Compounds interfering with Embryonic Lethal Abnormal Vision (ELAV) protein–RNA complexes: an avenue for discovering new drugs

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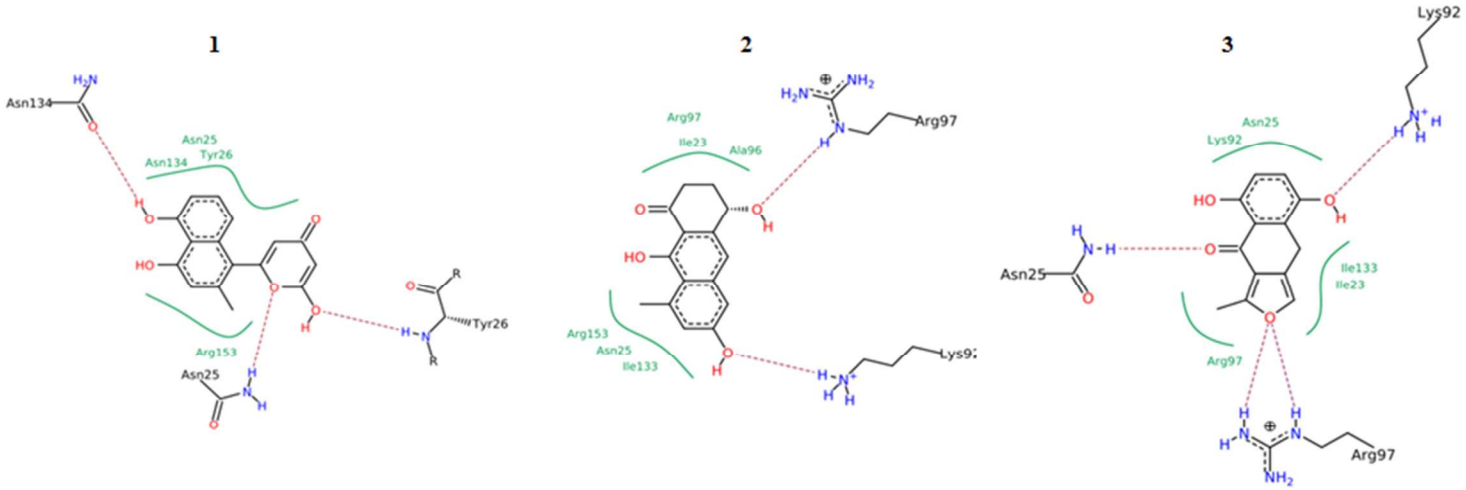
Supporting information

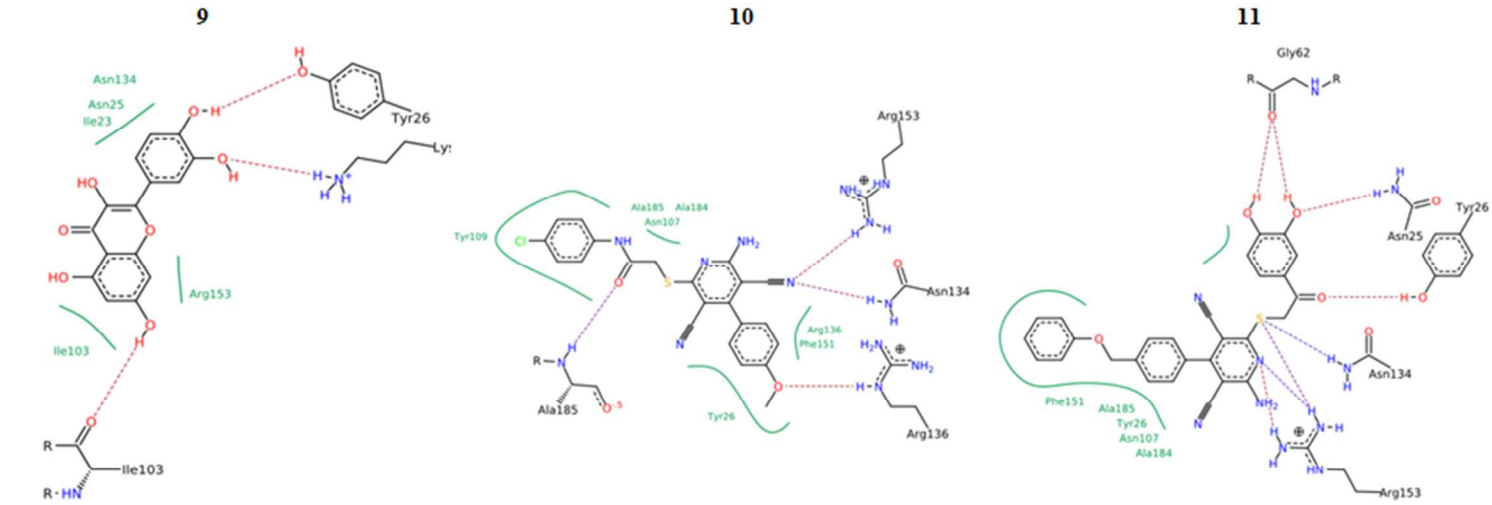
Predicted binding mode of compounds.

These binding modes are the result of a docking run using the FlexX docking module with 30 poses and represent the top-scoring pose after HYDE scoring with SEESAR and careful visual inspection to exclude poses with significant inter- or intra-molecular clash terms or unfavorable conformations. The figures were generated with PoseView as implemented in the LeadIT suite.

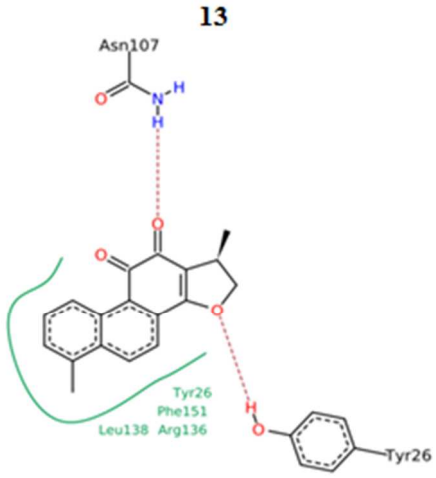
Compounds 1–3



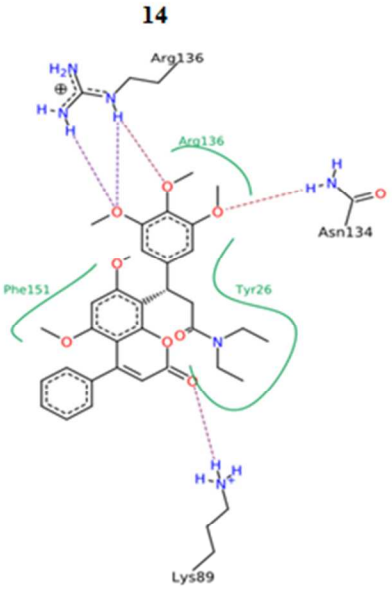
Compounds 9–11

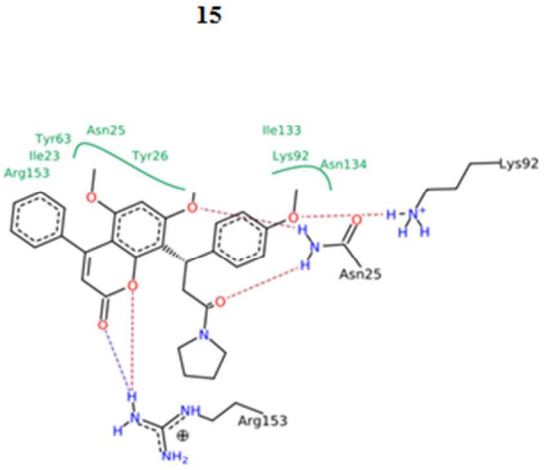


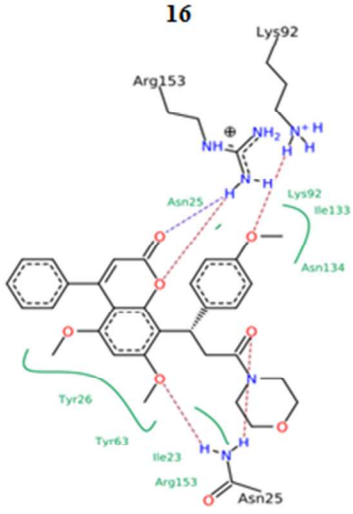
Compound 13



Compounds 14–19







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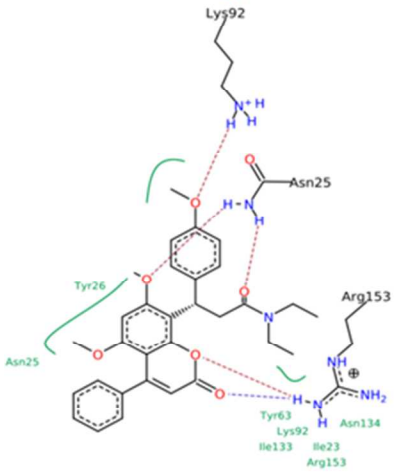

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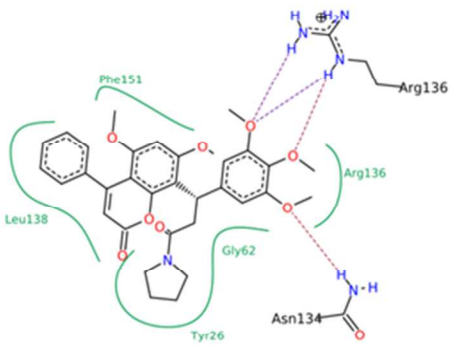

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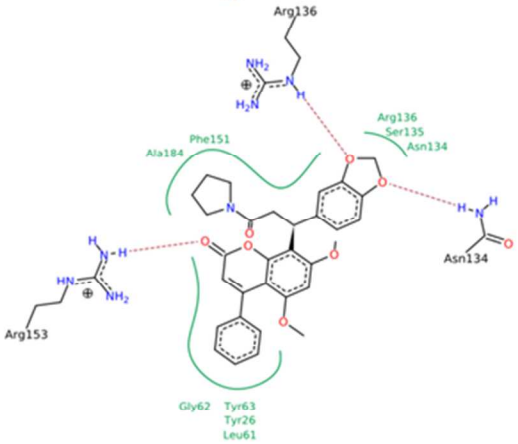

Compounds 20–22

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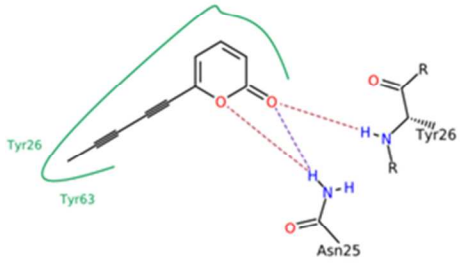

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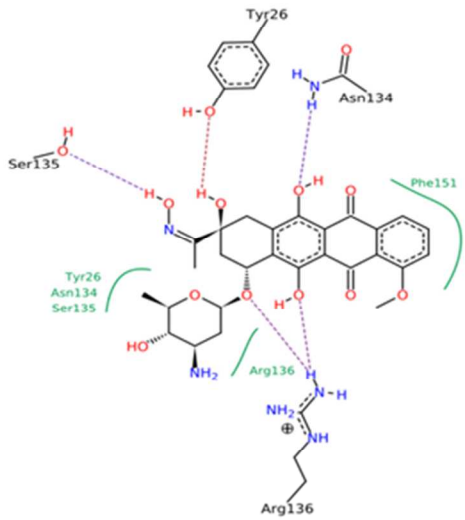

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