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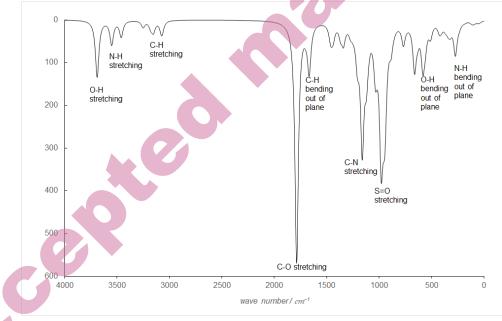
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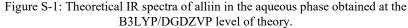
## SUPPLEMENTARY MATERIAL TO

## Chemical reactivity of alliin and its molecular interactions with the M protease<sup>pro</sup> of SARS-CoV-2

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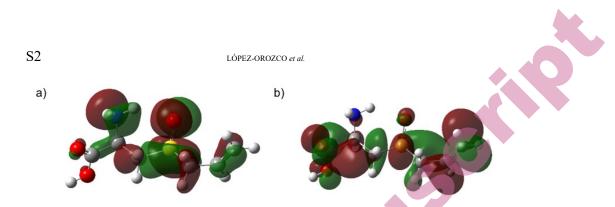
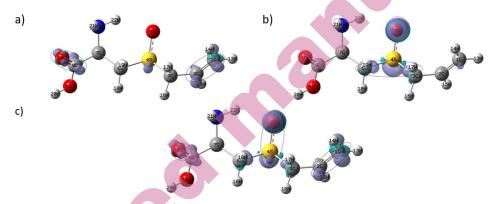
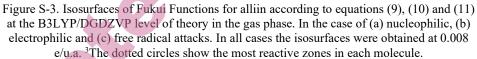


Figure S-2. HOMO and LUMO distributions on alliin obtained at the B3LYP/DGDZVP level of theory in the gas phase. In all cases the isosurfaces were obtained at 0.08 e/u.a.<sup>3</sup>.





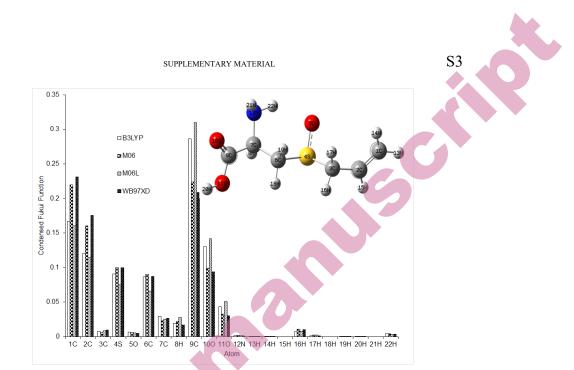


Figure S-4. Condensed Fukui function values for electrophilic attacks on alliin at the X/ DGDZVP level of theory (where X=B3LYP, M06, M06L and ωB97XD), in the aqueous phase employing the Hirshfeld population and equations (12)-(14), the dashed circles show the most reactive zones in each molecule.

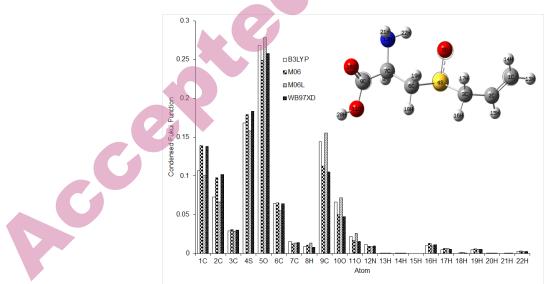


Figure S-5. Condensed Fukui function values for free radical attacks on alliin at the X/ DGDZVP level of theory (where X=B3LYP, M06, M06L and ωB97XD), in the aqueous phase employing the Hirshfeld population and equations (12)-(14), the dashed circles show the most reactive zones in each molecule.

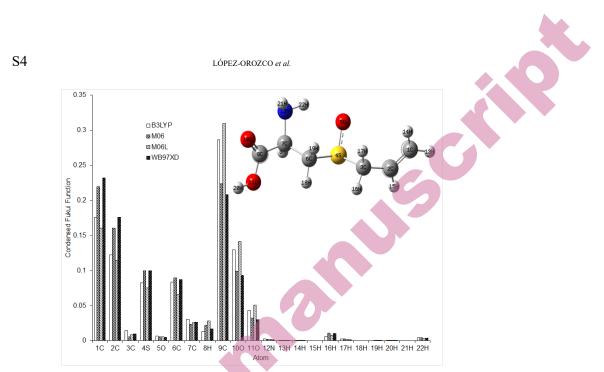


Figure S-6. Condensed Fukui function values for nucleophilic attacks on alliin at the X/ DGDZVP level of theory (where X=B3LYP, M06, M06L and  $\omega$ B97XD), in the gas phase employing the Hirshfeld population and equations (12)-(14), the dashed circles show the most reactive zones in each molecule.

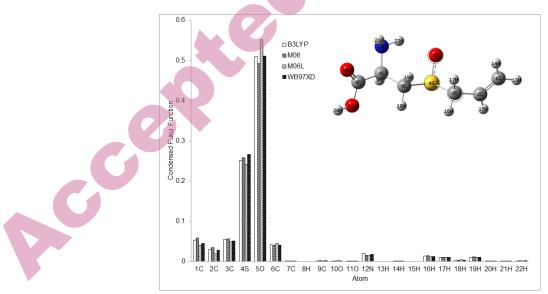


Figure S-7. Condensed Fukui function values for electrophilic attacks on alliin at the X/ DGDZVP level of theory (where X=B3LYP, M06, M06L and ωB97XD), in the gas phase employing the Hirshfeld population and equations (12)-(14), the dashed circles show the most reactive zones in each molecule.

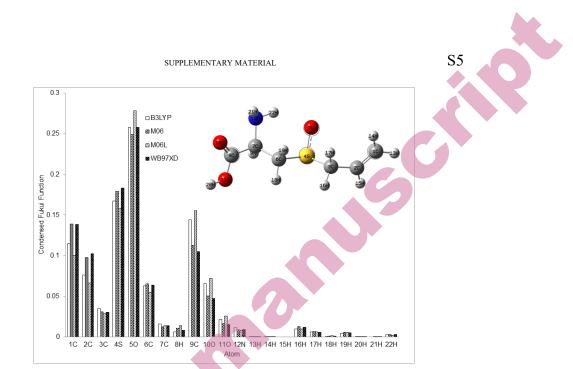


Figure S-8. Condensed Fukui function values for free radical attacks on alliin at the X/ DGDZVP level of theory (where X=B3LYP, M06, M06L and  $\omega$ B97XD), in the gas phase employing the Hirshfeld population and equations (12)-(14), the dashed circles show the most reactive zones in each molecule.

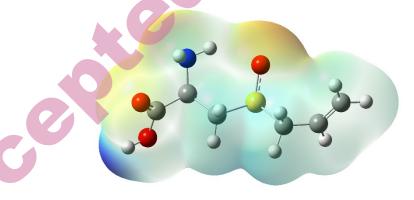


Figure S-9. Mapping of electrostatic potentials evaluated at the B3LYP/DGDZVP level of theory in the gas phase, over a density isosurface (value =0.002 e/a.u.<sup>3</sup>) for alliin.

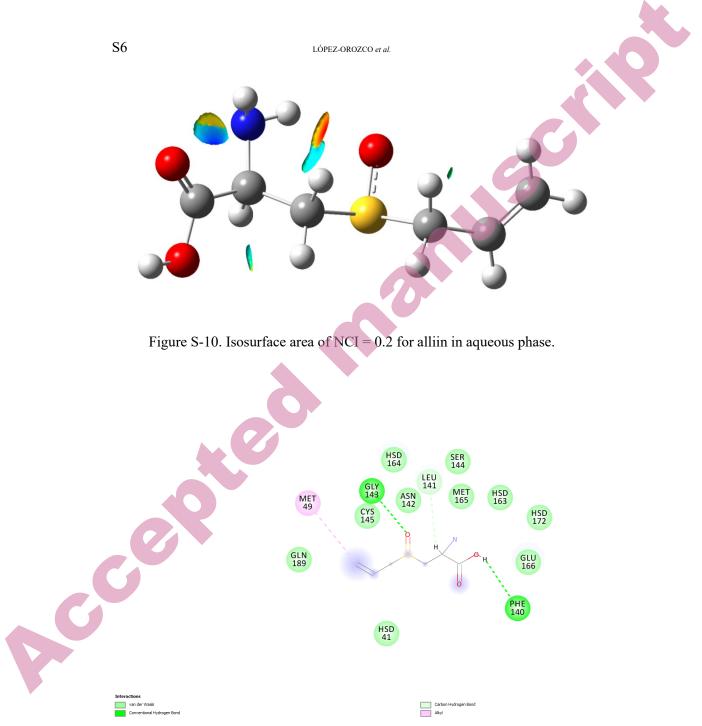


Figure S-11. 2D mapping of ligand/protein interactions for alliin.