

1. Keep up-to-date with relevant literature on this topic.
2. Compilation of a dataset of RNA selective binders and decoys.

3. Visualization and study of the RNA binders chemical subspace (PCA, tSNE ...).
4. Building of high-quality QSAR models to distinguish RNA selective binders.
5. Application of the QSAR models to automatically filter internal/commercial databases of compounds.
6. Present work results at group and department meetings.

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