How microfluidics ultrahigh-throughput screening enables the discovery of aptamers enzyme inhibitors.

Based on the work of Michaël Ryckelynck "Digital Biology of RNA" Research team from IBMC.

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Microfluidics consists in manipulating of microliters to picoliters volumes of fluid in lab-onchip devices, allowing to miniaturize and automate biological assays¹. Aptamers are singlestranded nucleic acids adopting a 3D structure conferring them the capacity to specifically interact with a target like a protein ^{2,3}. Aptamers have similar uses to antibodies, but antibodies development takes a long time and requires in vivo screening, whereas, aptamers can be developed in vitro, using ultrahigh-throughput methods. The "Digital Biology of RNA" research team of IBMC used SELEX and μIVC -seq methods together with bioinformatics to find out an aptamer that specifically inhibits the SPM-1 enzyme. This enzyme degrades several antibiotics and provides drug-resistance to strain expressing it. A specific inhibitory aptamer of SPM-1 could resensitize SPM-1 expressing strains to the antibiotic. The SELEX method isolates SPM-1 ligand aptamers sequences from 10^{15} sequences randomly generated. Then, the *µIVC-seq* method is used to select SPM-1 inhibitor aptamers from SPM-1 ligand aptamers. The specific SPM-1 inhibitor aptamers are then sequenced, to identify them and model their 3D structure. In addition, the aptamers were tested for RNase resistance and to be functional in a medium reproducing physiological ionic concentration. Microfluidic screening enables ultra-specific molecules to be selected rapidly and economically, thanks to the miniaturization of samples and automation of processes³.

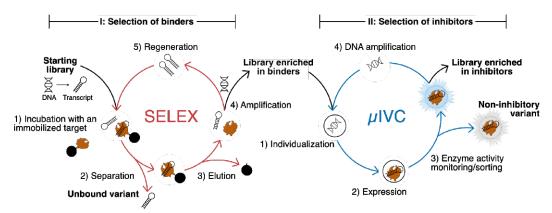


Figure: Ultrahigh-throughput microfluidic screening using SELEX and μIVC -seq methods³.

References

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