



*Citation for published version:*

Fienberg, S, Cozier, GE, Acharya, KR, Chibale, K & Sturrock, ED 2018, 'The Design and Development of a Potent and Selective Novel Diprolyl Derivative That Binds to the N-Domain of Angiotensin-I Converting Enzyme', *Journal of Medicinal Chemistry*, vol. 61, no. 1, pp. 344-359. <https://doi.org/10.1021/acs.jmedchem.7b01478>

*DOI:*

[10.1021/acs.jmedchem.7b01478](https://doi.org/10.1021/acs.jmedchem.7b01478)

*Publication date:*

2018

*Document Version*

Peer reviewed version

[Link to publication](https://doi.org/10.1021/acs.jmedchem.7b01478)

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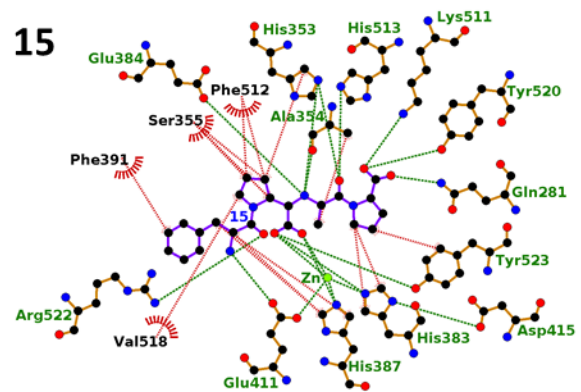
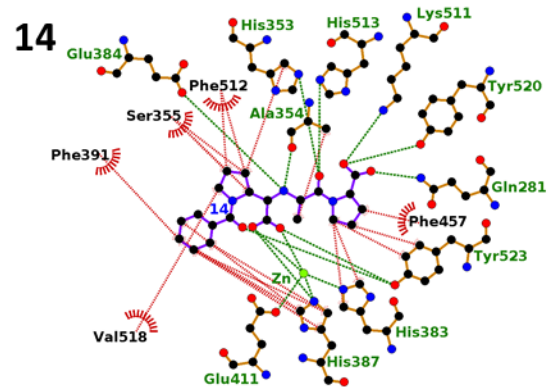
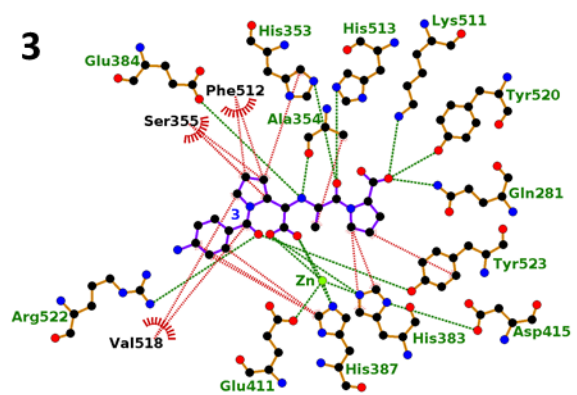
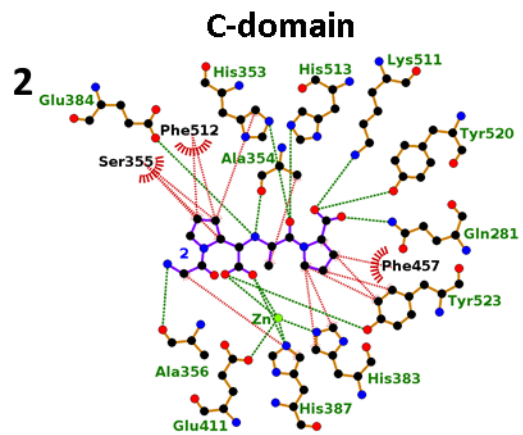
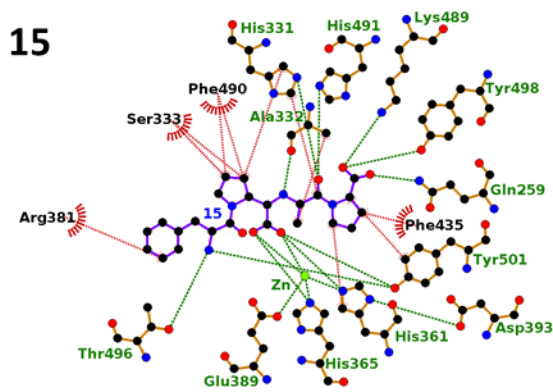
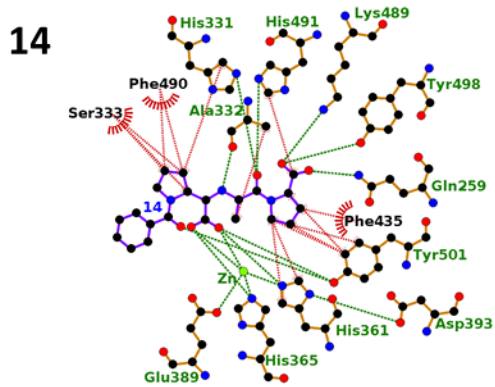
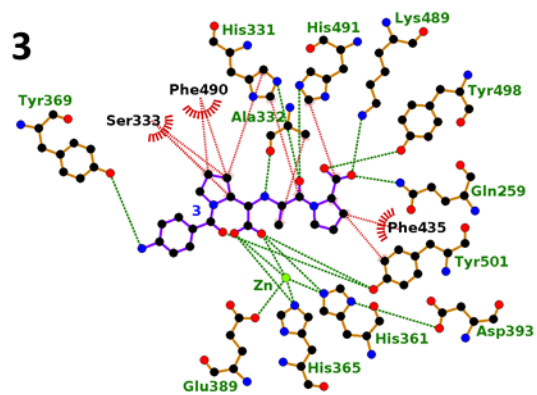
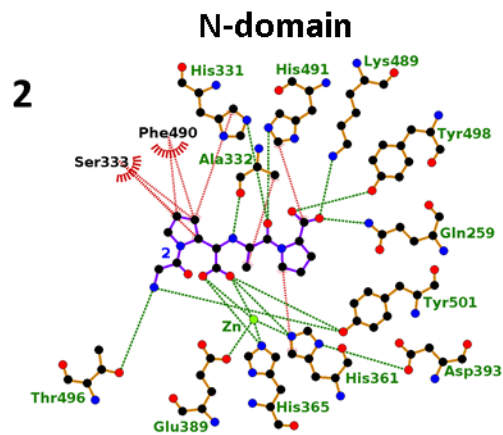
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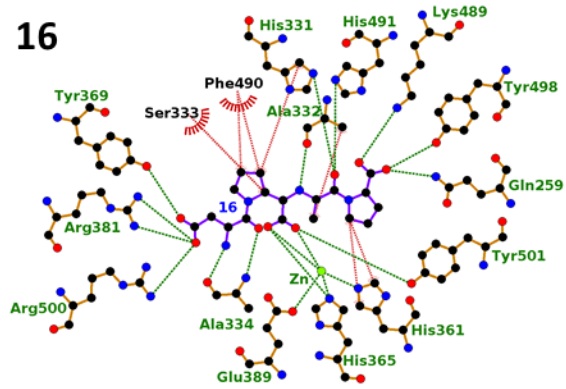
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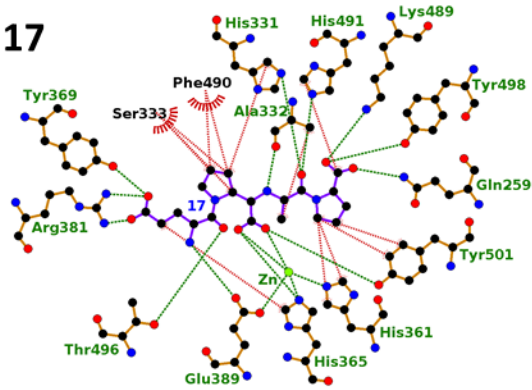


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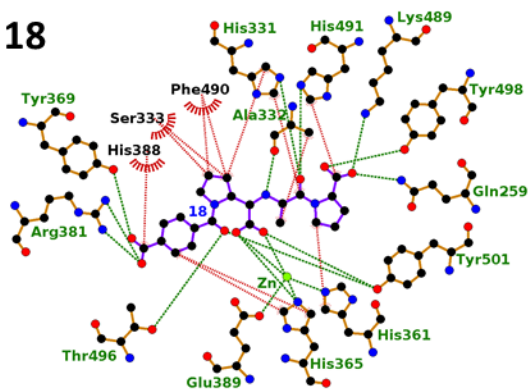
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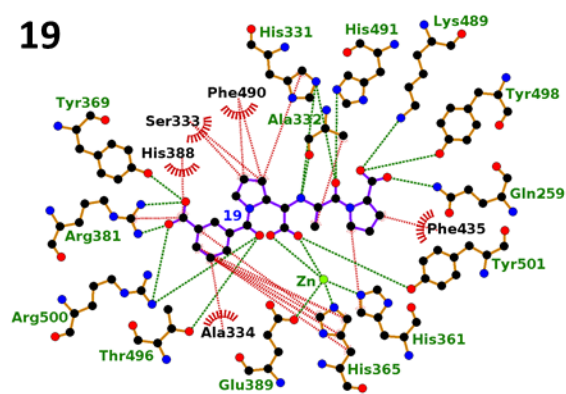
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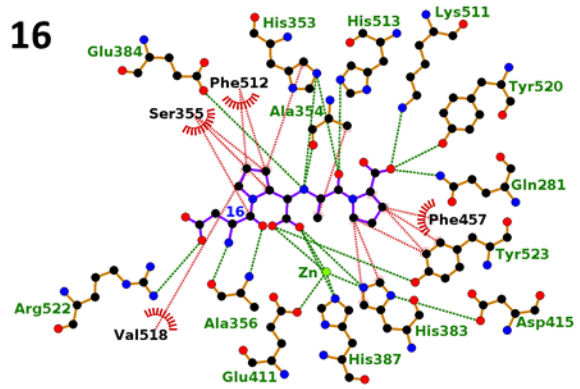


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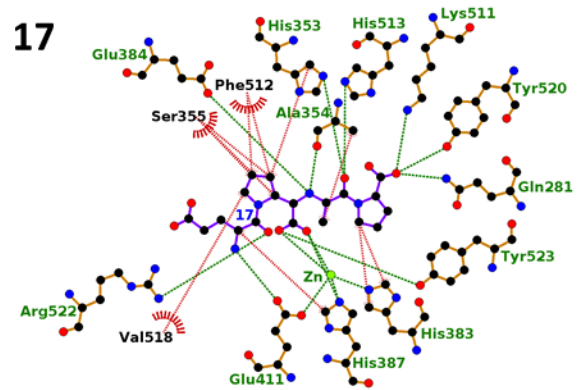


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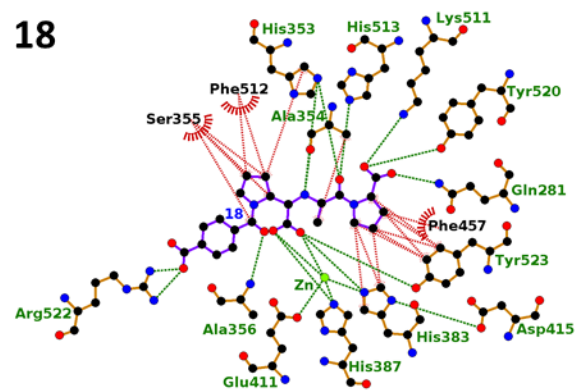
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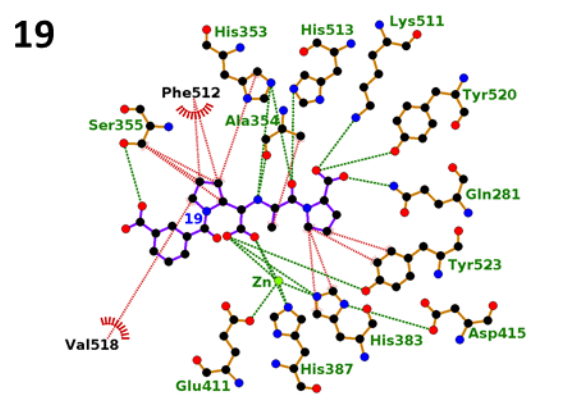
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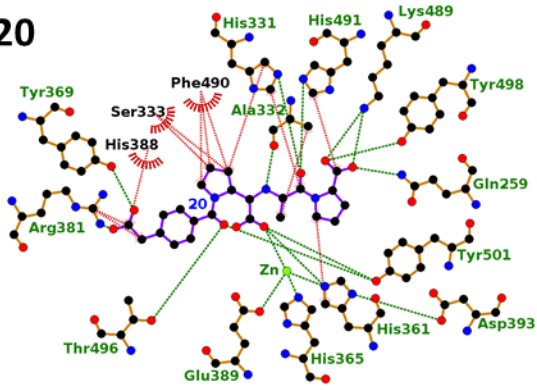


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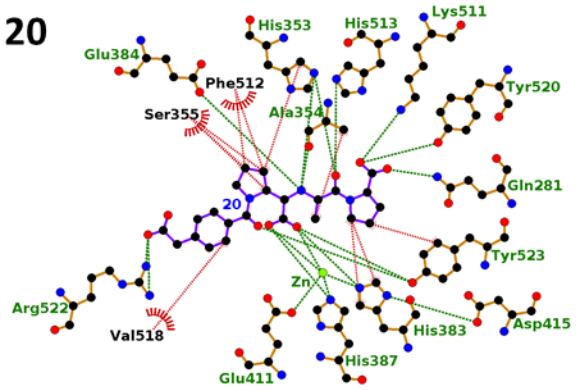
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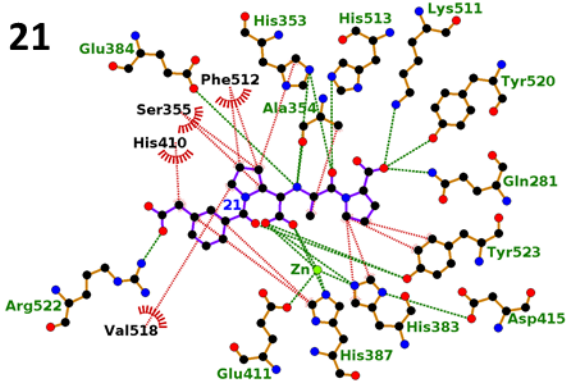


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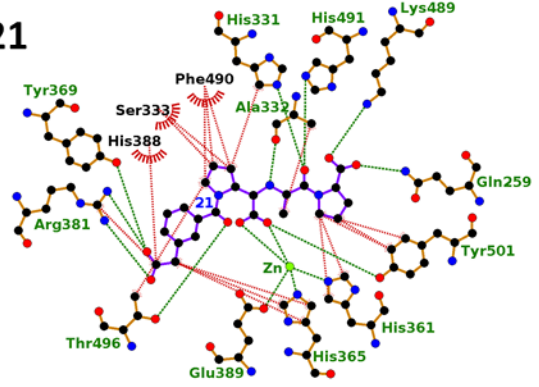
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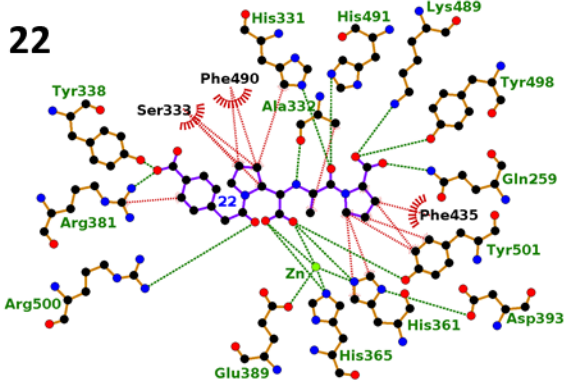
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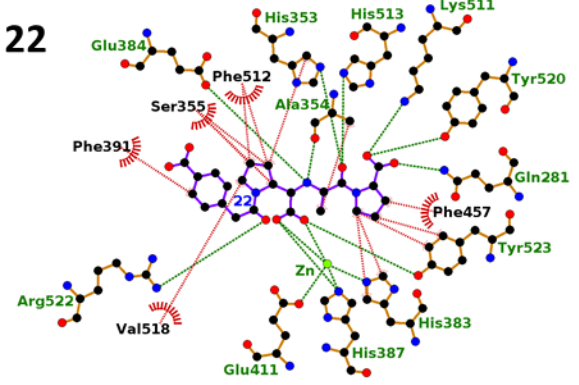
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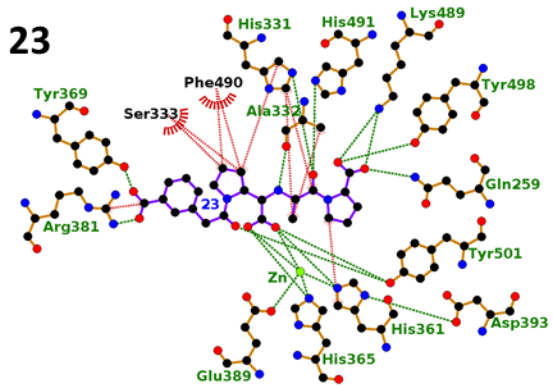
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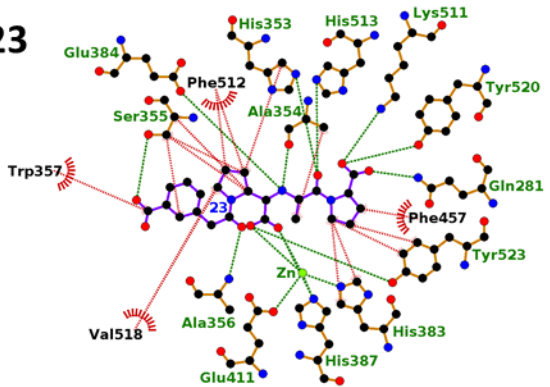
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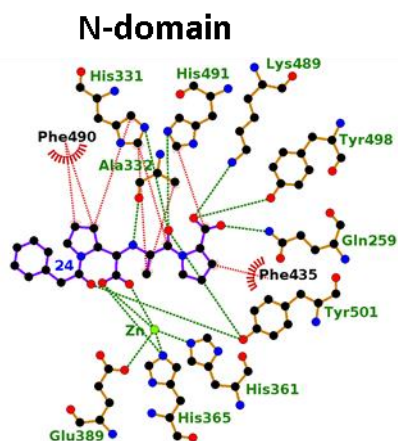
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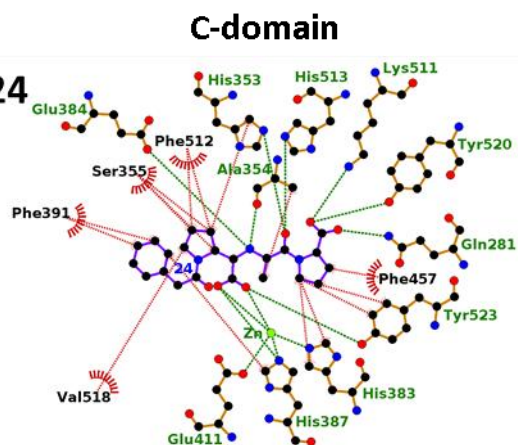
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**Figure SI1:** Ligplot representations of the binding site interactions resulting from the docking of the diprolyl series compounds in the ACE N- and C-domains. Hydrogen bonds and hydrophobic interactions are shown with green and red dotted lines respectively.