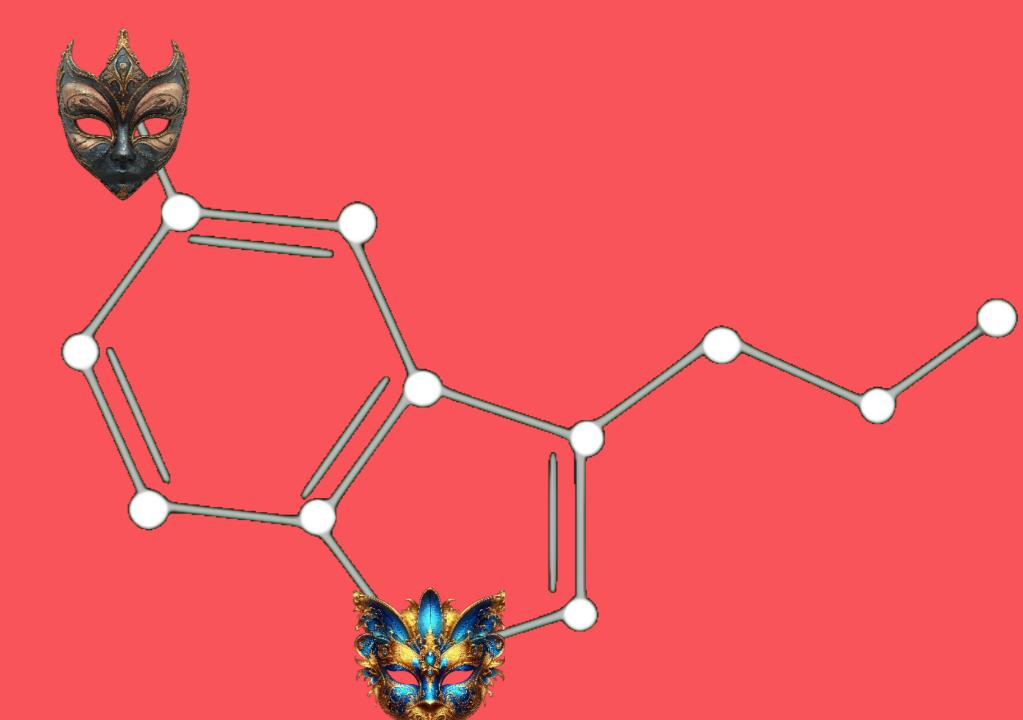


ENHANCING CHEMICAL EXPLAINABILITY THROUGH COUNTERFACTUAL MASKING



Łukasz Janisiów

Marek Kochańczyk

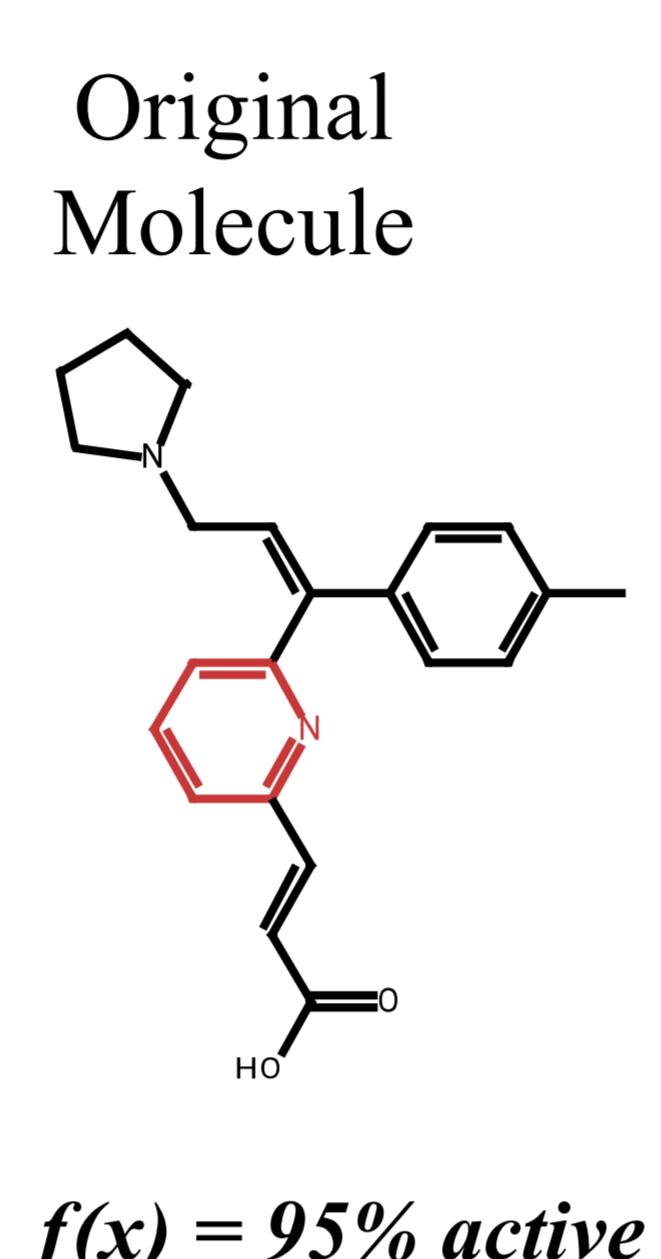
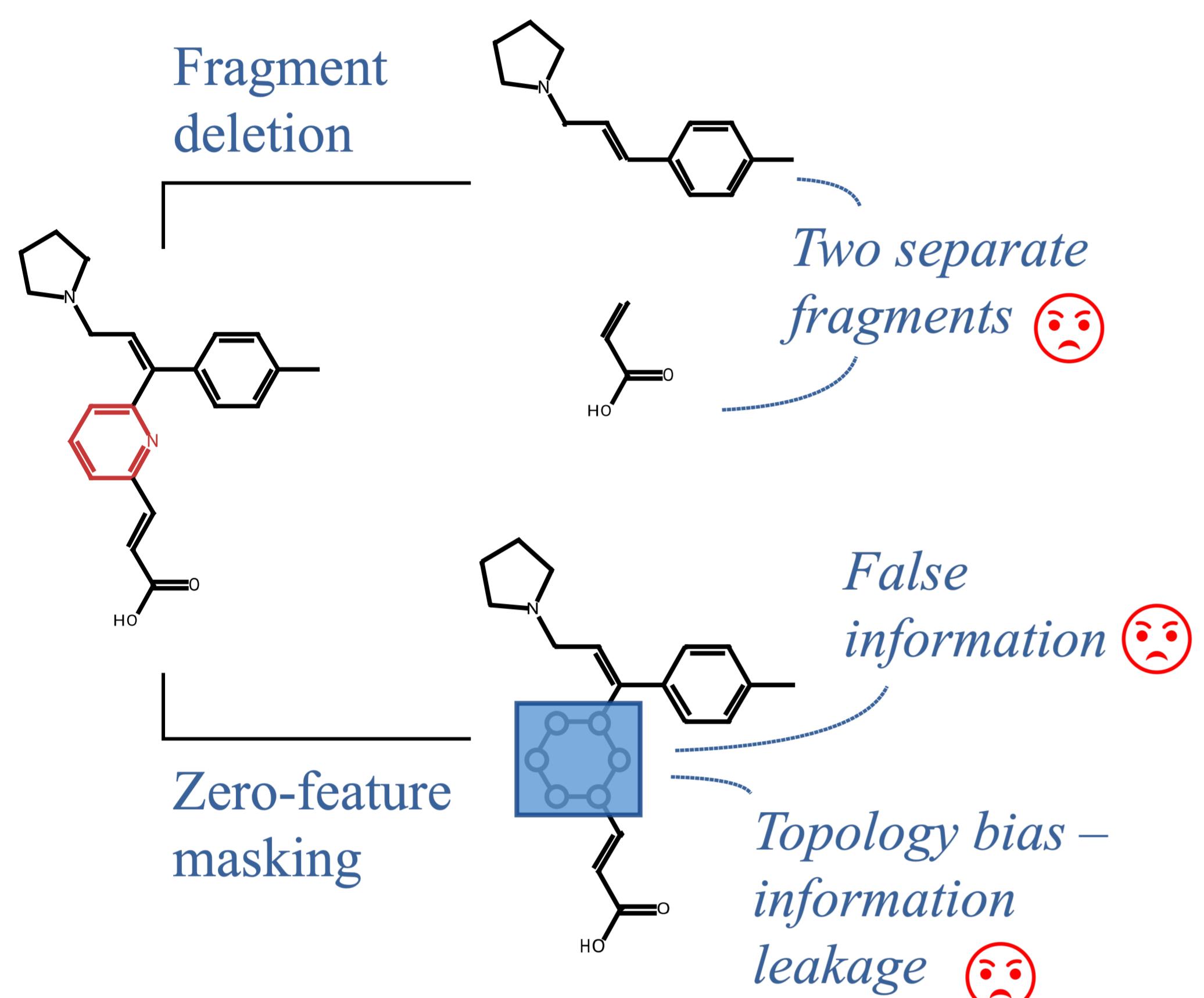
Bartosz Zieliński

Tomasz Danel

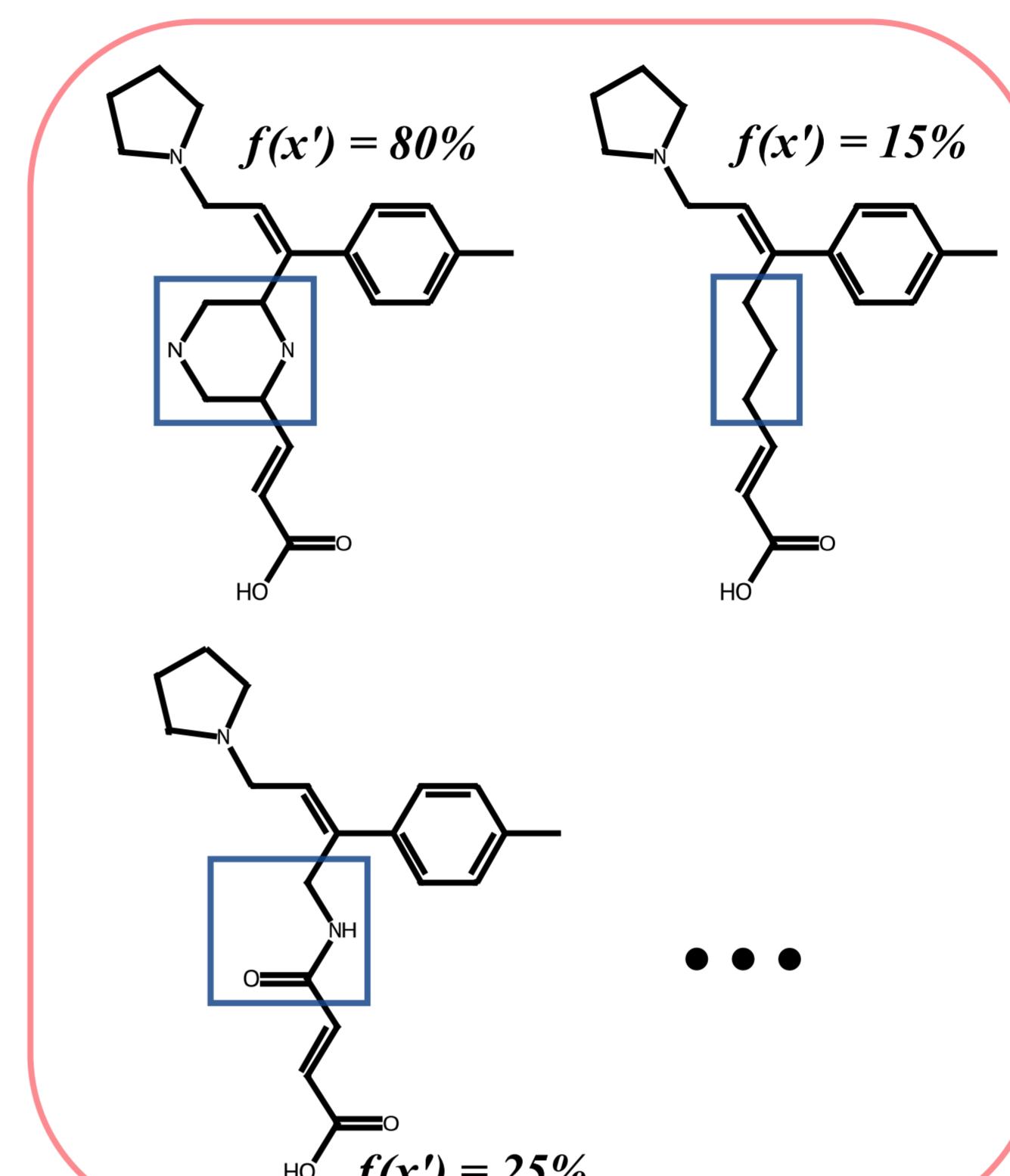
Jagiellonian University, Poland

MOTIVATION: Molecular Masking Is Broken

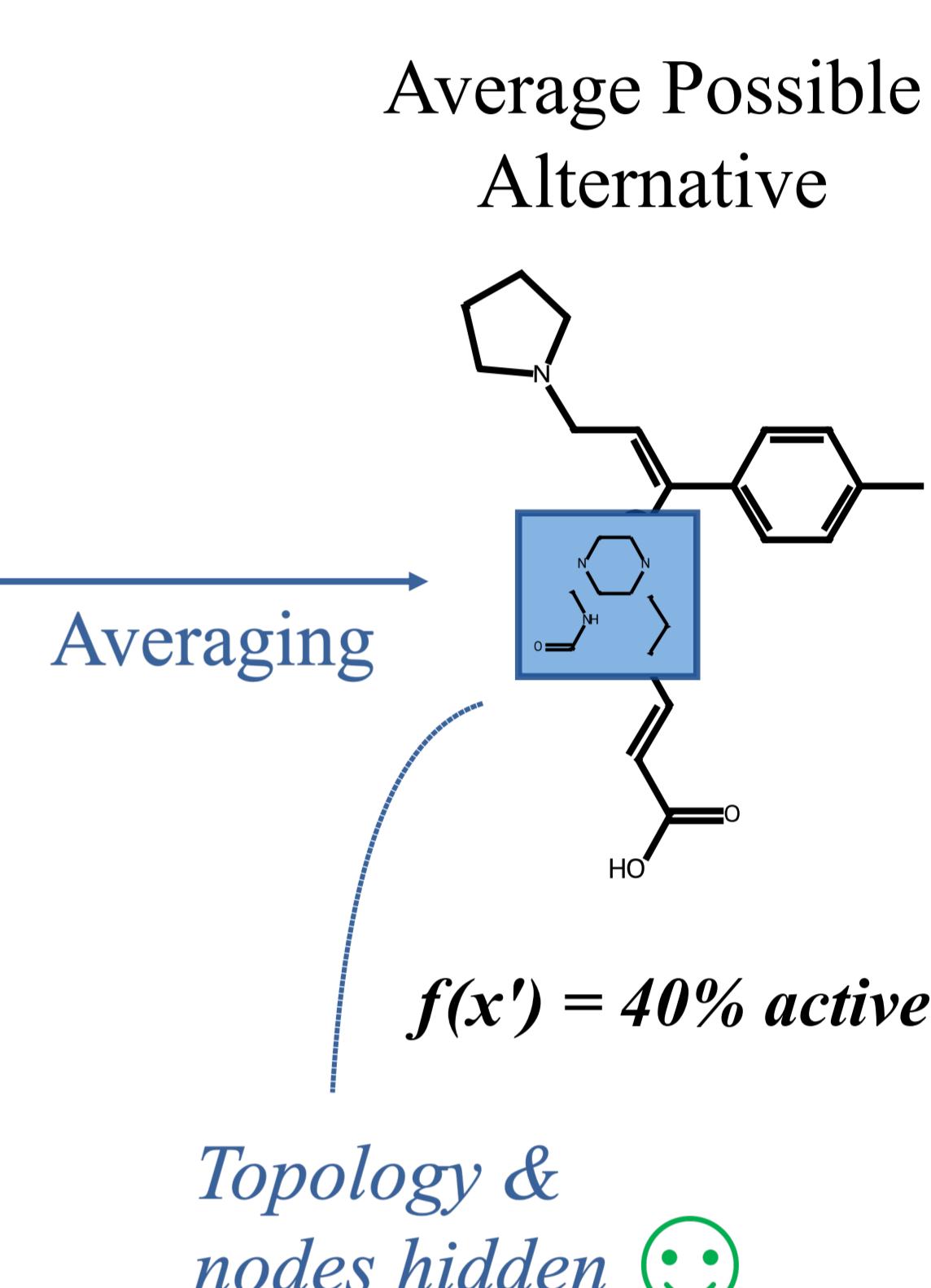
When atoms or bonds are naively masked (by zeroing features or deleting parts of molecule), the resulting molecules often become **chemically implausible or physically impossible**, producing examples that fall outside the training distribution. This **out-of-distribution problem** undermines the reliability of both the explanations and their evaluation metrics.



Deleting & Inpainting

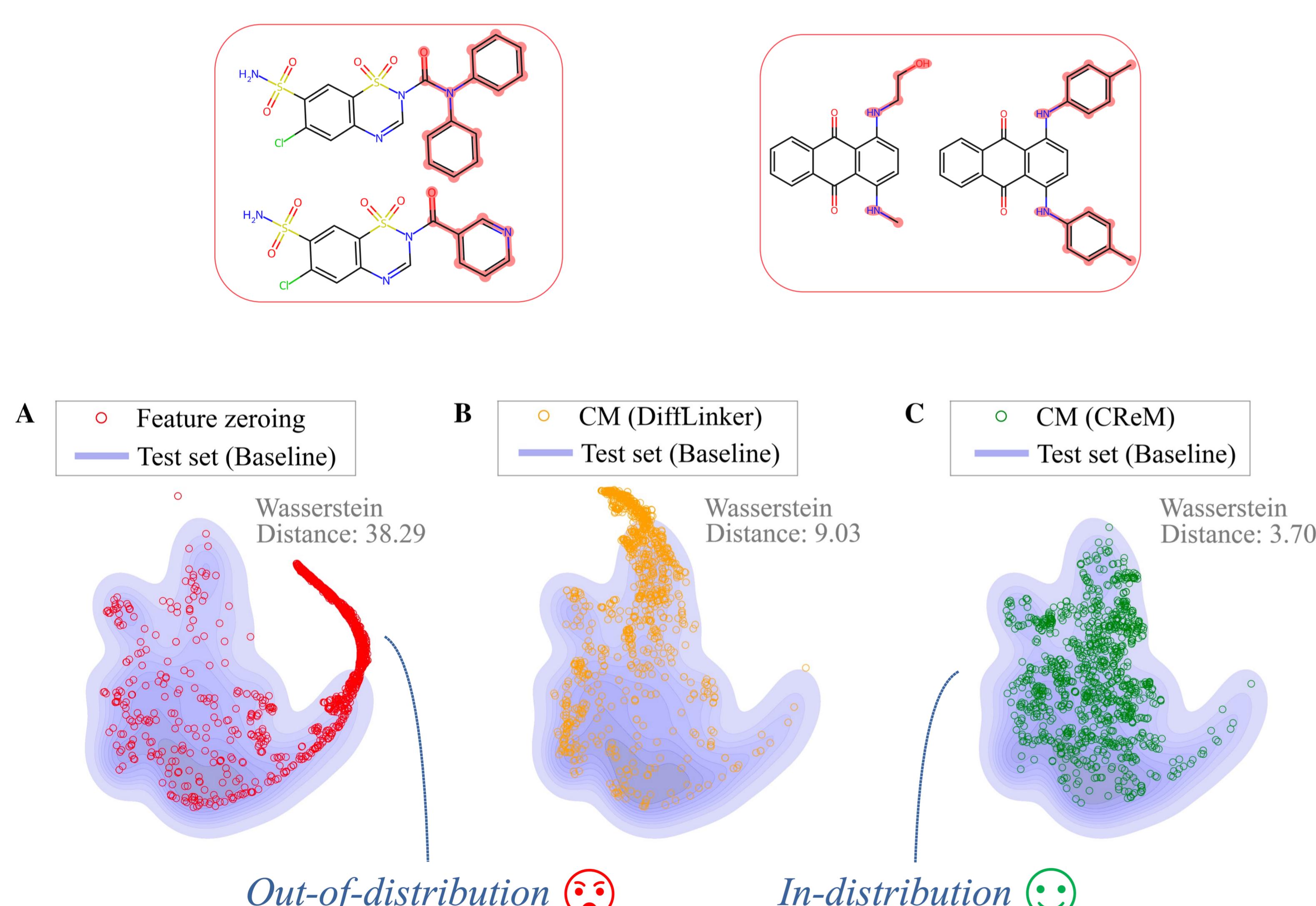


Context-fragment aware generation (DiffLinker & CReM)



Evaluation of Fragment Masking

To evaluate different masking techniques, we used the common substructure pair dataset, which contains pairs of molecules with the same core structure. In each pair, we masked the fragments that were not shared and measured the difference in the model's predictions. An ideal masking technique should completely obscure the masked fragment, resulting in identical predictions for both molecules in the pair.



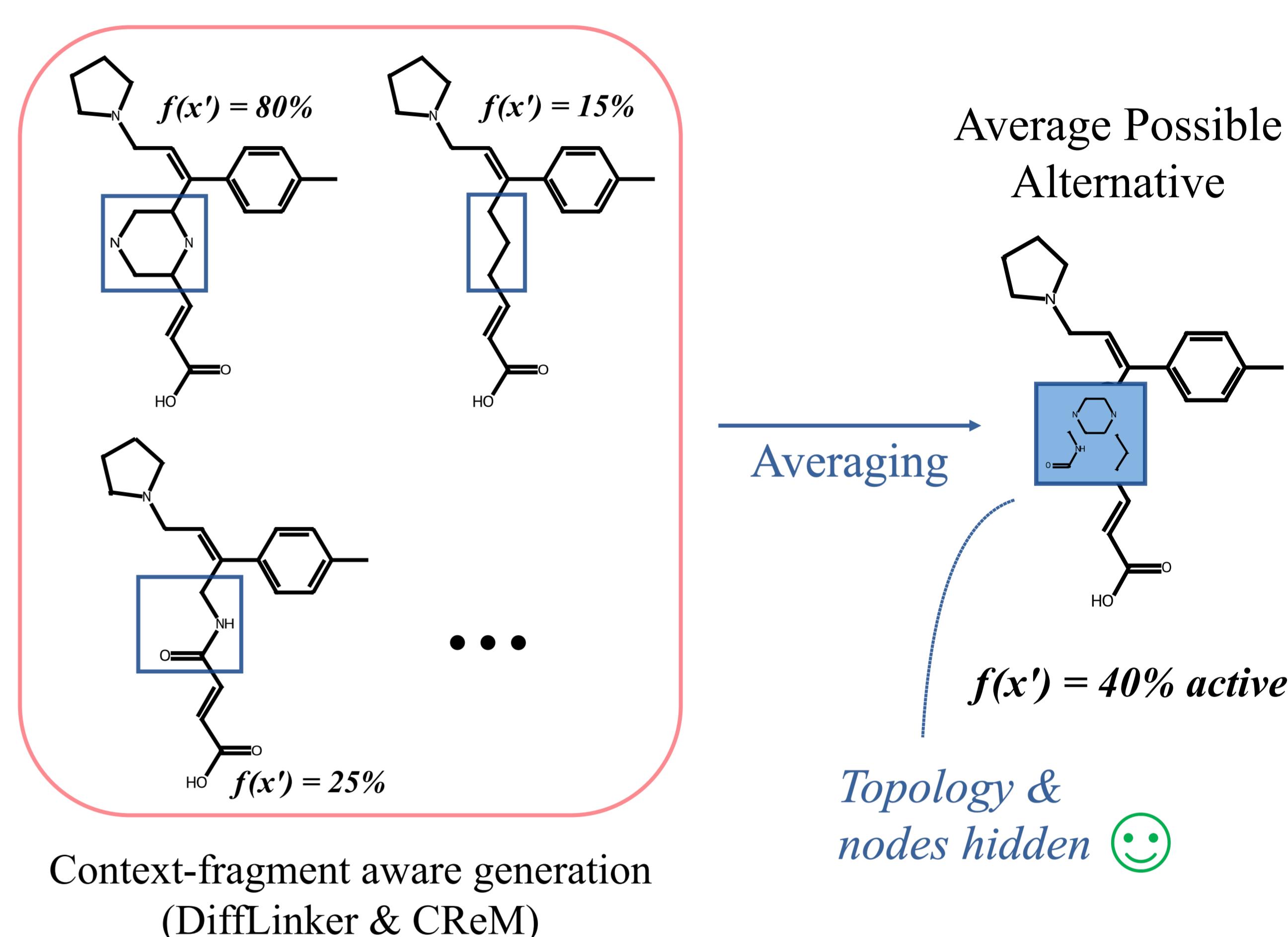
Masking	Single Anchor			Multiple Anchors			Both		
	$ \Delta\hat{y} \downarrow$	Validity \uparrow	Size \uparrow	$ \Delta\hat{y} \downarrow$	Validity \uparrow	Size \uparrow	$ \Delta\hat{y} \downarrow$	Validity \uparrow	Size \uparrow
No masking	2.81	100%	517	2.52	100%	638	2.91	100%	783
Feature zeroing	2.08	100%	517	1.60	100%	638	1.80	100%	783
CM (CReM)	1.68	88%	454	1.32	53%	341	1.72	65%	510
CM (DiffLinker)	1.81	67%	347	1.59	61%	386	1.55	65%	508

Acknowledgments

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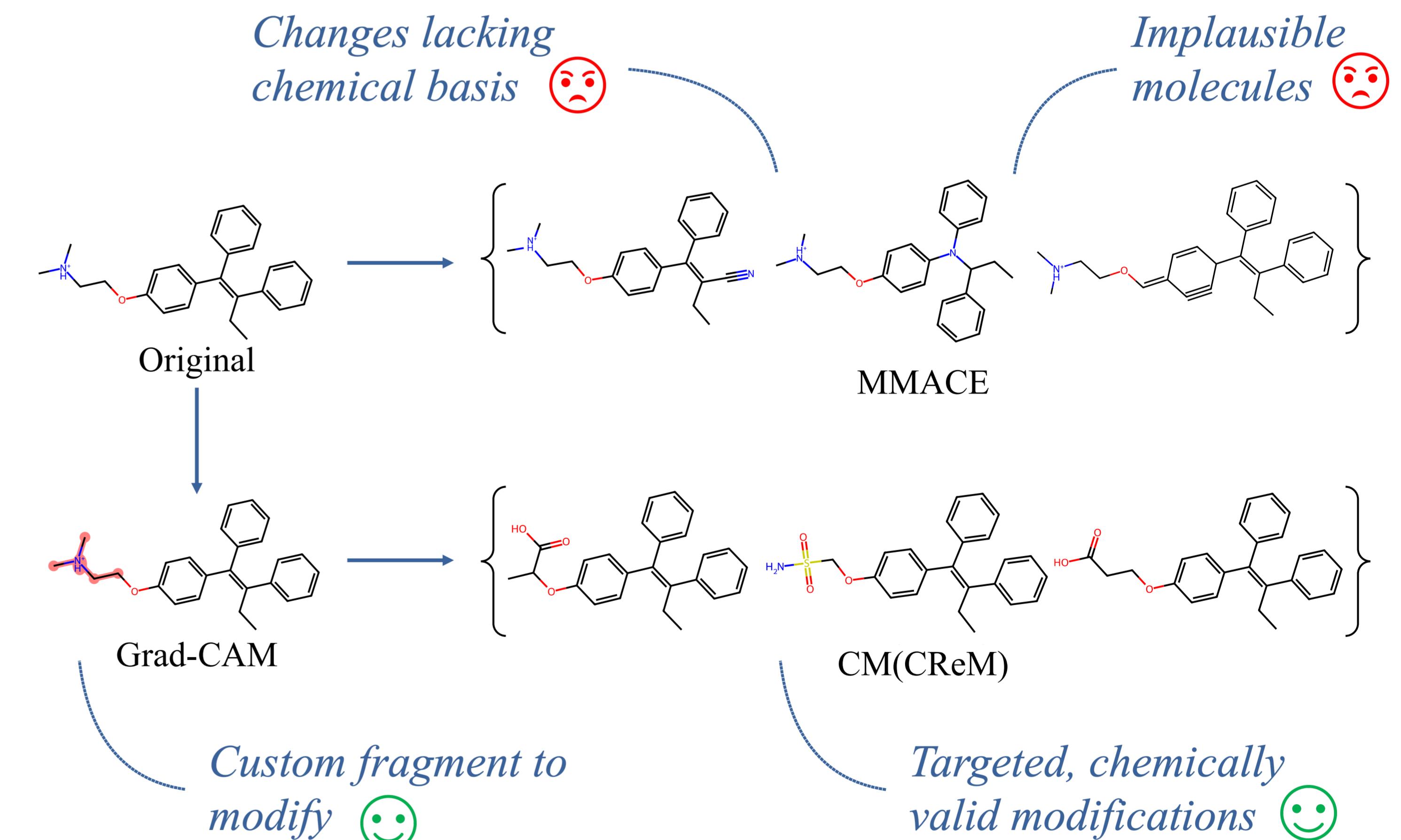
SOLUTION: Counterfactual Masking (CM)

Counterfactual Masking masks by replacing the important molecular fragments with chemically plausible alternatives. This enables a direct comparison between the model's prediction for the original molecule and the average prediction across valid counterfactuals (i.e., all possible alternatives).



Can We Generate Counterfactuals?

Although originally designed for masking tasks, CM can effectively generate realistic, chemically valid counterfactual examples by ensuring that molecules stay within the data distribution. In comparison to other counterfactual generation methods, it also enables targeted, local modifications to specific fragments, which can be easily interpreted by people with a chemistry background.



Dataset	Method	BSR \uparrow	Success Rate \uparrow	Similarity \uparrow	Diversity \uparrow	SA \downarrow
CYP 3A4	GNExplainer	0.16 ± 0.04	0.16 ± 0.04	0.33 ± 0.00	0.00 ± 0.00	3.67 ± 0.04
	NN (Nearest Neighbour)	1.00 ± 0.00	1.00 ± 0.00	0.37 ± 0.00	0.66 ± 0.00	2.53 ± 0.01
	MMACE	0.27 ± 0.00	1.00 ± 0.00	0.60 ± 0.01	0.52 ± 0.01	3.34 ± 0.02
	CM (CReM)	0.21 ± 0.01	1.00 ± 0.00	0.52 ± 0.04	0.46 ± 0.04	2.71 ± 0.08
	CM (DiffLinker)	0.41 ± 0.02	1.00 ± 0.00	0.41 ± 0.05	0.41 ± 0.01	3.91 ± 0.18

