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Diversity-Oriented Synthesis Approach to Macrocycles via Oxidative Ring Expansion

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Abstract

Macrocycles are key structural elements in numerous bioactive small molecules and are attractive targets in the diversity-oriented synthesis of natural product-based libraries. However, efficient and systematic access to diverse collections of macrocycles has proven difficult using classical macrocyclization reactions. To address this problem, we have developed a concise, modular approach to the diversity-oriented synthesis of macrolactones and macrolactams involving oxidative cleavage of a bridging double bond in polycyclic enol ethers and enamines. These substrates are assembled in only 4–5 synthetic steps and undergo ring expansion to afford highly functionalized macrocycles bearing handles for further diversification. In contrast to macrocyclization reactions of corresponding *seco*-acids, the ring expansion reactions are efficient and insensitive to ring size and stereochemistry, overcoming key limitations of conventional approaches to systematic macrocycle synthesis. Cheminformatic analysis indicates that these macrocycles access regions of chemical space that overlap with natural products, distinct from currently-targeted synthetic drugs.

Macrocycles are large ring structures found in myriad natural products, and over 100 of these molecules have been developed into approved drugs^{1,2}. Macrolactone and

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macrolactam structures are particularly prevalent (Fig. 1a), often derived from polyketide or peptide biosynthetic pathways. The macrocyclic constraint is a key structural element that organizes the overall molecular scaffold to present functional groups to biological targets in appropriate pharmacophoric conformations^{3–5}. This conformational restriction can also provide increased binding affinity⁶ (although not necessarily entropic in origin^{7,8}) and bioavailability⁹. Notably, a variety of macrocyclic natural products modulate challenging targets that are difficult to address with conventional drug-like molecules¹, which are often comprised of small heteroaromatic structures.

Accordingly, macrocycles are compelling targets in the diversity-oriented synthesis of natural product-based libraries for probe and drug discovery screening^{10,11}. However, macrocycles are severely underexploited in this regard due to challenges associated with their synthesis. Indeed, a recent substructure search of the >360,000-compound NIH Molecular Libraries Small Molecule Repository returned only 22 macrolactones and macrolactams with 10-, 11-, or 12-membered ring carbon scaffolds of the sort treated herein (Supplementary Fig. 1). Thus, while individual macrocycles can commonly be synthesized by macrocyclization of an appropriate linear precursor, such reactions are highly sensitive to substrate effects that impact precursor conformation, such as ring size, substituent pattern, and stereochemical configuration 12-15. As a result, this class continues to pose major challenges in the context of diversity-oriented synthesis, in which efficient, flexible, and ideally systematic access to a range of macrocyclic scaffolds is required. Indeed, previous efforts to synthesize macrocycle libraries by macrocyclization of diverse linear precursors have been hampered by the highly variable and unpredictable efficiency of these reactions^{16–19}. The high-dilution conditions typically required for macrocyclizations are also poorly suited for use in library synthesis. Thus, macrocycles remain underrepresented in current diversity libraries and new approaches to macrocycle synthesis must be developed to capitalize fully upon the biological potential of these molecules¹.

To address this chemical challenge, we posited that ring expansion reactions that are insensitive to substrate conformational effects would provide an attractive alternative to conventional macrocyclization-based strategies for systematic macrocycle library synthesis^{20–22}. Such approaches have received very limited attention in diversity-oriented synthesis, restricted to the synthesis of individual core scaffolds $^{23-25}$. We envisioned the general synthetic strategy outlined in Fig. 1b, which features rapid, modular substrate synthesis, leading to an oxidative ring expansion approach to macrolactones pioneered in the $1960s^{26-28}$ that builds upon the classic oxidative cleavage of 9,10 -octalin to 1,6cyclodecanedione²⁹. Cyclic 1,3-diketone starting materials $\mathbf{1}$, which are readily available with a variety of backbone functionalities and ring sizes, would first be converted to the corresponding 1,3-disiloxydienes 2. Diels-Alder cyclocondensation reactions with various dienophiles (e.g., aldehydes, imines) would then provide the bicyclic enones 3. Enantioselective versions of this transformation can be readily envisioned, and the resulting ketone functionality could undergo stereoselective transformations, leveraging cyclic stereocontrol, with introduction of additional diversity elements as desired. This would provide the key bicyclic precursors 4 having a bridging double bond for the pivotal oxidative ring expansion in only 4-5 synthetic steps. The ring expansion reaction would

then ideally proceed to macrocycles **5** efficiently and independent of substrate conformational effects that would impact conventional macrocyclizations of the corresponding linear precursors (*e.g.*, *seco*-acids). Subsequent transformations could be used for additional downstream functionalization of these macrocyclic scaffolds.

We report herein the development of a concise, modular approach to the systematic synthesis of functionalized macrolactone and macrolactam scaffolds related to those found in numerous natural products, using oxidative ring expansion of polycyclic substrates having a bridging double bond. We further demonstrate that this ring expansion approach provides superior reaction efficiency and independence from substrate conformational effects compared to conventional macrocyclizations of related *seco*-acid substrates. We have successfully applied this strategy to the synthesis of a first-generation library of macrocycles having diverse ring sizes and substituent patterns. Cheminformatic analyses of structural and physicochemical parameters using PCA and of three-dimensional molecular shapes using principal moment of inertia (PMI) analysis indicate that this library accesses regions of chemical space that overlap with natural products, including known macrolactones and macrolactams, and are complementary to those addressed by high-profile synthetic drugs currently targeted by the pharmaceutical industry and related drug-like libraries.

RESULTS

Modular synthesis of polycyclic ring expansion substrates

The requisite polycyclic enol ether and enamine substrates for ring expansion were synthesized as shown in Fig. 2. Cyclic ketones **6** were converted to β -diketones **7** by α -*C*-acetylation^{30,31} (**7a,d** are commercially available). Two sequential treatments with LDA and TMSCl in one pot provided the corresponding bis(trimethylsilyloxy)dienes **8** in 90% purity after filtration through basic alumina^{32,33}. Hetero-Diels–Alder reactions with aldehydes and *N*-nosyl imines afforded dihydropyrones **9** and dihydropyridones **10**, respectively³⁴. The linear Mukaiyama aldol adduct was observed by TLC prior to TFA-induced cyclodehydration in all cases, and was isolated in the case of **8a** \rightarrow **9a**. The intermediate Lewis acidity of Yb(OTf)₃³⁵ (vs. ZnCl₂ or BF₃·OEt₂) proved well-suited for the initial Mukaiyama aldol step and 4 Å MS were required for high efficiency³⁶, presumably by preventing hydrolysis of the diene substrate. The ketone functionality then underwent diastereoselective reductions to **11** and **12**³⁷, followed by silylation to afford ring expansion substrates **13** and **14** in excellent overall yields (see Table 2 below). Notably, these reactions could be carried out on multigram scale to enable scale-up of individual scaffolds as necessary.

Ring expansions to form macrolactones and macrolactams

Initial studies on the pivotal ring expansion reactions were carried out with bicyclic dihydropyran derivative **13a** (Table 1). Pioneering earlier work demonstrated the utility of *m*-CPBA and ozone for the oxidative cleavage of bridging olefins in ring expansion reactions of cyclic enol ethers^{26–28}, building upon the classic oxidative cleavage of 9,10 -octalin to 1,6-cyclodecanedione²⁹. Unfortunately, while *m*-CPBA was effective for **13a** (entry 1), it proved less efficient when applied to other substrates, such as enamine **14a**, for

which a complex mixture resulted. Initial attempts at ring expansion of **13a** using ozonolysis were also disappointing, yielding only 11% of the desired macrolactone **15a** (entry 2).

We were encouraged to find that a variety of RuO₄-mediated reactions provided promising results^{38,39}. Catalytic RuCl₃ with NaIO₄ as the stoichiometric oxidant was most effective at 3.5 mol% catalyst (entries 3-5)^{38–40}. Although use of Pb(OAc)₄ as the stoichiometric oxidant provided lower yields (entry 6), a buffered Oxone (DuPont)-based system was highly effective (entry 7)^{38,39}. Furthermore, RuO₂ could be substituted for RuCl₃ as the precatalyst (entry 8)^{41,42}.

Based on these results, we evaluated the most effective Ru-based oxidant systems across a wider range of substrates to produce both macrolactone and macrolactam scaffolds (Table 2). We were gratified to find that the ring expansion reactions proceeded with consistent yields across multiple ring sizes (entries 1–3, 4–6, 7–9), independent of ring size-based conformational effects that limit the utility of classical macrocyclization-based approaches in diversity-oriented synthesis. While slightly different oxidant systems proved optimal for the enol (**13a–c**), aryl enol (**13d–f**), and enamine substrate series (**14a–c**), this is not surprising given the electronically distinct nature of the reactive olefins in each series. More importantly, while substrate stereochemistry often has a large influence on the efficiency of classical macrocyclization reactions¹⁹, the ring expansion reaction was relatively insensitive to these effects, affording the diastereomeric methyl-substituted macrocycles **17** and **18** in similar efficiencies (entries 10,11; see Supplementary Methods for full details).

Related macrocycles **19–21** were synthesized by the analogous processes (entries 12–14; see Supplementary Methods for full details) and further highlight the tolerance of our ring expansion approach to latent functionalities that allow further downstream diversification. Macrocycle **19** includes an aryl iodide moiety suitable for subsequent palladium-catalyzed cross-coupling reactions. Macrocycles **20** and **21** contain protected alcohols and demonstrate the feasibility of including Ru-oxidation-sensitive benzyl ether moieties, albeit with somewhat reduced yield in the latter case. Macrocycle **21** also demonstrates the compatibility of the route with incorporation of aliphatic aldehydes at the Diels–Alder stage of the substrate synthesis.

Comparison to classical macrocyclization reactions

Next, to compare the efficiency of this ring expansion approach to conventional macrocyclizations, we investigated macrolactonization and macrolactamization reactions of the corresponding *seco*-acid substrates. We recognized that such substrates might be accessed systematically and conveniently for such studies by saponification of our macrocyclic ring expansion products. However, in initial experiments, we found that the *seco*-acid precursors that would lead directly to macrocycles **15a** and **16a** appear to be inherently unstable due to the reactive ketone functionality (see Supplementary Methods for full details). Accordingly, these macrocycles would be, at best, very difficult to access directly by classical macrocyclization.

To investigate *seco*-acid substrates that have closely related structures and might also reasonably be converted to our macrocyclic products after cyclization, the reactive ketone

functionality in **15a–d** was masked by methylenation (**23a–d**), after which saponification provided *seco*-acids **22a–d** (See Supplementary Methods for full details). We then attempted macrocyclizations of these substrates with Shiina's MNBA reagent, which is often effective for larger (13-membered) ring systems and is the preferred reagent in such macrolactonizations^{43,44}. Strikingly, none of these substrates underwent efficient macrocyclization under the standard conditions (high-dilution, slow inverse addition over 22 h), with large amounts of starting material remaining (Table 3). Attempted macrolactamization of a related *N*-nosyl amino acid substrate derived from **16a** under conditions that are effective for similar *N*-sulfonylamino acid substrates (DCC, 4pyrrolidinopyridine⁴⁵) led only to a linear byproduct (see Supplementary Methods for full details). Conditions that have been used in other macrolactamizations (HATU, DBU or Et₃N⁴⁶) led to complex mixtures. These results further highlight the unique effectiveness of our ring expansion approach in accessing these 10- to 12-membered macrolactones and macrolactams.

Further diversification of macrocyclic scaffolds

With a variety of macrocyclic scaffolds in hand, we investigated several downstream reactions to provide additional structural diversity and functional group handles for coupling reactions (Fig. 3). As hoped, aryl iodide **19** proved an effective partner in several palladiumcatalyzed cross-coupling reactions, including Sonogashira, Negishi, and Stille reactions (**24**–**26**). The nascent ketone functionalities in **15a–f** can also serve as handles for additional diversification, as highlighted by diastereoselective reductions to provide the corresponding alcohols **27a–f**. Alternatively, the ketone functionality of **15a–d** could be modified by Petasis methylenation to afford olefins **23a–d**. Subsequent hydroboration oxidation or epoxidation of **23a** provided alcohol **28** or epoxide **29**, respectively. In addition, the TBS-protected secondary alcohol **15a** could be desilylated to yield α-hydroxy ketone **30** and acylated to give α-acetoxy ketone **31**, illustrating the utility of this handle for further diversification.

Cheminformatic analysis of macrocycle library

To assess the ability of our macrocycle library to access underrepresented regions of chemical space, we carried out principal component analysis (PCA) on a collection of 32 macrocyclic structures (see Supplementary Fig. 2) derived from the scaffolds shown in Table 2 and Fig. 3 by *in silico* desilylation as necessary. Using PCA, we compared the macrocycle library with our established reference sets of 40 top-selling brand name drugs, 60 diverse natural products, and 20 drug-like library members¹⁰. The drug reference set was selected to illustrate the structural bias in high-profile synthetic drugs currently targeted by the pharmaceutical industry. The broad natural product reference set was selected to represent a diverse range of biological activities and biosynthetic origins. This set includes all 24 compounds identified previously as having led to an approved drug in 1981–2006⁴⁷, as well as 13 other clinically used drugs. The drug-like library reference set is comprised of 10 molecules in one scaffold class each from ChemBridge and ChemDiv, two of the earliest and most widely used commercial library suppliers. These drug-like compounds are all present in the NIH Molecular Libraries Small Molecule Repository. An additional reference

set of 24 macrocyclic natural products (see Supplementary Fig. 3) was also included to assess the ability of the macrocycle library to target a similar region of chemical space.

Each compound was analyzed for our established set of 20 calculated structural and physicochemical properties¹⁰ (see Supplementary Methods for full details on parameter selection), with PCA resulting in rotation of the complete 20-dimensional dataset onto three unitless, orthogonal axes that represent linear combinations of the original 20 parameters. Because several of the 20 properties are highly correlated with one another (see Supplementary Fig. 4 for loading plots and table), especially parameters that scale with molecular size, this three-dimensional plot is able to represent 75% of the information in the full 20-dimensional dataset (Fig. 4a–c and Supplementary Fig. 5). Similarly, >90% of the information in the full dataset is represented in the first six principal components.

As hoped, the macrocycle library accesses a distinct region of chemical space compared to our reference sets containing current, high-profile drugs and related drug-like libraries, overlapping to a significant extent with the macrocyclic natural products. Examination of the component loadings in the PCA (see Supplementary Fig. 4) indicates that molecular weight and other parameters associated with molecular size shift molecules to the left along the x-axis (PC1). Hydrophobicity and aromatic ring count shift molecules upward along the y-axis (PC2), while aqueous solubility shifts molecules downward. Notably, larger ring systems shift molecules rearward (negative) along the z-axis (PC3), and stereochemical complexity shifts molecules left, downward, and rearward. Because the macrocyclic natural product reference set and our macrocycle library are comprised of relatively smaller molecules having similar molecular sizes to those of the drug and drug-like library reference sets, these last two properties appear to be important in differentiating the macrocycles from our reference sets of current, high-profile synthetic drugs and related drug-like libraries (Fig. 4b).

We next carried out a molecular shape analysis using normalized ratios of principal moments of inertia (PMI) to visualize the three-dimensional shape diversity of the various compound datasets in a chemically intuitive manner, as previously reported^{19,48} (Fig. 4d and Supplementary Fig. 6). In this analysis, a stochastic conformational search is carried out for each molecule (see Supplementary Methods for full details). Low-energy conformations are identified for each molecule in the data set and three PMI values (I₁, I₂, and I₃; where I₃ I₂

 I_1) are calculated for each conformer. Normalized ratios of PMI (I_1/I_3 and I_2/I_3) are then calculated and plotted on a triangular graph, with the vertices (0,1), (0.5,0.5), and (1,1) representing a perfect rod, disc, and sphere, respectively⁴⁸. To account for the fact that molecules may not always bind to proteins in energy-minimized conformations⁴⁹, a collection of conformers within 3 kcal/mol of the minimum energy conformer was generated for each of the structures. For ease of visualization, the lowest energy conformers are plotted in Fig. 4d; additional PMI plots including all conformers up to various energy cutoffs above the minimum are shown in Supplementary Fig. 7.

While the distinctions between the different reference sets are less obvious in this PMI plot, the drug reference set tends to occupy rod-like space with some disc-like character (Fig. 4d). In contrast, our natural product reference set spans a much wider range of shapes, extending

well into sphere-like space. Notably, both the macrocyclic natural products, including several approved drugs, and our macrocycle library occupy similar regions of the PMI plot, displaying decreased rod-like character in comparison to the drug reference set. The structures of four macrocycle library members are shown in Fig. 4d to illustrate the differences in molecular shape between the lowest energy conformers. Structures **15f'**, **16c'**, **20'**, and **26'** are derived from macrocycles **15f**, **16c**, **20**, and **26**, respectively, via *in silico* desilylation (see Supplementary Fig. 2). Macrocycle **20'** falls closest to the rod vertex of the

triangular plot (0,1) and its elongated, rod-like shape is evident when compared to the more flat, disc-like shape of macrocycle **15f'**. In comparison, the substitution patterns of functional groups in the scaffolds of macrocycles **16c'** and **26'** impart more sphere-like character to these ring systems.

DISCUSSION

Despite their rich biological pedigree and distinctive molecular recognition capabilities, macrocycles remain severely underrepresented in current probe and drug discovery efforts¹. The influence of conformational effects based on ring size, stereochemistry, and backbone substitution patterns on the efficiency of macrocyclization reactions, as well as the need for long reaction times under high-dilution conditions, generally limit the utility of macrocyclization-based approaches in the synthesis of macrocycle libraries. Indeed, previous efforts to synthesize natural product-inspired macrocycle libraries using macrocyclization reactions have displayed considerable dependence on the substitution patterns of the linear precursors^{16–19}. For example, in a recent study, Marcaurelle and coworkers reported the synthesis of a collection of stereochemically diverse 8- to 14membered rings using an aldol-based 'build/couple/pair' strategy¹⁹. Several different cyclization reactions were used (nucleophilic aromatic substitution, Huisgen [3+2] cycloaddition, ring-closing metathesis), and all were influenced considerably by the stereochemistry of the corresponding linear substrates. In some cases, changing a single stereochemical configuration in the linear substrate led to as much as 43% variation in yields of isolated cyclic products.

To overcome these limitations of classical macrocyclization strategies in the context of diversity-oriented synthesis, we have developed a ring expansion approach to systematic macrocycle synthesis that provides efficient access to a variety of macrocyclic scaffolds that comprise the core structures of numerous natural products. Our polycyclic ring expansion substrates are synthesized via a concise, 4–5 step route in high overall yields, requiring only two chromatographic steps (Figs. 1b and 2). This modular synthetic approach provides a range of ring sizes and substitution patterns and also leverages cyclic stereocontrol during substrate assembly. While the carbonyl functionalities of 4 and 5 were reduced herein, a variety of other diastereoselective transformations can be envisioned to provide additional diversity in the future (Fig. 1b). The oxidative ring expansion reactions proceed efficiently and predictably, independent of ring size and stereochemistry-based substrate conformational effects. In addition, our ring expansion approach uses mild conditions that allow for the inclusion of a wide array of latent functionalities in the macrocyclic products. Thus, these macrocycles are poised for a variety of downstream transformations that provide

both additional structural diversity and functional group handles for coupling other building blocks in the future (Fig. 3).

Moreover, we have demonstrated explicitly that this ring expansion approach provides 10to 12-membered lactones and lactams that are not readily accessible via classical macrocyclizations of *seco*-acid substrates (Table 3). Macrolactonization reactions of *seco*acids **20a–d** using Shiina's MNBA reagent led to low conversion to the desired macrocycles after 22 h, with unreacted starting material and oligomeric byproducts observed. In contrast, the oxidative ring expansion reactions used to synthesize macrolactones **15a–d** proceeded to complete conversion in 2–24 h, affording the target molecules in good isolated yields. Notably, the ring expansion reactions are carried out in aqueous solution and do not require inert atmosphere, slow addition or high substrate dilution, overcoming major technical limitations of classical macrocyclization reactions in the context of library synthesis. Taken together, these aspects of the ring expansion approach illustrate the utility of our method for the diversity-oriented synthesis of macrocycle libraries.

Cheminformatic analyses using PCA suggest that the macrocyclic scaffolds generated via our ring expansion method address regions of chemical space that are distinct from those addressed by our reference sets of current, high-profile synthetic drugs and related drug-like libraries (Fig. 4a–c). The component loadings (Supplementary Fig. 4) can be used to understand the relative influence of the original 20 physicochemical parameters upon the positioning of molecules in the PCA plot. These data indicate that hydrophobicity and aromatic character are among the most important parameters contributing to the positioning of the drug and drug-like library reference sets. Meanwhile, stereochemical complexity and aqueous solubility are two of the key parameters that differentiate the macrocycle library from these reference sets. Insights gleaned from such analyses can now be leveraged in the design and planning of future libraries that may probe more deeply into natural product-like space.

While PCA highlights differences in structural and physicochemical properties between compound classes, PMI analysis illustrates differences in the three-dimensional molecular shapes of macrocycles compared to our reference sets of current, high-profile synthetic drugs and related drug-like libraries (Fig. 4d). The drug reference set spans the region of rod- and disc-like shapes, with a bias toward the rod vertex of the triangle. This observation is consistent with previous PMI analyses of drugs and drug-like molecules^{48,50}. Relative to the drug reference set, both the macrocyclic natural products and our macrocycle library members cluster farther from the rod vertex of the triangle and indicate a greater degree of sphere-like character. This difference in three-dimensional shapes may impact the abilities of these molecules to address challenging biological targets¹.

In conclusion, we have developed an efficient, modular ring expansion approach to the systematic synthesis of macrocyclic scaffolds similar to those found in numerous natural products. This chemistry has been applied to the production of a first-generation macrocycle library that has been submitted to the NIH Molecular Libraries Program (PubChem Substance search term: DST_MC1_*). These macrocyclic scaffolds are now being screened against a wide range of challenging biological targets to evaluate further the functional

METHODS

General procedures for key steps in the synthesis of macrolactones **15a–f** and macrolactams **16a–c** are outlined below. Unless otherwise noted, reactions were performed in flame-dried glassware under an anhydrous Ar atmosphere. See Supplementary Methods for additional protocols, detailed experimental and cheminformatic procedures, and complete analytical data for all compounds.

expansion of this synthetic approach will be reported in due course.

General protocol for hetero-Diels–Alder reactions

Ground 4 Å molecular sieves (660 mg/mmol dienophile) were placed in a roundbottom flask and flame dried under high vacuum. After cooling to rt, anhydrous THF (3.0 mL/mmol dienophile) was added, followed by the bis(trimethylsiloxy)diene **8a–f** (1.05–2.00 equiv) and benzaldehyde (1.00 equiv). Neat Yb(OTf)₃ (2.0 mol%) was added, the flask was sealed with a glass stopper, and the mixture was stirred overnight at rt. TLC analysis of a reaction aliquot (diluted with EtOAc) indicated complete consumption of the dienophile and formation of the intermediate Mukaiyama aldol product. The reaction mixture was then filtered through a pad of celite and the pad was rinsed thoroughly with EtOAc and CH₂Cl₂. The solvents were then removed by rotary evaporation.

The resulting residue was dissolved in CH_2Cl_2 (9.5 mL/mmol dienophile) in an Ar-flushed roundbottom flask. TFA (0.5 mL/mmol dienophile) was added and the mixture was stirred at rt until the reaction was judged complete by TLC. The reaction was quenched with satd aq NaHCO₃ and the mixture was stirred vigorously for 20 min. The layers were separated and the aqueous layer was extracted with CH_2Cl_2 (3×). The combined organic extracts were dried (MgSO₄), filtered, and concentrated by rotary evaporation. Purification by silica flash chromatography (hexanes/EtOAc) afforded the dihydropyrone **9a–f** or dihydropyridone **10a–c**.

General protocol for oxidative ring expansion using $RuCl_3$ with $NalO_4$ stoichiometric oxidant

TBS-protected alcohol **13a–c** was dissolved in 1,2-dichloroethane (5.50 mL/mmol substrate). Water (4.50 mL/mmol) was added, followed by a solution of RuCl₃ (0.035 M in water, 3.5 mol%). Neat NaIO₄ (2.00 equiv) was added in one portion, and the resulting biphasic mixture was stirred until the reaction was judged complete by TLC analysis of a reaction aliquot (quenched with satd aq Na₂S₂O₃, extracted with EtOAc). Satd aq Na₂S₂O₃ was added and the resulting mixture was stirred for 10 min, then extracted with CH₂Cl₂ (5×). The combined organic extracts were washed with water and brine, and dried (MgSO₄). To the desiccant slurry, a spatula of celite was added and the mixture was agitated for 5 min. The suspension was then filtered through a plug of celite and the solvents were removed by rotary evaporation. Purification by silica flash chromatography (hexanes/EtOAc, 0.5 vol% Et₃N) afforded macrolactone **15a–c**.

General protocol for oxidative ring expansion using RuCl₃ with Oxone (DuPont) stoichiometric oxidant

TBS-protected alcohol **13d–f** was dissolved in acetonitrile (16.0 mL/mmol). Water (10.0 mL/mmol substrate) was added, which in most cases resulted in the precipitation of substrate from solution. RuCl₃ (0.035 M in water, 3.5 mol%) was added and, after stirring for several minutes, the starting material partially redissolved. A solid mixture of Oxone (DuPont) (1.50 equiv) and NaHCO₃ (4.60 equiv, 1.02 equiv relative to Oxone protons) was then added in portions. The mixture was stirred until the reaction was judged complete by TLC analysis of a reaction aliquot (quenched with satd aq Na₂S₂O₃, extracted with EtOAc). Satd aq Na₂S₂O₃ was added, and the resulting mixture was stirred for 10 min, then extracted with EtOAc ($3\times$). The combined organic extracts were washed with water and brine, dried (MgSO₄), filtered, and concentrated by rotary evaporation. Purification by silica flash chromatography (hexanes/EtOAc, 0.5 vol% Et₃N) afforded macrolactone **15d–f**.

General protocol for oxidative ring expansion using ${\rm RuO}_2$ with ${\rm NalO}_4$ stoichiometric oxidant

TBS-protected alcohol **14a–c** was dissolved in CH_2Cl_2 (14.0 mL/mmol). Acetonitrile (6.00 mL/mmol substrate) and acetone (6.00 mL/mmol substrate) were added. In a separate flask, NaIO₄ (3.00 equiv) was dissolved in H₂O (5.3 mL/mmol substrate) and RuO₂ (17 mol%) was added. The activated catalyst was then added to the solution of TBS-protected alcohol in two portions over 5 min and stirred at until the reaction was complete as judged by TLC analysis (*ca.* 40 min). Satd aq Na₂S₂O₃ was added and the resulting mixture was extracted with CH₂Cl₂ (3×). The combined organic extracts were washed with brine, dried (MgSO₄), and filtered over a pad of celite covered by a thin layer of activated charcoal. The filtrate was concentrated by rotary evaporation. Purification by silica flash chromatography (hexanes/EtOAc) afforded the macrolactam **16a–c**.

Cheminformatic analyses

The PCA and PMI analyses were carried out on a total of 176 compounds: 40 top-selling brand-name small molecule drugs by revenue in 2006, 10 drug-like pyrrazolecarboxamides in the MLSMR from ChemBridge, 10 drug-like dihydrotriazolopyrimidines in the MLSMR from Chem Div, 60 natural products with diverse structures, 24 macrocycle natural products, and 32 macrocycle library members.

In the PCA, each molecule was analyzed for a set of 20 physicochemical properties using free online cheminformatics tools, ChemDraw, or manual inspection. These data were assembled in a spreadsheet with the molecules in rows and physicochemical descriptors in columns. The mean value and standard deviation of each column was calculated and mean values were also calculated within each compound series. PCA was performed on mean-centered, normalized data using the R open source statistical computing package. The results from R were used to generate PCA plots (see Supplementary Methods for full details).

In the PMI analysis, minimum energy conformers were generated for each compound using the stochastic conformational search algorithm, as implemented in the MOE molecular modeling software package. The Merck Molecular Force Field (MMFF94) combined with

the generalized Born (GB) solvation model of aqueous solvation were used. All conformers within 3 kcal/mol of the minimum energy conformation were recorded. Three principal moments of inertia (Ixx, Iyy, Izz) and normalized principal moments of inertia, npr1 (Ixx/Izz) and npr2 (Iyy/Izz) were calculated using MOE for each conformer. The npr1 and npr2 values were used to generate PMI plots (see Supplementary Methods for full details).

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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Figure 1. Macrocyclic natural products and overall strategy for the diversity-oriented synthesis of macrolactones and macrolactams

(a) Examples of macrocyclic natural products with diverse chemical structures and biological activities. (b) Overall approach to the diversity-oriented synthesis of functionalized macrocycles using oxidative ring expansion reactions. Polycyclic substrates are prepared using concise, modular reactions that allow for the introduction of chemical diversity both prior to and after ring expansion.





Figure 2. Modular approach to diverse polycyclic substrates for oxidative ring expansion Average yields en route to 13 and 14 are shown in brackets (*cf.* Table 2). a) LDA (inverse addition), THF, -78° C, 1 h; AcCl (inverse addition), -78° C, 1 h or Ac₂O, BF₃(AcOH)₂, 4.5 h; NaOAc, MeOH, H₂O, 85 °C, 2.5 h; [85%]. b) LDA, THF, -78° C, 1 h; TMSCl, -78° C \rightarrow rt, 1.5 h; LDA, -78° C, 1 h; TMSCl, -78° C \rightarrow rt, 1.5 h; [96%]. c) R¹CHX, Yb(OTf)₃, 4 Å MS, THF, 12 h; TFA, CH₂Cl₂, rt, 20 min; [86%]. d) NaBH₄, CeCl₃, CH₂Cl₂, MeOH, -78° C, 2 h or LiEt₃BH, toluene, -78° C, 40 min; [94%]. e) TBSCl, imidazole, DMF, 50°C; [87%]. LDA = lithium diisopropylamide; MS = molecular sieves; Ns = 2nitrobenzenesulfonyl; TBS = *tert*-butyldimethylsilyl; TFA = trifluoroacetic acid; TMS = trimethylsilyl. (See Supplementary Methods for full details).

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Figure 3. Functionalization of macrocyclic scaffolds

a) TBSOCH₂CH₂C=CH, Pd(PPh₃)₄, CuI, Et₃N, DMF, 60°C, 18 h, 81%. b) 4-MeOPhZnCl, Pd(PPh₃)₄, THF, rt, 16 h, 72%. c) Bu₃SnCH=CH₂, Pd₂(dba)₃, AsPh₃, THF, 50°C, 18 h, 81%. d) NaBH₄, CeCl₃, MeOH, CH₂Cl₂, -78° C, 2 h; **27a**: 86%, 98:2 dr, **27b**: 93%, 1:1 dr; **27c**: 99%, 90:10 dr; **27d**: 55%, 98:2 dr; **27e**: 73%, 96:4 dr; **27f**: 67%, 98:2 dr. e) Cp₂TiMe₂, toluene, 65°C, 12 h; **23a**: 55%; **23b**: 55%; **23c**: 50%; **23d**: 71%. f) 9-BBN, THF, 0°C \rightarrow rt, 12 h, then NaOH, H₂O₂, 0°C \rightarrow rt, 1.5 h, 70%, 98:2 dr. g) DMDO, CH₂Cl₂, 0°C \rightarrow rt, 12 h, 88%, 63:37 dr. h) TBAF, AcOH, THF, rt, 3.5 h, 72%. i) AcCl, pyridine, CH₂Cl₂, 0°C, 10 min, 79%. 9-BBN = 9-bora-bicyclo[3.3.1]nonane; DMDO = dimethyldioxirane; DMF = *N*,*N*-dimethylformamide; TBAF = tetrabutylammonium fluoride; TBS = *tert*-butyldimethylsilyl; THF = tetrahydrofuran.

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Figure 4. Cheminformatic analyses of macrocycle library

Principal component analysis (PCA) and principal moment of inertia (PMI) plots of 32 macrocycle library members, 24 macrocyclic natural products, and established reference sets of 40 top-selling brand name drugs, 60 diverse natural products, and 20 ChemBridge and ChemDiv drug-like library members. (a) PCA plot of PC1 vs. PC2 generated from 20 structural and physicochemical parameters. (b) PCA plot of PC3 vs. PC2. (c) PCA plot of PC1 vs. PC3. (d) PMI plot and representative macrocycle library members with lowest energy conformations having characteristic shapes. Structures 15f', 16c', 20', and 26' are derived from macrocycles 15f, 16c, 20, and 26, respectively, via *in silico* desilylation (see Supplementary Fig. 2). See Supplementary Figs. 5 and 6 for expanded PCA and PMI plots and Supplementary Datasets 1 and 2 for complete data. See Supplementary Fig. 7 for PMI plots with additional conformers within 3 kcal/mol of the minimum.

Table 1

Oxidants for ring expansion of 13a to 15a.^[a]

Entry	Oxidant	Conditions	Yield [%]
1	m-CPBA	CH ₂ Cl ₂ , rt, 1h	66
2	O ₃	CH ₂ Cl ₂ , -78 °C, 5 min; PPh ₃	11
3	0.5 mol% RuCl ₃ , NaIO ₄	H ₂ O/1,2-DCE, rt, 24 h ^b	62
4	3.5 mol% RuCl ₃ , NaIO ₄	H2O/1,2-DCE, rt, 20 h	71
5	10 mol% RuCl ₃ , NaIO ₄	H ₂ O/1,2-DCE, rt, 20 h	37
6	3.5 mol% RuCl ₃ , Pb(OAc) ₄	H ₂ O/1,2-DCE, rt,1h	35
7	3.5 mol% RuCl ₃ , Oxone	NaHCO ₃ , H ₂ O/CH ₃ CN, rt, 2 h	82
8	17 mol% RuO ₂ , NaIO ₄	H ₂ O/CH ₂ Cl ₂ /acetone/CH ₃ CN, rt, 1 h	69

 $[a]_{1,2-\text{DCE}} = 1,2-\text{dichloroethane}; m-\text{CPBA} = 3-\text{chloroperbenzoic acid}; \text{Oxone (DuPont)} = \text{potassium peroxomonosulfate};$

^b conversion not complete after 1 day.

Table 2

Synthesis of diverse macrolactones and macrolactams by oxidative ring expansion

Entry	Macrocycle P	roduct	Precursor Yield (4 steps) [%] ^[a]	Product Yield [%]
1	O Ph	15a (n = 1)	81 (13a)	71 ^[b]
2	\wedge	15b (n = 2)	78 (13b)	70 ^[b]
3	OTES	15c (n = 3)	73 (13c)	71 ^[b]
	0			
4	O Ph	15d (n = 1)	68 (13d)	56 ^[c]
5		15e (n = 2)	52 (13e)	69[c]
6	CTBS	15f $(n = 3)$	67 (13f)	59[c]
	U			
7	O Ph II I	16a (n = 1)	73 (14a)	68 ^[d]
8	N.	16b (n = 2)	75 (14b)	65 ^[d]
9	The OTBS	16c (n = 3)	66 (14c)	66 ^[d]
	U O			
10	O Ph	17	60	69 ^[c]
	OTBS			
11	O Ph	18	56	60 ^[c]
	OTBS 0			
12		19 (R = <i>m</i> -I)	71	70 ^[c]
13	R	20 (R = p -CH ₂ -OMOM)	45	71 ^[c]
	ÎÝ			
	OTBS			
	U O			

[a] Overall yield of ring expansion substrates (*e.g.*, **13,14**) over four steps from diketones **7**, which were commercially available (**7a,d**) or synthesized from **6** as shown in Fig. 2. (See Supplementary Methods for full details).

[b] 3.5 mol% RuCl3, NaIO4, 1:1 H2O/1,2-DCE, rt, 20–24 h.

[c] 3.5 mol% RuCl3, Oxone (DuPont), NaHCO3, 1:1.45 H2O/CH3CN, rt, 2–18 h

^[d]17 mol% RuO₂, NaIO₄, 1:2.3:1:1 H₂O/CH₂Cl₂/acetone/CH₃CN, rt, 1–2 h.

Table 3

Macrocyclization of corresponding seco-acid substrates.[a]

Entry	Substrate	Unreacted 22 [%][b]	Lactone 23 [%][b]	Corresponding Ring Expn Yield (15) [%] ^[c]
1	22a (n = 1)	63	14	71 (15a)
2	22b (n = 2)	63	5	70 (15b)
3	22c (n = 3)	56	14	71 (15c)
4	22d (n = 1, <i>benzo</i>)	$_{44}[d]$	31	56 (15d)

[a] DMAP = 4-dimethylaminopyridine; MNBA = 2-methyl-6-nitrobenzoic anhydride.

^[b]Yields determined by ¹H-NMR analysis of crude products after aqueous workup, based on an internal standard.

[c] Isolated yields of **15a–d** from Table 2.

 $^{[d]}$ Corresponding mixed anhydride observed intact after aqueous workup by ¹H-NMR and MS.