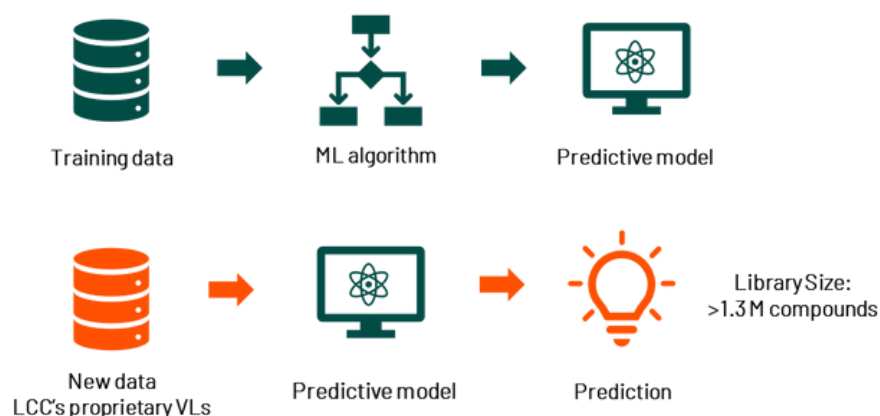




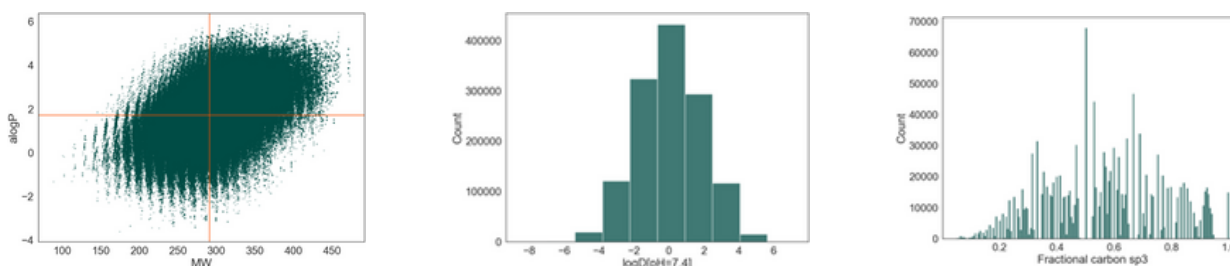
LCC'S MOLECULES OF THE MONTH

RNA Virtual Library

While classical drug discovery efforts have focused on targeting the proteome, it is estimated that only ~15% is druggable. Whereas, transcription of the human genome into RNA stands at around 75%, with only ~1% encoding for proteins.¹ This paves the way for the use of **RNA-targeting molecules** to target the encoded mRNA of 'undruggable' proteins, or non-coding RNA in relevant disease-regulatory mechanisms.¹ While other RNA-targeting modalities, such as antisense oligonucleotides, have been used extensively to target RNA, these can often present challenges when it comes to cell permeability and distribution.² Comparatively, **small molecules** can be advantageous due to their well-established synthetic methodologies and their tunability, enabling easier modification to improve delivery or uptake.³



LCC has designed and enumerated an **RNA-focused virtual library**. The library is based on LCC's novel, chirally pure, multi-functional scaffolds and was created by using a machine learning algorithm, trained with experimentally derived sets of RNA binders (ROBIN).² By design, the library is diverse and optimised for Hit-ID, while near-neighbour analogues can be found in the parent virtual space and rapidly synthesised in LCC's **parallel synthesis laboratory**.



If you are interested in accessing LCC's RNA Virtual Library, **please get in touch!**



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¹ Physicochemical principles driving small molecule binding to RNA, 2024.

² Machine learning informs RNA-binding chemical space, 2022.

³ Chemical Communications, 2020, 56, 14744–14756.

