

Catalytic asymmetric synthesis of *meta* benzene isosteres

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Mingkai Zhang¹, Matthew Chapman¹, Bhagyesh R. Sarode², Bingcong Xiong¹, Hao Liang¹, James K. Chen^{2,3,4}✉, Eranthie Weerapana¹✉ & James P. Morken¹✉

Although aromatic rings are common elements in pharmaceutically active compounds, the presence of these motifs brings several liabilities with respect to the developability of a drug¹. Nonoptimal potency, metabolic stability, solubility and lipophilicity in pharmaceutical compounds can be improved by replacing aromatic rings with non-aromatic isosteric motifs². Moreover, whereas aromatic rings are planar and lack three-dimensionality, the binding pockets of most pharmaceutical targets are chiral. Thus, the stereochemical configuration of the isosteric replacements may offer an added opportunity to improve the affinity of derived ligands for target receptors. A notable impediment to this approach is the lack of simple and scalable catalytic enantioselective syntheses of candidate isosteres from readily available precursors. Here we present a previously unknown palladium-catalysed reaction that converts hydrocarbon-derived precursors to chiral boron-containing nortricyclanes and we show that the shape of these nortricyclanes makes them plausible isosteres for *meta* disubstituted aromatic rings. With chiral catalysts, the Pd-catalysed reaction can be accomplished in an enantioselective fashion and subsequent transformation of the boron group provides access to a broad array of structures. We also show that the incorporation of nortricyclanes into pharmaceutical motifs can result in improved biophysical properties along with stereochemistry-dependent activity. We anticipate that these features, coupled with the simple, inexpensive synthesis of the functionalized nortricyclane scaffold, will render this platform a useful foundation for the assembly of new biologically active agents.

Bioactive compounds, from natural products to clinically relevant active ingredients, often contain aromatic rings. The reasons for incorporation of this motif are multifold¹: (1) the rigidity of aromatic rings often restricts available molecular conformations thereby lessening the entropic cost for the aromatic ligand to bind to a target receptor; (2) aromatic rings participate in unique non-covalent interactions that can provide enhanced attraction to receptor binding pockets; and (3) the methods used to attach substituents to aromatic rings include efficient cross-coupling reactions³ that facilitate construction and screening of diverse compound collections. These notable benefits are counter-balanced by the ready oxidation of π -electron-rich aromatic rings that can severely diminish metabolic stability⁴. Moreover, relative to their $C(sp^3)$ counterparts, aromatic molecules have decreased aqueous solubility⁵ and increased $\log D$ (ref. 6). A particular challenge arising from the increased lipophilicity of an aromatic compound is that this feature increases off-target promiscuity⁷ that is problematic for drug development. To address this limitation, bicyclic hydrocarbon frameworks have been advanced as $C(sp^3)$ -based isosteric replacements for aromatic rings⁸, with the expectation that the saturated bicyclic analogues might offer enhanced biophysical properties while retaining

biological activity⁹. When appropriately designed, these molecular scaffolds can retain the conformational restriction of the benzenoid ring while simultaneously avoiding associated liabilities. Following this model, an array of small bicyclic motifs have been developed that position substituents at a similar distance and with similar exit vectors compared with substituted arenes, with selected examples of *meta* benzene isosteres shown in Fig. 1a (refs. 10–22).

In the context of ligand development, two important features arise when planar $C(sp^2)$ -based aromatic rings are replaced with $C(sp^3)$ -based bicyclic isosteres. The first is that of retaining the efficient, reliable, modular and scalable synthesis techniques that accompany modern catalytic cross-coupling methods. The second issue is that of three-dimensional shape. Even if the isostere positions its substituents in a manner that would superimpose them onto those of an aromatic nucleus, the tetrahedral nature of the saturated carbon centres in the isostere automatically results in a three-dimensionality that is absent from the flat aromatic ring system. As the preponderance of biologically relevant receptors are chiral, the handedness of a nonsymmetric isostere may either enhance or impede the binding of the derived ligand to its cognate receptor^{18,23}. The way MK-5108 binds to the inactive DFG-out

¹Department of Chemistry, Boston College, Chestnut Hill, MA, USA. ²Department of Chemical and Systems Biology, Stanford University, Stanford, CA, USA. ³Department of Developmental Biology, Stanford University, Stanford, CA, USA. ⁴Department of Chemistry, Stanford University, Stanford, CA, USA. ✉e-mail: jameschen@stanford.edu; eranthie.weerapana@bc.edu; morken@bc.edu

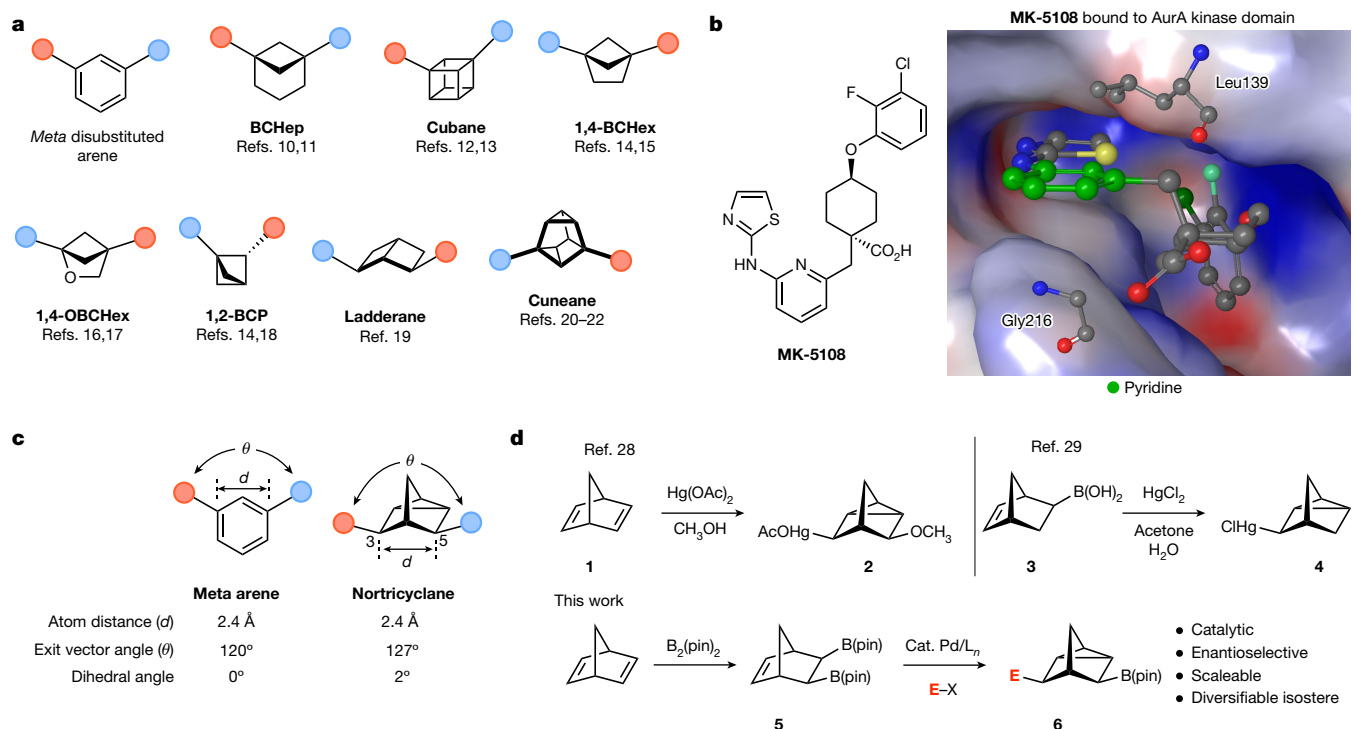


Fig. 1 | Saturated bicycloalkanes as isosteric replacements for aromatic rings. **a**, Selected recent examples of *meta* benzene isosteres. **b**, The kinase inhibitor MK-5108 bound to the chiral binding pocket in the kinase domain of AurA. **c**, Structural comparison of a *meta* disubstituted arene and

state of the mitotic kinase Aurora A is illustrative^{24,25}. When bound, the *meta* disubstituted arene in MK-5108 is sandwiched in a hydrophobic pocket with Leu139 positioned 3.9 Å above the arene (carbon-to-carbon distance) and Gly216 3.5 Å below (Fig. 1b). Thus, when a nonsymmetric benzene isostere occupies this binding pocket, it would be expected to fit better if it occupies more volume in the upper region of the receptor rather than the lower. Such a lock-and-key fit of ligands to their complementary receptors can result in enhanced specificity and ligand promiscuity has been shown to diminish as the number of chiral centres in a compound increases⁷. From this vantage point, it is clear that methods are needed to control the absolute configuration of the overall molecular ensemble. In this report, we describe a reaction that retains all the practical advantages of contemporary cross-coupling and that allows conversion of simple starting materials into enantiomerically enriched 3,5-disubstituted nortricyclane frameworks (Fig. 1c). Of note, comparison of the metrical parameters of the nortricyclane with those of benzene show that carbons 3 and 5 are equidistant with the *meta* carbons of benzene. Moreover, the exit vectors for connection to substituents at carbons 3 and 5 have an angle between them that is nearly the same as in benzene (127° versus 120°), and the vectors are essentially co-planar as in the aromatic framework. Last, the process that we describe for the production of 3,5-disubstituted nortricyclanes occurs by a heretofore unknown catalytic reaction. This reaction occurs with low loadings of a chiral palladium complex and delivers malleable boron-containing isosteres, holding marked potential for further modification.

The bicyclic norbornadiene skeleton (**1**; Fig. 1d) has provided an instructive framework for studying the nature of non-classical carbocations and for probing the ability of homoconjugation to stabilize organic structures and influence the course of their chemical reactions^{26,27}. Owing to the through-space interaction between the two π -bonds, the conversion of the bicyclo[2.2.1] framework to nortricyclanes (that is, **2**) readily occurs on the addition of suitable

a 3,5-disubstituted nortricyclane. **d**, Metal-promoted reactions for the construction of nortricyclanes from simple bicycloheptane precursors. pin, pinacolato group.

electrophilic reagents²⁸. Similar electrophilic activation of bicyclic organometallics (that is, **3**) can result in ring-closing C–C bond formation to furnish the nortricyclane skeleton (**4**) (ref. 29). In contrast to these stoichiometric reactions, we herein demonstrate that a catalytic synthesis of nortricyclanes operates when the mercury salts used in the synthesis of **2** and **4** are replaced with substoichiometric quantities of palladium complexes in conjunction with an organic electrophile. The reaction that occurs results in ring closure with the concomitant addition of the organic electrophile to the bicyclic framework. An important finding for the application of this reaction to the modular construction of enantiomerically enriched *meta* benzene isosteres is that the cyclization occurs with *meso* diboron substrate **5**, a compound that is shelf-stable and readily available on a multigram scale. Reaction of **5** delivers a nortricyclane **6** that retains one boronic ester functional group, apart from the newly appended organic substituent (E). After optimization of the reaction (conditions, ligand structure), it was found that a wide array of enantiomerically enriched boron-containing nortricyclanes can be prepared in excellent yield and enantioselectivity.

Development of the Pd-catalysed reaction

In the presence of catalytic quantities of Pd(OAc)₂ and chiral biarylmonophosphine ligand **L1** (ref. 30) (Fig. 2), diboron substrate **5** and organic electrophiles engage in catalytic enantioselective ring closure to nortricyclanes. With simple substituted aryl bromides, the reaction occurs in high yield and enantioselectivity. The products from both electron-deficient (**11**, **12**) and electron-rich (**13**, **15**) electrophiles can be readily obtained, although selectivity is diminished with a strong electron-withdrawing nitroarene electrophile (**16**). Also of note, heterocyclic electrophiles were found to engage in the reaction (products **20–23**). With respect to practical features, it merits mention that unprotected anilines and phenols do not seem to interfere

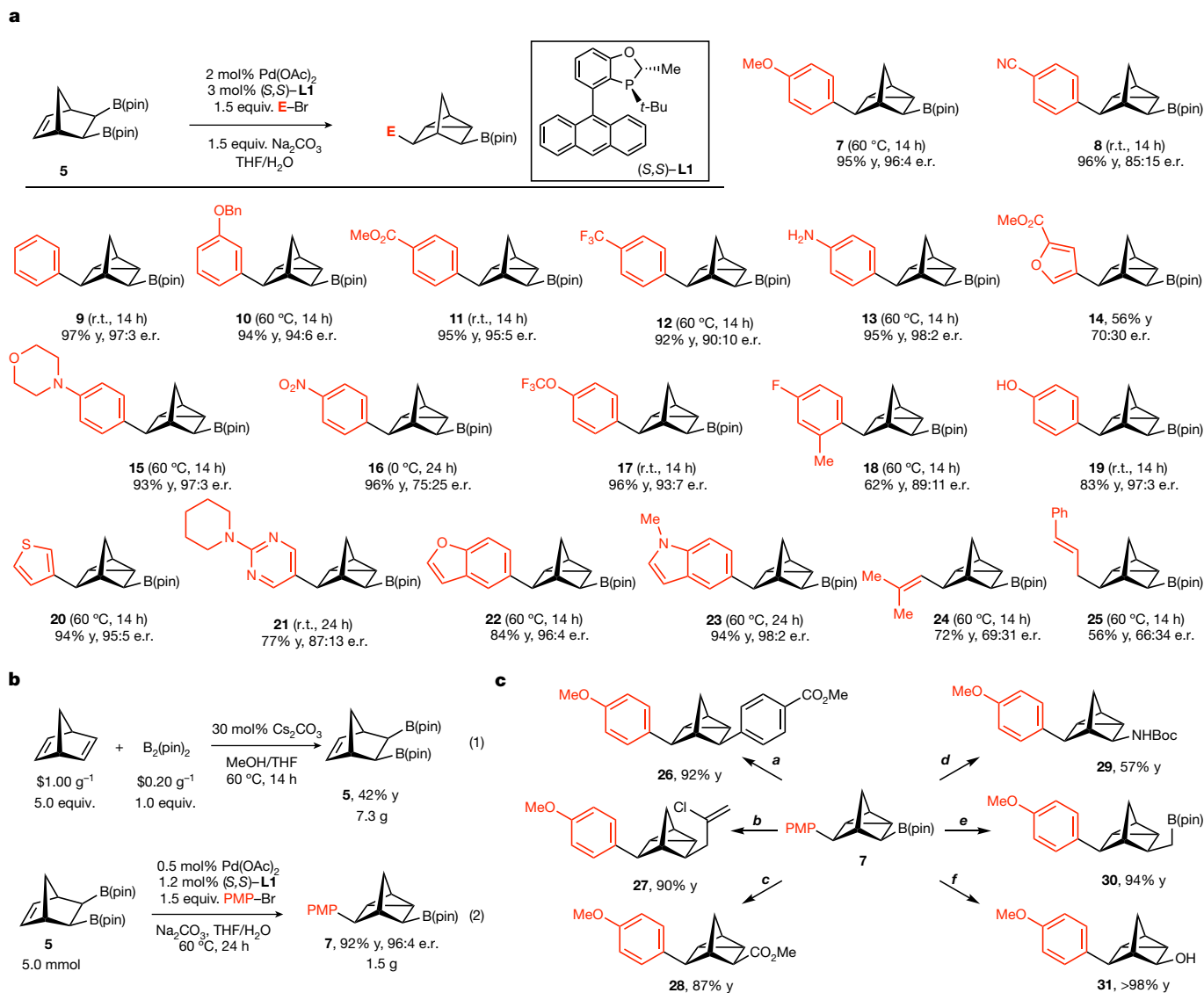


Fig. 2 | Pd-catalysed enantioselective construction of chiral nortricyclanes and their applications. a, Enantioselective coupling of **5** with electrophiles.

Yields (y) are isolated yields of the purified products. The enantiomeric ratio (e.r.) was determined by supercritical fluid chromatography. Product **21** required 5 mol% catalyst; product **25** from cinnamyl chloride. Absolute configuration determined by X-ray analysis of compound **20**. Bn, benzyl; Ph, phenyl; PMP, *para*-methoxyphenyl; r.t., room temperature. **b**, Single-step

with the process and that extension to non-aromatic electrophiles (**24**, **25**) seems promising. Practical application of the nortricyclane synthesis necessitates procedures that operate on a large scale as well as functional group transformations that effectively replace the remaining boronic ester in the catalytic reaction product. With regards to the first aspect, it was found that the bis(boronate) substrate **5** could be prepared on a multigram scale directly from inexpensive commercially available norbornadiene and B₂(pin)₂ (Fig. 2b, (1)). The reaction of **5** could be conducted on a 5 mmol scale and, so long as the reaction time is extended to 24 h, could be accomplished in good yield with only 0.5 mol% of palladium complex (Fig. 2b, (2)). As the transformations in Fig. 2c indicate, the boronic ester in product **7** can be replaced stereospecifically by Zn-based catalytic cross-coupling³¹ (**26**), Cu-catalysed allylation and carboxylation³² (**27** and **28**), direct amination³³ (**29**), homologation³⁴ (**30**) or conversion to the derived alcohol (**31**).

multigram-scale synthesis of *meso* bis(boronate) substrate **5** and its conversion to **7**. The prices shown are from Ambeed for B₂(pin)₂ and Sigma-Aldrich for norbornadiene. **c**, Transformations of boron-containing nortricyclane **7**. Conditions: a: *t*-BuLi then Zn(OAc)₂; G3-PdCPhos, Ar-Br. b: *t*-BuLi, then 20 mol% CuCN, 2,3-dichloropropene. c: *t*-BuLi, then 20 mol% CuCN, methylchloroformate. d: MeONH₂, *n*-BuLi; then Boc₂O. e: *n*-BuLi, CH₂BrCl. f: NaOH, H₂O₂.

Study of reaction mechanism

Mechanistic experiments provide insight into a likely pathway that leads to the product from substrate-derived cyclic ate complex **32** (Fig. 3a), a compound that has precedent from previous studies in our lab^{35,36}. The density functional theory calculations were used to determine the free energy profile of reaction paths originating from a palladium–olefin complex (**GSO**) that would be obtained by oxidative addition between LPd(0) and an electrophile, followed by association with **32**. In one route (path A, red), carbopalladation of the alkene (**TS1a**) delivers Pd(II) intermediate **GS1a**, which might then undergo an uncommon displacement of palladium(II) (refs. 37,38) by **TS2a** in which the organoboron serves as a nucleophile to simultaneously expel and reduce the metal. Alternatively (path B, blue), on binding to Pd(II), the alkene may become sufficiently electrophilic that nucleopalladation³⁹ by **TS1b** may generate **GS1b**, with direct reductive

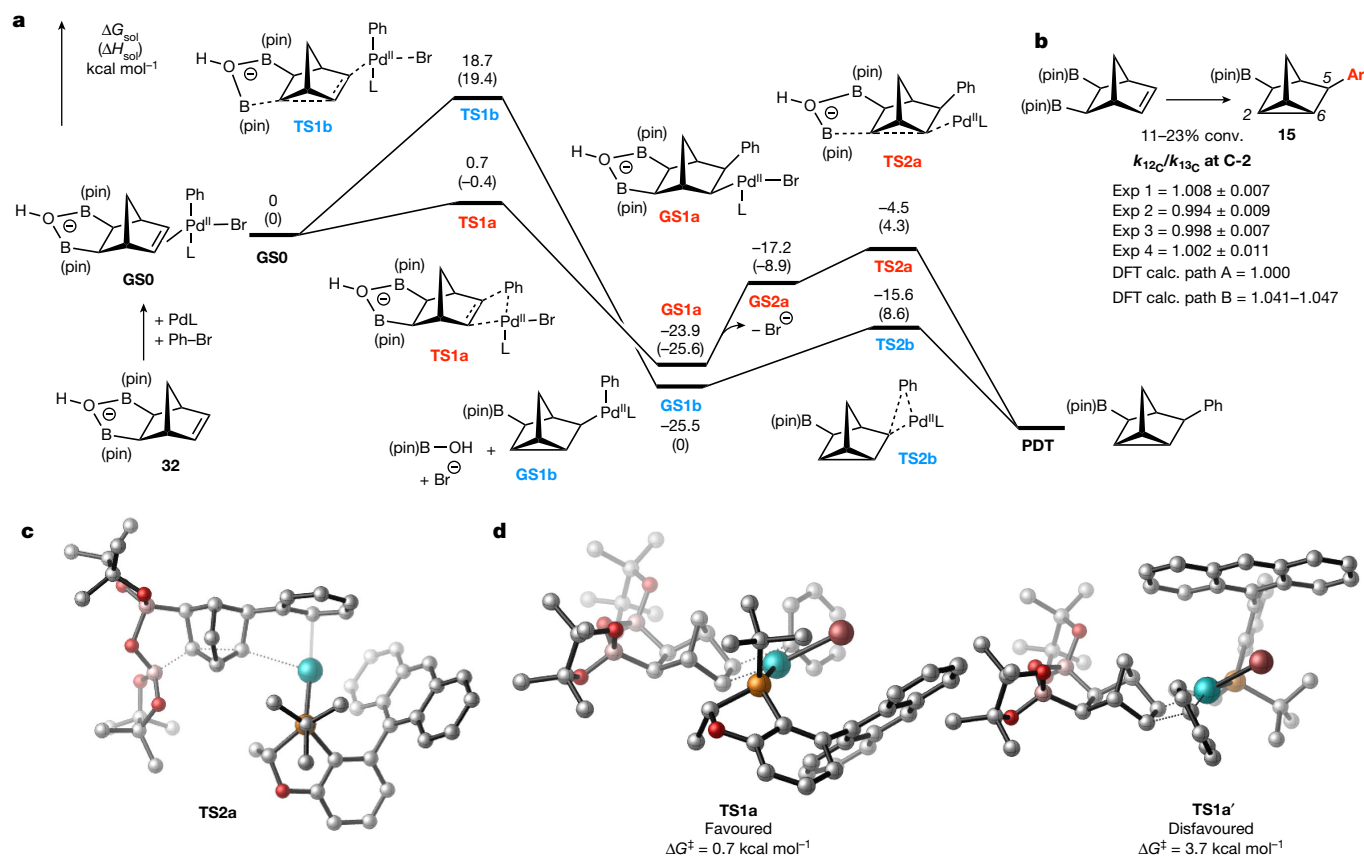


Fig. 3 | Mechanistic aspects of the Pd-catalysed nortricyclane synthesis. **a**, Calculated ground states and transition states for a carbopalladation-based mechanism (path A, red) and a nucleopalladation-based mechanism (path B, blue) for the Pd-catalysed synthesis of nortricyclanes. Methods: B3PW91(D3)/def2-TZVP/def2-TZVPP(Pd)/smd(THF)//B3LYP(D3BJ)/def2-SVP/def2-TZVP(Pd).

elimination then delivering the product. Calculations indicate that the carbopalladation pathway is lower in energy, with the olefin migratory insertion to give **GS1a** occurring by a much lower barrier compared with the nucleopalladation step (0.7 kcal mol⁻¹ versus 18.7 kcal mol⁻¹). From **GS1a**, dissociation of bromide allows the ring-closing reductive displacement of Pd(II) to occur with an accessible 19.4 kcal mol⁻¹ barrier by **TS2a** (Fig. 3c). Of note, calculation of the ¹³C kinetic isotope effects that would arise from **TS1a** (ref. 40) matches well with those determined experimentally by the Singleton method⁴¹ (Fig. 3b). Most notably, palladium-catalysed coupling of **5** exhibits a negligible KIE at C2, which is consistent with the carbopalladation-based mechanism but not the nucleopalladation pathway. Further mechanistic evidence in support of the carbopalladation pathway can be found in the Supplementary Information. According to these experiments, the carbopalladation step is enantio-determining and computational studies were undertaken to elucidate the origin of enantioinduction. As shown in Fig. 3d, the calculated structure of the transition state leading to the minor product stereoisomer places the CH₂ bridge of the bicycloheptane motif directly under the sterically encumbered anthracene ring system, whereas the conformation of the transition state for the major isomer avoids this interaction.

Biophysical and biochemical properties

To determine whether nortricyclane frameworks are suited for use in bioactive molecules, both enantiomers of **33** were prepared as chiral isosteric analogues of the fatty acid amide hydrolase (FAAH)

b, Selected ¹³C kinetic isotope effects as determined by the natural abundance isotope method of ref. 41. **c**, Calculated transition state geometry for **TS2a**. **d**, Calculated stereochemistry-determining transition states leading to the major and minor enantiomers of carbopalladation product. ΔG[‡], energy difference between the transition state and the reactants.

inhibitor URB597 (refs. 42,43) and subjected to analysis (Fig. 4a). Of note, although the isosteric analogues have similar molecular weight as URB597, they have 10-fold increased solubility in the aqueous buffer while maintaining comparable lipophilicity (α log *D*). Apart from these positive attributes, the isosteric analogues exhibit measurably increased metabolic stability relative to URB597 as determined by mouse liver microsomal assay. To probe for the effect of chirality on biological activity, compounds (*R,S*)-**33** and (*S,R*)-**33** were compared by quantitative activity-based protein profiling experiments (for complete details, see the Supplementary Information). Mouse-brain homogenates were treated with varying concentrations of each compound before exposure to the serine hydrolase-directed activity-based probe fluorophosphonate (FP)-biotin. FP-labelled proteins were enriched on streptavidin beads and analysed by quantitative tandem mass tagging (TMT)-based mass spectrometry⁴⁴. Of note, although (*R,S*)-**33** and (*S,R*)-**33** were not as potent as URB597, the isosteric analogues retained the ability to inhibit FAAH and show different potencies: (*R,S*)-**33** exhibits an IC₅₀ of 0.50 μM, whereas that for (*S,R*)-**33** is more than 4 μM. Furthermore, the incorporation of the isosteres did not alter the proteome-wide selectivity of URB597, as both isosteres maintained high selectivity for FAAH over 23 other detected serine hydrolases.

In a second comparative analysis, the enantiomeric tricyclanes (*S,R*)-**34** and (*R,S*)-**34** were prepared as analogues of the Hedgehog (Hh) pathway⁴⁵ inhibitor, sonidegib⁴⁶ (Fig. 4b). Aberrant Hedgehog signalling has been implicated in certain types of cancer, and sonidegib inhibits activation of this pathway by interacting with the associated signal transducer, Smoothened (SMO). Comparison of stereoisomeric

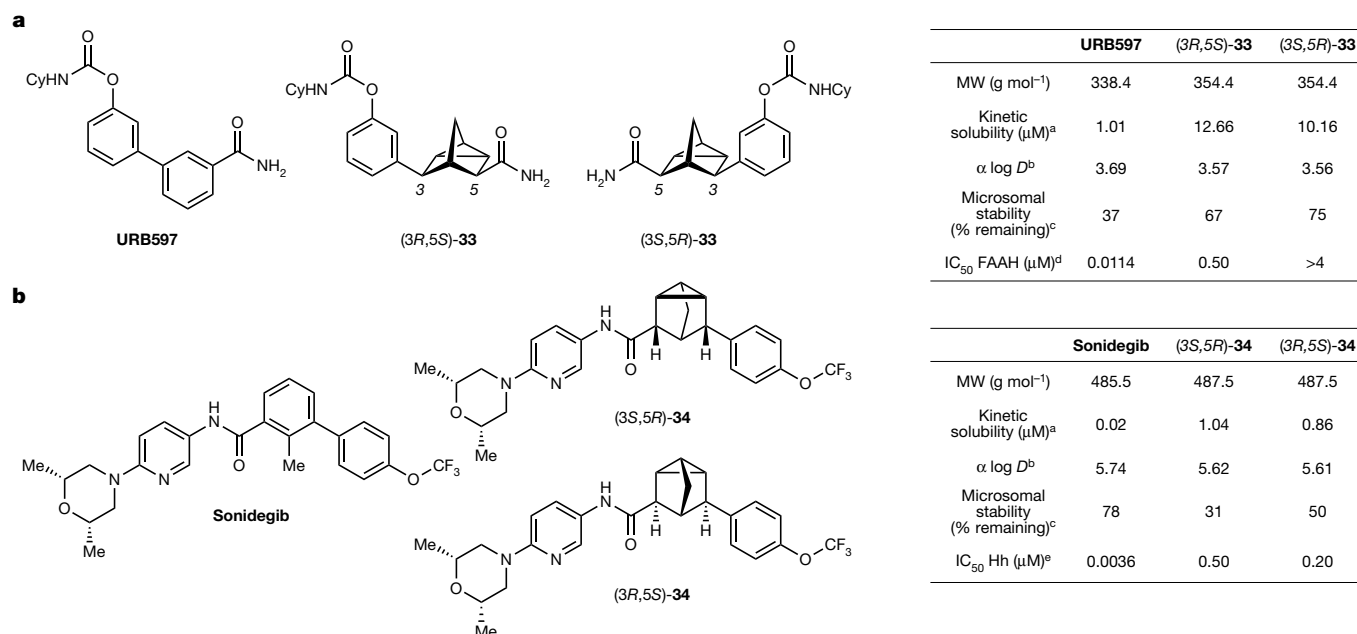


Fig. 4 | Pharmacokinetic analysis of nortricycane-based analogues and their inhibitor activity against FAAH and Hedgehog signalling. a, Analysis of the FAAH inhibitor URB597 and analogues (*R,S*)-**33** and (*S,R*)-**33**. **b**, Analysis of sonidegib and analogues (*S,R*)-**34** and (*R,S*)-**34**. ^aDetermined in PBS buffer at pH = 7.4 after 24 h. ^bDefined as log *D* at pH = 7.4; determined by HPLC analysis.

isosteres **34** was conducted by three independent IC₅₀ determinations using a dual luciferase assay in Shh-LIGHT2 cells⁴⁷, and it was found that the enantiomeric sonidegib analogues inhibit Hh signalling with submicromolar potencies (IC₅₀ = 0.50 μM and 0.20 μM for the (*S,R*)-**34** and (*R,S*)-**34**, respectively), and with a modest, at best (*P* = 0.26), difference in activity for the two stereoisomers. Overall, the preliminary analysis of the nortricycane isosteres is in line with expectations: replacement of arenes with C(*sp*³)-based isosteres results in a 10-fold to 50-fold improvement in solubility, probably as a result of increased solvation of the C(*sp*³) analogues as suggested in refs. 48,49. Of note, the stereochemistry-dependent activity of nortricycane isosteres (**33** and **34**) seems to reinforce the notion that shape matters when binding to biological targets, and it is notable that differences in activity are detected even with isosteric analogues that have not been reoptimized to bind their target receptors. We can anticipate that as chiral ligands are fine-tuned for affinity, the absolute stereochemistry of the isosteres is likely to become increasingly important.

In summary, catalytic enantioselective nortricycane synthesis provides an efficient entry into chiral *meta* benzene bioisosteres, and the chirality of the tricyclic motif plays a part in the shape-dependent binding of the derived ligands with their target receptors. The ease and robustness of the catalytic synthesis, combined with the biophysical properties of nortricyclanes should provide opportunities for pharmaceutical development. Although further exploration of the tricycloheptane skeleton seems warranted, further study of the unique catalytic process that leads to its formation—especially the C–C bond-forming reductive displacement of Pd(II)—may provide new strategies for reaction development.

Online content

Any methods, additional references, Nature Portfolio reporting summaries, source data, extended data, supplementary information, acknowledgements, peer review information; details of author contributions and competing interests; and statements of data and code availability are available at <https://doi.org/10.1038/s41586-024-07865-4>.

^cDetermined at 30 min in the presence of male CD-1 mouse liver microsomes with NADPH. ^dInhibition of FAAH determined by quantitative mass spectrometry profiling. See the Supplementary Information for a complete dataset. ^eInhibition of Hedgehog signalling in Shh-LIGHT2 cells. Data are an average of five replicates.

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Article

Reporting summary

Further information on research design is available in the Nature Portfolio Reporting Summary linked to this article.

Data availability

Crystal structure data for compound **20** have been deposited at the Cambridge Structure Data Centre (CCDC 2325329). Mass spectrometry proteomics data have been deposited to the ProteomeXchange Consortium through the PRIDE⁵⁰ partner repository with the dataset identifier PXD051400. All other data are available in the main text or in the Supplementary Information.

50. Perez-Riverol, Y. et al. The PRIDE database resources in 2022: a hub for mass spectrometry-based proteomics evidences. *Nucleic Acids Res.* **50**, D543–D552 (2022).

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Author contributions M.Z. and J.P.M. conceptualized the study; M.Z. and M.C. conducted the synthetic investigation; H.L. performed the density functional theory calculations; B.X. and E.W. studied the hydrolase inhibition; B.R.S. and J.K.C. investigated the Hh inhibition; J.P.M. assisted with the writing of the initial draft; and all authors contributed to the review and editing of the paper.

Competing interests J.P.M., M.Z. and H.L. declare that provisional patent applications have been filed on boron-containing cyclic molecules (US provisional application 63/509,173 and international patent application PCT/US24/34873). All other authors have no competing interests.

Additional information

Supplementary information The online version contains supplementary material available at <https://doi.org/10.1038/s41586-024-07865-4>.

Correspondence and requests for materials should be addressed to James K. Chen, Eranthie Weerapana or James P. Morken.

Peer review information *Nature* thanks Antonia Stepan and the other, anonymous, reviewer(s) for their contribution to the peer review of this work.

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Reporting Summary

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Statistics

For all statistical analyses, confirm that the following items are present in the figure legend, table legend, main text, or Methods section.

n/a Confirmed

- The exact sample size (n) for each experimental group/condition, given as a discrete number and unit of measurement
- A statement on whether measurements were taken from distinct samples or whether the same sample was measured repeatedly
- The statistical test(s) used AND whether they are one- or two-sided
Only common tests should be described solely by name; describe more complex techniques in the Methods section.
- A description of all covariates tested
- A description of any assumptions or corrections, such as tests of normality and adjustment for multiple comparisons
- A full description of the statistical parameters including central tendency (e.g. means) or other basic estimates (e.g. regression coefficient) AND variation (e.g. standard deviation) or associated estimates of uncertainty (e.g. confidence intervals)
- For null hypothesis testing, the test statistic (e.g. F , t , r) with confidence intervals, effect sizes, degrees of freedom and P value noted
Give P values as exact values whenever suitable.
- For Bayesian analysis, information on the choice of priors and Markov chain Monte Carlo settings
- For hierarchical and complex designs, identification of the appropriate level for tests and full reporting of outcomes
- Estimates of effect sizes (e.g. Cohen's d , Pearson's r), indicating how they were calculated

Our web collection on [statistics for biologists](#) contains articles on many of the points above.

Software and code

Policy information about [availability of computer code](#)

Data collection Tecan Infinite M1000/M1000 PRO microplate reader. Mass spectrometry data was collected on an Orbitrap Exploris 240 running Thermo Scientific Xcalibur v4

Data analysis Microsoft Excel (version:16.81), Prism 9 (version:9.3.1) Mass spectrometry data was analyzed using Thermo Proteome Discoverer v2.4 running SequestHT and Percolator

For manuscripts utilizing custom algorithms or software that are central to the research but not yet described in published literature, software must be made available to editors and reviewers. We strongly encourage code deposition in a community repository (e.g. GitHub). See the Nature Research [guidelines for submitting code & software](#) for further information.

Data

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All manuscripts must include a [data availability statement](#). This statement should provide the following information, where applicable:

- Accession codes, unique identifiers, or web links for publicly available datasets
- A list of figures that have associated raw data
- A description of any restrictions on data availability

Any data generated or analyzed during this study, associated protocols, and materials are available from the corresponding author on reasonable request. Mass spectrometry data for this study will be deposited in ProteomeXchange via the PRIDE partner repository. Protein databases employed for MS/MS searches were obtained from the third party UniprotKB protein database.

Field-specific reporting

Please select the one below that is the best fit for your research. If you are not sure, read the appropriate sections before making your selection.

Life sciences Behavioural & social sciences Ecological, evolutionary & environmental sciences

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Life sciences study design

All studies must disclose on these points even when the disclosure is negative.

| | |
|-----------------|--------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| Sample size | To determine the inhibitory potency of each compound in the SHH signaling assays, at least five biological replicates were used for each compound concentration. Three independent experiments were conducted to obtain average IC50 and s.e.m. values. These sample sizes were based on our previous experience with these assays and the number of replicates required to obtain reliable and reproducible results. For hydrolase assays, mouse brains were purchased from commercial sources, and brain homogenates from a single mouse brain were treated with the indicated concentrations of compounds |
| Data exclusions | None. |
| Replication | Descriptions of the number of replicates and error bars are included in the main text and method section. Student's t-test (two-tailed, unpaired) analysis in Prism were used to compare results and calculate P values. For hydrolase assays, the number of technical replicates has been indicated and the average ratio of each entry has been reported in supplementary data. |
| Randomization | Randomization is not relevant to our SHH signaling experiments. All samples in these assays were treated uniformly and subjected to the same data analysis procedures. |
| Blinding | The SHH signaling experiments were not blinded because the quantitative results are not subject to human biases. For hydrolase assays, blinding was not performed in this study as the proteomics measurements were collected and analyzed by computer software. |

Reporting for specific materials, systems and methods

We require information from authors about some types of materials, experimental systems and methods used in many studies. Here, indicate whether each material, system or method listed is relevant to your study. If you are not sure if a list item applies to your research, read the appropriate section before selecting a response.

Materials & experimental systems

| n/a | Involvement in the study |
|-------------------------------------|-----------------------------------------------------------------|
| <input checked="" type="checkbox"/> | <input type="checkbox"/> Antibodies |
| <input type="checkbox"/> | <input checked="" type="checkbox"/> Eukaryotic cell lines |
| <input checked="" type="checkbox"/> | <input type="checkbox"/> Palaeontology and archaeology |
| <input type="checkbox"/> | <input checked="" type="checkbox"/> Animals and other organisms |
| <input checked="" type="checkbox"/> | <input type="checkbox"/> Human research participants |
| <input checked="" type="checkbox"/> | <input type="checkbox"/> Clinical data |
| <input checked="" type="checkbox"/> | <input type="checkbox"/> Dual use research of concern |

Methods

| n/a | Involvement in the study |
|-------------------------------------|-------------------------------------------------|
| <input checked="" type="checkbox"/> | <input type="checkbox"/> ChIP-seq |
| <input checked="" type="checkbox"/> | <input type="checkbox"/> Flow cytometry |
| <input checked="" type="checkbox"/> | <input type="checkbox"/> MRI-based neuroimaging |

Eukaryotic cell lines

Policy information about [cell lines](#)

| | |
|----------------------------------------------------------------------|-----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| Cell line source(s) | Shh-LIGHT2 cells: This NIH-3T3 cell-derived line was provided by Phil Beachy (Stanford University). The generation of this line is described in Taipale et. al., Nature (2000) 406:1005-1009. |
| Authentication | We confirmed SHH ligand-dependent activation of the firefly luciferase reporter in this line. No further authentication was performed. |
| Mycoplasma contamination | The cells were not tested for mycoplasma contamination. |
| Commonly misidentified lines (See ICLAC register) | None of the misidentified lines were used in this study. |

Animals and other organisms

Policy information about [studies involving animals](#); [ARRIVE guidelines](#) recommended for reporting animal research

| | |
|-------------------------|-----------------------------------------------------------------------------------------------------------------------------------------------|
| Laboratory animals | One C57Bl/6 strain, 12-week-old female mouse was used in this study. Frozen C57BL/6J mouse brains were purchased from The Jackson Laboratory. |
| Wild animals | Wild animals are not involved in this study. |
| Field-collected samples | The study did not involve any field-collected samples. |
| Ethics oversight | The use of the animal was approved by Stanford University Administrative Panel on Laboratory Animal Care (Protocol Number 14145). |

Note that full information on the approval of the study protocol must also be provided in the manuscript.