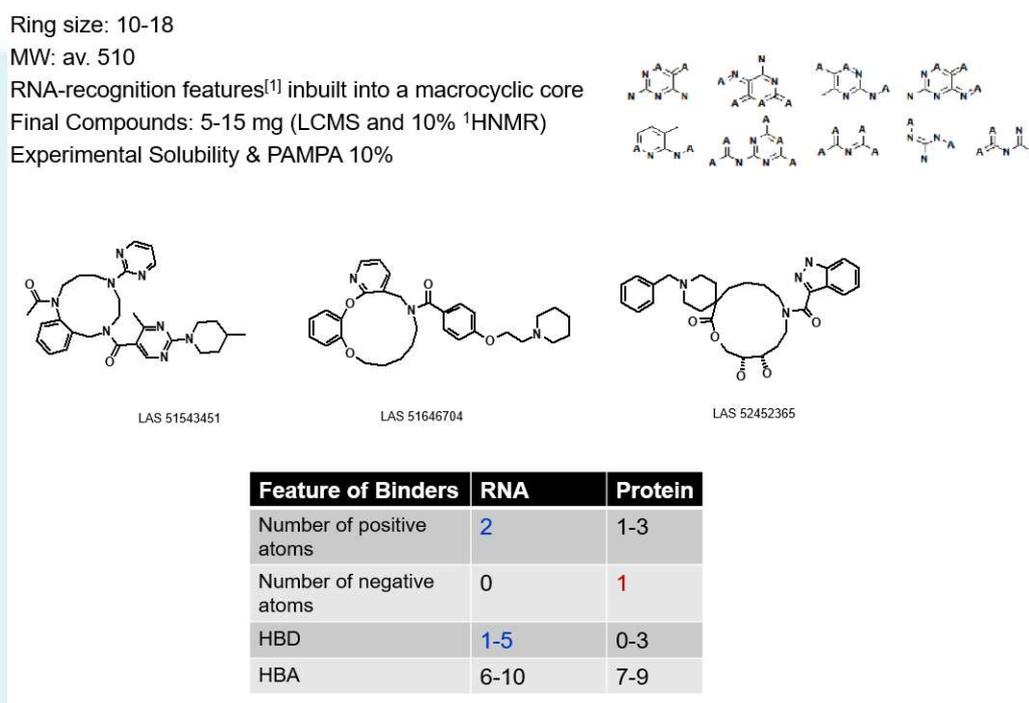


Macrocycles for RNA Drug Discovery

RNA is an emerging class of therapeutic targets that require identification and development of potent small molecule drugs. RNA-relevant chemical matter is deemed to be somewhat different from existing protein-targeting modalities and this area of research requires the integration of new ideas, design concepts, and efficiency in the production of new molecules. Comparative analyses of binding sites and ligands of RNAs and proteins have revealed that RNA pockets are less buried and more polar, and often adopt ligands that have elongated “rod”-like molecular shapes [1]. The incorporation of known RNA recognition pharmacophores [2-4] into the design of novel RNA-binding molecules is beneficial as this increases the corresponding success rate in finding new RNA-targeted hits. In addition to promising target recognition features, an RNA-binding molecule should also demonstrate reasonable intracellular bioavailability. There is a growing scientific record that properly functionalized macrocyclic scaffolds could generate very ligand-efficient molecules with advantageous cell permeability properties. These and other factors make macrocycles promising scaffolds for the design of novel RNA-targeting molecules.

Our collection of macrocycles for RNA consists of very diverse, drug-like molecules which incorporate certain known RNA-recognition elements (e.g. nucleobase ring systems and analogs) distributed within macrocyclic rings or peripheral fragments. As macrocyclic molecules tend to be larger than traditional screening molecules, it is vital to carefully assess and control their physicochemical properties. All our macrocycles have been tested for **aqueous and DMSO solubility** with cut-offs applied at 10 mM in DMSO and 50 μ M in PBS (pH 7.4); **PAMPA** permeability has also been tested for representative set of macrocycles.

Figure 1



For any requests please contact: lsadovenko@asinex.com

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