene (**9**) (Merten et al., 2017). It has three other diastereomers besides **9a** (6S,7S,11R): **9b** (6R,7R,11R), **9c** (6S,7R,11R), and **9d** (6R,7S,11R) (Figure 2C).

After comprehensive analysis, diastereomer **9a** is preferred. As the authors indicated, "The DP4 analysis should not be used as a black-box method by simply evaluating the calculated and experimental ^{13}C NMR shifts in the order of appearance (e.g., lowest calculated with lowest experimental shift, and so on). In case of FCdiene, neglect of a correct connectivity-based assignment leads to different results for both methods, as shown in Merten et al. (2017): Using the unassigned ^{13}C chemical shifts in a DP4 analysis, diastereomer **9b** becomes favored over diastereomer **9a**, with a probability of 94.9%. It is therefore important to point out the importance to include the connectivity information obtained from 2D NMR spectroscopy in the assignment of the experimental to the calculated ^{13}C NMR shifts in order to ensure a reliable, unambiguous assignment of a relative configuration".

In addition, comparing the calculated ^{13}C NMR with the experimental results determines that the organic compounds are not only used in AC but also applied for planar structure determinations since ^{13}C NMR calculations involve the position of all atoms in space and the chemical environment. In some cases, it can be used to determine the *cis* or *trans* structure for a compound with C=C bonds. For example, cyclopeptide anthranilic acid (**10**) was assigned as a *cis* structure by comparing its ^{13}C NMR with the calculated data (Figure 3A) (Zheng et al., 2009). In the first case (*Z*), all of the maximum relative errors of chemical shift were below 8.0 ppm. However, in the second case (*E*-), the maximum errors at C-21 were 8.9 ppm. Thus, it was assigned as (*Z*)-**10a**. As another method, ECD, which will be referred to later, it is also a good way to assign the *E/Z* structures (Rode and Frelek, 2017).

By comparing the relative chemical shifts of ^1H and/or ^{13}C NMR, valid information in the chiral compound's configuration assignments can be provided. An example is zosteraphenol A (**11**) from the seagrass *Zostera marina* (Grauso et al., 2020). One minor conformer (**11b**) and major (**11a**) rotamer were at equilibrium in solution due to the repulsion between the -MeO in ring A and the -OH in ring B, the rotation of the single bond C2-C2' being restricted. This barrier is big enough at room temperature to block the single-bond rotation and leads the exchange signals at 4.65, 4.80 and 5.20 ppm to appear as a broad peak. Once the temperature decreased to 258 K (-15°C), the signal became very sharp (Yu et al., 2015). After computing its rotamers, ^{13}C NMR at the chemical shifts were subsequently calculated at the PBE0/6-311+G(2d,p) level using B3LYP/6-31+G(d,p)-optimized geometries *via* the PCM model in chloroform (Figure 3B). The computational results supported the hypothesized conformational equilibrium for **11**.

Because the shift of nuclei is sensitive to their position in the molecule, we can compare the differences of the same nuclei in different diastereoisomers. This is very useful when some epimers were obtained. In this case, structural assignments could be made by comparing the experimental shift difference between two epimers with the experimental shift differences between the two diastereoisomers. For example, glaucumolides B (**12**) and bistrochelides A (**13**) were a pair of structurally similar epimers (Figure 3C) (Sun et al., 2019). MAEs between the investigated diastereoisomers were used to assign the relative configuration of C11 by comparing the error pattern and referring the X-ray crystalline of key compound **14**. The AC of glaucumolide B (**12**) was assigned as (1S,2R,3R,4R,5S,21S,26S,27R) in the assistance of ECD spectra comparison of experimental and theoretical results.

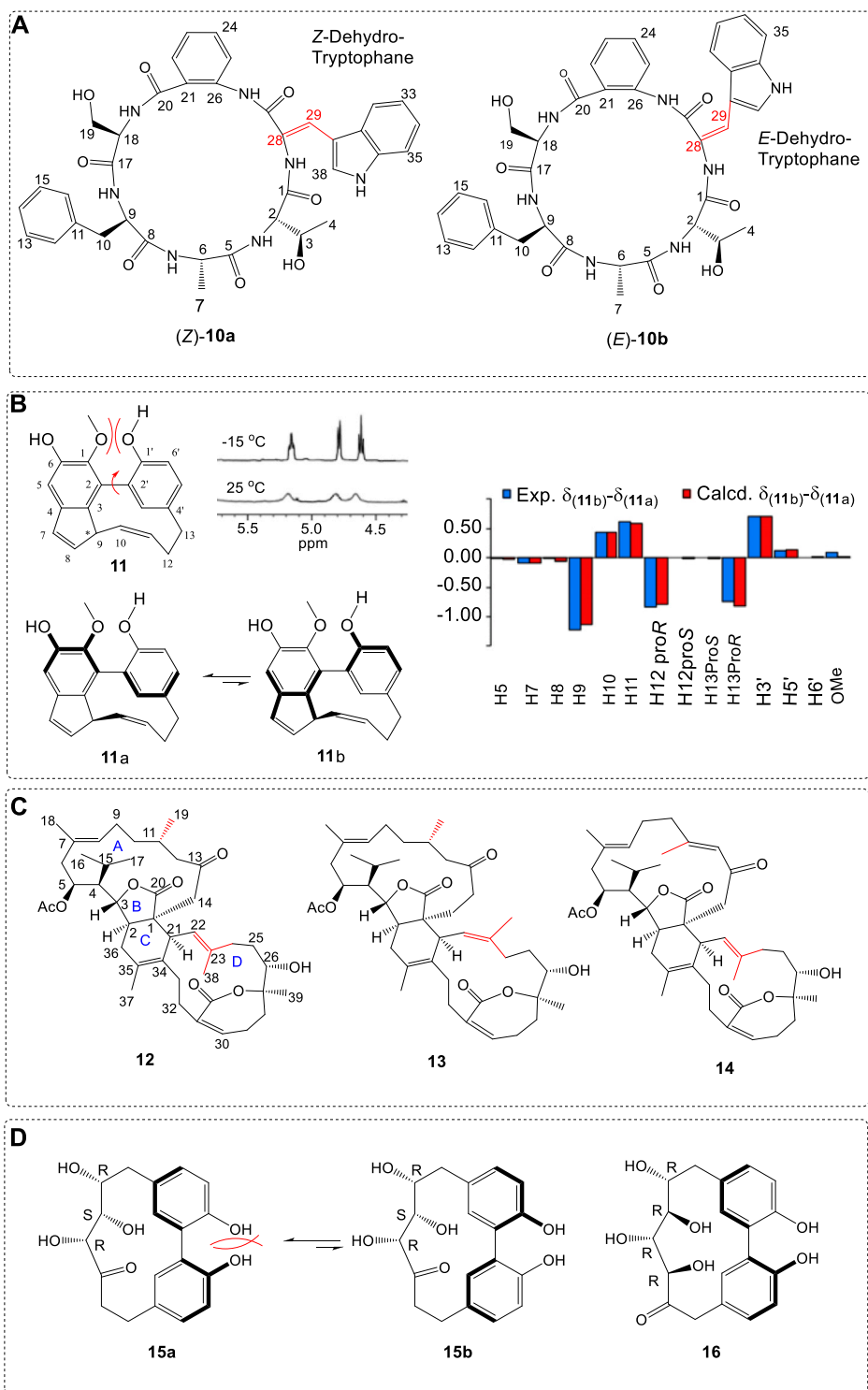
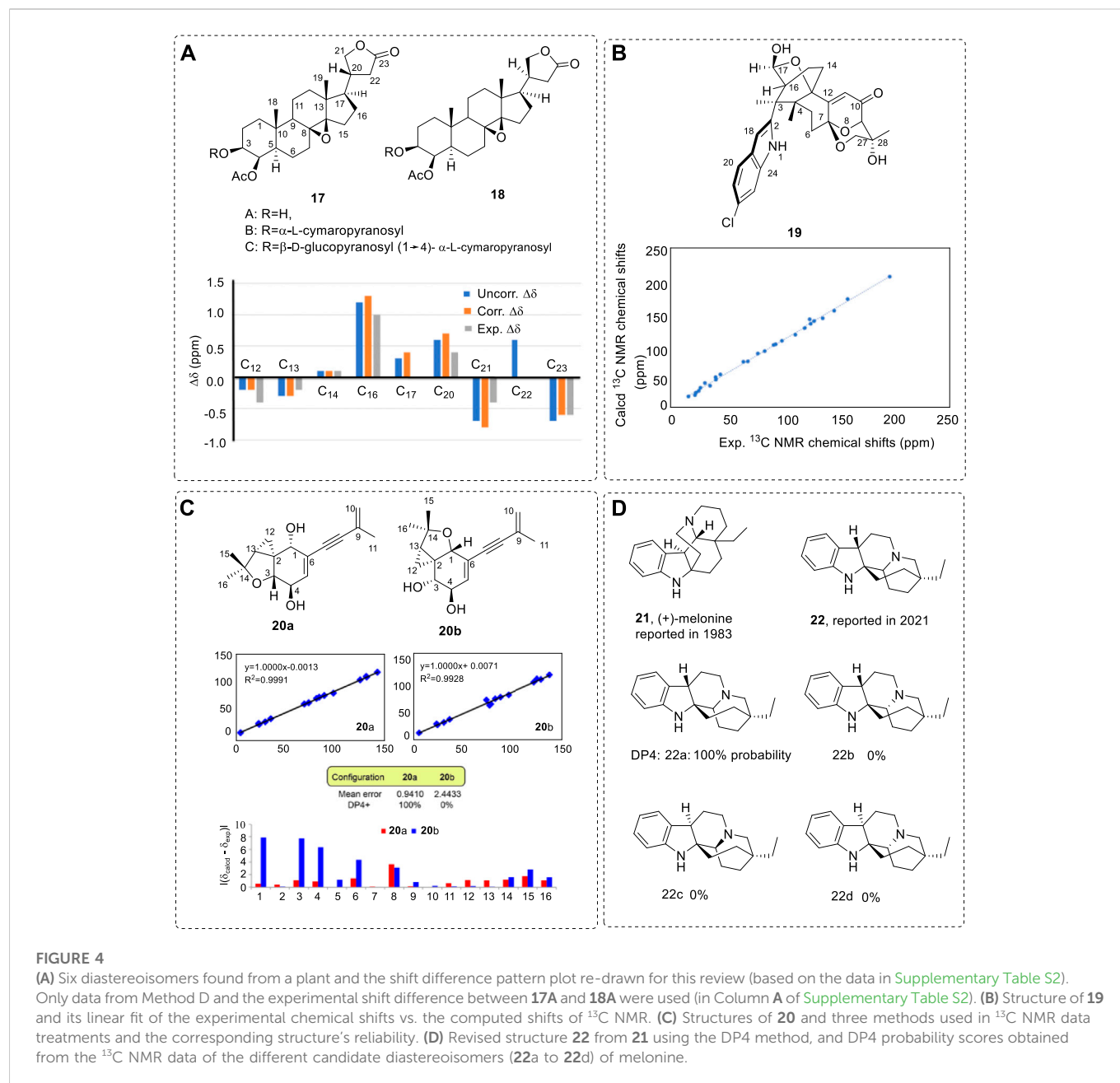


FIGURE 3

(A) (*Z*) and (*E*)-structures of cyclopeptide anthranilic acid (10). (B) Structure of zosteraphenol A (11) and its two rotamers (11a and 11b) in equilibrium, together with the experimental and DFT-calculated differences of ^1H NMR between chemical shifts of the minor (11b) and the major (11a) rotamers. All shielding and de-shielding effects shown are reproduced within 0.08 ppm. (C) The structures of 12 to 14 used in NMR calculations. (D) Structures 15 and 16 and the two conformers 15a and 15b of atropisomer 15.

Similarly, the relative configuration of stereoisomers 15 and 16 can also be assigned by comparing the experimental and theoretical $^{13}\text{C}/^1\text{H}$ NMR spectra (Figure 3D) (Cerulli et al., 2017). Eight isomers for 15 and 16 were used in the computations. After the ^{13}C NMR shifts were

obtained by a Boltzmann statistics sum, the DP4+ method was applied for comparison. Their relative configurations were assigned by a combined quantum mechanical/NMR approach, comparing the experimental $^{13}\text{C}/^1\text{H}$ NMR chemical shift data and the related



predicted values. It was found that they are atropisomers due to the restricted single-bond rotation where the resistance came from the repulsion of the two hydroxyl groups on the phenyl rings. The predicted transition state (TS) barrier between **15a** and **15b** was $\Delta G^\ddagger = 20.7$ kcal/mol. This barrier may be high enough to block its rotation $r = \text{freely}$ at room temperature.

There is another method for assigning the relative configurations for diastereoisomers by using the *chemical shift error patterns* of the experimental and computed ^{13}C NMR data. For example, there are two stereoisomers **17** and **18** from *Parepigynum funingense* (Hua et al., 2004); their configurations were assigned by comparing the theoretical ^{13}C NMR with the experimental data using the current configurations in Figure 4A (Hua et al., 2007). Only C20 had a different configuration. In this case, if we listed the shift differences from top to bottom, it could be

seen that the sign and shift difference values change pattern almost similarly (Supplementary Table S2). Therefore, the real relative configurations—for example, **17A** via **18A**—should have the same relative configurations with those that were used in ^{13}C NMR computations of **17A** and **18A**. If the sign is reversed, the real relative configurations of **17A** and **18A** should be the configurations of **18A** and **17A** that used in the ^{13}C NMR computations. This can be re-drawn as shift difference pattern plot as illustrated in Figure 4A. If the error pattern was used, it would be found almost the same between the linear fit corrections. Thus, in this case, using the error pattern is relatively convenient. Importantly, if the HF/6-31G(d)//HF/6-31G(d) was used in the computations, the shift error differences would almost be the same between the experimental and computational results. This is a cheap method that can be applied to this study.

Note that a deeper truth that can be found in the compounds. The different stereogenic center was C20; however, the shift difference of C20 was only 0.4 ppm between the **17** and **18** in their corresponding **A**, **B**, and **C** compounds. The largest shift difference was 1.0 ppm at C16. This is a typical example, and one must be very careful in handling experimental data. Generally, the different chiral center may have small to medium chemical shifts among the neighboring carbon atoms in the diastereoisomers.

A novel indole-diterpenoid **19** was obtained and its relative configuration was assigned based on NOESY spectra, with assistance provided by comparing the experimental ¹³C NMR with that computed (Figure 4B) (Zhou et al., 2021). Its coefficient was 0.9968 after the linear correlations. In this case, its AC can be determined using ECD.

One more example shows that several methods can reach the same conclusion if the structure is reliable. For example, the following two possible structures were identified using the method mentioned above. Diisoprenyl-cyclohexene/ane-type meroterpenoids are obtained from *Biscogniauxia* sp. 71-10-1-1 w. Compound **20** may have either a **20a** or **20b** structure (Figure 4C). As a typical example, it affords us a general viewpoint: cross-testing using several methods can provide more confident conclusions, and, if the structure is correct, these methods must give the same conclusion.

Finally, a structure revision example is illustrated here using DP4 and other assistance. The original structure melonine (+)-**21** was reported early (Baassou et al., 1983). After nearly 40 years, it was corrected as (2R,3S,7S,20R)-**22** using NMR data (Figure 4D) (Kouamé et al., 2021). This is easily understood since the structure elucidation mostly depended on 1D NMR data from 1983. This structure correction also led to its different biogenetic pathway.

4 Chiroptical spectroscopy

Specific optical rotation, which has the longest history in AC study, has played an important role in AC research. Even in today, it is still a useful method. Along with progress in electronic circular dichroism (ECD), vibrational circular dichroism (VCD), and Raman optical activity (ROA)—including some derivative technologies like magnetic circular dichroism (MCD)—these modern technologies have become increasingly important tools in stereochemistry. As a valid method, OR has been widely used as a necessary datum in many reports of chiral compounds.

4.1 OR and ORD methods

Specific optical rotation ($[\alpha]_D$) is simply called “optical rotation” (OR) in this text. Different OR values measured at different wavelengths are termed “optical rotatory dispersion” (ORD). In the absence of an ORD spectrometer, OR at several wavelengths is used in ORD study—633, 589, 546, 436, and 365 nm. Quantum method theories are widely applied for AC study (Grauso et al., 2019). Some typical chiral molecules were used in the study (Egidi et al., 2019). As one of the simplest and cheapest tools in AC determination, calculating chiral molecular OR and then comparing it with the experimental value is a valid and effective method for OR values that are generally not less than 20 degrees (Raghavan and Polavarapu, 2017). OR and ORD have

been widely used in the study of AC (Reinscheid and Reinscheid, 2016; Saito and Schreiner, 2020). For example, the chiral compounds **23–25** were assigned using OR methods; they are listed in Figure 5A (Li et al., 2011a; Li et al., 2011b; Ding et al., 2016).

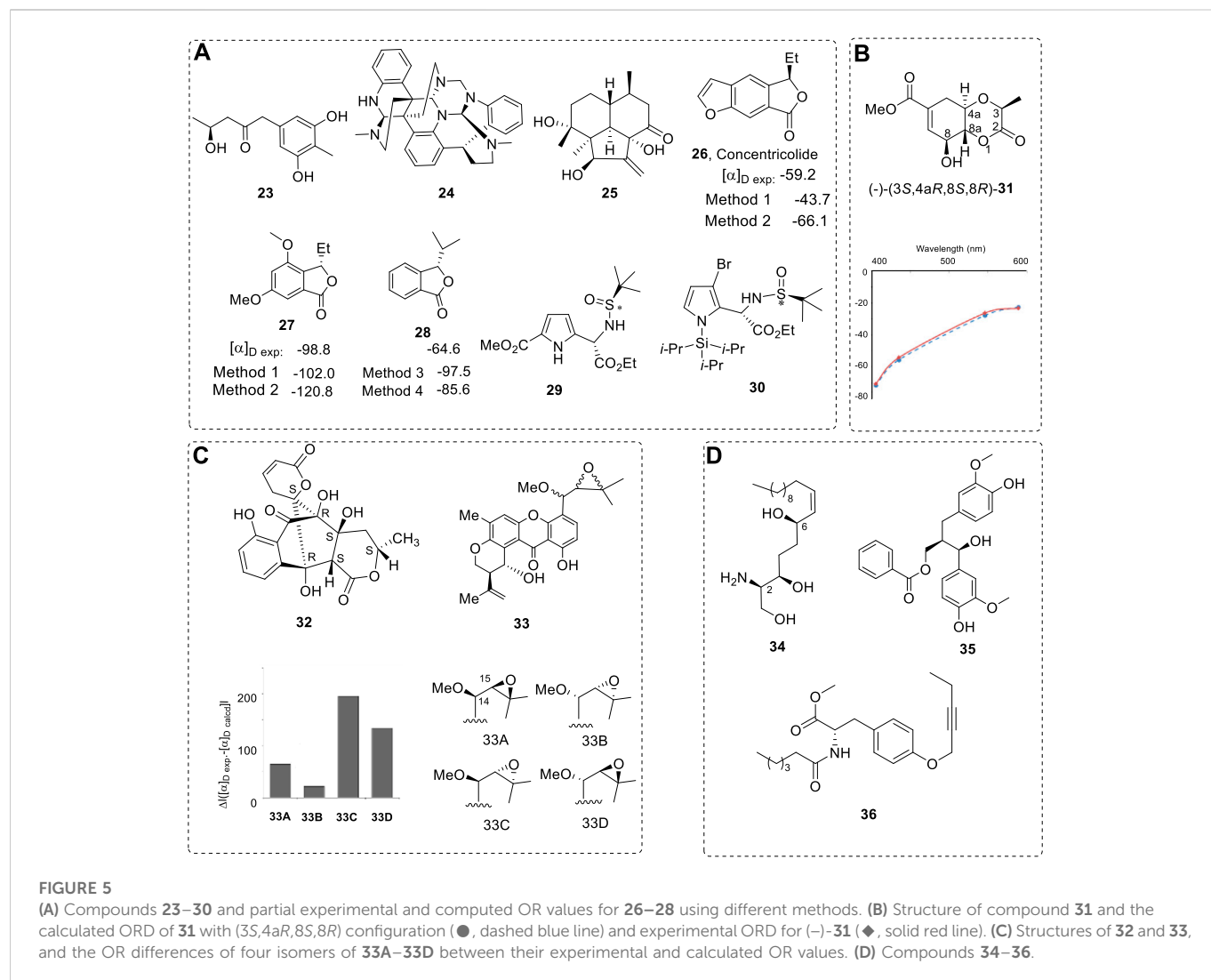
Compound **27** was obtained because natural concentricolide (**26**) has an anti-HIV property; it was assigned as (*R*) configuration with a -59.2 of OR (Qin et al., 2006). Quantum methods were used in the AC study (Ren et al., 2009). Method 1: B3LYP/aug-cc-pVDZ//B3LYP/6-31+G(d), and Method 2: B3LYP/aug-cc-pVDZ//MP2/6-311+G(d) were used for OR computations of (*S*)-**26**. Both methods predicted -43.7 and -66.1 . Thus, the configuration of **26** must be (*S*). The other compound **28** with OR value of -64.6 (Li and Si, 2012) provided other evidence to support this conclusion (Method 3: (B3LYP/6-31+G(d,p))/PCM//B3LYP/6-31+G(d,p) and Method 4: PCM/B3LYP/6-31++G(d,p)//B3LYP/6-1+G(d,p)) (Figure 5A). Other kinds of chiral compounds, such as sulfone (**29**) or Si-, Br-atom-containing compound (**30**) could be used in OR computations using current theoretical methods (Figure 5A) (Haghdani et al., 2016). The interesting chirality of $>S=O$ in **29** and **30** have been well investigated. Like CAM-B3LYP functional and the second-order approximate coupled cluster singles and doubles (CC2) method. Currently, OR values for any chiral compounds that contain most elements of the periodic table could be computed using quantum methods.

Firstly, it should be clear that the assigned AC concluded from OR must be the same as the other quantum theories or experimental results. The bioactive metabolite phyllostin (**31**) was isolated from *Phyllosticta cirsii* fungus (Evidente et al., 2008); its relative configuration was established as (3*R*,4*aS*,8*R*,8*aS*) by NMR and X-ray analyses (Tuzi et al., 2010) while its AC remained unknown. Total synthesis (Muralidharam et al., 1990) and the ECD exciton chirality method (Isogai et al., 1985; Mu et al., 2021) permitted the determination of the AC of its C8 epimer. The ORD study gave the same conclusion: (-)-(3*S*,4*aR*,8*S*,8*R*)-**31** (Figure 5B) (Mazzeo et al., 2013).

It is simple to assign one compound's AC if one chiral compound is relatively rigid and its relative configuration is known. It is only necessary to compute its OR directly and then compare its OR value with the experimental results to assign its AC. Compound **32** was obtained with a known relative configuration *via* X-ray (Mo radiation) (Ding et al., 2010); its OR was $+57.4$ in MeOH. The computed OR for (*R*)-**32** was $+74.3$ in the gas phase and $+66.2$ in methanol using the PCM model (Figure 5C). This means the (+)-**32** must have the configuration as illustrated in Figure 5C. Indeed, ideal situations are rare.

In some cases, there are several diastereoisomers—for example, **33** (Figure 5C) (Figueroa et al., 2009). Its side-chain configurations with ethylene oxide moiety were not assigned. By comparing the $\Delta[\alpha]_D$ values among the four diastereoisomers, isomer **33B** with the smallest $\Delta[\alpha]_D$ value was assigned as the most likely structure.

Many chiral compounds may have flexible structures and their AC assignment is not easy due to many conformations in solution. For example, with chiral compound halisphingosine A (**34**) (Molinski et al., 2013), a very flexible linear chiral compound with three stereogenic centers, it is hard to compute its OR to assign its AC. A simpler model is needed—to be discussed later—or making its derivatives for ECD exciton chirality spectra. Compounds (7*R*,8*R*)-**35**



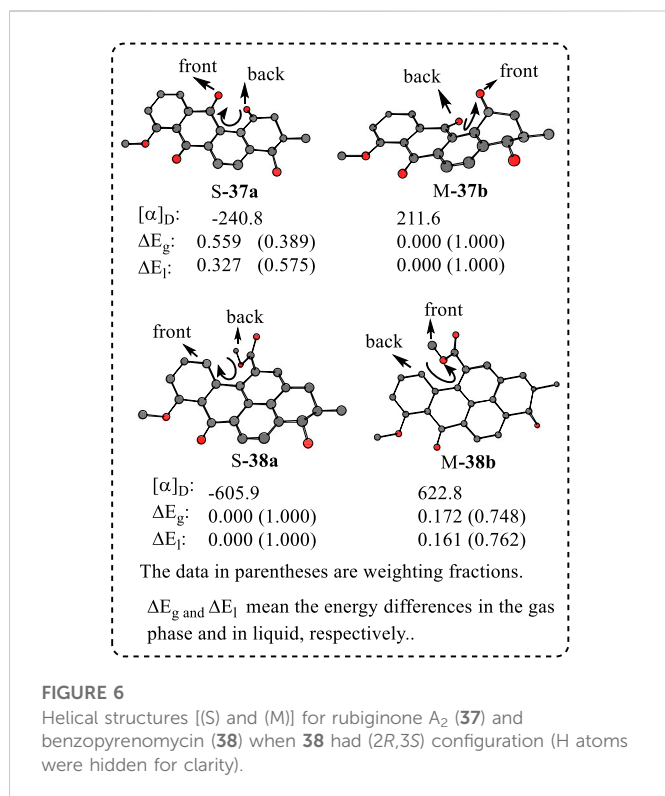
(Zhang et al., 2021a) and **36** (Zaman et al., 2021) are not as flexible as those of **34** (Figure 5D); they can be used directly in computations of OR or ECD.

It has been reported that the structurally similar compounds rubiginone A₂ (**37**) and benzopyrenomycin (**38**) have close OR values; **37** had an OR of +50 and **38** had +38 in CHCl₃ (Figure 6). However, the computed OR values were 14 to 87 for **37** and -12 to -50 for **38**, using different quantum methods (Li et al., 2013). Therefore, **37** had the (2*R*,3*S*) configuration and **38** must have (2*S*,3*R*) configurations. Why is there such a big difference from the viewpoint of structural features? The key point is that these two molecules formed different helical chirality, as illustrated below (Figure 6). When both had (2*R*,3*S*) configuration, their conformation analyses were performed. The minor conformers S-**37a** had an OR value of -240.8 and major conformer M-**37b** had 211.6. After the Boltzmann statistics, the OR value depended on the major conformer M-**37b**'s OR sign. In contrast, the major conformer S-**38a** had -605 and the minor M-**38b** had 622.8. Finally, the simulated OR value and sign were decided by the major conformer S-**38a**. Thus, if both had (2*R*,3*S*) configuration, they should have a reversed OR sign. Consequently, when both had the same OR sign, they must have different ACs.

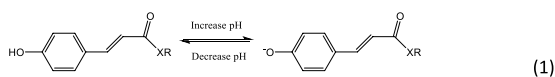
This example provided an important hint: when two structures are very similar, the real conformer structures must be carefully examined before identifying their AC using their OR values. Helical structures exist widely and, importantly, generally have large OR values. This example again indicates that conformational study is very important, as we have emphasized previously.

ORD provided more information in other wavelengths for various structures. Since different structures have different sensitivity at different wavelengths, different structures may have obvious and different ORD curves. The AC of C8 of jonquiline (**39**) was assigned as (*R*) (Vergura et al., 2018) based on the computations of ORD at the level of B3LYP/aug-cc-pVDZ//B3LYP/TZVP level in the gas phase. The (8*S*)-**39** had a large difference in ORD from the (8*R*)-**39** and the experimental ORD (Figure 7). The ORD values of **39** in methanol were larger than those in chloroform.

Because, for some chiral phenol compounds, the -OH is acidic, its ORD or other properties are dependent on the pH values of the solution. In this case, different solvents may have varying effects on the OR. For example, different ORD curves were recorded for the diesters of (1*R*,2*R*)- and (1*S*,2*S*)-cyclohexanediols and diamides of (1*R*,2*R*)- and (1*S*,2*S*)-diaminocyclohexane (**40** and **41**) with *p*-hydroxycinnamic acid (Figure 7) (Mazzeo et al., 2016). This is



due to the equilibrium under different pH values (Eq. 1). The compounds also exhibited different chiroptical spectroscopies in solution.



In addition, the ORD of any chiral compounds, such as the reaction atropisomer **45** from **42**, could be also used in the AC assignment (Figure 8A) (Mirco et al., 2019). The enantiomers (+)-**45** and (-)-**45** exhibited a very good mirror image as expected. The (S)-**45** was used in the ORD study.

Finally, an important phenomenon should be noted. Many linear chiral compounds have only very small OR values (Lu et al., 2020). In these cases, it is common to use quantum methods to compute OR values to assign AC unless there is a so-called known analogue's OR (theoretical and experimental OR values used as reference). For more about this study, please see the simplified model section. The OR analysis of four additional chiral compounds (**46**–**49**) are summarized below (Figure 8B) (Taniguchi et al., 2009; Ding et al., 2012; Yang et al., 2013; Sun et al., 2021a).

A notable case is the computed OR value being generally larger than the experimental datum. In some cases, the computed OR is two or more times larger than the experimental OR values using current DFT methods. Therefore, it is better to refer to other methods in AC determinations for complex chiral compounds, particularly if any of the OR values are small, such as less than 20°.

OR has been widely used for a long time and it will continue to be, as has the ORD method. However, its application for AC study is not as frequent as OR or ECD.

4.2 ECD and its uses

Electronic circular dichroism (ECD) spectra have been used for AC determination since the 1960s. ECD was originally known more simply as CD, and, more recently, many reports use the term ECD for AC assignment (Guo et al., 2020; Zhang et al., 2021b; Kim et al., 2021; Nhoek et al., 2021; Quan et al., 2021). As mentioned above, a detailed conformational search should first be performed (Mazzeo et al., 2014). Many known chiral compounds have been used this way in chiroptical spectroscopic study (Molteni et al., 2015; Stepanek and Bour, 2015; Pescitelli and Bruhn, 2016; Rossi et al., 2016; Johnson et al., 2018) while new synthetic chiral compounds have also been studied (Paolino et al., 2021; Yajima et al., 2021). Chiral compounds require one or two UV-vis chromophores in order to use ECD in their configuration assignments.

The DFT theory has been widely used in ECD computations for various chiral organic compounds (Polavarapu and Covington, 2014; Jiang et al., 2021a; Ji et al., 2021; Niu et al., 2021), including metallic chiral compounds (Enamullah et al., 2016; Saha et al., 2017; Ravutsov et al., 2021). This method provides not only a good balance in treating different molecular systems but also gives relatively accurate computational results when applied to real molecules. For example, using the functional B3LYP at the 6-311+G(d) set level basis can obtain relatively good agreement between theoretical prediction and experimental results. An improved theory, such as cam-DFT (Yanai et al., 2004), is also used in ECD computation. Other methods, like simplified TD-DFT, may give fast and accurate assignment for some chiral compounds (Cerra et al., 2019; Martynov et al., 2019; Ianni et al., 2022). For some chiral molecules, this method can provide more accurate results, although this varies depending on different molecules. A general mixed quantum/classical method for the calculation of the vibronic ECD has been reported as valid tests (Cerezo et al., 2018). Compared to DFT calculations, its computational cost is relatively larger, but it is useful in cases where the ECD shape arises from a subtle balance between vibronic effects and conformational varieties. Indeed, as an active research area, many developed methods have been reported (Liu et al., 2016a; Hodecker et al., 2016; Hong et al., 2016; Nørby et al., 2017; Jimenez et al., 2019; Scott et al., 2021; Wibowo et al., 2021), including the efficient simplified time-dependent density functional theory (sTD-DFT) that has been used to compute large molecular systems with more than 100 atoms (María-Magdalena and Jorge, 2015) such as helicene (Bannwarth et al., 2016), which consists of 102 atom and 16 annulated benzene rings. Simulations of ECD for nucleic acids have also effectively used quantum theory (Zhu et al., 2014; Padula et al., 2016a; Zou et al., 2021).

DFT methods should first be used and, if the predicted ECD curves are greatly different from the experimental results after UV correction, cam-DFT, such as cam-B3LYP theory, can be used in computations to compare both measured and predicted ECD spectra.

DFT is an efficient tool used in AC determinations. Some typical compounds and their ECD spectra are summarized below, such as an unprecedented C₉-spiro-fused 7,8-seco-ent-abietane, decandrinin (**50**) (Figure 9A) (Wang et al., 2014a). The ECD signals had almost no red or blue shifts. The ECD of organic salts, such as clavulanate potassium, can also be calculated using DFT method shift (Li et al., 2018). The structure of some chiral compounds, such as anacosins A (**51**), contain some isolated C=C or C=O groups, and ECD signals

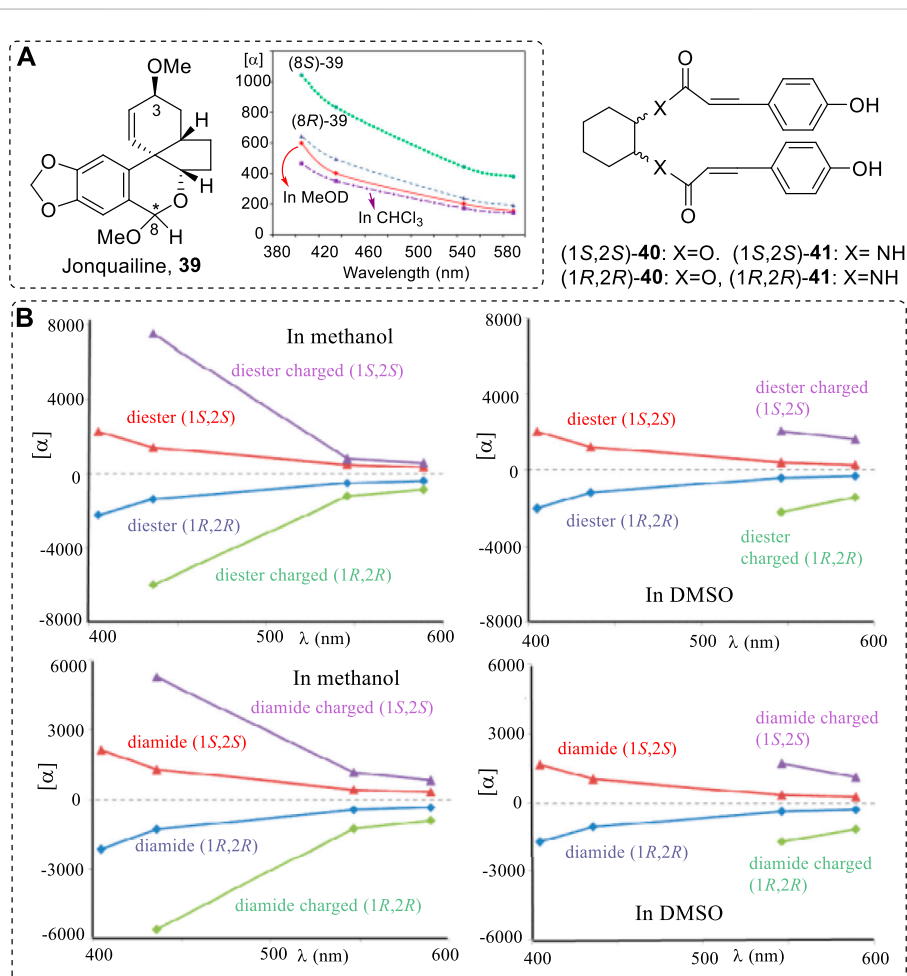


FIGURE 7

(A) Experimental ORD of **39** (methanol, solid red curve, chloroform, dash-dotted purple curve) and computed spectra (gas phase) for (8S)-**39** (dotted green curve) and (8R)-**39** (dashed blue curve). (B) Experimental ORD curves of neutral and charged diesters of (1S,2S)- and (1R,2R)-**40** (up panels) and (1R,2R)- and (1S,2S)-**41** (down panels) in MeOH and DMSO, respectively.

mostly appear from 200–250 nm (Cai et al., 2019). The best evidence is its match of the predicted VCD with experimental results.

In ECD study generally, chiral molecules have one or more vis-UV chromophores in the UV wavelength range of 200–400 nm for organic compounds. The short wavelength range, such as 180–210 nm, can also provide valid information for aliphatic compounds, such as terpenoids, peptides, or proteins (Farkas et al., 2016; Banerjee and Sheet, 2017). However, signals of 180–210 nm are more difficult to measure without distortions. Most ECD study involves the range of 210–400 nm for organic compounds (Li et al., 2014; Nicu et al., 2014; Podlech et al., 2014; Yu, 2014; Han et al., 2021; Omar et al., 2021; Saetang et al., 2021). The ECD wavelength range for chiral metallic ion-containing organic compounds or poly-conjugated compounds may cover 500 nm or more (Wu et al., 2014; Saleh et al., 2015; Enamullah et al., 2016; Vesga and Hernandez, 2016).

One configurationally stable chiral dithia-bridged hetero (Marty et al., 2014) helicene radical cation (**52**) was studied (Figure 9B) (Gliemann et al., 2017) involving nitrogen *N*-chirality. This molecule had a bulky negative ion SbF₆⁻, which is located close to the *N* atom to fix its conformation. This provides stability to the *N* chirality

formation—natural compounds containing *N*-cation in natural products are not rare.

Some novel types of chiral compounds were recently obtained from natural sources. Anti-inflammatory cassane-type diterpenoids from the seed kernels of *Caesalpinia sinensis* have a novel framework formed with the new ring D. This rigid framework (+)-**53** was confirmed by X-ray, and its analogues (+)-**54** must have the same AC; this has been confirmed by comparing the computed ECD for (5S,6R,8R,9S,10R,14S)-**54** with the experimental ECD of (+)-**54** as (5S,6R,8R,9S,10R,14S) (Figure 9C) (Wang et al., 2021a). The ACs confirmed by X-ray structure and ECD spectra are the same.

The dimeric form of axial carboline compound **55** has been isolated and identified using ECD (CAM-B3LYP/6-31G(d)//B97D/TZVP) (Jin et al., 2021), which was obtained from the culture broth of a cold-seep derived actinomycete, *Streptomyces olivaceus* OUCLQ19-3 (Figure 9D). Its other axial diastereoisomer was also obtained and the predicted ECD spectrum agreed well with the experimental result. The N-C single bond rotation was seriously restricted; if the rotation barrier was provided there, we could more clearly understand the atropisomer's properties of NMR in solution.

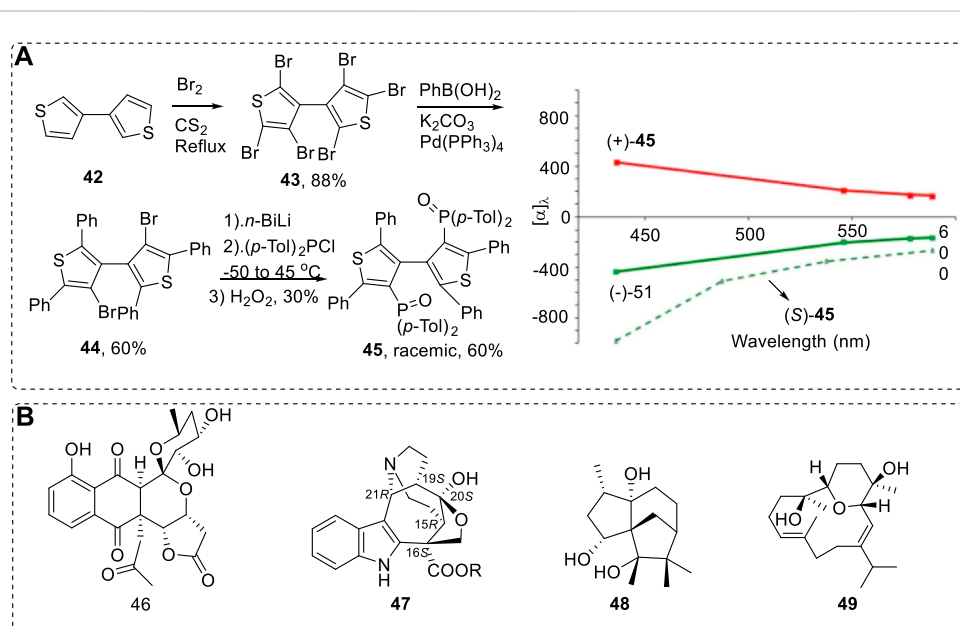


FIGURE 8

(A) Left is the reaction scheme for the synthesis of racemic; right is the experimental and calculated (dashed lines) ORD spectra of enantiomers (**45**) in chloroform. (B) Structures of compounds **46–49**.

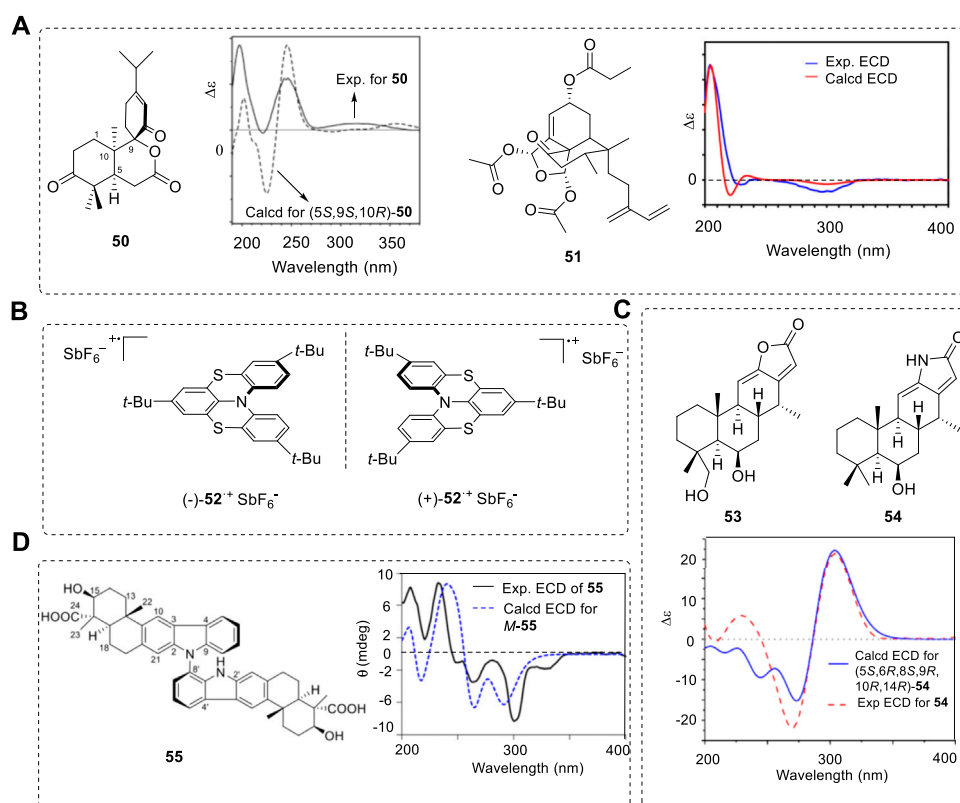
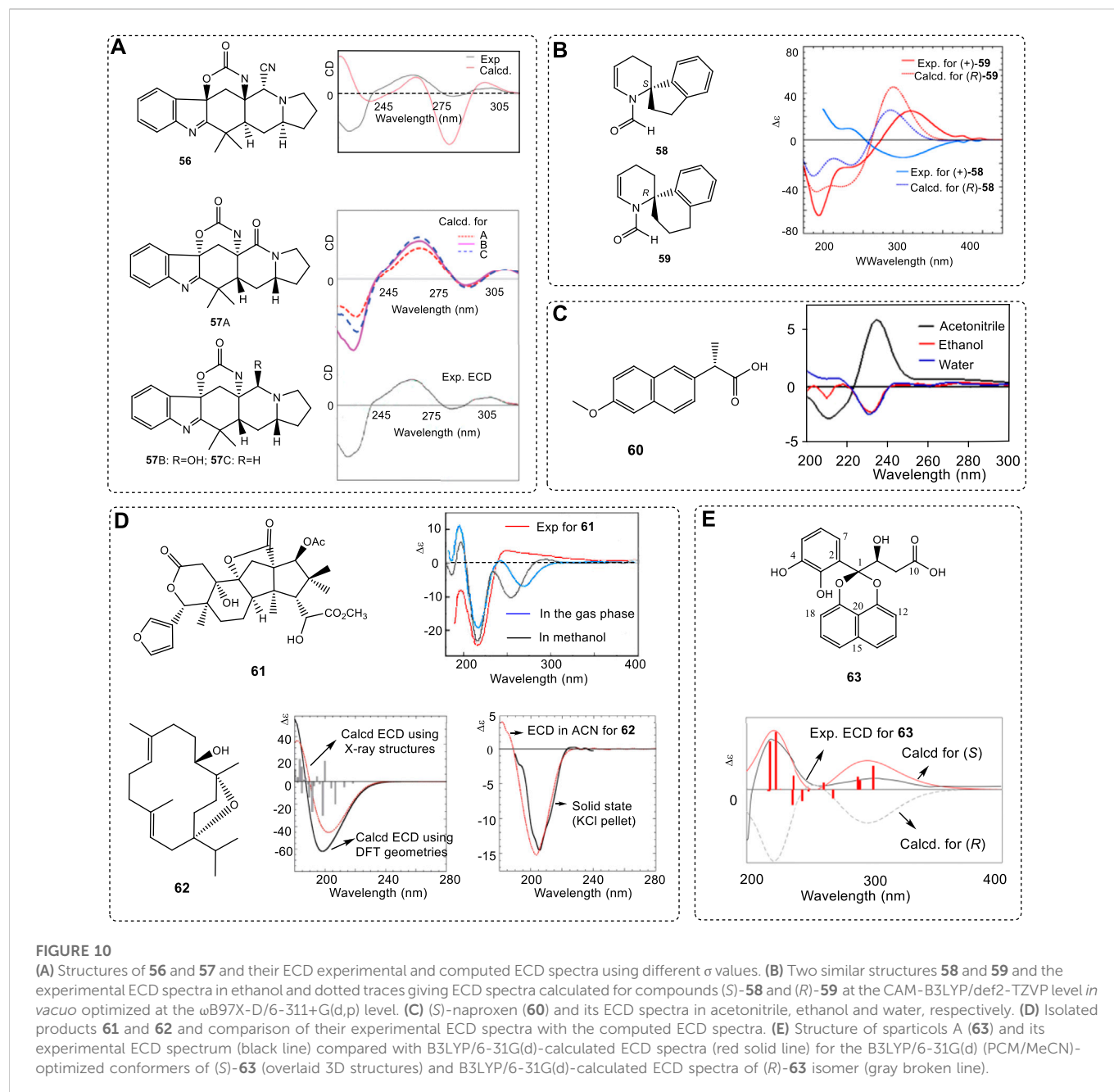


FIGURE 9

(A) Structures of **50** and **51** and their ECD spectra. (B) Chemical structures of investigated dithia-bridged hetero[4]helicene radical cations **(-)-52⁺ SbF₆⁻** and **(+)-52⁺ SbF₆⁻**. (C) Structures **53** and **54** and experimental and theoretical ECD spectra. (D) Dimeric form of carbolines (**55**) and its ECD obtained at the CAM-B3LYP/6-31G(d)/B97D/TZVP.



Finally, we consider the standard deviation (σ) factor of the data. Among the various factors that affect the AC determinations, one problem that has been ignored in the teaching process is the data of σ . The wrong assignment of AC was caused by a wrong use of the sigma. Thus, this factor must be used carefully by beginners to this method.

The data of σ could be 0.2 to 0.5 eV, but 0.3 and/or 0.4 eV has mostly been used in ECD simulations. The general principle of using σ depends on the ECD signals recorded in experiments. Unless the experimental ECD resolution is very high, 0.3 or 0.4 eV can be used in most cases. Once the signal resolution is very high, where the signal's half-height is small, 0.2 or 0.25 eV can be used to simulate the ECD. In contrast, 0.5 eV may be used in simulation of ECD spectra. Generally, the smaller the σ value, the sharper the signals of ECD will be and the resolution of the ECD will be high.

However, when the ECD signals are complex, it is not easy to select their σ values. For example, one wrong assignment of chiral compound **56** was reported (Ji et al., 2013). The conformational search for all geometries was performed correctly, and the excitation state numbers were covered from 200 nm to 400 nm in the calculations. Its ECD simulation was performed at the B3LYP/6-31G(d) level. The wrong σ value of 0.2 eV was used. After its analogues A–C were obtained, the ACs were assigned again. The original AC of **56** was corrected to **57** by using 0.35 eV (Figure 10A) (Li et al., 2016b). Obviously, the small σ value of 0.2 eV in the report led to the wrong assignment of the compound **56**.

The default sigma value of 0.4 eV is used in ECD data treatment in GaussView software; there is much no-cost software can be used for ECD simulations. For more details, please refer to the corresponding websites and download the appropriate software. The corresponding

website for ECD spectra is at <https://github.com/jbloino/estampes>. Readers who need to use it may contact the corresponding person there. Other software may exist on different websites.

As mentioned above, correct conformer structures are extremely important for ACs. This procedure must use quantum theory. Due to the computational limits from before 2000 or earlier, the use of a high basis set, such as 6-311+G(d) for a large chiral molecule, was then difficult. In 1997, two spiro compounds **58** and **59** with very similar skeletons exhibited almost identical ECD spectra (Figure 10B) (Ripa et al., 1997). Compound **59** just had one more CH₂ than **58**; both had the same two chromophoric moieties. It was expected that both (*S*)-**58** and (*S*)-**59** would have the same ECD spectra since their spiro-center structure was identical. However, an early report argued that both would have mirror-image ECD spectra using Schellman matrix methods (Sandström, 2000a)—a popular approach in 1990s based on independent system approximation (ISA). For more details about ISA, refer the corresponding materials (Raabe et al., 2012; Sandström, 2000b).

The two structures were then carefully examined using DFT theory (Padula et al., 2016b). The new study "...demonstrates that the main source of the error in the original investigation lies in the fact that the aromatic and N-formyl enamine chromophores are not independent in the sense implied by the ISA; therefore, the ECD spectra of both compounds **58** and **59** cannot be correctly treated by the matrix method approach (Padula et al., 2016b)." Obviously, the early conclusion was not correct. After using all the new conformers in the ECD simulations, the correct results were obtained (Figure 10B). The key reason was that a TDDFT-based fragmentation approach (ISA) led to very different results from full TDDFT calculations. All theories of B3LYP, CAM-B3LYP, and M06-2X with the def2-TZVP basis sets predicted the same conclusion. With the development of computers, full chiral molecules are now used in computations unless the chiral molecules are extremely large.

In general, solvents may have some effect on ECD spectra. In a very unusual case, naproxen (**60**) had completely different ECD in different solvents (Figure 10C) (Ximenes et al., 2018). It had a (+)-Cotton signal at near 235 nm in acetonitrile and one (−)-Cotton signal at 210 nm. In contrast, it had a (−)-Cotton signal at 230 and 231 nm in ethanol and water and an unremarkable (+)-Cotton effect from 200 to 220 nm. However, the simulated ECD spectra in the three solvents gave very similar curves. All predicted ECD had a (+)-Cotton effect at a 240–245 nm range and a (−)-Cotton effect of 202–205 nm. This is an extremely unusual case, in that the predicted ECD did not match the experimental results in different solvents. Its importance may be that

This finding deserves great attention because polar protic and aprotic solvents are widely used for HPLC analysis in quality control of pharmaceutical drugs. If we consider the importance of discrimination between the active drug [(*S*)-naproxen, anti-inflammatory] and the toxic one [(*R*)-naproxen, hepatotoxic] using ECD detector coupled to HPLC, the findings reported here could represent an important source of error in these analyses (authors' statement).

The reason was not analyzed until now. It is possible that there was a very strong solvation effect on the formation of different geometries with different energies.

The use of ECD to assign a chiral molecule's AC requires the chiral center(s) to be close to the chromophore. If the stereogenic center is too far from the chromophores, it can be only used when this molecular relative configuration is reliable. An example is

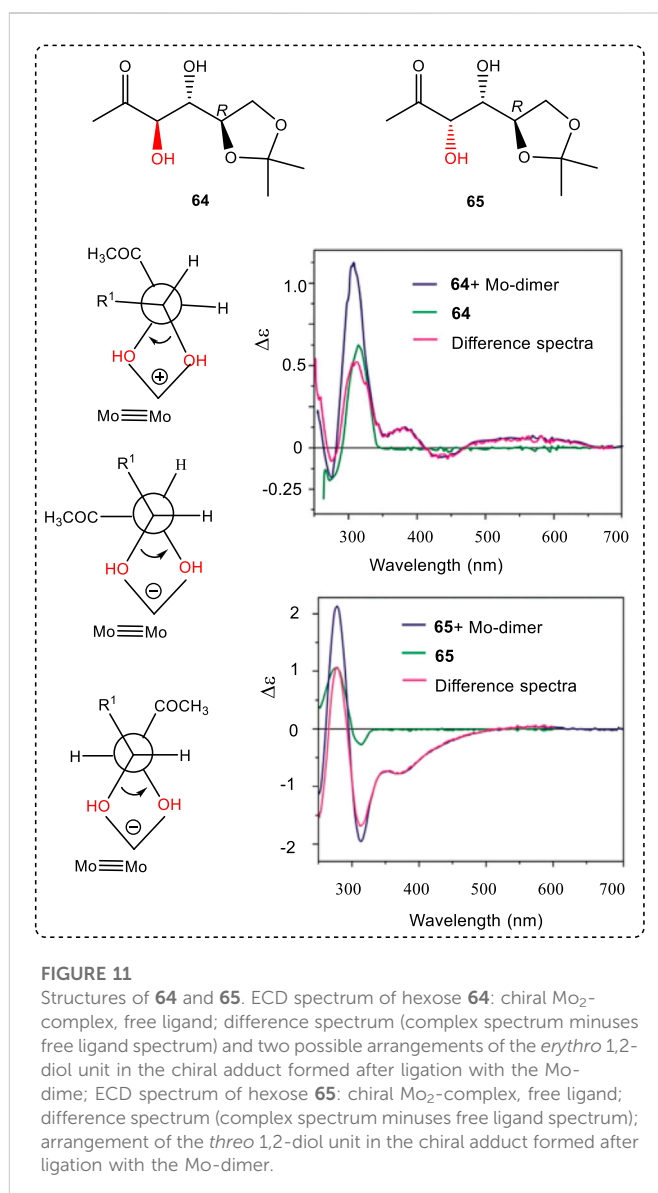


FIGURE 11
Structures of **64** and **65**. ECD spectrum of hexose **64**: chiral Mo₂-complex, free ligand; difference spectrum (complex spectrum minus free ligand spectrum) and two possible arrangements of the *erythro* 1,2-diol unit in the chiral adduct formed after ligation with the Mo-dimer; ECD spectrum of hexose **65**: chiral Mo₂-complex, free ligand; difference spectrum (complex spectrum minus free ligand spectrum); arrangement of the *threo* 1,2-diol unit in the chiral adduct formed after ligation with the Mo-dimer.

compound **62** (Yuan et al., 2010), which has very limited chromophores. If one had one to two different configurations from the current one (**61**) illustrated below, it is impossible to record the obvious ECD spectra difference from the **62s** ECD. Only when its relative configuration is reliable, such as the X-ray information, then can its AC using ECD be significant and correct (**62**, Figure 10D) (Avula et al., 2016).

Once again, it must be emphasized that, although compound **61** has one more sugar moiety with five chiral centers, it has no chromophores and its stereogenic centers have very little effect on its ECD spectra. This raises another question: if a chiral molecule has polychiral centers but its relative configuration is not identified, then, if anyone wants to use ECD to identify its AC, it is impossible, since its ECD curves would hardly change, no matter which stereogenic center changes its configuration.

When the chiral center is located on the side chain, then it is the favored relative to the neighboring ring structures. For example, sparticolos A (**63**) has a chiral center at C8 connected to a hydroxyl group, and its computed ECD and experimental ECD spectrum

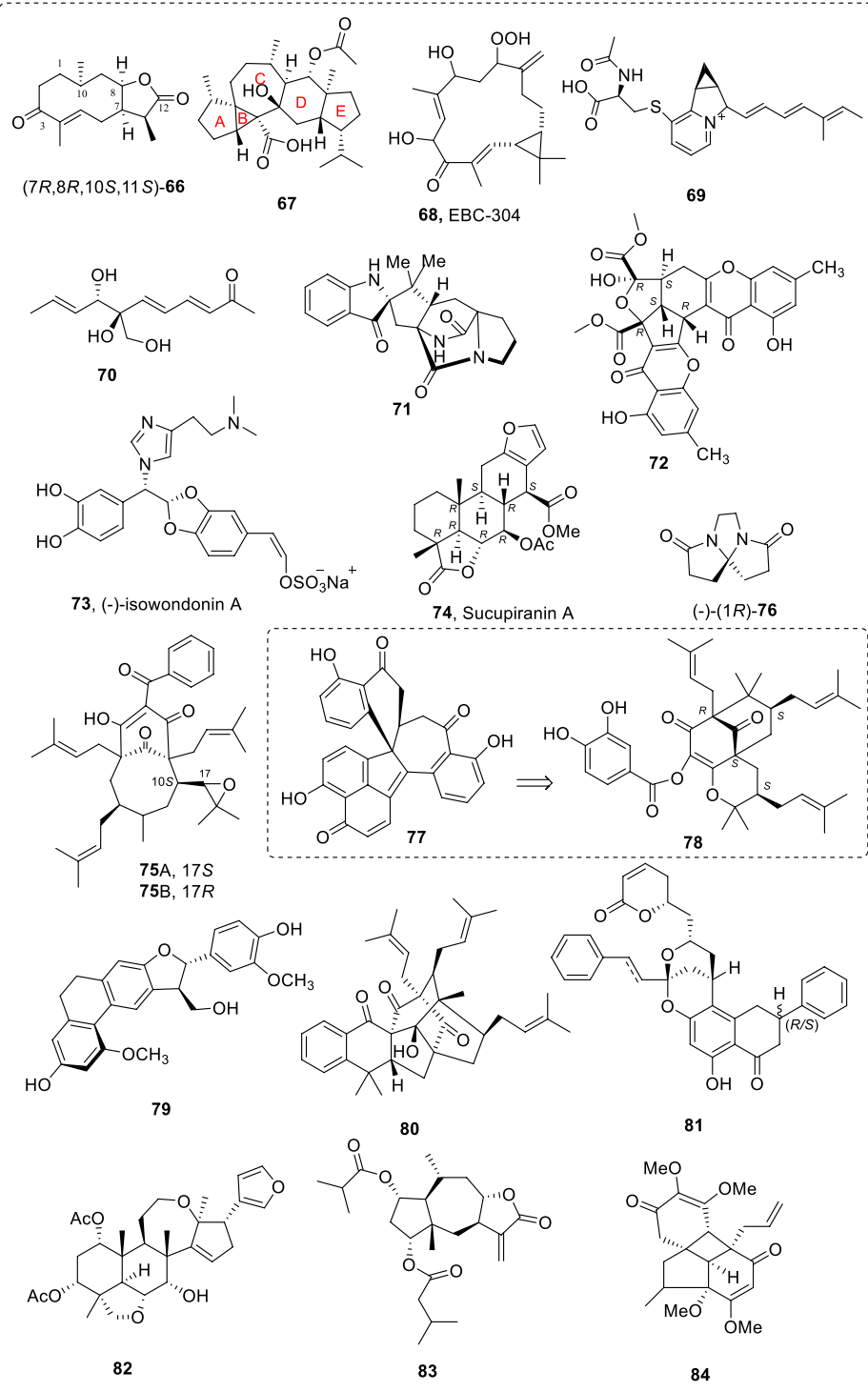


FIGURE 12
Structures 66–84.

matched well (Figure 10E) (Garcia et al., 2021). Due to the through space interaction of the -OH with the phenyl ring and the intramolecular H-bond, it has limited flexibility. In this case, its major contribution to the conformer generally played a key role in ECD spectra. Other examples, like asperglauclin A, have been reported (Lin et al., 2021).

Diols with a diol moiety often have biological activities, such as antibiotics (Fischbach and Walsh, 2009) and steroidal hormones (Norman et al., 2004; Niwa and Niwa, 2014). In organic chemistry, a diol fragment can be used as a chiral building block (Hartung and Grubbs, 2014) and promoter used in asymmetric synthesis (Huo et al., 2014). On the other hand, many diol-containing compounds are

transparent in the UV-vis spectral range. Thus, they need to be converted into a derivative with a chromophoric group. Chiroptical methods can be used to identify the ACs of some diols (Jawiczuk et al., 2015). It is convenient to convert diols, which may be located on one ring moiety or on a linear carbon chain, into a new cyclic moiety to reduce the conformers in various sugar-derivatives or poly-alcohol-containing compounds in AC study (Pardo-Novoa et al., 2016).

The method of determining the AC of diols is to use dimolybdenum methodology with dimolybdenum tetracarboxylates which act as auxiliaries (Jawiczuk et al., 2014; Masuda et al., 2021). This consists of the *in situ* formation of chiral complexes (Mo₂-core). The measured ECD spectra of the chiral complexes minus the free-ligand spectrum afford a different spectrum. The determination is strictly associated with the dependence of the signs of the Cotton effect (Górecki et al., 2007). Identical signs occurred respectively at near 400 nm and 310 nm, which is consistent with the sign of the O-C-C-O torsion angle. Examples are introduced here using **64** and **65** (Popik et al., 2014). Hexose **65** negative signs of decisive ECD band at 310 nm and 380 nm conform to the negative sign of the O-C-C-O torsion angle. The second requirement—the antiperiplanar orientation of both hydroxyl groups versus 1,2-O-isopropylidene ring and acetyl group—is also met (Figure 11). On this basis, the assignment of 3S,4S AC in compound **65** was directly possible from the level of the ECD spectrum. However, it is difficult to assign the AC of **64** because it has two possible Newman projections with two states (+) and (-); when two hydroxy groups and two connected Mo atoms were located at the bottom, the large groups had two different positions in space. Thus, it is important to give the proper Newman projection for a given chiral molecule using this method.

A popular method of analysis is to calculate ECD and compare it with the experimental spectrum. Many examples have been studied and the predicted ECD curves agreed well with the experimental results (Junior et al., 2014; Wang et al., 2014b; Fleming et al., 2015; Pescitelli and Bruhn, 2016; Masnyk et al., 2016; Lee et al., 2018; Zhu et al., 2018; Liu et al., 2019a; Liu et al., 2019b; Guo et al., 2021a; Arreaga-González et al., 2021; Guo et al., 2021b; Wang et al., 2021b; Rao et al., 2021; Tan et al., 2021; Xie et al., 2021); this includes the compounds in which the ECD were compared with the compound's configuration that was assigned using quantum theory methods (Socolsky et al., 2016). Some examples (**66–84**) are summarized in Figure 12 (Zhang et al., 2008; Pazderkova et al., 2014; Zhang et al., 2014; Bultinck et al., 2015; Sherer et al., 2015; Liu et al., 2016b; Xu et al., 2016b; Evidente et al., 2016; Nakashima et al., 2016; Yu et al., 2016; Endo et al., 2017; Maslovskaya et al., 2017; Xu et al., 2019; Zhou et al., 2019; He et al., 2021a; Sun et al., 2021b; Lu et al., 2021; Xue et al., 2021; Yang et al., 2021).

Another CD method is CD exciton chirality (Pescitelli, 2018). This method generally requires two chromophores to be present in the molecule (could be obtained by synthesis). It is different from the ECD calculations using Gaussian or another package *via* quantum equations, since the CD exciton method does not give a specific ECD curve or a series of excitation state energy for any chiral compounds. Instead, it uses the relative position of dichromophores, such as the dibenzoates of acyclic 1,2-glycols in space to determine the ACs. This may entail uncertainties for some chiral compounds where it is not easy to correctly orientate the chromophores' position. It is useful to determine the AC of the acyclic 1,2-glycols, acetylene alcohols (Naito et al., 2005), or similar chiral compounds. *As emphasized above, a key step is to locate the*

relative positions of two chromophores for a specific chiral compound by conformational considerations.

The Mosher ester method is valid for the AC assignment of secondary alcohols or others (Dale and Mosher, 1973; Seco et al., 2004). This method provides very valid assistance in the study of ACs. In most cases, this method requires the substituents adjacent to the stereogenic center to be small. A previous example clearly stated this point. *If there is a very bulky group, it may change the most and second-most stable conformers' distribution and will finally lead to a wrong conclusion.* This point must be clearly remembered.

Both methods are out of the scope of the present review; therefore, the two useful methods are just mentioned here rather than described in detail. Readers interested in the two methods can look for the corresponding reports (Gao et al., 2021), including other experimental methods (Fujii et al., 1998).

4.3 VCD methods

Generally, vibrational circular dichroism (VCD) spectroscopy can be used for the AC determinations of any chiral compounds. It differs from ECD in that ECD spectrometers were quickly commercialized in the 1970s and were widely applied to the study of AC. VCD was first applied to AC determinations in 1990s by using magnetic field perturbation equations (Nafie, 1983; Stephens, 1985; Nafie, 1992; Stephens et al., 1994), especially when quantum theory was developed and computational methods became available by use of the Gaussian package (commercially available from 1998, and then more widely for Gaussian 03 in 2003). Modern VCD instruments provide important assistance for AC study (Zhu et al., 2015; Keiderling and Lakhani, 2018; Ren et al., 2019) while computational methods (Polavarapu and Covington, 2014), including anharmonic VCD methods or other theoretical methods (Barone et al., 2012; Domingos et al., 2015; Fuse et al., 2019), make it possible for scientists to perform calculations and comparisons with a wide range of experimental results (Rossi et al., 2017; Taniguchi, 2017; Kohout et al., 2016; Góbi et al., 2015; Abbate et al., 2014; Frelek et al., 2014; Batista et al., 2015; Giovannini et al., 2016; Nieto et al., 2010; Bösel et al., 2019; Demarque et al., 2020; Ren and Zhu, 2009; Monde et al., 2003; Cichewicz et al., 2005; Batista et al., 2011a).

VCD spectroscopy is a valid method used in ACs determination for various chiral compounds (Polavarapu and Santoro, 2020). Since it uses IR light as the source for covering vibrational transitions in all parts of a molecule, it does not require any chromophores in the vis-UV region. It has been widely used in the past decade in the AC assignment of various natural products (Nafie, 2008; Batista et al., 2011b; Debie et al., 2011; Felipe et al., 2012; Lourenço et al., 2012; Cheng et al., 2020; Nafie, 2020).

Raman optical activity (ROA) spectroscopy is hampered by lower sensitivity than VCD for small molecules; however, there are some possibilities for enhancing its signal. Here, we note a recently developed mechanism wherein chirality is enhanced using the resonance that results from supramolecular aggregation. This mechanism is aggregation-induced resonance Raman optical activity (AIRROA). As an example, J-aggregates of astaxanthin (AXT) strongly absorb circularly polarized light in the range of ROA excitation at 532 nm. The implications of aggregation-induced signal enhancement for chiroptical spectroscopy are discussed by Nafie (1996), Vargék et al.

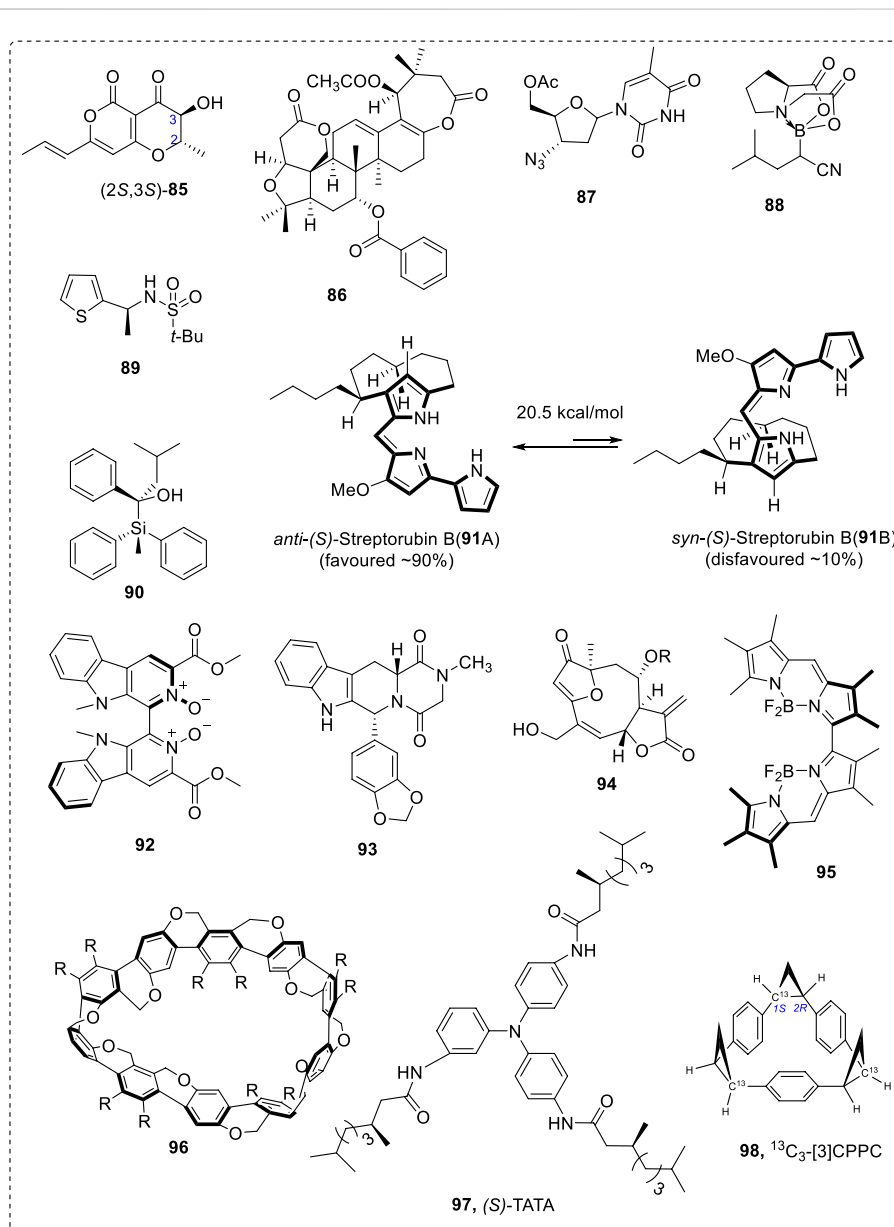


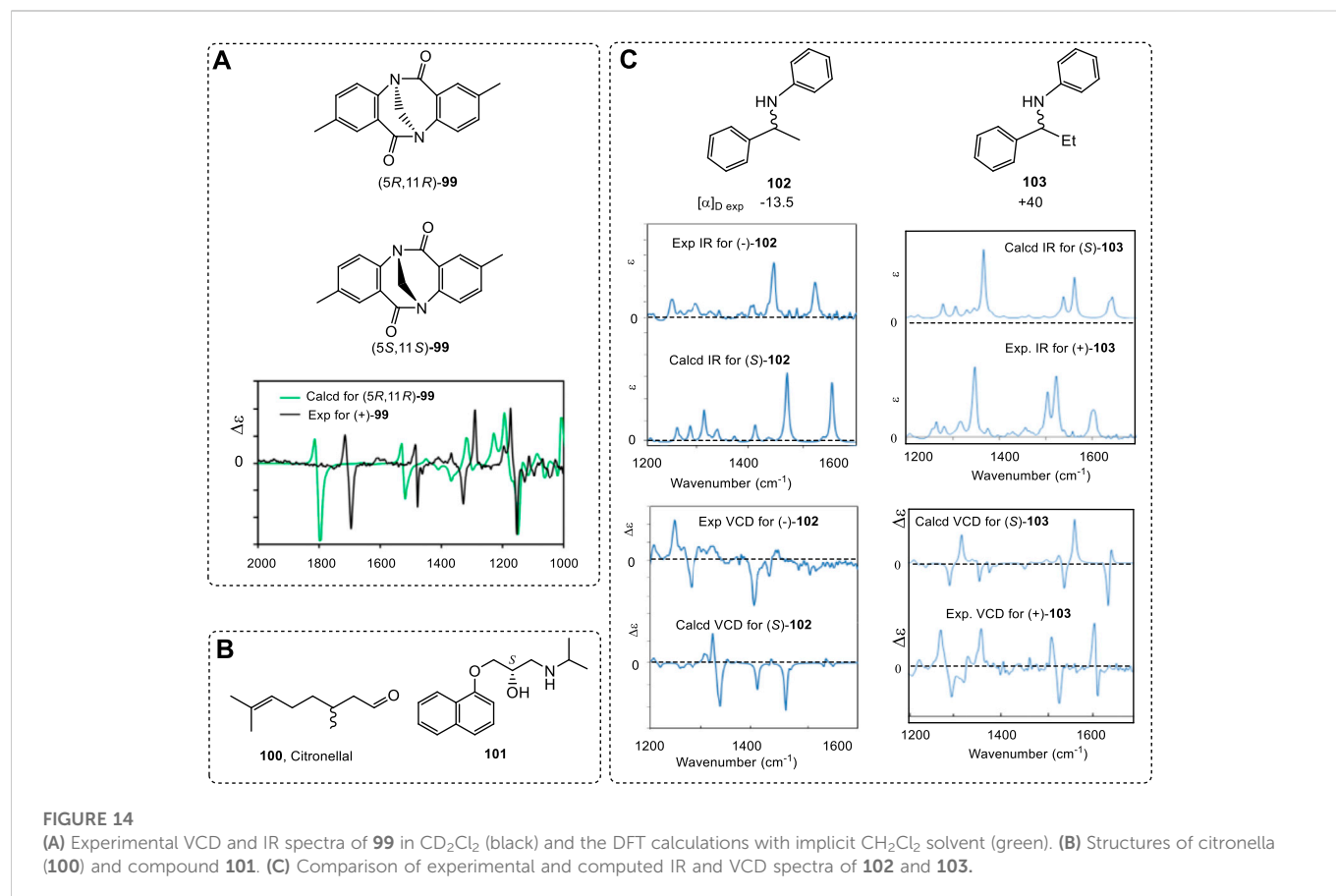
FIGURE 13
AC assigned for some typical structures 85–98.

(1998), and Zajac et al. (2016). Some ROA studies have been reported (Daugey et al., 2014), including the AC determination of riboflavin (Pinto et al., 2015), which is a kind of cyclopeptide with moderate effectiveness ($IC_{50} = 42 \mu\text{M}$) against the *Plasmodium falciparum* strain 3D7. However, this method has not been widely used in AC study until now, and will not be further discussed in this review.

It has been theoretically shown using nuclear velocity perturbation theory that crystal packing induces the enhancement of VCD or other methods (He et al., 2001; Nafie, 2004; Domingos et al., 2014a; Jahnigen et al., 2018). A so-called “switchable amplification of VCD” for chiral structures was reported to amplify VCD signals (Domingos et al., 2014b). Another way is to use lanthanide tris(β -diketonates) as useful probes in the stereogenic center determination of biological amino alcohols near 1500 cm^{-1} (β -diketonate IR absorption region) due to

the formation of ternary complexation with racemic lanthanide tris(β -diketonates) (Miyake et al., 2014).

In some cases, flexible chiral compounds, such as β -peptides and foldamers, could be analyzed by careful conformational study (Farkas et al., 2019)—for example, compounds 85 (Santoro et al., 2019) to 98 (Mazzeo et al., 2017a; Taniguchi et al., 2017; Gimenes et al., 2019). Typical structures are 86 (Gimenes et al., 2019) and 87 (Taniguchi et al., 2017), boron-containing chiral compound 88 (Mazzeo et al., 2017a), chiral thiophene sulfonamide 89 (Rode et al., 2017), chiral Si-containing compound 90 (Xia et al., 2018), pseudoenantiomeric atropisomers *anti*-(S)-streptorubin B (91A) and *syn*-(S)-streptorubin (91B) (Andrade et al., 2015), axial chiral ligand 92 (Zhu et al., 2015), 93 (Qiu et al., 2013), bioactive compound 94 (Junior et al., 2015), axial N-BF₂ compound 95 (Abbate et al., 2017), Möbius-shaped cycloparaphenylenes 96 (Nishigaki et al., 2019), and



axial 3,3'-bithiophene atropisomeric scaffold **97** (Gabrieli et al., 2016). The chirality of isotope of ¹²C and ¹³C-containing compound (**98**) can be assigned using VCD (Miura et al., 2019). These compounds nearly cover most important organic compound types (Figure 13). Many other natural products have been well investigated using VCD methods (Nafie, 2011; Ji et al., 2014; Kessler et al., 2014; Mazzeo et al., 2015; Taniguchi et al., 2015; Cheng et al., 2016; Covington et al., 2016; Zhang et al., 2016).

As a useful tool, VCD has been widely examined experimentally and theoretically by theoretical chemists (Ortega et al., 2015a; Nafie et al., 2018; Nafie and Laane, 2018). An interesting example is the *N*-chirality molecule, the analogue (**99**) of Tröger's base. It can be obtained using HPLC with mixtures of CH₂Cl₂ and *n*-heptane as eluents (Runarsson et al., 2015). A frequency-scaling factor was not applied to the predicted spectra for correcting the calculated spectra appearing at a higher wave number (Figure 14A). In this case, the calculated VCD/IR spectra were calculated at the B3LYP/6-311++G(d,p) level of theory, and the calculation results matched the experimental results well.

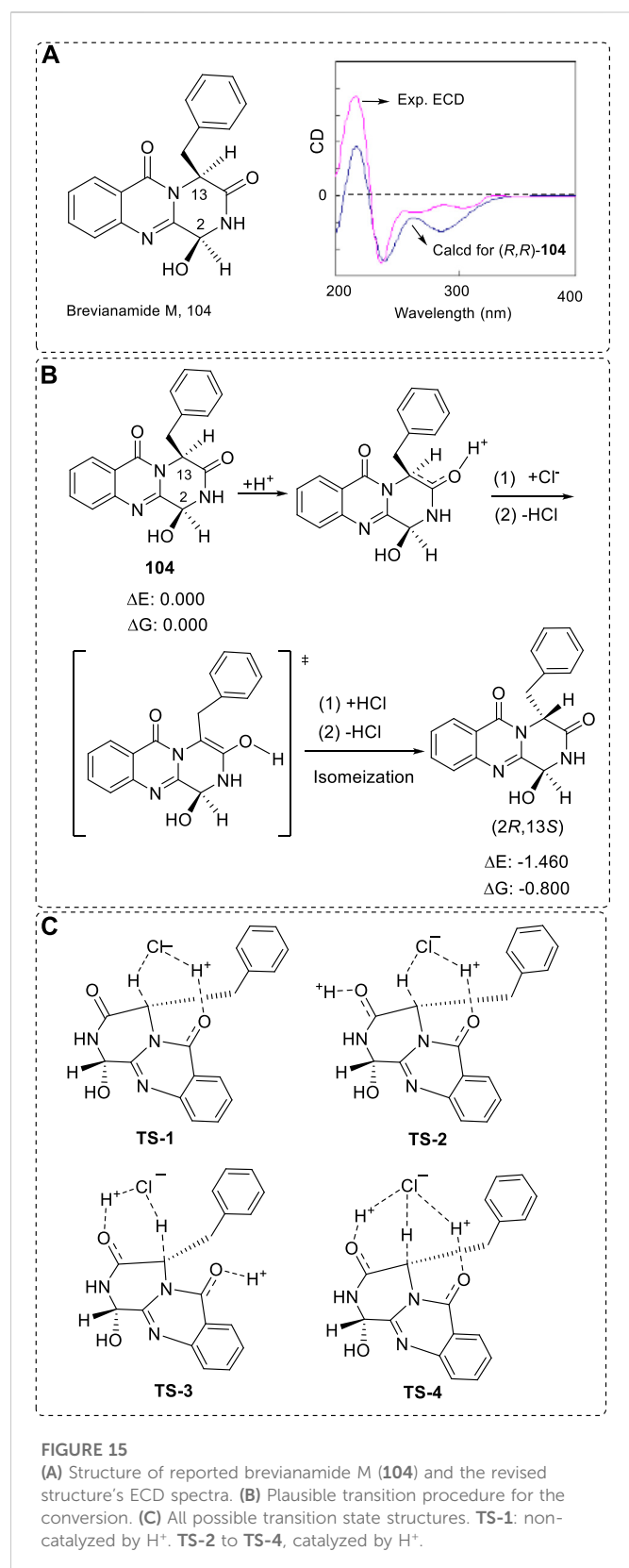
As mentioned previously, it is hard to find all conformers for linear chiral compounds. It is also difficult to compute all found geometries in simulations of VCD. The conformer numbers increased quickly with the free rotated single bonds in linear chiral compounds. In one example, the conformers of citronellal (**100**, Figure 14B), a kind of flavor, were 162 in liquid (Koenis et al., 2019). The total geometries were used in VCD simulations with good agreement between the experimental and computed spectra. Similar examples included

compound **101** (Górecki et al., 2017) and others (Zinna and Pescitelli, 2016).

In some cases, chiral molecules are quite large. As with ECD study, significant progress in the study of VCD using quantum mechanical theory for large molecules has been made in the past decade using fragment-based methods (Gordon et al., 2012). The molecules-in-molecules (MIM) fragment-based method is now suggested for handling large molecular VCD treatment (Mayhall and Raghavachari, 2011; Jose et al., 2015). It employs a multilayer method with multiple levels of theory using a generalized hybrid energy expression, or when considering many solvent molecules mixed with chiral molecules using QM/MM methods (Ghidinelli et al., 2018). This is similar to the ONIOM methods used in optimization for large molecules. For more details, readers can refer to the corresponding reports.

For 1,2-diol compounds, VCD can use the di-Mo-intermediates as used in ECD (Jawiczuk et al., 2015). On the other hand, converting the molecule into a five-ring structure is also helpful when the two hydroxyl groups are located on a linear carbon chain, such as pseudotriin A₃ with an unusual hetero-spirocyclic system (Xu et al., 2016c).

To compute a VCD spectrum requires calculating both atomic polar and atomic axial tensors based on the force constant matrix, which is generally time-consuming. Density functional-based tight binding (DFTB) theory was recently developed by Visscher (Teodoro et al., 2018), making it possible to obtain accurate VCD spectra with a much lower computational demand than standard DFT methods.



As mentioned above, the ECD spectrum is sensitive to chromophores. In a series of chiral analogues, these molecules have very similar ECD spectra. However, the corresponding VCD spectra are quite different. Even though two chiral compounds have extremely

similar structures, they may have totally different VCD spectra. For example, compounds **102** and **103** are two extremely similar analogues (Zhu and Zhao, 2015; Cao et al., 2017), but, while their ECD spectra are similar, their VCD spectra are extremely different (Figure 14C).

Therefore, it is impossible to compare the VCD spectra among a similar series of chiral compounds to assign their AC by comparing the unknown compound's VCD with the known structure's VCD. One must compute the VCD using the quantum methods and then compare this with the experimental result.

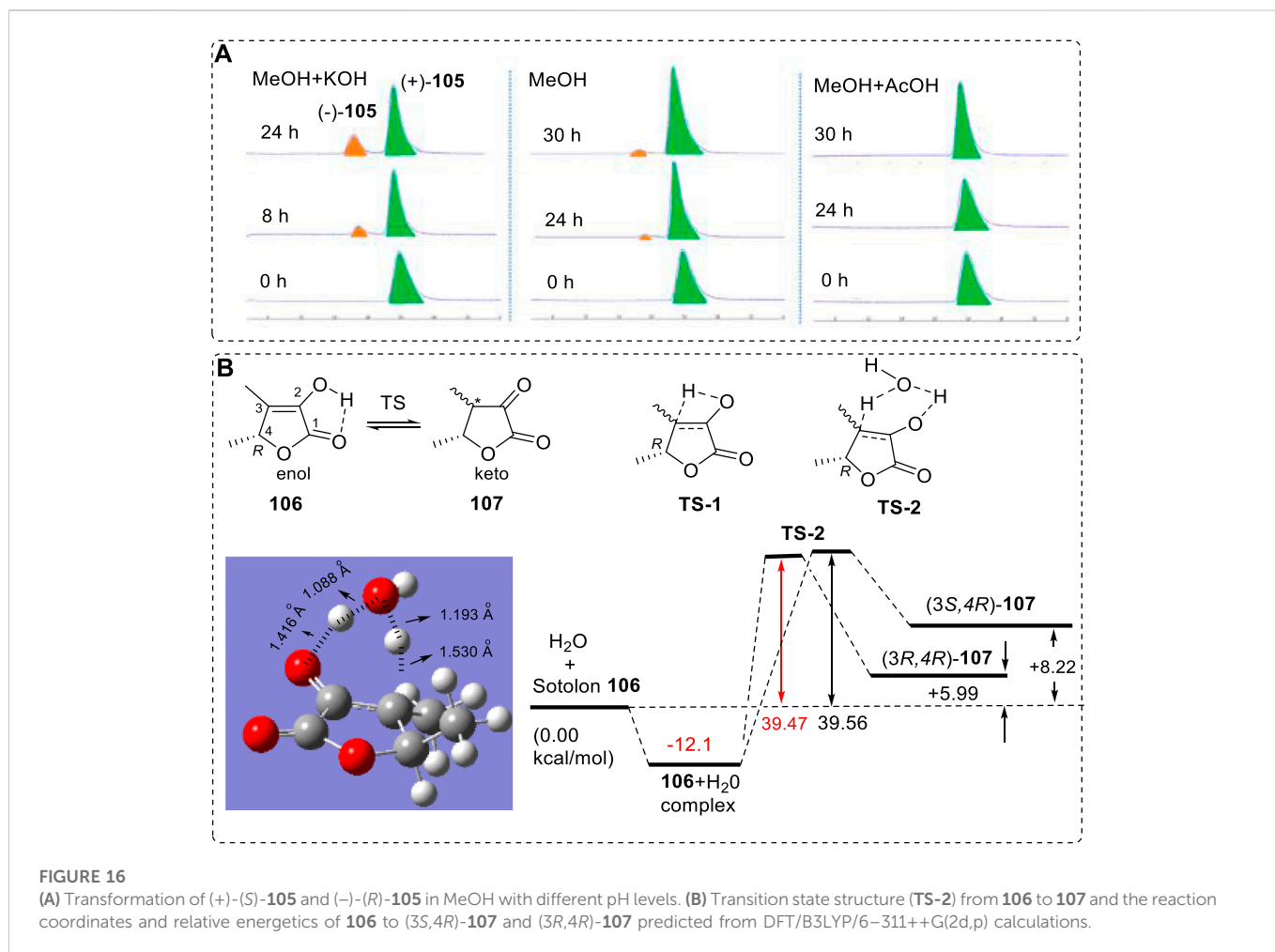
An advantage of VCD is that it can be used to treat the interaction of two molecules in space. One early report studied the structural arrangement of four molecules of 6,6'-dimethyl-[1,1'-biphenyl]-2,2'-dicarboxylic acid in space (Urbanová et al., 2005). It is now possible to study the self-assembly of supramolecular polymers of *N*-centered triarylamine trisamides (Koenis et al., 2020). In a catalytic reaction, solvent-induced conformational changes of the thiourea and the evidence for an unexpected binding topology were discussed between the thiourea and an acetate anion interaction in space (Kreienborg et al., 2016). In real systems, the interaction of water with chiral molecules such as dipeptides is quite strong. Careful investigation of the water-molecule interaction has been performed. For example, there are six water molecules around the alanine dipeptide using the VCD probe (Mirtic et al., 2014). Many similar cases, like conformational equilibrium in (-)-*S*-nicotine, have been reported (Ortega et al., 2015b).

5 Transition state (TS) computation for assistance in AC determination

When determining the AC of various chiral compounds, other methods, such as transition state barrier calculation, may prove useful for understanding the assignments of some special compounds. For example, the alkaloid brevianamide M (**104**) was assigned as a (2*S*,13*S*) configuration using X-ray (Mo radiation) and hydrolysis experiments to afford L-phenylalanine (Figure 15A) (Li et al., 2009). This assignment is very solid. The recorded experimental OR was -146; however, the correct AC was (2*R*,13*R*) instead of (2*S*,13*S*). This re-assignment was performed using the OR, ECD (using Gibbs free energy data in Boltzmann statistics), and VCD methods (Ren et al., 2013). Computations were performed at the B3LYP/6-311++G(2d,p)//B3LYP/6-311++G(2d,p) level. The correct structure was finally confirmed by X-ray using Cu-radiation. It is a very interesting question as to why using the AC of L-phenylalanine from the hydrolysis of (-)-**104** led to a wrong conclusion.

This may possibly be because of the configuration conversion of C13 from *R* to *S* during the hydrolysis of (-)-**104** (Figure 15B) via four possible TS structures (**TS-1** to **TS-4**), since the two groups of -OH and benzyl ring are *cis* orientated (Figure 15C). In this procedure, if the final product's energy is higher than the starting material, this conversion would not happen. Thus, firstly, the relative energy of (2*R*,13*R*)-**104** and (2*R*,13*S*)-**104** must be compared before the TS analysis. Fortunately, the (2*R*,13*S*)-**104** was lower by 0.8 kcal/mol than the (2*R*,13*R*)-**104** using Gibbs free energy at the B3LYP/6-311++G(2d,p) level in the gas phase after conformational searches. Hence, if the TS barrier is not high enough, 92% of the (2*R*,13*R*)-**104** using total electronic energy or 80% using Gibbs free energy would convert into (2*R*,13*S*)-**104**.

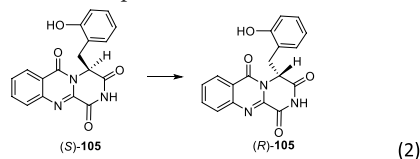
The calculated barrier was about 25.1 kcal/mol via **TS-1** in water without any acidic promotion. The acidic catalysis procedures via **TS-2** and **TS-4**



were 16.5 and 17.9 kcal/mol, respectively. This barrier of 25.1 kcal/mol may mean that the reaction may possibly happen when the water temperature increases to about 65°C. After the configuration conversion, the next step is the hydrolysis of (2R,13S)-brevianamide M in water—this barrier was only 21.3 kcal/mol in water. Thus, after the configuration conversion, it immediately decomposed into the corresponding L-phenylalanine.

It is very popular to hydrolyze one complex compound into one or two simple known compounds in order to assign the complex compound's structure. Although there are not many examples with configuration conversion, the possibility does exist in some examples. To raise vigilance in daily study may help reduce the possibility of wrong conclusions.

Interestingly, another similar example (**105**) was reported (Eq. 2). This compound racemized in room temperature under a strong base KOH condition (Figure 16A) (Xu et al., 2021). However, it is stable under acetic acid at room temperature. This shows that acidic promotion needs about 25 kcal/mol (HCl). It would be better if there were a TS barrier calculation for this conversion; in this case, the barrier could be compared with the barrier under acidic catalysis.



In contrast to the above case, another example can exclude the possibility of a natural product to form the other structure. For example, Sotolon (**106**) is a naturally occurring chiral furanone with a well measured VCD. However, the predicted VCD was little different from the experimental results at 1080 and 1060 cm^{-1} (Nakahashi et al., 2011). Should it then form the corresponding enol structures and should the enol structure bring the corresponding contribution to the VCD spectra? Before investigating the OR, ECD, and VCD spectra of **106**, the possible conversion of enol **106** to its keto tautomer **107** was theoretically investigated. (*R*)-**106** could transform into either (3R,4R)-**107** or (3S,4R)-**107** (Figure 2), depending on the values of the different transition state (TS) barriers in the pathway. Direct conversion *via* TS-1 would face a large barrier since it involves a four-membered ring in TS. Therefore, only TS-2 was considered. The barrier to generate (3R,4R)-**107** was 39.48 Kcal/mol while the formation of (3S,4R)-**107** required overcoming a 39.56 Kcal/mol barrier (Yang et al., 2017). Both barriers are high, suggesting that this conversion is impossible. Furthermore, even if this barrier could be overcome, the keto forms (3R,4R)-**107** or (3S,4R)-**107** were predicted to be 5.99 or 8.22 Kcal/mol less stable, respectively, than the sum of the starting material **106** and a water molecule (Figure 16B). Hence, **106** is predicted to greatly dominate over either (3R,4R)-**107** or (3S,4R)-

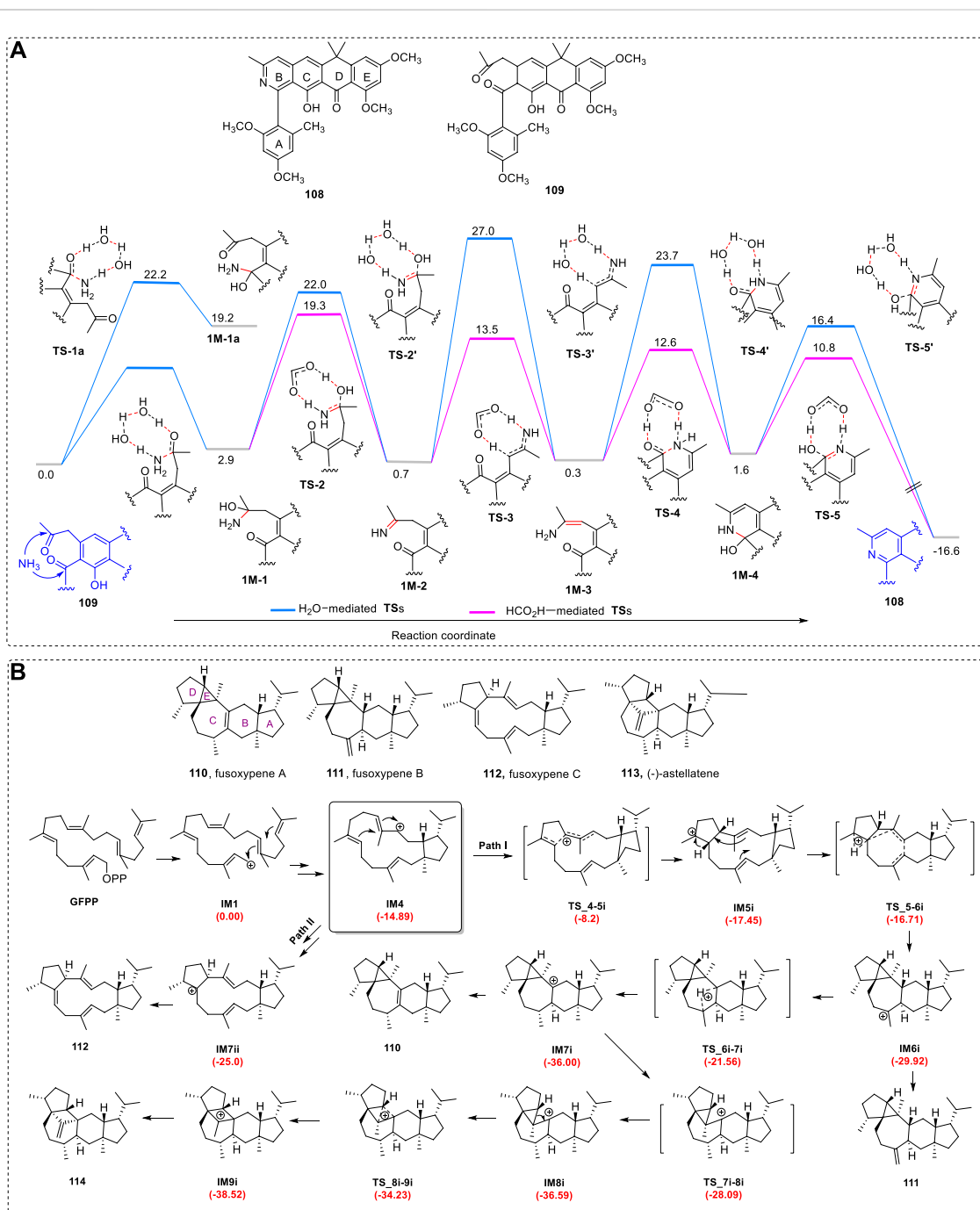


FIGURE 17

(A) Reaction coordinate of 1,5-dicarbonyl condensation with ammonia for conversion of **109** to **108** (data represent ΔG values in kcal/mol). (B) Simplified computed reaction pathways of **110**–**114**. IM4 is the bifurcation point to generate Path I and Path II. IM = InterMediate. The potential energy profiles relative to IM1 are shown in red.

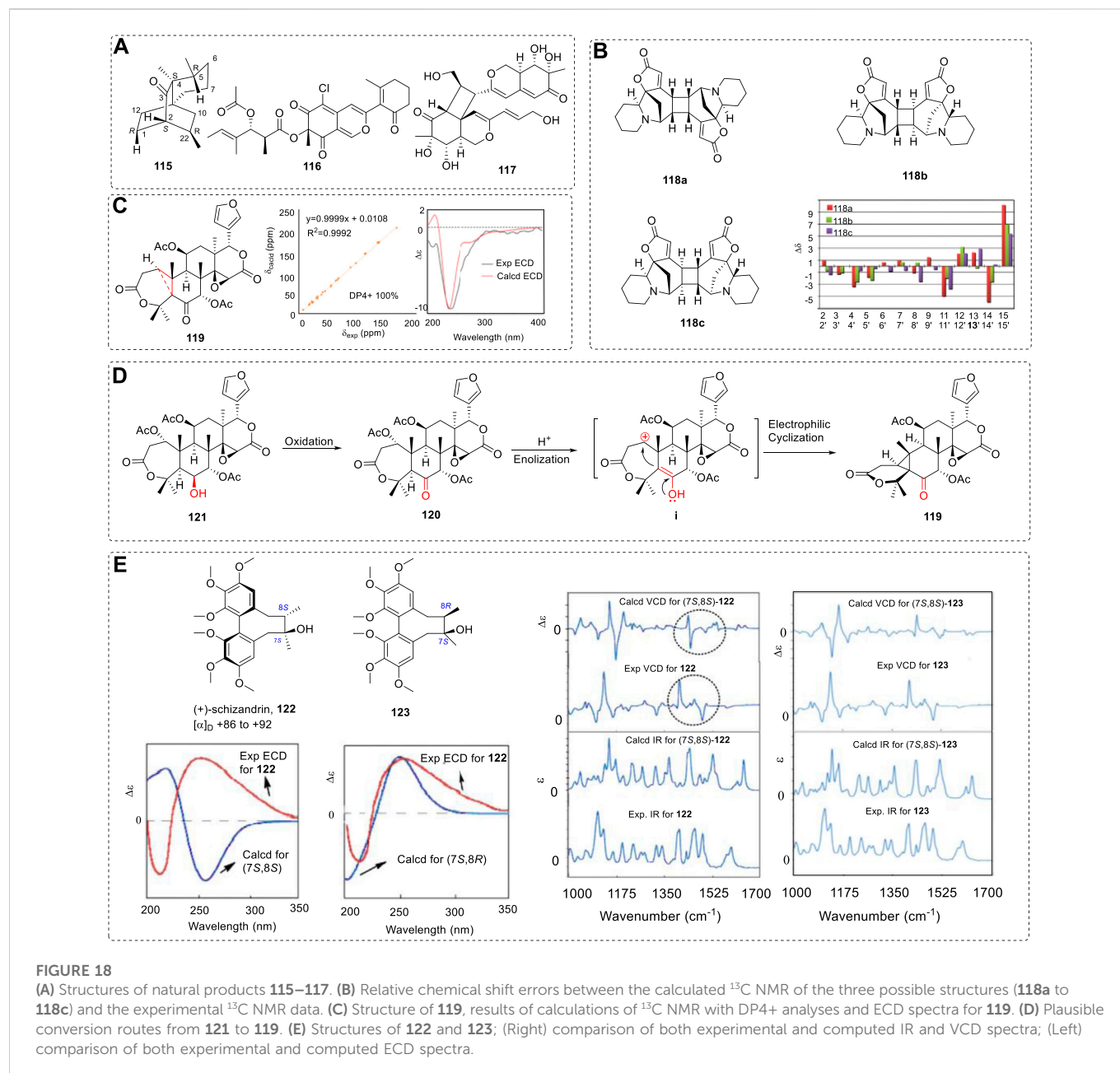
107 at equilibrium and it or its other structural form may be considered in its OR or CD calculation.

Obviously, using TS barrier data can provide many benefits for understanding the AC assignment procedure. It is also obvious that computing TS barriers is not only the job of synthetic chemists. Correct use of TS barriers can accelerate the understanding of AC study.

In some cases, the biogenetic pathway is an important way of performing the organic synthesis of some bioactive compounds. Theoretical methods can provide, to a certain degree, direct

knowledge of whether the pathway is possible. For example, after obtaining the diastereoisomers **108** and **109**, compound **108** may be converted from **109** via an ammonia molecule present in the reaction (Figure 17A). The computed TS coordinates for the conversion were fully investigated and the largest TS barrier was only 19.2 kcal/mol, using the B3LYP-D3BJ/def2-SVP level of theory with the PCM for H₂O.

The genome-based discovery of two previously unreported fungal bifunctional terpene synthases (BFTSs) from phytopathogenic fungi



has been reported. The formation of fusoxypenes A–C (**110**–**112**) and (–)-astellatene (**113**) has been investigated using quantum theory (Figure 17B) (Jiang et al., 2021b). The Gibbs free energies were calculated from the electron energies at the mPW1PW91/6–31+G(d,p) level plus the thermal correction at the M06-2X/6–31G(d,p) level.

6 Comprehensive methods in AC determination

Generally, a single method may settle some chiral molecular ACs. However, for some chiral compounds, one method cannot provide enough evidence to support the AC assignments. It is thus better if two, three, or all of the OR, ECD, VCD and ^{13}C NMR methods are used. This is more important for determining the AC of chiral

medicines, where many drugs have been investigated using various methods (Polavarapu, 2016). Example **6** used the ^{13}C NMR and OR methods. There are many examples used with various combinations of the tools—for example, rare sesquiterpene (+)-3-ishwarone (**115**) was investigated using NMR, ORD, ECD, and VCD (Junior et al., 2014; Hu et al., 2021; Lee et al., 2021), and other examples include **116** and **117** (Cao et al., 2019; Cao et al., 2020b) (Figure 18A).

Flueggidine, isolated from *Flueggea virosa* (Zhao et al., 2013), has one set of NMR but two units of molecular weight—a very rigid compound. During the conformational searches, one conformer was manually obtained. The theoretical ^{13}C NMR data of **118a**, **118b**, and **118c** were calculated at the B3LYP/6–311++G(2d,p)//B3LYP/6–31+G(d) level in the gas phase. Structure **118a** must be excluded due to the relatively high chemical shift at up to 10.2 ppm at C15 (Figure 18B). Structure **118c** had the smallest relative errors at C15; its linear coefficient should be the largest. The calculated OR value of

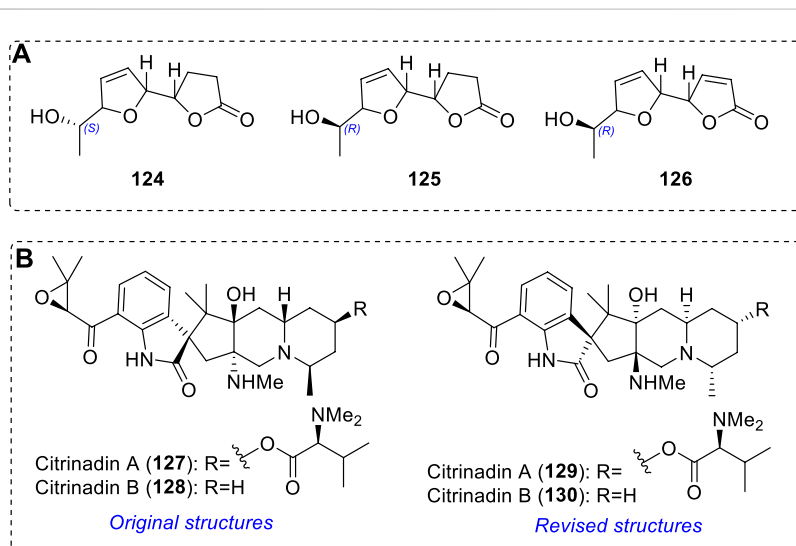


FIGURE 19
(A) Structures of 124 to 126. (B) Original structures 127 and 128, and revised structures 129 and 130.

118c (-38.8) was close to the recorded one (-33.5). In contrast, the calculated OR magnitude for **118b** was +62.5. Thus, the stereochemistry of **118** was determined to be **118c**—2(2′)*S*, 7(7′)*R*, 9(9′)*R*, 14(14′)*S*, and 15(15′)*S*.

For an unusual aglatestine **A 1** (**119**) with a cyclopropenyl moiety, its relative and AC was assigned *via* ^{13}C NMR and ECD methods (Figures 18B, C) (Sun et al., 2021c). In the series of analogues, it was found that the three compounds obtained may have biosynthetic routes (Figure 18D). The relative configuration of the three compounds matched well and this confirmed the relationship among them (**119**–**121**).

However, the use of ECD for some specific chiral compounds for their AC assignment may not be enough. For example, schizandrin **122** was assigned as (7*S*,8*S*) in a previous report but was corrected as (7*S*,8*R*). Its ECD spectra is illustrated below (Figure 18E), predicting that its original configuration assignment was wrong. Based on the calculated ECD spectrum curve shape, it is tempting to predict its real configuration as (7*R*,8*R*), the opposite of the original assignment. However, VCD study shows that its real configuration is (7*S*,8*R*) (Figure 18E). In this case, the predicted ECD for **123** matched well with the experimental results in both ECD and VCD studies (He et al., 2014). The DFT calculations were performed at the B3LYP/6-311+G(d)//B3LYP/6-311 + G(d) level in the gas phase.

In some cases, the chiral molecules are not large and rigid but are inflexible, such as diplobifuranylonones **A–C** (**124–126**) (Figure 19A) (Mazzeo et al., 2017b). In the assignment procedures, the Mosher method was used while the ECD, VCD, and ORD were applied for the AC study.

It is now very popular to use two or more chiroptical spectroscopies in AC determinations (Reinhardt et al., 2019). Some chiral compounds have their stereogenic centers on a linear carbon chain. In these cases, Mosher methods may provide good assistance during AC assignment—for example, the compounds **1** and **4** (El-Kashef et al., 2019). It can be seen that the groups near the stereogenic center (on chain) are not as large as in compound **5**.

Two bioactive spirooxindole alkaloids, (–)-citrinadin **A** (**127**) and (+)-citrinadin **B** (**128**), were isolated and reported (Tsuda et al., 2004; Mugishima et al., 2005). Their AC assignment used the ROSEY, ECD,

and VCD methods. However, after a total synthesis of both compounds, it was found that the correct structures should be **129** and **130** (Figure 19B) (Bian et al., 2014). It is unresolved why the study of combinational uses of NMR, ECD, and VCD do not afford the correct ACs.

Organic synthesis is still a good way of testing the stereochemistry of some compounds accompanied by quantum calculation. The revision of the absolute stereochemistry of pycnidione was achieved *via* the NMR and ECD methods (Bemis et al., 2021).

With the development of computations and synthetic methods, the AC assignments of many different chiral compounds have used two more chiroptical spectroscopies (Zhang et al., 2021c). Some typical examples are listed in Table 1.

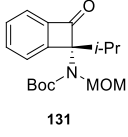
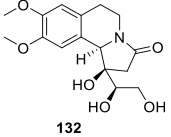
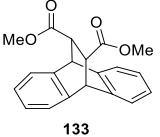
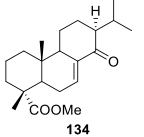
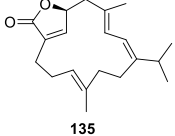
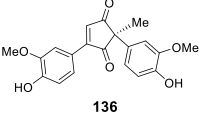
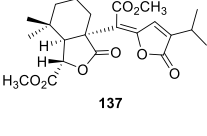
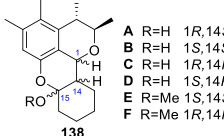
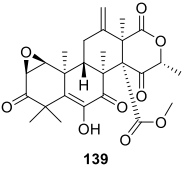
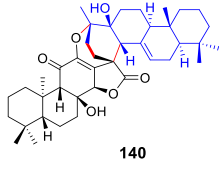
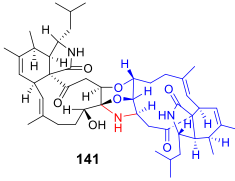
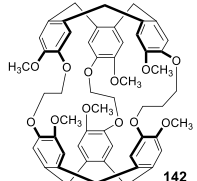
7 Simplified models

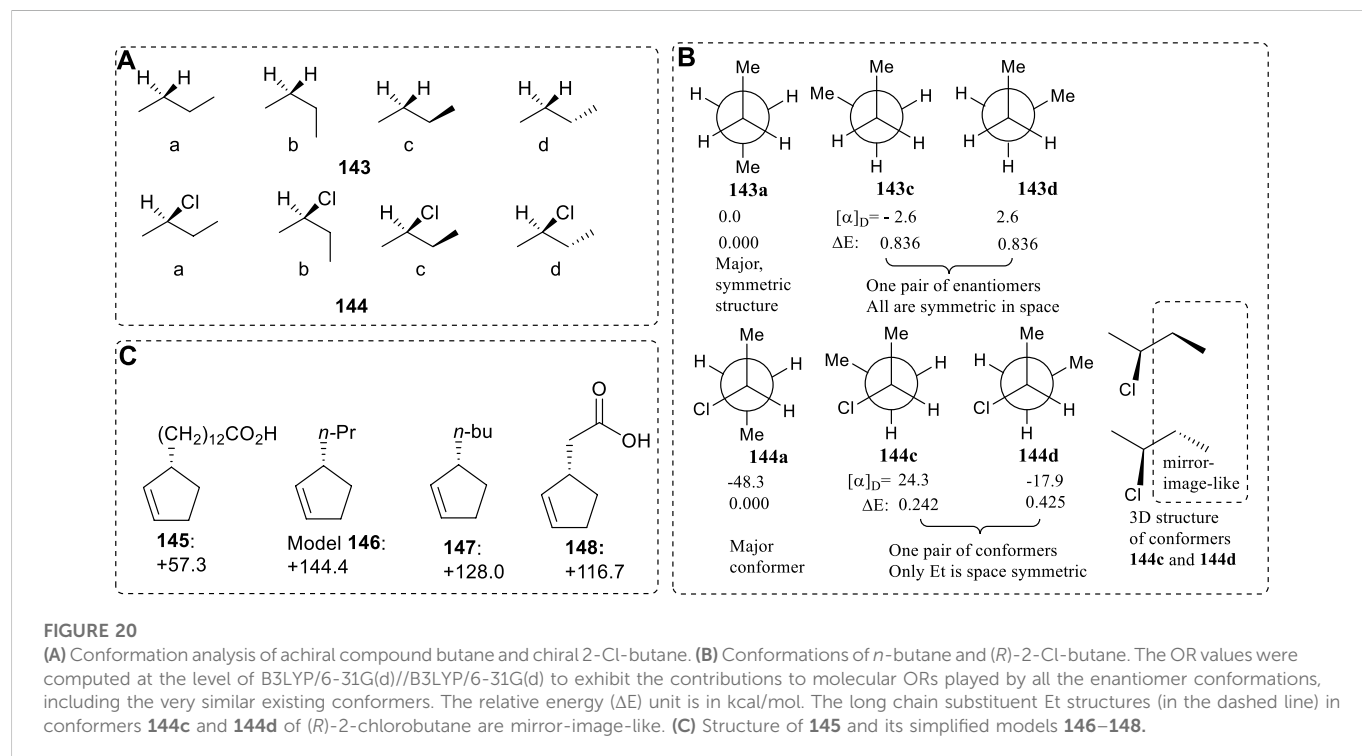
Use of a simplified model is popular in the interpretation of chiroptical spectroscopies or in TS barrier calculations. For example, researchers frequently use a vinyl group or allyl to replace a benzyl group in a TS calculation. A methyl, an ethyl, or a propyl is used to represent a long carbon chain which has no stereogenic center. There are two possible reasons for researchers to collect in this way. One is that current computational methods, even DFT methods, are not the most accurate: they are all approximate calculations which introduce computational errors in energy and chiroptical spectroscopy calculations. Another reason is that there are fundamental reasons to shorten the long carbon chain to a short chain, or a methyl. Although the concept has been suggested in OR computations (Zhao et al., 2019; Zhu, 2022b), this principle could be extended to ECD or VCD calculations using simplified models.

7.1 Fundamental basis for using models

The aim of using simplified models in chiroptical calculations is to reduce computational time without greatly reducing the accuracy of

TABLE 1 AC assignment for structures of 131 to 142 using various methods.

Molecule	Molecule	Molecule	Molecule
 131	 132	 133	 134
ECD, VCD Kasamatsu et al. (2017)	ORD, ECD, and VCD Johnson et al. (2019)	ECD, VCD Passarello et al. (2014)	ECD, VCD Masnyk et al. (2018)
 135	 136	 137	 138 A R=H 1R,14S,15S B R=H 1S,14S,15R C R=H 1R,14R,15S D R=H 1S,14R,15R E R=Me 1S,14S,15R F R=Me 1R,14R,15S
ECD, X-Ray Li et al. (2019b)	ECD, VCD Du et al. (2012)	VCD, X-Ray Yun et al. (2014)	NMR, ECD Cheng et al. (2021)
 139	 140	 141	 142
ECD, X-Ray Fang et al. (2021)	ECD, X-Ray He et al. (2021b)	¹³ CNMR, ECD, X-Ray Wang et al. (2021c)	ECD, VCD, ROA Brotin et al. (2015)

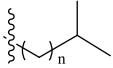
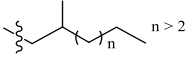



the calculation. This involves a new concept: a conformation pair (or conformer pair) (Zhao et al., 2019; Zhu, 2022b). For example, butane (*S*) has four conformers: **a**, **b**, **c**, and **d** (Figure 20A). Conformation **c** has no symmetric element, so its OR will not be 0, as with **d**. Since **c** and **d** are mirror-image symmetric, their relative energy and OR

magnitudes will be same, but the OR signs will be reversed. Thus, the net OR value from **c** and **d** is 0. Therefore, the total OR for 143 will be 0.

In the next example, (*R*)-2-chlorobutane 144 (Figure 20A) has one pair of conformers (144c and 144d) that have symmetric Et

TABLE 2 Suggested simplified groups used to replace the big groups in the calculations.

Entry	Real group	Simplified group
1	$-(\text{CH}_2)_n\text{CH}_3$, n is odd	$-\text{CH}_2\text{CH}_3$
2	$-(\text{CH}_2)_n\text{CH}_3$, n is even	$-\text{CH}_2\text{CH}_2\text{CH}_3$
3	 $n > 2$	$-\text{CH}_2\text{CH}_3$
4	 $n > 2$	
5	Phenyl ring	2-propenyl
6	Poly-substituted phenyl ring	One-substituted phenyl ring
7	Poly-phenyl rings	Single phenyl ring
8	$-(\text{CH} = \text{CH})_n\text{CH}_3$, $n \geq 2$	$-\text{CH} = \text{CH}_2$
9	$-(\text{CH} = \text{CH})_n\text{CH}_2\text{-(CH} = \text{CH})_m\text{CH}_3$, $n \geq 2$, $m \geq 2$	$-\text{CH} = \text{CH}_2$ or $-\text{CH} = \text{CHCH}_3$
10	$-\text{CH}_2\text{C} = \text{CH}(\text{CH}_2)_n\text{CH}_3$	$-\text{CH}_2\text{C} = \text{CH}_2$

substituent structures (Figure 20B); this pair has opposite OR signs. This is very similar to the case in **143c** and **143d**, which are complete mirror-images. The symmetric structures that appear in **144c** and **144d** exist within their Et substituents. Therefore, the sum of the OR contributions of conformations **144c** and **144d** have a very small magnitude of only 7.4 degrees, calculated from their individual OR values at the B3LYP/6-31G(d)//B3LYP/6-31G(d) level and based on their relative conformer populations derived from their relative energies. Thus, the calculated OR ($[\alpha]_D$) contribution (7.4°) of this **144c** and **144d** pair of conformers was only 15% of the OR value of -48.3 from conformation **144a**.

Conformer pairs like **143c** and **143d**, or **144c** and **144d**, are expected in any chiral linear molecules existing in solution. The final contribution from a summation of all the enantiomeric conformer pairs and the closely symmetric pairs will only represent a small fraction of the total measured molecular $[\alpha]_D$ value. The major $[\alpha]_D$ contribution is from the most stable conformer, as illustrated by conformer **144a** in (*R*)-2-chlorobutane.

As the number of a chain's carbons increase, the number of stable conformations increases by $3^{(n-2)}$ (here n is the number of single bond rotations and must be larger than 3). This formula is applicable to many chiral linear compounds like the above molecules. No matter how long the chain is, only a unique conformer with the lowest energy exists; it has the largest OR values. The conformer pairs must exist in double numbers, and their Boltzmann statistic fraction will not be larger than the most stable conformation's OR values. Theoretically, the major OR contribution is from this OR: the most stable conformer's OR sign closely represents the chiral molecular AC. Thus, the fundamental basis for using simplified models is the existence of conformer pairs for chiral compounds with a long chain.

When the carbon chain numbers are more than six, it is difficult to find all conformations. This may lead to computational errors since some conformer pairs, where there is not another conformer to form a pair to the conformer pair, are not found in the conformational search. At the same time, every theoretical method must have its accuracy level in energy and OR computations. This point again entails

computation errors. Therefore, both sides entail such errors. Thus, when using a short carbon chain to representative a long chain, it is easier to simulate a large OR value than use a long carbon chain. To use an ethyl to representative a long chain with an even carbon number or a propyl to represent an odd carbon chain number may be a good choice. For example, the chiral compound **145** was assigned based on the following the OR values of three model molecules (models **146–148**) (Figure 20C) (Wang et al., 2010). The computed OR of **147** with four carbons was almost 16 degrees smaller than the compound **146** with a three-carbon chain. If the side chain was simplified as an acetyl (**148**), its OR became 8 degrees less than **147**. All of the models have over-estimated OR values compared to the OR of the actual molecule of $+57.3^\circ$. Some suggested simplified substituents to replace the corresponding large and complex substituents are summarized in Table 2.

7.2 Examples using simplified models in OR, ECD, and VCD study

Although the use of simplified models is determined above from the OR, it can also be used in related calculations for ECD and VCD. Many natural products are too complex or flexible to use their large number of low energy conformations in calculations of ECD, OR, or VCD spectra. Thus, model molecules must be selected that are representative of the original structure. The choice of satisfactory model structures is the focus of this section. For example, the compounds **149** and **150** can be simplified as the models **151** and **152** in the ECD study (Figure 21A) (Tanaka et al., 2017), including ^{13}C NMR simulations.

The highlighted red is the core framework of cyclohelminthols in **149**. It was thus separated as two parts: **151A** and **151B** and X (Figure 21A) for **149**. The **152** part and X part were used for **150**. Conformational searches were performed using semiempirical AM1 MMFF provided for only a limited number of candidate conformers for some models. The DFT theory of B3LYP/def2-

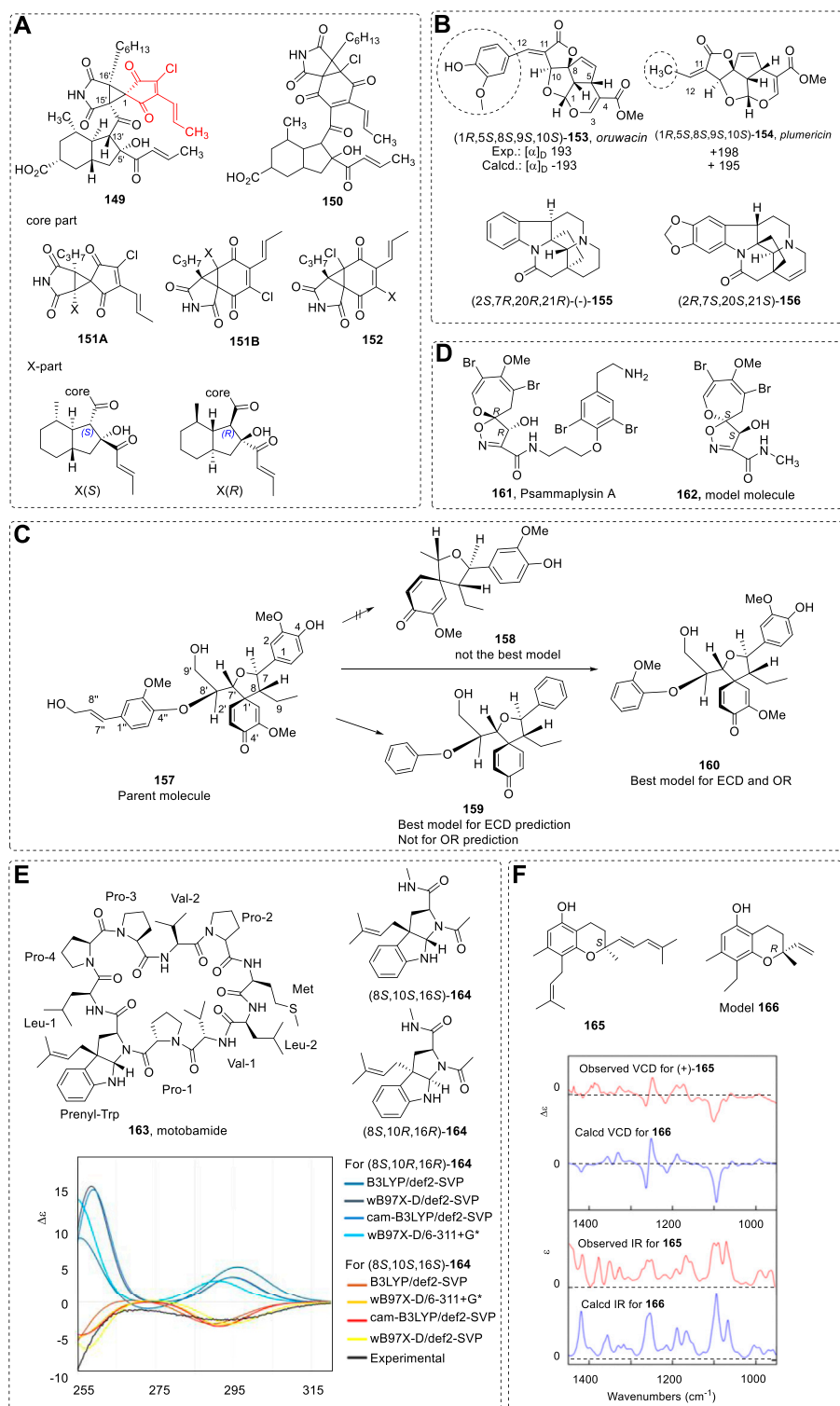


FIGURE 21

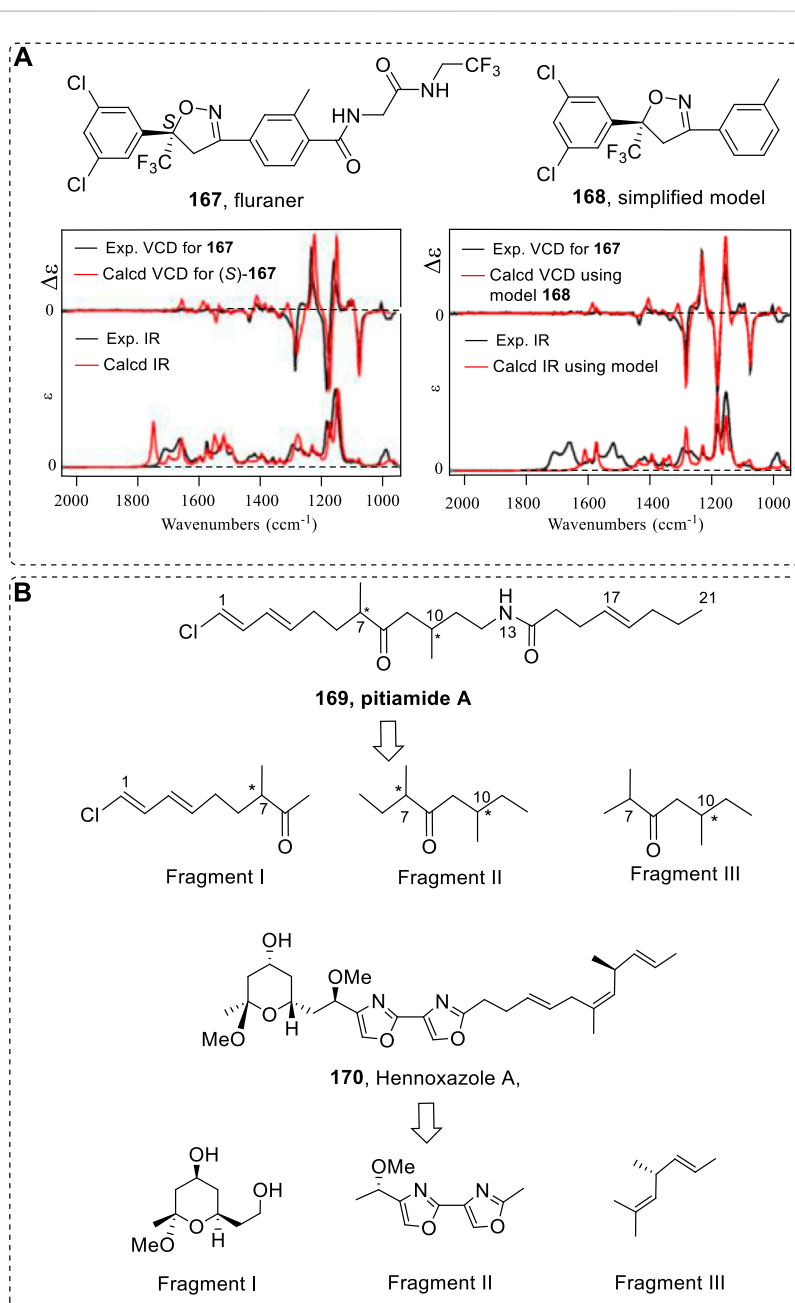
(A) Structures of cyclohelminthol X (**149**) and AD0157 (**150**) and their simplified model parts. (B) Two pairs of similar analogues of **153** vs. **154** and **155** vs. **156**. (C) Natural product **157** and its three simplified models used in OR and ECD study. (D) Psammamplysin A (**161**) and its simplified model compound (**162**) used for ECD prediction. The model is smaller and, more importantly, much less conformationally mobile than the natural product. (E) Experimental ECD spectrum of motobamide (**163**) and theoretical ECD spectra of the model compounds (8S, 10S, 16S)-**164** and (8S, 10R, 16R)-**164**. (F) Structure **165** and its simplified model **166** and their experimental and computed VCD spectra.

TZVP was used for ECD calculations, and ω B97X-D17 was used at the 6-31G* basis set in the ^{13}C NMR calculation.

The two parts were independently computed. Thus, different combinations were used for the calculations (Supplementary Table

S3); good agreement between the theoretical results and the experimental data was achieved.

However, use of a wrong simplified group may result in the wrong assignment. For example, ourwacin and plumerichi are very similar

**FIGURE 22**

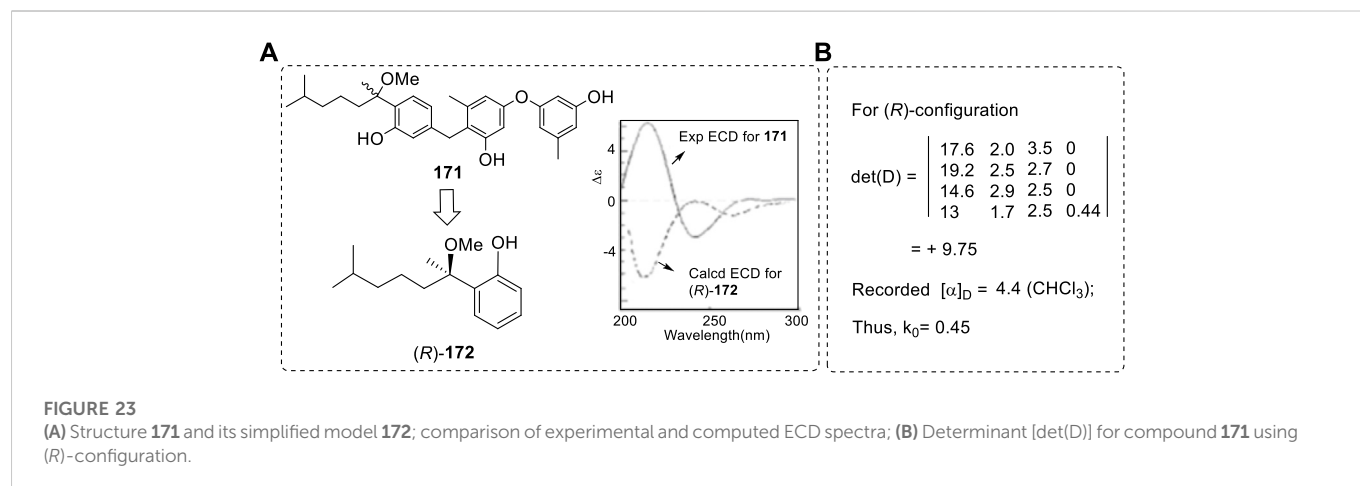
(A) Structures of **167** and **168**; the experimental VCD for (+)-fluralaner (black line) and the calculated VCD (red line) for (S)-fluralaner (left). Comparison of the experimental VCD and the computed VCD using the simplified (S)- model (right). (B) Original structure **169** and **170** and the corresponding separated fragments.

natural products (Stephens et al., 2007a). Ourwacin's (**153**) experimental OR is 193. (1*R*,5*S*,8*S*,9*S*,10*S*)-plumerichi (**154**) had a +198. It is easy to assign (+)-ourwacin as (1*R*,5*S*,8*S*,9*S*,10*S*). If the (1*R*,5*S*,8*S*,9*S*,10*S*)-ourwacin was used in OR calculations, -193 was obtained. However, if (1*R*,5*S*,8*S*,9*S*,10*S*)-plumerichi was used in the calculations, the +195 of OR value was recorded. This clearly shows that a phenyl ring cannot be simplified as a methyl group in calculations. As an opposite example, the compound **155** had -25.4 of OR values (Hajček and Trojanek, 1981; Laguna et al., 1984), and its structural similar compound **156** had $[\alpha]_D$ of +15.5 in CHCl₃ (Renner and Kernweisz, 1963). Its AC was assigned as the (2*R*,7*S*,20*S*,21*S*) using ECD, VCD, and ORD

(Stephens et al., 2007b). It has no phenyl group. The same configuration had the same OR sign.

In this case, it is easy to see that simplifying a phenyl group as a non-aromatic substitute like methyl leads to a wrong conclusion. This can be observed again in another example: **157** (Figure 21C) (Zhao et al., 2016). The model molecule **158** was not the best in the simulations of ECD or OR.

In this model study, consider for computing OR that the -OMe on the phenyl ring may restrict some stable conformation formations. Thus, it is necessary to keep this OMe at its ortho-position instead of replacing it as a H atom since OR is more sensitive to geometries and spatial locations while ECD is more sensitive to the positions of double bond or phenyl ring chromophores.



The structure psammaphysin A (**161**) has a large side chain (Mandi et al., 2015). It was simplified as the model molecule **162** in ECD computations (Figure 21D). In this way, the chromophore is a long distance from the stereogenic centers. Its effect on ECD curves is not large enough to affect the assignment of AC.

A cyclic peptide (motobamide **163**) was simplified as **164** below and used in ECD study (Figure 21E) (Takahashi et al., 2021). This simplification is quite bold, since the >C=O is also a chromophore, although it has a weak contribution to entire molecular ECD spectra. It should be noted that the model is a linear structure and the peptide is cyclic. Although conformer structures of the linear model are expected to be different from cyclic ones, it remains enigmatic whether this difference should yield a ECD difference or not.

As mentioned above, the effect of the carbon chain length on the OR has its limitation, which means that, even if the chain is very long, its OR value will not increase significantly. This clearly leads to the question of why the chain increase does not have an effect on its OR value.

Although this conclusion is obtained from OR calculations, it can be used in other calculations as mentioned above, such as VCD. For example, chiral compound **165** can be simplified as **166** in its VCD calculations (Figure 21F) (Batista et al., 2009; Mota et al., 2009; Batista et al., 2010).

Fluralaner (**167**), a racemic animal health product used to prevent fleas and ticks, was used in a VCD study that used the entire molecule and its simplified model (**168**) (Kong et al., 2017). Six conformers were found using the entire molecule and only two for the simplified model molecule. There is good agreement using the simplified model in simulations of its VCD spectra (Figure 22A).

If a chiral molecule has several stereogenic centers on its carbon chain, such as pitamide A (**169**) (Nagle et al., 1997), one possible method is to cut the structure into several small molecules that have one or two stereogenic centers in OR calculations (Ribe et al., 2000). Another example hennoxazole A (**170**) had been treated similarly (Figure 22B) (Kondru et al., 1997; Kondru et al., 1998). These are only model molecules and it is not easy to cut similar molecules into small molecules in practice if one has had no experience in this study.

Consider a chiral molecule with a single stereogenic center: molecule **171** (Lu et al., 2010). If the entire molecule was used in ECD calculation, the number of conformers would be large. To reduce

the computation time, it was simplified as the model **172** for ECD computations. In this case, the ortho-OH cannot be cut off since it restricts the rotation of the phenyl ring and thereby affects the geometry's overall structure and energy distribution. Simplification is possible by using an ethyl to replace the isopentyl group. Considering that the chiral molecule has only three isolated phenyl rings, its ECD maximum of CD signal must appear at the 220–240 nm range. Here, the long chain may have a relatively large effect on the observed CD. Indeed, to use model B in the ECD simulation agrees well with the experimental results obtained (Figure 23A). This conclusion also agrees with the matrix method results.

The matrix may thus be another approach to this study. For example, molecule **171** yielded a very small OR value of 4.4°. If the entire molecule is used in OR calculations, the large number of conformers needs to be investigated. On the other hand, use of the matrix model is relatively quick. For the current group order (S) configuration, its determinant was +9.75. Based on the positive k_0 value (0.45) for a tertiary-alcohol derivative, it is concluded that it has S configuration. This conclusion agrees with that from the ECD calculations using simplified model molecule **172**.

Many other chiral compounds have been similarly reported. For example, the compound **173** is bioactive; its AC was assigned using Mosher and ECD analysis (Liu et al., 2015). If a simplified model can be used, it may be easily assigned since it can form a six-membered ring structure *via* an intramolecular H-bond if using the model **174** as illustrated below (Figure 24A).

A total of 46 conformers were found for **174** using the Gaussian package. The lowest energy conformer which had an intramolecular H-bond was about 7.4 kcal/mol lower than the lowest energy conformer without this bond (Figure 24B). It is surprising that the energy of the conformer with a side chain located on an *axial*-bond is lower by about 5.4 kcal/mol than the conformer that has its H-bond as an *equatorial*-bond. If the large group is on *equatorial*-bond, the intramolecular conformer is only about 2.0 kcal/mol lower than the one without H-bond and also has the large group located on *equatorial*-bond. To clearly state the conformer question, the conformational search was performed using a MMFF94S force field. All the conformers were then computed at HF/3-21G*. Both kinds of conformer with or without H-bonds with relative energy from 0–2.0 kcal/mol were then re-optimized at B3LYP/6-31G* level in the

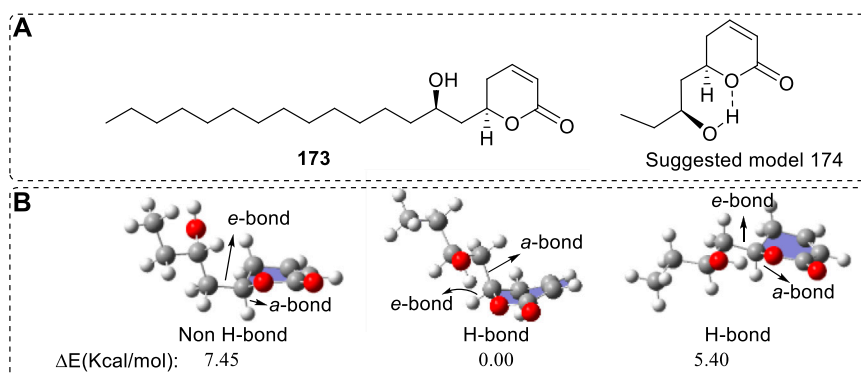


FIGURE 24

(A) Structure of **173** and its simplified model **174**. (B) Energy difference of the conformers with or without intramolecular H-bond (blue section is the ring structure moiety).

TABLE 3 OR simulations of sotolon (**106**) and furanone (**175**) and their dimer structures (**106a** and **175a**).

Form	M1 (deg.)	M2 (deg.)	M3 (deg.)	M4 (deg.)	M5 (deg.)
Monomer of 106	+0.9	-1.8	-8.7	-16.5	-14.5
Dimer of 106a	-34.7	-42.7	-24.6	-20.0	-33.1
Monomer of 175	+74.2	+81.8	+66.7	+62.9	+50
Dimer of maple 175a	+89.7	+79.3	+103.4	+69.0	+75

It is to emphasize the computed data are very close to the experimental values.

gas phase. These B3LYP/6-31G*-optimized conformers with relative energy from 0.0 to 2.0 kcal/mol were finally optimized at the B3LYP/6-311G* level. All optimizations were performed in the gas phase. Since the conformers with low energy are very limited, they could be used for ECD or other spectroscopic computations.

8 Dimeric forms of some chiral compounds in solution

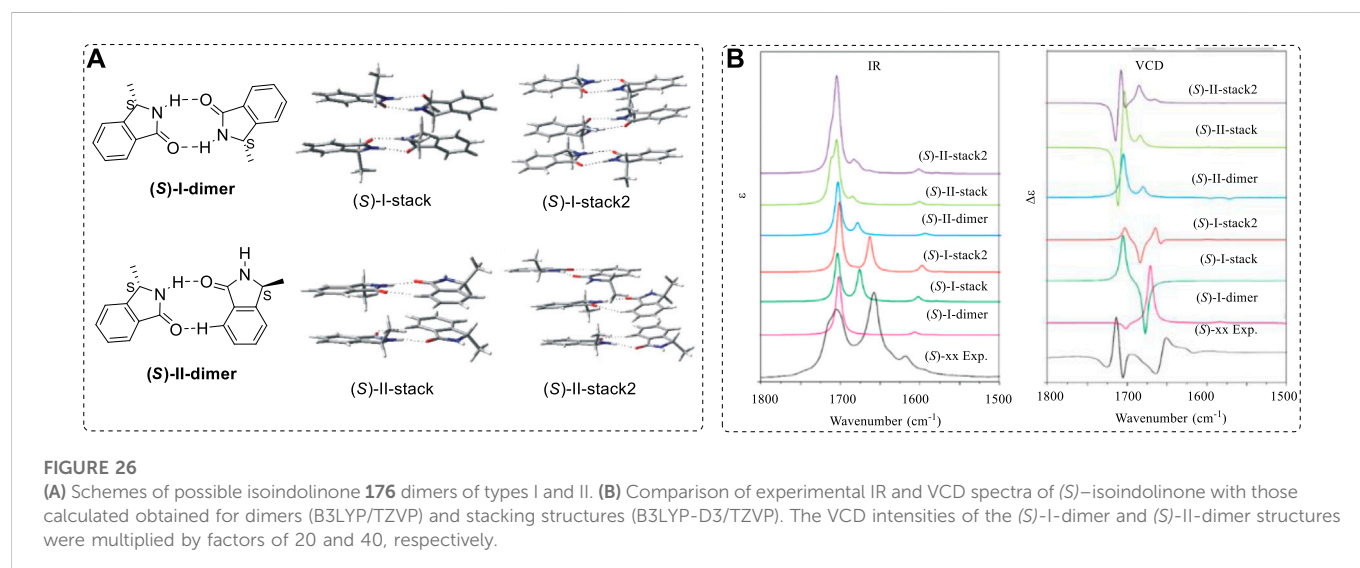
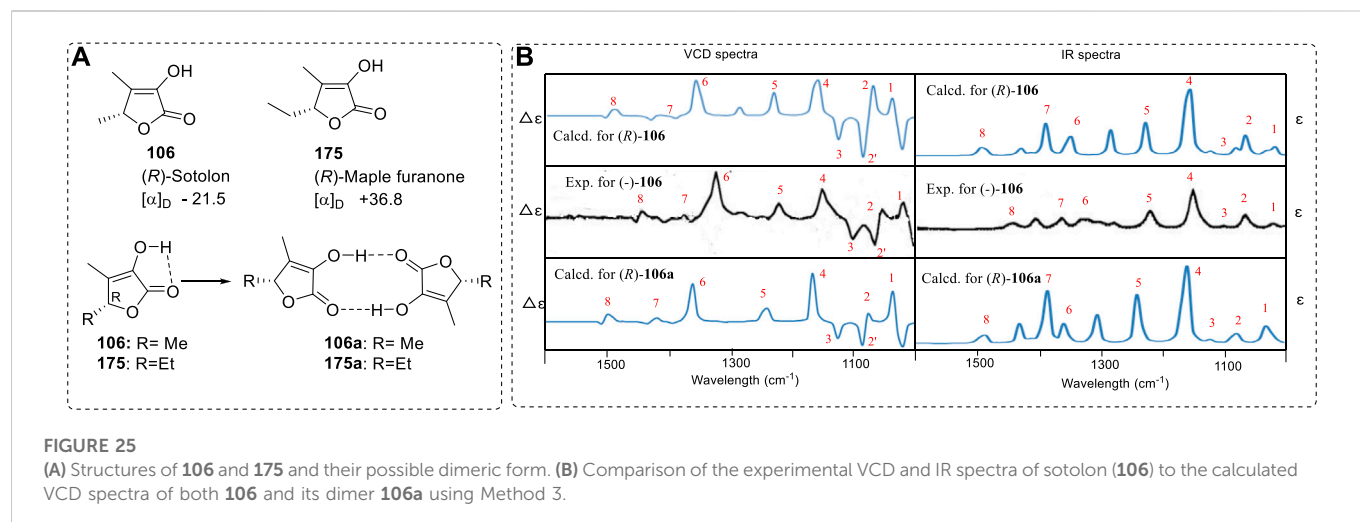
Finally, a possible case to consider is that of chiral flavor molecules that may form dimers (Slaughter, 1999; Collin et al., 2012; Yang et al., 2017). Generally, the four bases A, G, T, C are found in DNA structures where the largest substituent is methyl group-base thymine (T). This contains a Me group instead of a larger substituent such as Et among the four base pairs C, G, T, and A in DNA. Very stable H-bonds can form among the four base pairs. Could the derivative of T, with an ethyl group instead of T in DNA, form a strong H-bond in living systems? Why is only methyl involved instead of Et or a larger group in DNA evolution? In reference to this interesting question, compound **106**, which has two methyl groups, can form dimers in solution, but compound **175**, which has one ethyl group, cannot. This result might provide some hints to understanding such questions. Since OR values are very sensitive to their conformers in solution, monomer versus a dimer structure may have a large calculated difference in OR values (Jiang et al., 2021b). Five different methods (B3LYP/6-31+G(d)//B3LYP/6-31+G(d) (Method 1, M1), B3LYP/6-311++G (2d,p)//B3LYP/6-311++G(2d,p) (Method 2, M2),

B3LYP/6-311++G (2d,p)(PCM-CHCl₃)//B3LYP/6-311++G(2d,p) (PCM-CHCl₃) (Method 3, M3), MPW1PW91/6-311+G(d)//MPW1PW91/6-311+G(2d,p) (Method 4, M4), and MPW1PW91/6-311+G(d) (PCM-CHCl₃)//MPW1PW91/6-311+G(d) (PCM-CHCl₃) (Method 5, M5)) were used in OR calculations. Only the OR values of **106a** approached the experimental OR value range (reported OR is -21.5). Method 3 gave a much more accurate estimation of the OR value (-24.6) (Table 3).

The ECD of **106** and **106a**, and **175** and **175a** were little different. However, the predicted VCD had a very interesting difference in strength at the vibration numbers of 1020 and 1040 cm⁻¹ (Figure 25B). In experiments, mode 1 is stronger than mode 2. In the monomer's VCD signals, mode 1 is weaker than mode 2. However, signal 1 is stronger than signal 2 in dimeric structure **106a**.

The VCD of **175** and **175a** had a very large difference between the vibration numbers from 1300–1420 cm⁻¹. Their OR, ECD, VCD, or ROA calculations should involve the dimeric structure in the calculations. However, if there is Et or a larger group on the structure, a dimeric structure may not be possible for consideration in computations.

The VCD study of isoindolinones (such as **176**) gave more information about the 3D structures organized by dimeric structures, forming poly-unit connected structures (polymeric form) (Rode et al., 2018). The most stable conformers were obtained by conformational searches for each isoindolinone, and each of the most stable conformations were used to organize the corresponding poly-unit structure layer by layer ((S)-I-stack and (S)-I-stack2 via (S)-II-stack and (S)-II-stack2), as illustrated below



(Figure 26A). The IR and VCD ranges were near 1700 cm^{-1} for focusing on the C=O vibration modes.

The IR results clearly show that the (*S*)-I-stack and/or (*S*)-I-stack2 were the possible structures for (*S*)-**176**. Furthermore, the VCD spectra showed that the (*S*)-I-stack2 was preferred in spatial arrangement (Figure 26B). Clearly, for the rigid structures, it is easy to form the dimeric structures in solution.

When the methyl of isoindolinone was replaced by other large substituents, they may assume polymeric form in solution by H-bonds. Their 3D structures were quite complex and are not easily computed in VCD study. Amides can easily form dimeric or polymeric structure, like diastereomer diketopiperazine peptides derived from phenylalanine in VCD study (Pérez-Mellor and Zehnacker, 2017).

9 Summary

This review summarizes the OR (ORD), matrix, ECD, VCD, TS, and simplified models used in AC study for natural products. The

monomer and dimeric forms of the chiral compounds were also introduced to readers. Some guidance is given to prevent wrong conclusions in the study of organic stereochemistry.

Author contributions

HZ is responding for almost writing the entire review. YW is responding for partial compound structure preparation and handling references. LN is responding for the VCD outline and English proofreading of the whole review.

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Conflict of interest

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Supplementary material

The Supplementary Material for this article can be found online at: <https://www.frontiersin.org/articles/10.3389/fntpr.2022.1086897/full#supplementary-material>

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