


 Cite this: *RSC Adv.*, 2024, 14, 4345

# Discovery of new tetrazines for bioorthogonal reactions with strained alkenes *via* computational chemistry†

 Michal Májek \* and Matej Trtúšek

Tetrazines are widely employed reagents in bioorthogonal chemistry, as they react readily with strained alkenes in inverse electron demand Diels–Alder reactions, allowing for selective labeling of biomacromolecules. For optimal performance, tetrazine reagents have to react readily with strained alkenes, while remaining inert against nucleophiles like thiols. Balancing these conditions is a challenge, as reactivity towards strained alkenes and nucleophiles is governed by the same factor – the energy of unoccupied orbitals of tetrazine. Herein, we utilize computational chemistry to screen a set of tetrazine derivatives, aiming to identify structural elements responsible for a better ratio of reactivity with strained alkenes vs. stability against nucleophiles. This advantageous trait is present in sulfone- and sulfoxide-substituted tetrazines. In the end, the distortion/interaction model helped us to identify that the reason behind this enhanced reactivity profile is a secondary orbital interaction between the strained alkene and sulfone-/sulfoxide-substituted tetrazine. This insight can be used to design new tetrazines for bioorthogonal chemistry with improved reactivity/stability profiles.

Received 21st December 2023

Accepted 25th January 2024

DOI: 10.1039/d3ra08712c

[rsc.li/rsc-advances](https://rsc.li/rsc-advances)

## Introduction

Biological systems are an important target of molecular science research, containing biomacromolecules, and being the host of fast, selective enzymatic reactions and precise molecular sensing systems. To comprehensively study these complex systems, the capability to perform chemical reactions selectively and in real-time within these systems is essential. However, this is a difficult task due to the inherent hostility of the biological environment towards typical organic reagents. Cells contain high concentrations of free reactive nucleophilic centers like thiols and amines from amino acid residues. This is combined with an aqueous reaction medium containing oxygen and different electrolytes. Bioorthogonal chemistry introduced new reactions that enable selective chemistry in such complex systems.<sup>1–6</sup>

For bioorthogonal reagents, robustness under physiological conditions is crucial. Yet, they need to provide the required reaction at a very high rate, as the local concentrations of the reagents are usually relatively low when compared to a typical organic reaction. Having these requirements in mind, researchers have been successful in developing multiple classes of bioorthogonal reactions. The first-generation reactions, such

as Staudinger ligation<sup>7,8</sup> and copper-catalyzed alkyne–azide cycloaddition,<sup>9,10</sup> served as a proof of concept but exhibited significant limitations in real systems. Staudinger ligation relies on phosphorus(III) reagents, unstable in the presence of oxygen, while copper-based catalysts have only limited compatibility with living organisms. The breakthrough came with the second generation of bioorthogonal reactions – strain-promoted reactions like azide–alkyne cycloadditions (SPAAC),<sup>11,12</sup> alkyne–nitrene cycloadditions (SPANAC),<sup>13,14</sup> and the tetrazine (*s*-tetrazine) ligation.<sup>15–17</sup> These reactions proceed rapidly while having superior stability of the required reagents at physiological conditions in comparison with the first-generation reactions. Tetrazine ligation has multiple attractive properties concerning the development of systems applicable to living organisms. The reactivity of the tetrazine can be modulated by modifying the substitution pattern on the tetrazine core to achieve the desired reaction rates.<sup>18</sup> Moreover, the tetrazine system undergoes significant electronic structure reorganization during the ligation reaction, allowing the design of switchable fluorescent tags, where fluorescence is extinguished or enhanced after the ligation process.<sup>19</sup>

When designing a new tetrazine reagent suitable for bioorthogonal ligation reactions a balance between their reactivity and stability is crucial. Such tetrazines need to have a high rate constant for the required ligation reaction. Mechanistically, tetrazine ligation reactions are usually inverse-electron demand Diels–Alder reactions (IEDDA).<sup>20</sup> Tetrazines serve as the diene component in these reactions and therefore, an efficient interaction of one of their unoccupied orbitals with the HOMO

Comenius University Bratislava, Faculty of Natural Sciences, Department of Organic Chemistry, Mlynská Dolina, Ilkovičova 6, 842 15 Bratislava, Slovakia. E-mail: [michal.majek@uniba.sk](mailto:michal.majek@uniba.sk)

† Electronic supplementary information (ESI) available. See DOI: <https://doi.org/10.1039/d3ra08712c>



orbital of the dienophile is required to enhance the reaction rate. This can be accomplished by introducing electron-withdrawing functional groups on the tetrazine core.<sup>21</sup> Typically ester- and pyridyl-substituents can be used for this purpose. The presence of electron-withdrawing groups lowers the energy of unoccupied orbitals and promotes the desired iEDDA reaction.

Unfortunately, this also leads to decreased stability of the tetrazine reagent in physiological conditions, making electron-poor tetrazines susceptible to attack by nucleophiles, leading to their decomposition in aqueous conditions.<sup>22</sup> Even worse, such tetrazines react readily with exposed nucleophilic centers present on biomacromolecules, leading to unselective labeling of biomacromolecules with the tetrazine reagent. Due to their high reactivity, thiol groups present in molecules containing cysteine are the biggest issue.<sup>23</sup>

Multiple strategies have been employed to increase the reactivity of dienes for bioorthogonal applications in iEDDA reactions while preserving their stability in biological systems. Due to the low stability of electron-poor tetrazines, this often necessitated shifting away from tetrazines to alternative scaffolds, such as triazines, where sluggish reactivity in iEDDA reactions becomes a problem.<sup>24,25</sup> In this case, the reactivity could be successfully increased by alkylating one of the nitrogens to generate a positively charged species – *N*1-alkyl-1,2,4-triazinium salts.<sup>26</sup> Another approach is to decrease the electron density on the triazine core by the coordination of heavy metal ions to the triazine core.<sup>27,28</sup> Yet the reactivity of triazines in iEDDA reactions with strained alkynes and *trans*-alkenes is generally significantly lower than that of similar tetrazines. Thus, the search for an optimal diene system for bioorthogonal iEDDA reactions continues (Scheme 1).

Computational chemistry has emerged as an efficient tool in this pursuit, enabling rapid screening of potential structures and providing insights into the operative reaction mechanism.<sup>29</sup> DFT computational models, extensively applied by Houk and coworkers, have identified multiple new promising reagents.<sup>30–32</sup> The simplest approach to the theoretical study of iEDDA reactions relies only on the energies of FMO (frontier molecular orbitals). Unfortunately, while computationally very

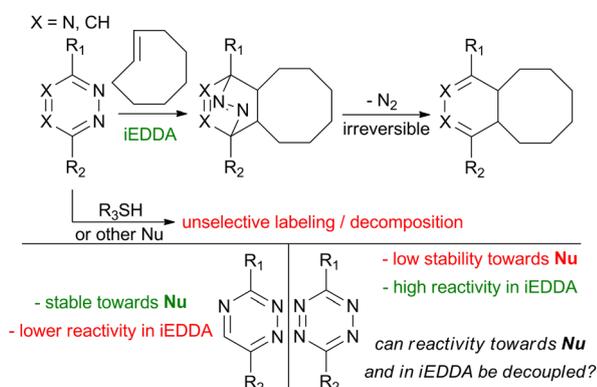
simple to achieve, this approach does not take into account all of the interactions that occur during the reaction – such as steric effects or electrostatic forces. To obtain a precise prediction of reactivity of the new iEDDA reagents, activation energy has to be calculated, which requires the localization of the transition state(s). This can be coupled with more advanced methods, such as distortion–interaction analysis, which allows us to gain insight into the steric and electronic effects during the reaction, or molecular dynamics simulations.<sup>33</sup>

The current pool of tetrazines tested in bioorthogonal iEDDA reactions included systems having various substituents on the tetrazine core, notably esters, pyridines, alkyl chains, and amines. Yet, there is still a significant number of substituents that have not been tested for their suitability in this type of application. Our aim is to identify new suitable candidates within the previously unexplored substituent space. To help with this task, we composed a set of different tetrazines to be subjected to our study. We aim to find a new system in which increased reactivity in the desired iEDDA reaction would not translate into increased susceptibility towards nucleophilic attack and thus degradation in biological environments. So far, there have been few reports of substituted tetrazines where the reactivity in iEDDA reactions has been uncoupled from the reactivity towards nucleophiles. Recently, Svatunek and Mikula have uncovered the underlying mechanism responsible for this feature using distortion–interaction analysis.<sup>34</sup> We followed a similar approach on our set of tetrazines.

## Results

Firstly, we identified which substitution patterns on tetrazine will be the target of this study. When using tetrazine derivatives for biomolecule tagging, one of the substituents on the tetrazine core has to act as a linker. These substituents allow for connection with labels or biomacromolecules. This is accomplished by carboxylic acids and hydroxyl- and amino-groups, which are then used to form an ester- or amide-bond. Importantly, such groups are usually not directly introduced onto the tetrazine core, but they are linked to it *via* a phenyl- or alkyl-linker.<sup>22</sup> In this study, we simplify these linkers as a phenyl- or methyl-substituent. The other substituent on the tetrazine can then be used to modulate its reactivity. We have selected a series of 14 different substituents to study, composed from both electron-withdrawing (EWG) as well as electron donating (EDG) groups (for quantification of the nature of the selected substituents by Hammett constants see ESI†) and atoms from different periods so that also a variance in the size of the substituents is achieved in our set (Fig. 1a).

Our strategy involves identifying potential candidates *via* the correlation of the activation barriers of iEDDA reactions with the energies of the relevant unoccupied orbitals on the tetrazines. To select the appropriate orbitals to be considered in our study, we calculated their shapes (Fig. 1b and c). Previous literature indicates that both carbon atoms and the nitrogen atoms of tetrazine are susceptible to nucleophilic attack.<sup>35,36</sup> Given these previous reports on the reactivity of the tetrazines, both LUMO and LUMO+1 orbitals are relevant for the tetrazine



**Scheme 1** Bioorthogonal iEDDA reactions using tetrazines or triazines as dienes with strained alkenes as dienophiles.



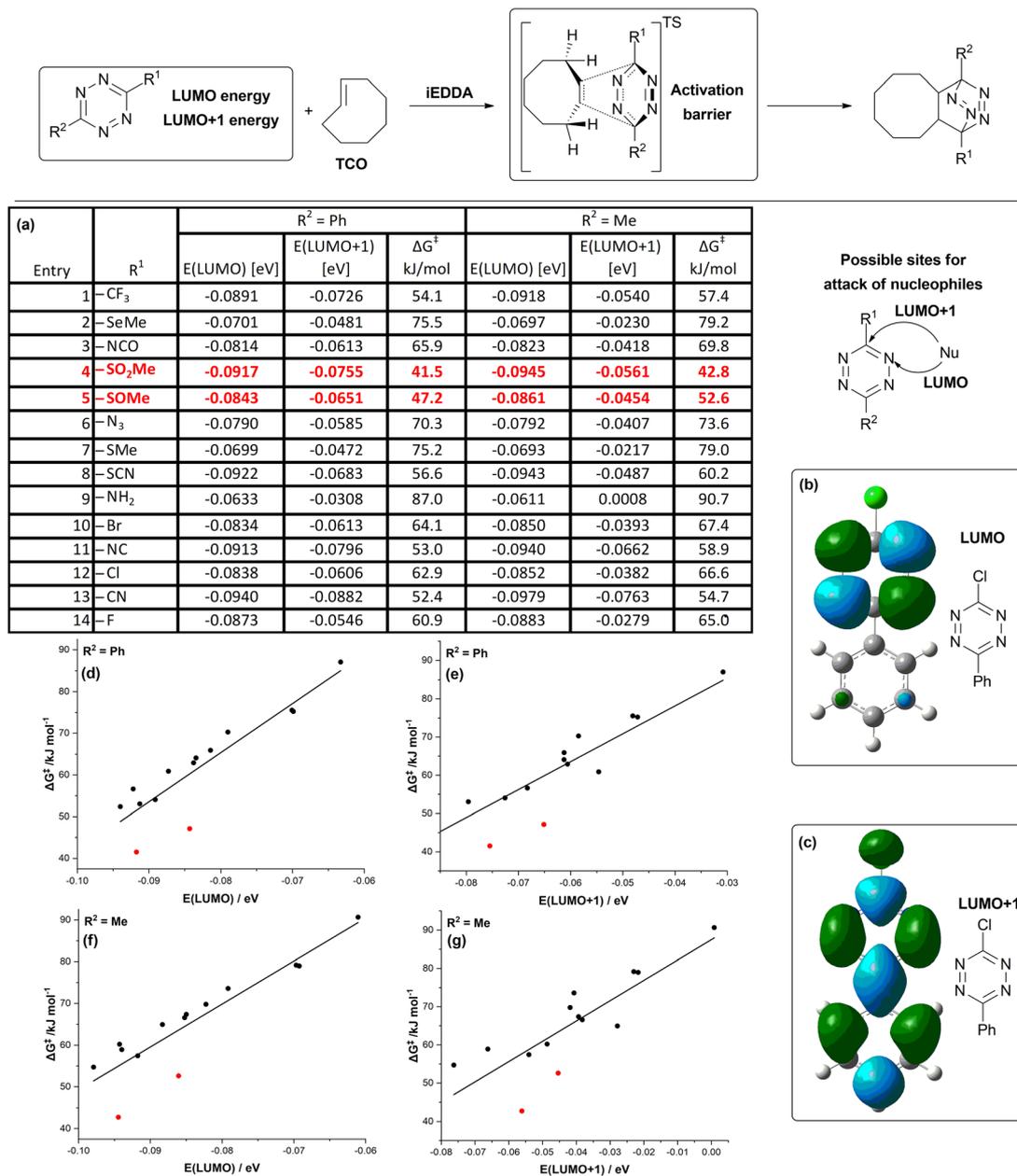


Fig. 1 (a) Results of DFT calculations for the set of tetrazine derivatives. Energies of LUMO and LUMO+1 of respective tetrazines are given in eV, free energies of activation for reaction with TCO are given in kJ mol<sup>-1</sup>. Calculations details are in the ESI;† (b) shape of LUMO orbital of tetrazine TCl; (c) shape of LUMO+1 orbital of tetrazine TCl; (d–g) plots of free energies of activation for reaction of tetrazines with TCO against LUMO and LUMO+1 energies of tetrazines. Derivatives TSO and TSO2 are marked red.

decomposition pathway – as the LUMO orbital can be implicated in the mechanism, where an attack on nitrogen occurs, whereas the LUMO+1 orbital is responsible for the reactivity of carbon atoms towards nucleophiles. As the model iEDDA reaction, we chose a reaction with *trans*-cyclooctene (TCO), a commonly used type of reagent for bioorthogonal applications.<sup>37</sup>

Across all cases studied, phenylated (R<sup>2</sup> = Ph) tetrazines exhibited a lower activation barrier with TCO compared to respective alkylated (R<sup>2</sup> = Me) tetrazines. This is in line with previously reported data, where the tetrazines bearing alkyl

linker generally reacted more sluggishly than the tetrazines with phenyl linker.<sup>22</sup> As anticipated, the introduction of electron-donating groups on tetrazines increased the activation barriers of the iEDDA reaction. This was most pronounced with amino-alkylsulfanyl- and alkylselenyl-compounds (Fig. 1a; entries 2, 7 and 9). Conversely, electron-withdrawing substituents had the opposite effect. Among these were trifluoromethyl- and nitrile-substituents (Fig. 1a; entries 1 and 13), but interestingly, also thiocyno- and isonitrile-substituents (Fig. 1a; entries 8 and 11). The most activated compounds were sulfoxides and sulfones (Fig. 1a; entries 4 and 5). With this data at



hand, we proceeded to plot the activation energy of the iEDDA reaction of tetrazines with TCO against the energies of LUMO orbitals (Fig. 1d and f) and LUMO+1 orbitals (Fig. 1e and g) of tetrazines. This was done in order to identify compounds where uncoupling of their reactivity in iEDDA reactions from their reactivity towards nucleophiles can be expected. For most of the compounds, a linear relationship between reactivity in the iEDDA reaction with TCO with the energy of the LUMO+1 orbital of tetrazines should be observed, as this is the orbital interacting with the dienophile during the course of the reaction. Indeed, this relationship holds true for most of the studied compounds. The same can be said about the energy of the LUMO orbital of tetrazines – the linear relationship between its energy and the predicted reactivity is again seen for most of the derivatives. There are two classes of compounds which do not follow this trend – sulfoxides and sulfones (Fig. 1a; entries 4 and 5). Interestingly, this behaviour of sulfones and sulfoxides is exhibited at all of the four studied relationships (Fig. 1d–g). The predicted reactivity of sulfones and sulfoxides is significantly higher than expected, based on their energies of LUMO and LUMO+1 orbitals. In order to gain further insight into this phenomenon, we subjected these derivatives to a more thorough study *via* the distortion–interaction analysis. The distortion–interaction model allows for the decomposition of the activation barrier into two components – distortion energy, which is the energy necessary for the deformation of the reactants from their equilibrium geometry into the geometry required by the transition state, and interaction energy, which has a stabilizing effect and comes from the interaction of orbitals of the two reactants in the transition state. This model has been successfully used before to provide explanation for the unexpected reactivity of several bioorthogonal reagents.<sup>13,38</sup>

For the analysis *via* the distortion–interaction model, we have chosen three different derivatives from our set. All of them have methyl as one of the substituents ( $R^2 = \text{Me}$ ). Reason behind this choice is that methyl group is likely to influence the transition states less than phenyl, which may influence the electronic distribution due to conjugation of its  $\pi$ -electrons. We chose chlorine, sulfoxide and sulfone as the other substituents (Fig. 1a, entries 4, 5 and 12). Sulfone TSO and sulfoxide TSO2 were the substituents on tetrazines, which exhibited the abnormally low activation barrier of the iEDDA reaction.

Chlorine-bearing tetrazine TCL, on the other hand, behaved in line with the trend of reactivity predicted from the energies of LUMO+1 orbitals. It is known that distortion–interaction analysis at transition state geometries can lead to skewed results when used to compare transition states with very different geometries. In our set of compounds, bond lengths in the transition state of the sulfone system (Fig. 2c) vary somewhat from the ones of chloro compound (Fig. 2a), but the difference between the geometry of the transition state of the sulfoxide-tetrazine (Fig. 2b) is similar, so the analysis of this pair (Fig. 2a *vs.* b) should give meaningful results.<sup>39</sup> The outcome of the distortion–interaction analysis shows that both sulfone TSO2 as well as sulfoxide TSO have significantly stronger interaction with the TCO than the chlorinated tetrazine TCL ( $-71.1 \text{ kJ mol}^{-1}$  and  $-70.5 \text{ kJ mol}^{-1}$  *vs.*  $-62.1 \text{ kJ mol}^{-1}$ ) in the

transition state. The distortion energy of the tetrazine moiety is also lower ( $42.2 \text{ kJ mol}^{-1}$  and  $45.2 \text{ kJ mol}^{-1}$  *vs.*  $51.9 \text{ kJ mol}^{-1}$ ). This hints that interaction other than the bond-forming process is happening during the course of the reaction of the compounds TSO2 and TSO with TCO, stabilizing the transition state. Lower distortion energy in combination with somewhat longer bonds in transition show state ( $2.40 \text{ \AA}$ ;  $2.341 \text{ \AA}$  for TSO2 and  $2.35 \text{ \AA}$ ;  $2.37 \text{ \AA}$  for TSO *vs.*  $2.32 \text{ \AA}$ ;  $2.29 \text{ \AA}$  for TCL), that the transition state occurs earlier in the reaction for sulfone and sulfoxide than for the TCL.

In order to identify the source of this effect, we have calculated the shape of the orbitals interacting together in the reaction – *i.e.* HOMO orbital of the TCO (Fig. 3a) and the LUMO+1 orbitals of the tetrazines TSO and TSO2 (Fig. 3b and c). Due to the strained nature of the double bond in TCO, its HOMO orbital is not a pure  $\pi$ -bonding orbital centered above and below the C=C double bond, but it also contains lobes on adjacent C–H bonds. This leads to a stabilizing interaction with lobes of LUMO+1 orbitals centered on the sulfone- and sulfoxide moiety of the respective tetrazine derivatives TSO2 and TSO (Fig. 3b and c), explaining the increased interaction energy of compounds TSO and TSO2 with respect to the compound TCL. Geometry of the transition state suggests that such secondary orbital interaction indeed occurs, as the C–H bond of TSO moves towards the SO moiety of TSO and TSO2 ( $2.63 \text{ \AA}$  for sulfoxide and  $2.53 \text{ \AA}$  for sulfone) during the reaction. Such additional stabilization of the transition state explains the unexpectedly low activation barrier for the derivatives TSO and TSO2 in iEDDA cycloaddition with TCO. Secondary orbital interactions have been reported as the driving force behind unexpected reactivity patterns of tetrazines before.<sup>40</sup> As this lowering of the activation barrier was not achieved by decreasing the energy of LUMO and LUMO+1 orbitals, we can predict that sulfone- and sulfoxide-substituted tetrazines will exhibit a better ratio between reactivity in iEDDA reactions with TCO to their lability towards nucleophiles, making them an interesting candidate for incorporation in bioorthogonal reagents. Importantly, from the point of future applications, the synthesis of sulfoxide/sulfone-substituted tetrazines is already known.<sup>41</sup> In addition to the found secondary orbital interaction, it is possible that the unusual reactivity of TSO and TSO2 is also affected by differences in Pauli repulsion of the reactants, as this has been proposed recently for similar derivatives where reactivity is not driven by differences in orbital energy.<sup>42</sup> Our group is currently actively working on the application of sulfoxide/sulfone-substituted tetrazines in bioorthogonal chemistry and further analysis of their reactivity by experimental as well as theoretical means.

## Computational details

Conformation analysis was performed using Spartan program package.<sup>43</sup> quantum chemical calculations were performed using Gaussian G16 RevC.01 software package.<sup>44</sup> Geometry optimization of minima involved were performed at M062X level of theory and 6-31G(d) basis set for optimisation of geometries. This combination has been successfully used



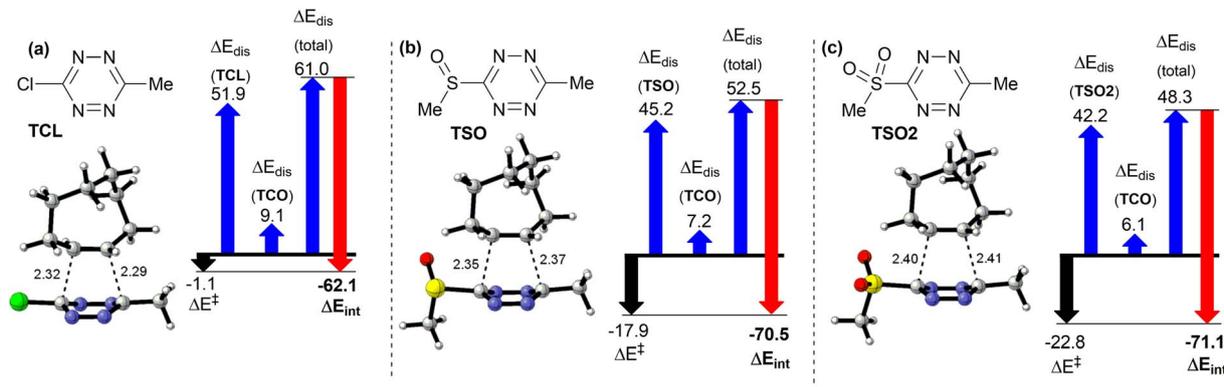


Fig. 2 Distortion–interaction analyses of reaction of TCO with (a) TCL; (b) TSO; (c) TSO2. Electronic activation energy  $\Delta E^\ddagger$  is shown with black arrows, distortion energy  $\Delta E_{\text{dis}}$  is shown with blue arrows and interaction energy  $\Delta E_{\text{int}}$  is shown with red arrows.

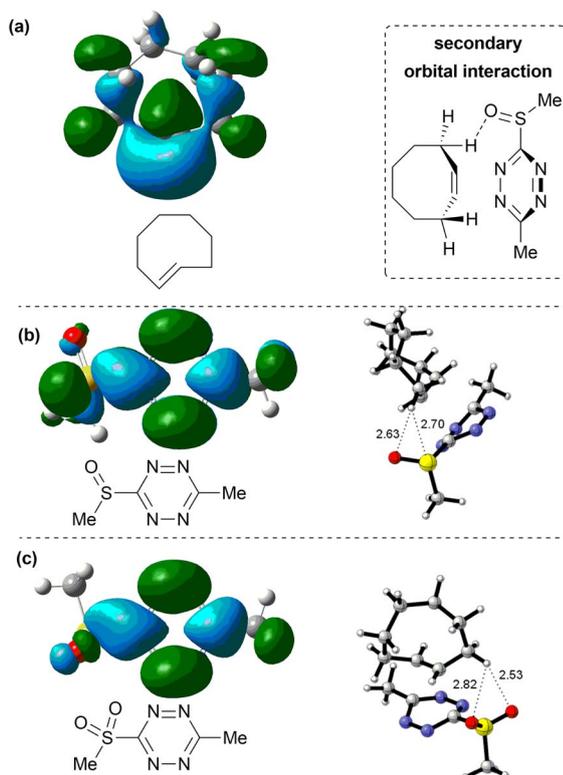


Fig. 3 (a) HOMO orbital of TCO; (b) LUMO+1 orbital of TSO; (c) LUMO+1 orbital of TSO2.

before to obtain transition state geometries of pericyclic reactions of tetrazine.<sup>15,45</sup> Single point calculations were performed using 6-311+g(2d,p) basis set. All calculations were done *in vacuo*. Frequency calculations were performed on the minima in order to ascertain the type of stationary point as well as to obtain thermochemistry. The nature of the transition states was investigated by running an IRC calculation, in order to prove that the correct transition state was found. Thermodynamic properties were calculated at 298 Kelvin. A quasiharmonic correction was applied during the entropy calculation by setting

all positive frequencies that are less than  $100 \text{ cm}^{-1}$  to  $100 \text{ cm}^{-1}$  using script GoodVibes.<sup>46</sup>

## Conclusions

We have predicted the reactivity of a series of substituted tetrazines in iEDDA reaction with *trans*-cyclooctene. For most of the tetrazines, their reactivity in such reaction was linearly correlated with the energy of their LUMO and LUMO+1. Tetrazines bearing sulfoxide and sulfone moiety were an exception to this trend, as their calculated activation barrier for the iEDDA reaction with *trans*-cyclooctene was significantly lower than what would be expected based on the energy of their LUMO and LUMO+1 orbitals. This unexpected enhancement of reactivity could be explained by distortion–interaction analysis. The underlying reason behind this phenomenon is a stabilizing secondary orbital interaction between the sulfoxide/sulfone moiety and the *trans*-cyclooctene. This makes sulfoxide/sulfone-substituted tetrazines an interesting candidate for applications in bioorthogonal chemistry, as they should exhibit increased reactivity in the iEDDA reactions with *trans*-cyclooctene without being more susceptible to decomposition by nucleophiles (*e.g.* thiols).

## Author contributions

MM designed the project, analysed the data, prepared and edited the manuscript; MT performed the DFT calculations.

## Conflicts of interest

There are no conflicts to declare.

## Acknowledgements

This work has been supported by VEGA grant no. 1/0633/21. Funded by the European Union (ERC, CAPELE, 101078608). Views and opinions expressed are however those of the author(s) only and do not necessarily reflect those of the European Union or the European Research Council. Neither the



European Union nor the granting authority can be held responsible for them.

## Notes and references

- Q. Fu, S. Shen, P. Sun, Z. Gu, Y. Bai, X. Wang and Z. Liu, *Chem. Soc. Rev.*, 2023, **52**, 7737–7772.
- R. E. Bird, S. A. Lemmel, X. Yu and Q. A. Zhou, *Bioconjugate Chem.*, 2021, **32**, 2457–2479.
- S. S. Nguyen and J. A. Prescher, *Nat. Rev. Chem.*, 2020, **4**, 476–789.
- S. L. Scinto, D. A. Bilodeau, R. Hincapie, W. Lee, S. S. Nguyen, M. Xu, C. W. am Ende, M. G. Finn, K. Lang, Q. Lin, J. P. Pezacki, J. A. Prescher, M. S. Robillard and J. M. Fox, *Nat. Rev. Methods Primers*, 2021, **1**, 30.
- N. K. Devaraj, *ACS Cent. Sci.*, 2018, **4**, 952–959.
- E. M. Sletten and C. R. Bertozzi, *Angew. Chem., Int. Ed.*, 2009, **48**, 6974–6998.
- C. Bednarek, I. Wehl, N. Jung, U. Schepers and S. Braese, *Chem. Rev.*, 2020, **120**, 4301–4354.
- E. Saxon and C. R. Bertozzi, *Science*, 2000, **287**, 2007–2010.
- L. Li and Z. Zhang, *Molecules*, 2016, **21**, 1393.
- Q. Wang, T. R. Chan, R. Hilgraf, V. V. Fokin, B. Sharpless and M. G. Finn, *J. Am. Chem. Soc.*, 2003, **125**(11), 3192–3193.
- E. G. Chupakhin and M. Y. Krasavin, *Chem. Heterocycl. Compd.*, 2018, **54**, 483–501.
- N. J. Agard, J. A. Prescher and C. R. Bertozzi, *J. Am. Chem. Soc.*, 2004, **126**, 15046–15047.
- M. Nakajima, D. A. Bilodeau and J. P. Pezacki, *RSC Adv.*, 2020, **10**, 29306–29310.
- C. S. Craig, J. Moran and J. P. Pezacki, *Chem. Commun.*, 2010, **46**, 931–933.
- H. Wu and N. K. Devaraj, *Acc. Chem. Res.*, 2018, **51**, 1249–1259.
- N. K. Devaraj, R. Weissleder and S. A. Hilderbrand, *Bioconjugate Chem.*, 2008, **19**, 2297–2299.
- M. L. Blackman, M. Royzen and J. M. Fox, *J. Am. Chem. Soc.*, 2008, **130**, 13518–13519.
- M. R. Karver, R. Weissleder and S. A. Hilderbrand, *Bioconjugate Chem.*, 2011, **22**, 2263–2270.
- G. Beliu, A. J. Kurz, A. C. Kuhleemann, L. Behringer-Pliess, M. Meub, N. Wolf, J. Seibel, Z.-D. Shi, M. Schnermann, J. B. Grimm, L. D. Lavis, S. Doose and M. Sauer, *Commun. Biol.*, 2019, **2**, 261.
- J. Sauer and D. Lang, *Angew. Chem.*, 1964, **76**, 603.
- B. L. Oliveira, Z. Guo and G. J. L. Bernardes, *Chem. Soc. Rev.*, 2017, **46**, 4895–4950.
- A. Maggi, E. Ruivo, J. Fissers, C. Vangestel, S. Chatterjee, J. Joossens, F. Sobott, S. Staelens, S. Stroobants, R. Van Der Veken, L. wyffels and K. Augustyns, *Org. Biomol. Chem.*, 2016, **14**, 7544–7551.
- H. E. Murrey, J. C. Judkins, C. W. am Ende, T. E. Ballard, Y. Fang, K. Riccardi, E. R. Guilmette, J. W. Schwartz, J. M. Fox and D. S. Johnson, *J. Am. Chem. Soc.*, 2015, **137**, 11461–11475.
- F.-G. Zhang, Z. Chen, X. Tang and J.-A. Ma, *Chem. Rev.*, 2021, **121**, 14555–14593.
- D. N. Kamber, Y. Liang, R. J. Blizzard, F. Liu, R. A. Mehl, K. N. Houk and J. A. Prescher, *J. Am. Chem. Soc.*, 2015, **137**, 8388–8391.
- Z. V. Šlachtová, S. Bellová, A. La-Venia, J. Galeta, M. Dračínský, K. Chalupský, A. Dvořáková, H. Mertlíková-Kaiserová, P. Rukovanský, R. Dzijak and M. Vrabel, *Angew. Chem., Int. Ed.*, 2023, **62**, e202306828.
- M. Sims, S. Kyriakou, A. Matthews, M. E. Deary and V. N. Kozhevnikov, *Dalton Trans.*, 2023, **52**, 10927–10932.
- V. N. Kozhevnikov, M. E. Deary, T. Mantso, M. I. Panayiotidis and M. T. Sims, *Chem. Commun.*, 2019, **55**, 14283–14286.
- T. Deb, J. Tu and R. M. Franzini, *Chem. Rev.*, 2021, **121**, 6850–6914.
- A. Sengupta, B. Li, D. Svatunek and K. N. Houk, *Acc. Chem. Res.*, 2022, **55**, 2467–2479.
- F. Liu, Y. Liang and K. N. Houk, *Acc. Chem. Res.*, 2017, **50**, 2297–2308.
- F. Liu, Y. Liang and K. N. Houk, *J. Am. Chem. Soc.*, 2014, **136**, 11483–11493.
- P. Yu, Z. Yang, Y. Liang, X. Hong, Y. Li and K. N. Houk, *J. Am. Chem. Soc.*, 2016, **138**, 8247–8252.
- D. Svatunek, M. Wilkovitsch, L. Hartmann, K. N. Houk and H. Mikula, *J. Am. Chem. Soc.*, 2022, **144**, 8171–8177.
- S. D. Schnell, M. Schilling, J. Sklyaruk, A. Linden, S. Lubert and K. Gademann, *Org. Lett.*, 2021, **23**, 2426–2430.
- Q. Zhou, P. Audebert, G. Clavier, F. Miomandre and J. Tang, *RSC Adv.*, 2014, **4**, 7193–7195.
- R. Selvaraj and J. M. Fox, *Curr. Opin. Chem. Biol.*, 2013, **17**, 753–760.
- E. M. Bickelhaupt and K. N. Houk, *Angew. Chem.*, 2017, **56**, 10070–10086.
- T. A. Hamlin, D. Svatunek, S. Yu, L. Ridder, I. Infante, L. Visscher and F. M. Bickelhaupt, *Eur. J. Org. Chem.*, 2019, **2–3**, 378–386.
- B. J. Levandowski, D. Svatunek, B. Sohr, H. Mikula and K. N. Houk, *J. Am. Chem. Soc.*, 2019, **141**, 2224–2227.
- A. Hamasaki, R. Ducray and D. L. Boger, *J. Org. Chem.*, 2006, **71**, 185–193.
- N. Houszka, H. Mikula and D. Svatunek, *Chem. Eur. J.*, 2023, **29**, e202300345.
- Spartan 14v114*, Wavefunction, Inc., Irvine, 2014.
- M. J. Frisch, G. W. Trucks, H. B. Schlegel, G. E. Scuseria, M. A. Robb, J. R. Cheeseman, G. Scalmani, V. Barone, G. A. Petersson, H. Nakatsuji, X. Li, M. Caricato, A. V. Marenich, J. Bloino, B. G. Janesko, R. Gomperts, B. Mennucci, H. P. Hratchian, J. V. Ortiz, A. F. Izmaylov, J. L. Sonnenberg, D. Williams-Young, F. Ding, F. Lipparini, F. Egidi, J. Goings, B. Peng, A. Petrone, T. Henderson, D. Ranasinghe, V. G. Zakrzewski, J. Gao, N. Rega, G. Zheng, W. Liang, M. Hada, M. Ehara, K. Toyota, R. Fukuda, J. Hasegawa, M. Ishida, T. Nakajima, Y. Honda, O. Kitao, H. Nakai, T. Vreven, K. Throssell, J. A. Montgomery Jr, J. E. Peralta, F. Ogliaro, M. J. Bearpark, J. J. Heyd, E. N. Brothers, K. N. Kudin, V. N. Staroverov, T. A. Keith, R. Kobayashi, J. Normand, K. Raghavachari, A. P. Rendell, J. C. Burant, S. S. Iyengar, J. Tomasi, M. Cossi, J. M. Millam, M. Klene, C. Adamo,



- R. Cammi, J. W. Ochterski, R. L. Martin, K. Morokuma, O. Farkas, J. B. Foresman, and D. J. Fox, *Gaussian 16*, Revision C.01, Gaussian, Inc., Wallingford CT, 2016.
- 45 S.-E. Suh, S. Chen, K. N. Houk and D. M. Chenoweth, *Chem. Sci.*, 2018, **9**, 7688–7693.
- 46 *GoodVibes Version 3.2*, Zenodo, DOI: [10.5281/zenodo.595246](https://doi.org/10.5281/zenodo.595246).

