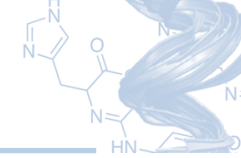


Combining ETKDG and NOE-derived Distance Bounds

Prof. Sereina Riniker
10th RDKit UGM, October 15, 2021

What is ETKDG?



ETKDG = Experimental torsions – knowledge (terms) – distance geometry

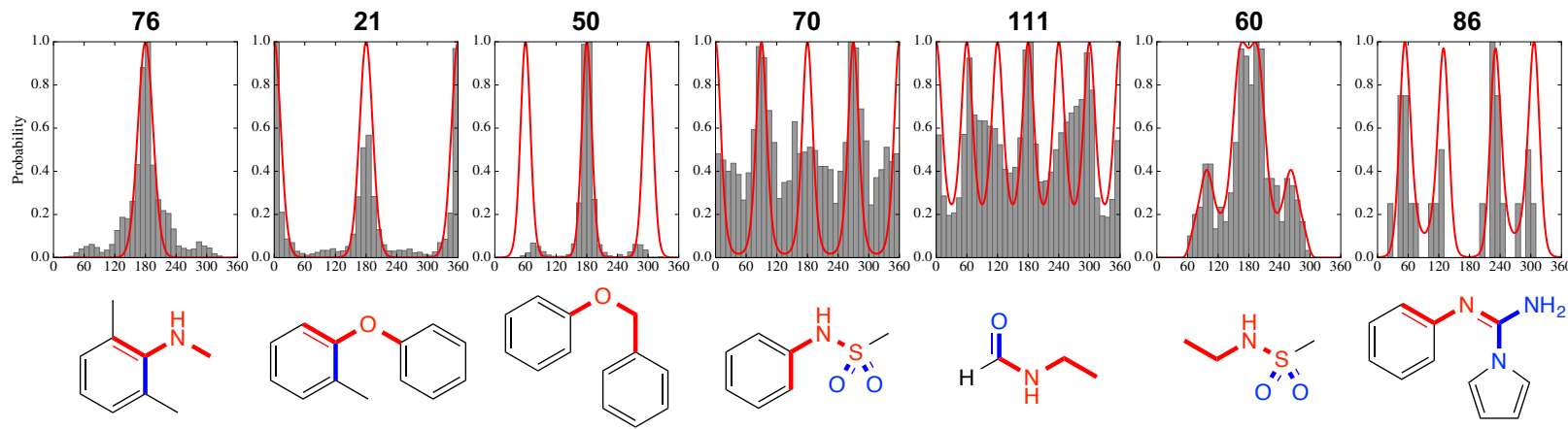
Experimental torsional-angle preferences of *acyclic* bonds from CSD

Examples of fitted torsion potentials:

$$V_{tors}(x) = \sum_{i=1}^6 K_i (1 + \cos(\delta_i) \cos(m_i x))$$

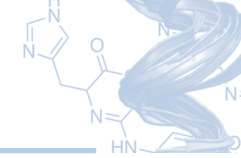
387 SMARTS patterns from:

- Version 1:** Schärfer et al., *J. Med. Chem.*, 56, 2016 (2013).
- Version 2:** Guba et al., *J. Chem. Inf. Model.*, 56, 1 (2016).



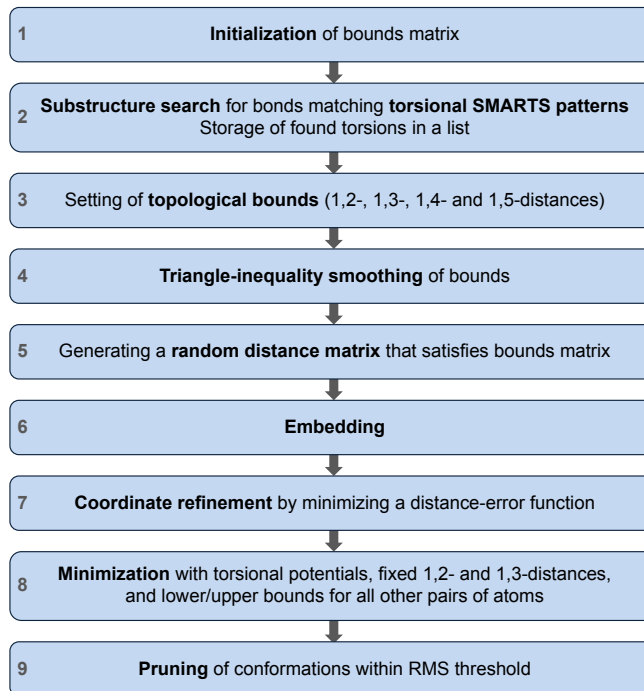
Knowledge terms: Flat sp² carbons, straight triple bonds

What is ETKDG?

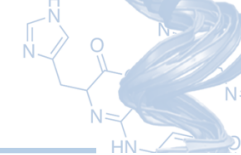


ETKDG = Experimental torsions – knowledge (terms) – distance geometry

Conformer generation workflow



ETKDG version 3



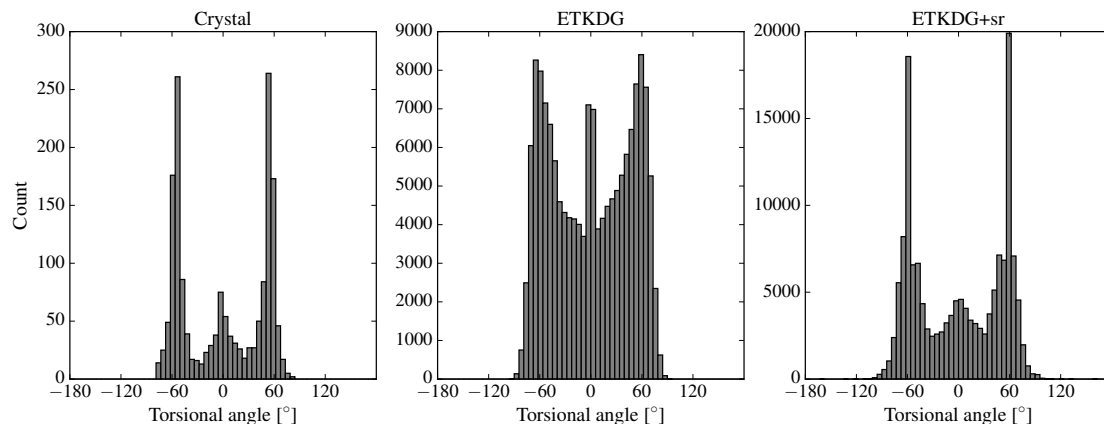
What about aliphatic cyclic bonds?

Distinguishing between small rings and macrocycles

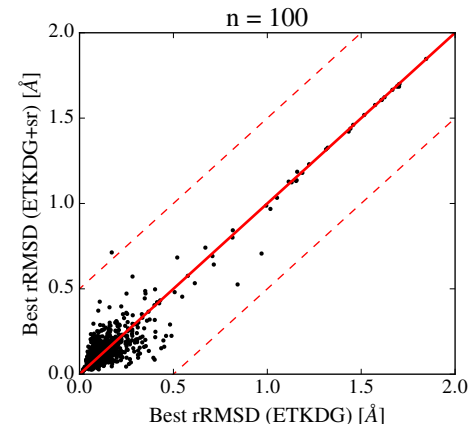
Small rings:

- New torsion SMARTS pattern derived (105 in total)
- Maximum ring size = 8
- Bridged systems: Bonds in >1 SSSR ring excluded, if ring shares >2 bonds with another ring → excluded

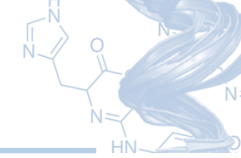
Six-membered rings



Best rRMSD on CSD test set



ETKDG version 3



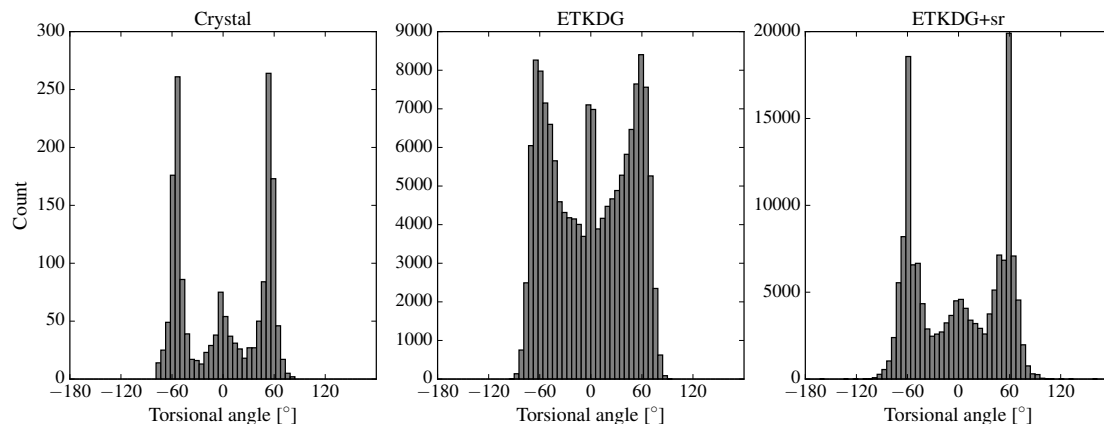
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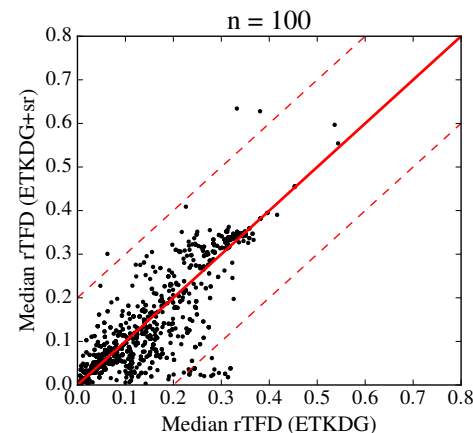
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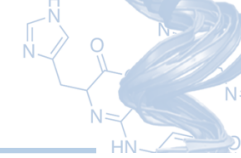
Six-membered rings



Median rTFD on CSD test set



ETKDG version 3



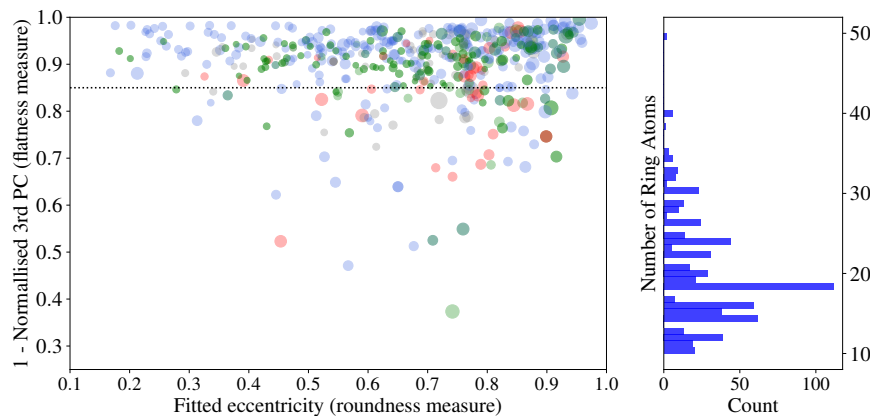
What about aliphatic cyclic bonds?

Distinguishing between small rings and macrocycles

Macrocycles:

- Cyclic bonds in rings > 8 behave similar to acyclic bonds (in terms of torsions)
- SMARTS patterns from version 2 adapted for macrocyclic bonds
- Changes to existing 1,4-bonds subroutines to treat macrocyclic bonds as acyclic

Flatness and ring size in dataset



Size of markers = macrocyclic ring size

Colours = source of the macrocycles:

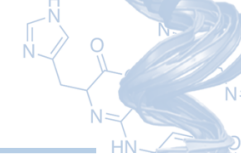
blue = CSD

green = Prime

red = BIRD

grey = Mac10

ETKDG version 3



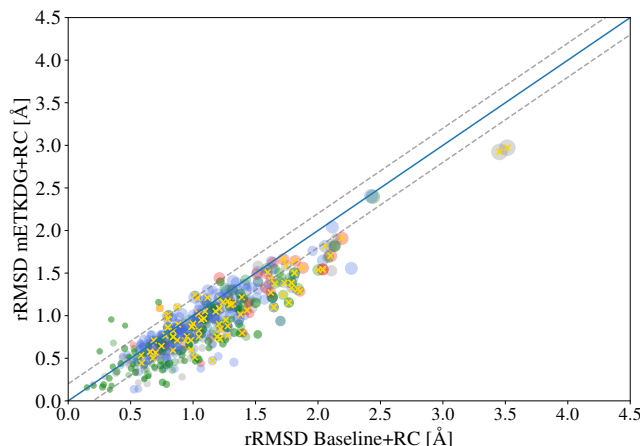
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mETKDG versus ETKDGv2 (random coordinates)



Size of markers = macrocyclic ring size

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blue = CSD

green = Prime

red = BIRD

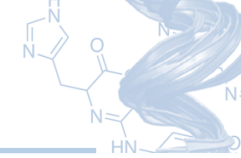
grey = Mac10

Blue solid lines: $y = x$

Gray dotted lines: ± 0.20 Å deviation

Golden crosses on a marker = compounds for which the experimental structure has a flatness value of <0.85

ETKDG version 3

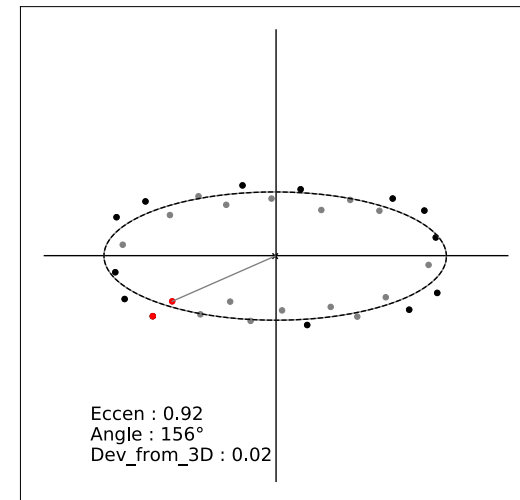
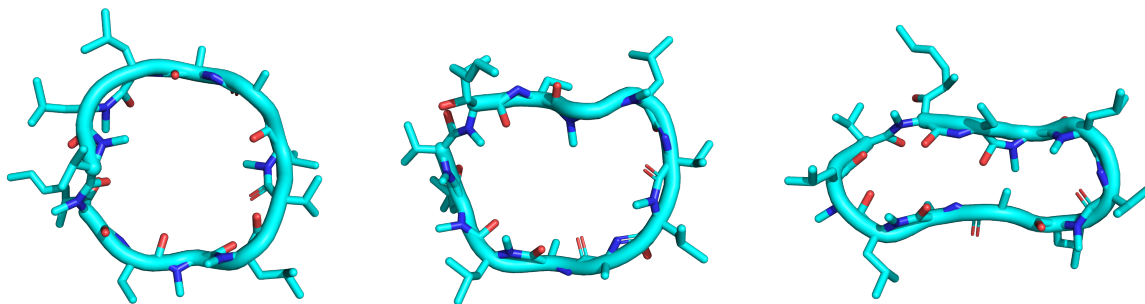


What about aliphatic cyclic bonds?

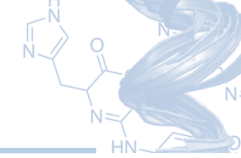
Distinguishing between small rings and macrocycles

Macrocycles: Eccentricity

- Crystal conformations of macrocycle backbone are often ellipsoid



ETKDG version 3



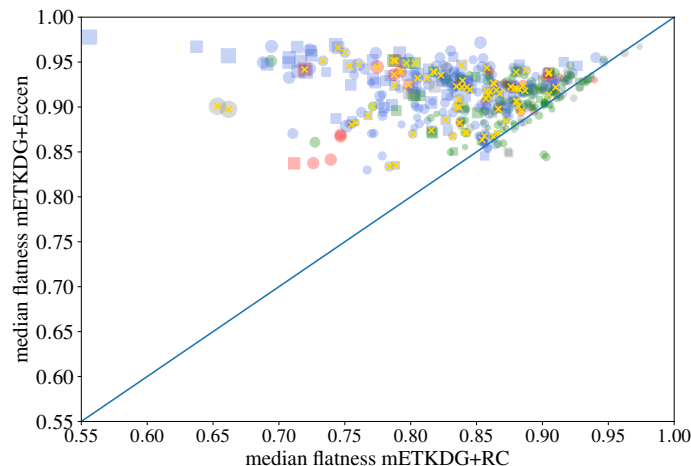
What about aliphatic cyclic bonds?

Distinguishing between small rings and macrocycles

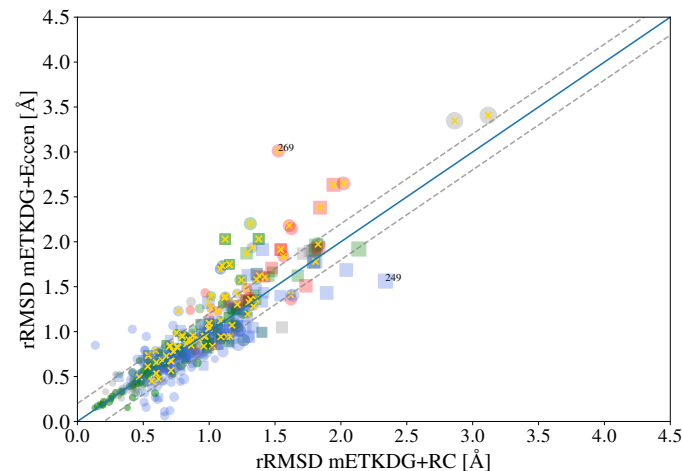
Macrocycles: Eccentricity

- Crystal conformations of macrocycle backbone are often ellipsoid
- Eccentricity constraints to target search in conformational space

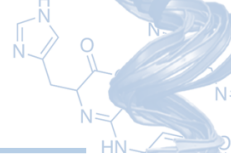
Median flatness



Best rRMSD



ETKDG + NOE



Can we use NOE-derived distance bounds to direct conformational search?

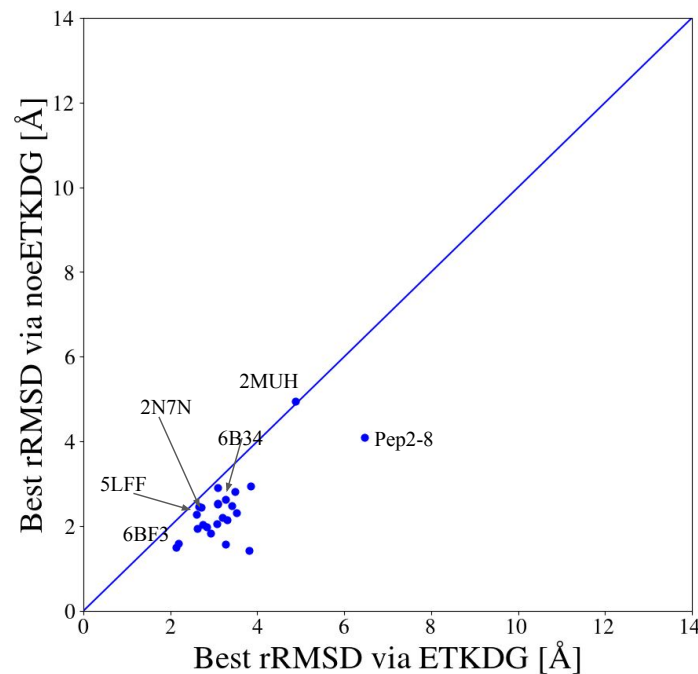
Goal: Conformer bundles fulfilling experimental restraints

ETKDG workflow:

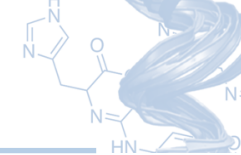
- Modification of upper bounds in the molecular bounds matrix
- Advantage: No force field involved (easy to handle unnatural amino acids, non-peptidic motifs)

Dataset + pipeline:

- 24 peptides (23 cyclic + 1 linear)
- Parsing: XPLOR-type files
- 2000 conformer generated with noeETKDG
- Bundles: Subselection of 9-20 conformers
 - Random
 - Most diverse macrocycle conformations
 - Lowest energy
 - Least NOE violations (LICUV)
 - NAMFIS [JACS, 117, 1027 (1995)]



ETKDG + NOE

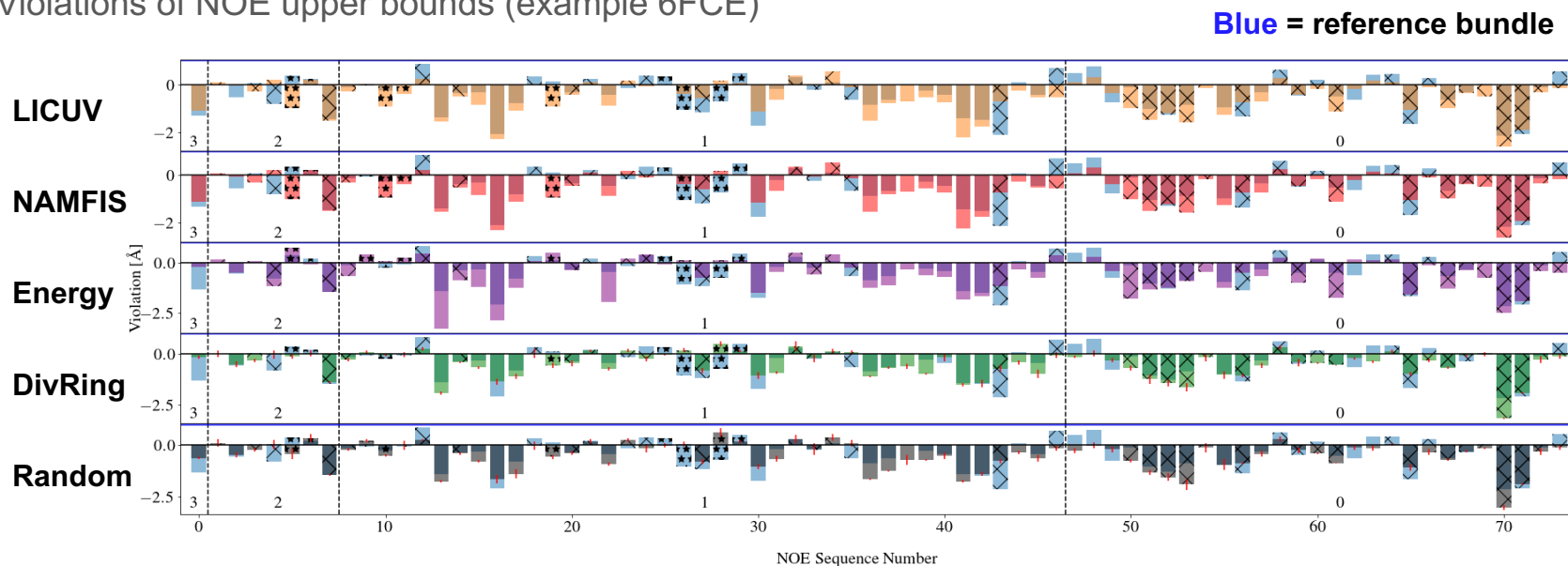


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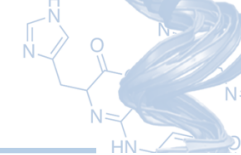
Goal: Conformer bundles fulfilling experimental restraints

Comparison of bundles:

Violations of NOE upper bounds (example 6FCE)



ETKDG + NOE



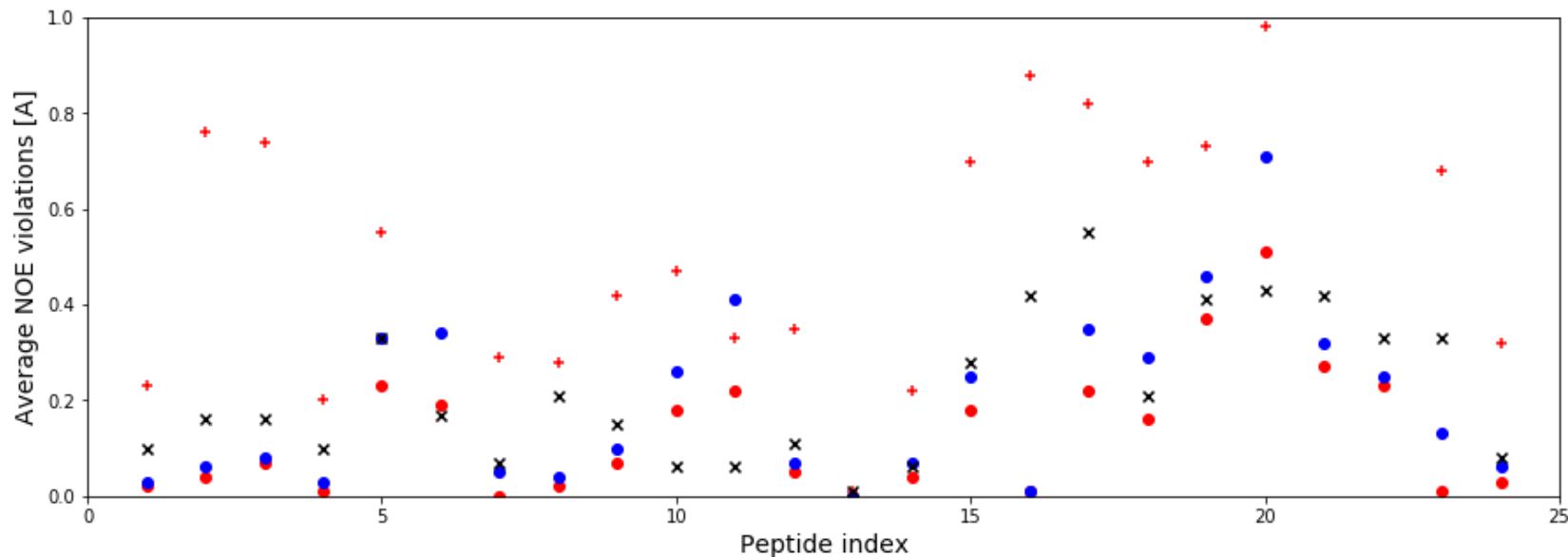
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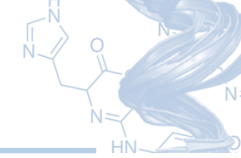
Comparison of bundles:

Average NOE violations

+ ETKDG / LICUV
o noeETKDG / LICUV
o noeETKDG / NAMFIS
x reference bundle



ETKDG + NOE



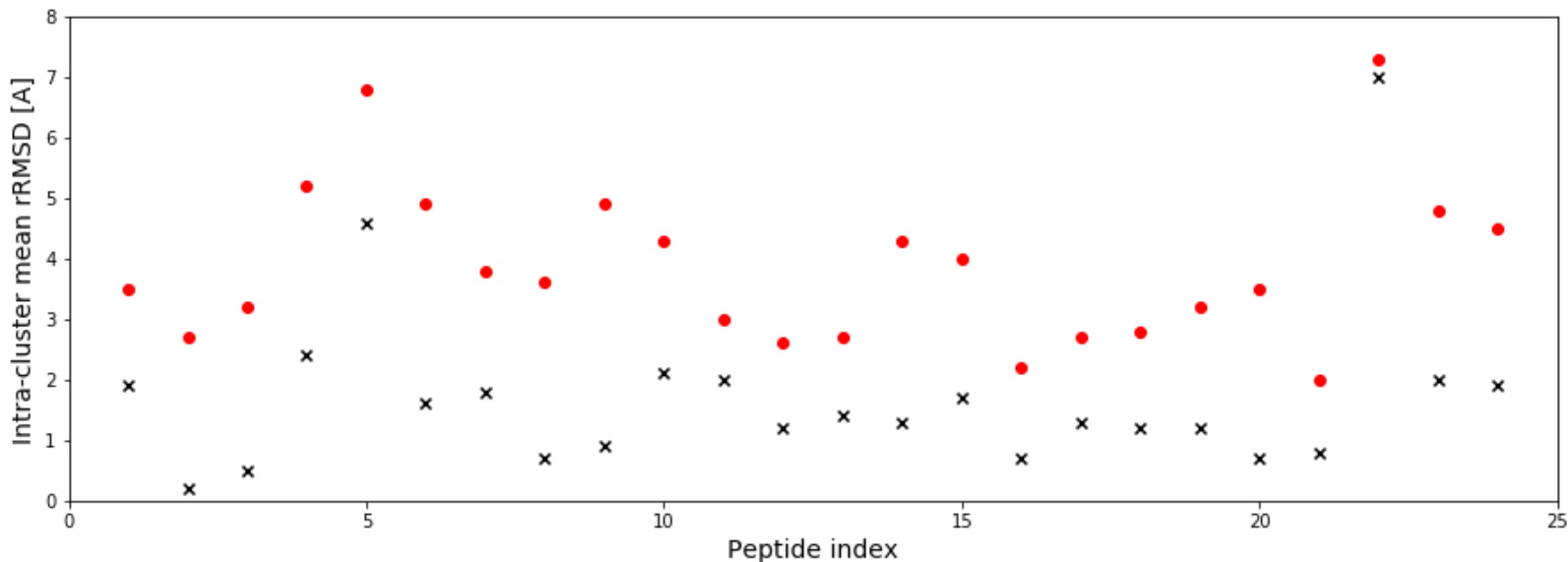
Can we use NOE-derived distance bounds to direct conformational search?

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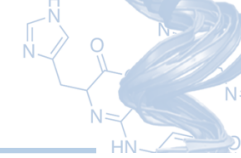
Comparison of bundles:

Intra-bundle macrocycle diversity (average rRMSD)

o noeETKDG / LICUV
x reference bundle



ETKDG + NOE



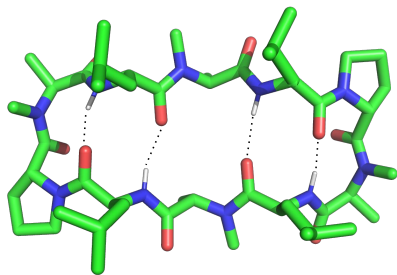
Can we use NOE-derived distance bounds to direct conformational search?

Goal: Conformer bundles fulfilling experimental restraints

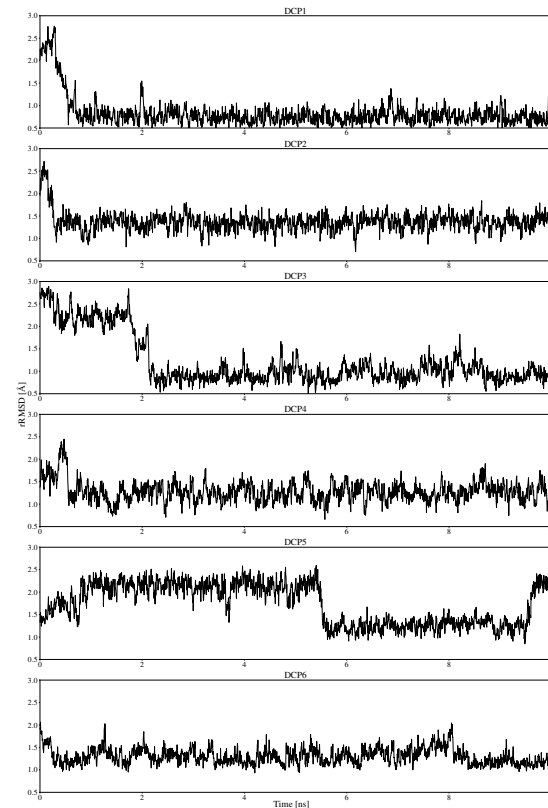
Post-processing with time-averaged NOE-restraining MD:

Use noeETKDG conformers as good starting points

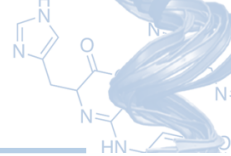
Example: Six cyclic decapeptides with 4 transannular hydrogen bonds



[A]	noeETKDG LICUV		Restraining MD Low-energy frames		Ref. bundle
	Mean NOE viol.	Intra- bundle rRMSD	Mean NOE viol.	Intra- bundle rRMSD	Intra- bundle rRMSD
DCP1	0.2	4.0	0.4	1.5	1.7
DCP2	0.01	2.2	1.1	1.3	0.7
DCP3	0.2	2.7	0.5	1.7	1.3
DCP4	0.2	2.8	0.3	1.5	1.2
DCP5	0.4	3.2	0.6	2.1	1.2
DCP6	0.5	3.5	0.6	1.7	0.7



ETKDG + NOE



Availability

External Python package using RDKit to modify conformer generator

customETKDG on Github:

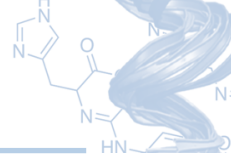
<https://github.com/rinikerlab/customETKDG>

Wang, et al., *J. Chem. Inf. Model.*, submitted (2021).

Example Jupyter notebook:

https://github.com/rinikerlab/customETKDG/blob/api/examples/demo_rdkit_ugm_2021.ipynb

Conclusions



Macrocyclic ETKDG (version3)

- Cyclic bonds behave as acyclic ones
- Tools for restricting the conformational search space:
 - Eccentricity constraints
 - Custom pairwise Coulombic interactions (CPCIs)

ETKDG + NOE-derived distance restraints

- Modification of upper distance bounds in ETKDG
- noeETKDG conformers more similar to structures in reference bundles (compared to ETKDG)
- Bundle selection: least NOE violations appears to be a good strategy
- noeETKD bundles contain more diverse ring structures (compared to reference bundles)
- Post-processing with time-averaged NOE-restraining MD simulations