

Screening Multiple PPI Targets in Parallel: Accelerating Portfolio-level Decisions

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Overview: HTS projects for multiple targets were developed in collaboration with our client and performed in parallel in a cross-site collaboration. Up to 370,000 compounds from a variety of chemical libraries were screened in a 1536-well plate format using TR-FRET technology. During screening plate effects were observed and investigated, the cause was attributed to shipment of assay ready plates on dry-ice.

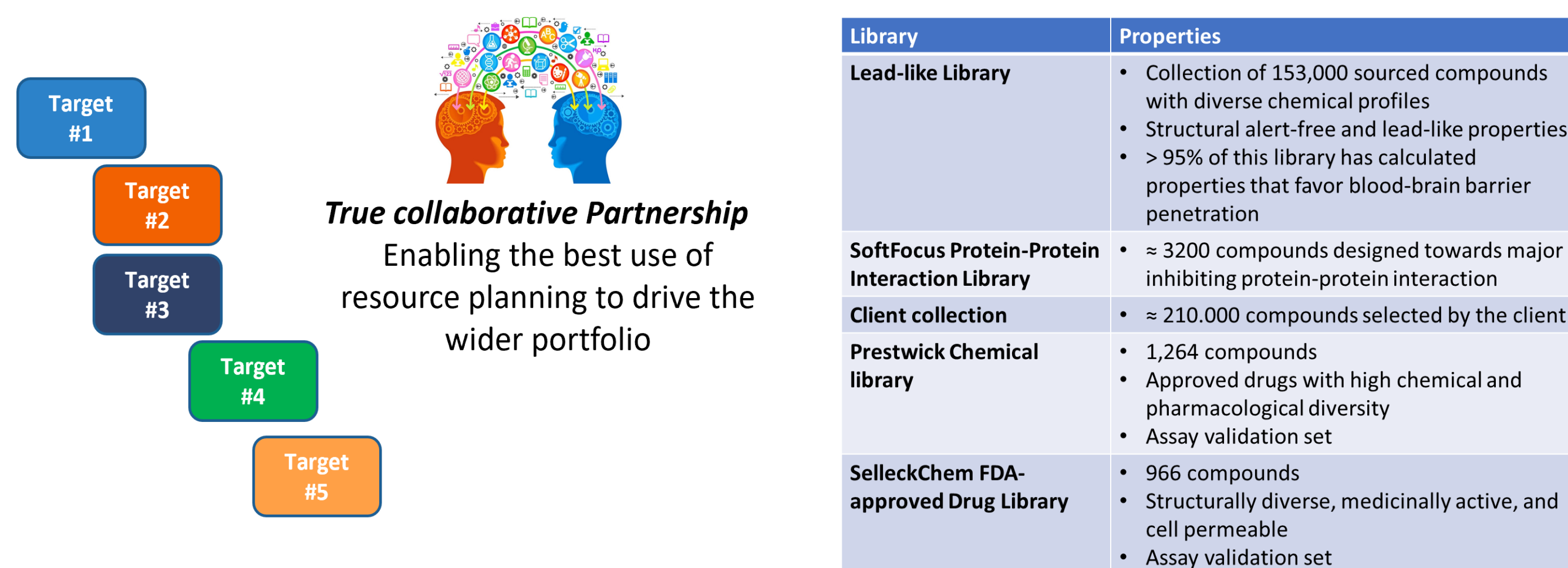


1 INTRODUCTION

In a fast-paced evolving world, high quality data and trustworthy partnerships are the key components to success. At Charles River, we quickly developed multiple high throughput screening projects in collaboration with our client to detect small molecule inhibitors of protein-protein interaction (PPI) targets. The assays were performed in parallel in a unique cross-site collaboration between HTS groups in the UK and the Netherlands.

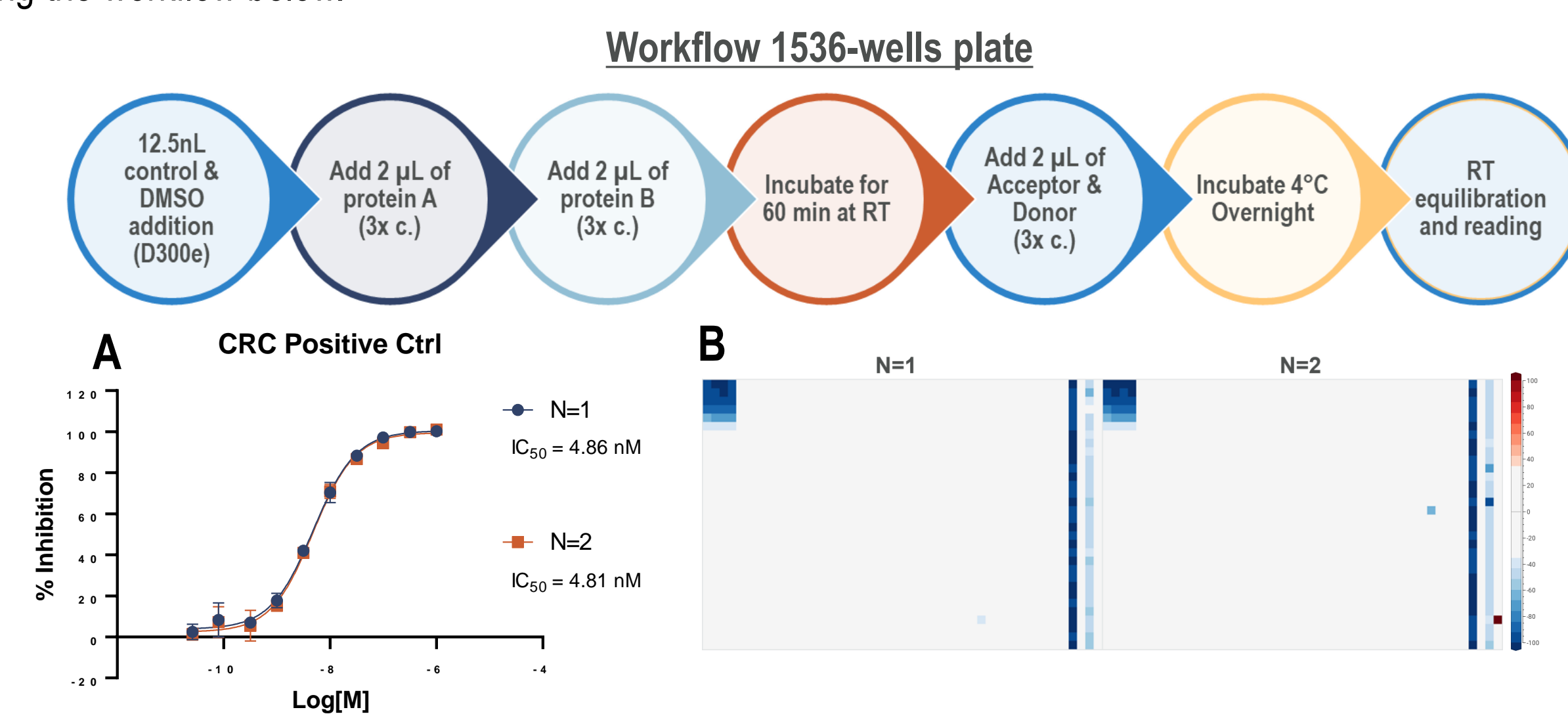
Assays were developed successively in 1536-well format using TR-FRET technology. With each successive target timelines were reduced by around 50% from assay-development to the final data packages, this was achieved by refining the assay development process and taking learnings from developed methods.

Compounds from both Charles River and Client libraries were screened simultaneously in a collaborative partnership. HTS scientists and medicinal chemists from both parties worked together to analyse the data and generate hit lists for progression through the HTS workflow and clients screening cascade.



2 METHODS

Assays for all targets were developed using the TR-FRET technology and the processes aligned for maximum efficiency ensuring robust assays were developed for HTS purposes. Optimisation experiments were performed for all detection reagents including testing a variety of detection pairings as well as K_d determinations for binding of the protein partners. Assays were validated using tool compounds when available and the tolerance to EDTA and DMSO were assessed. To assess the stability of each assay, plate uniformity studies were conducted followed by a pilot screen of 10,000 compounds tested in duplicate in the TR-FRET format before proceeding to the HTS. All assays produced a robust assay, acceptable plate statistics with Z' >0.7 and consistent signal to background following the workflow below.



3 RESULTS

Once final conditions were identified and following successful pilot screens for each target, the TR-FRET assays were used to screen approximately 370,000 compounds comprising of the entire Discovery UK Lead-Like Library of approximately 155,000 compounds and a Client library screened using assay-ready plates containing 12.5nL of compounds and final assay compound concentration of 20 μM.

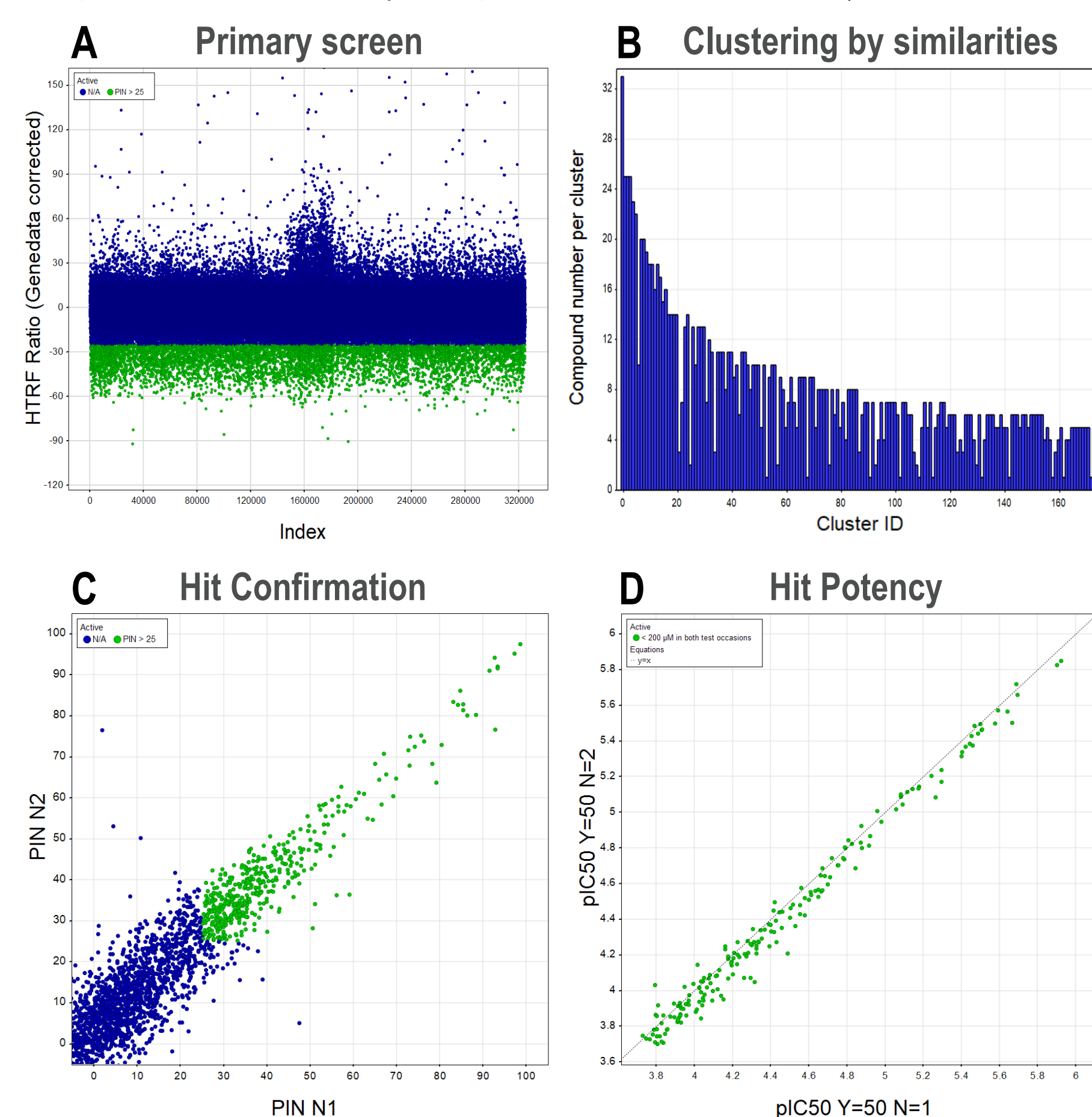
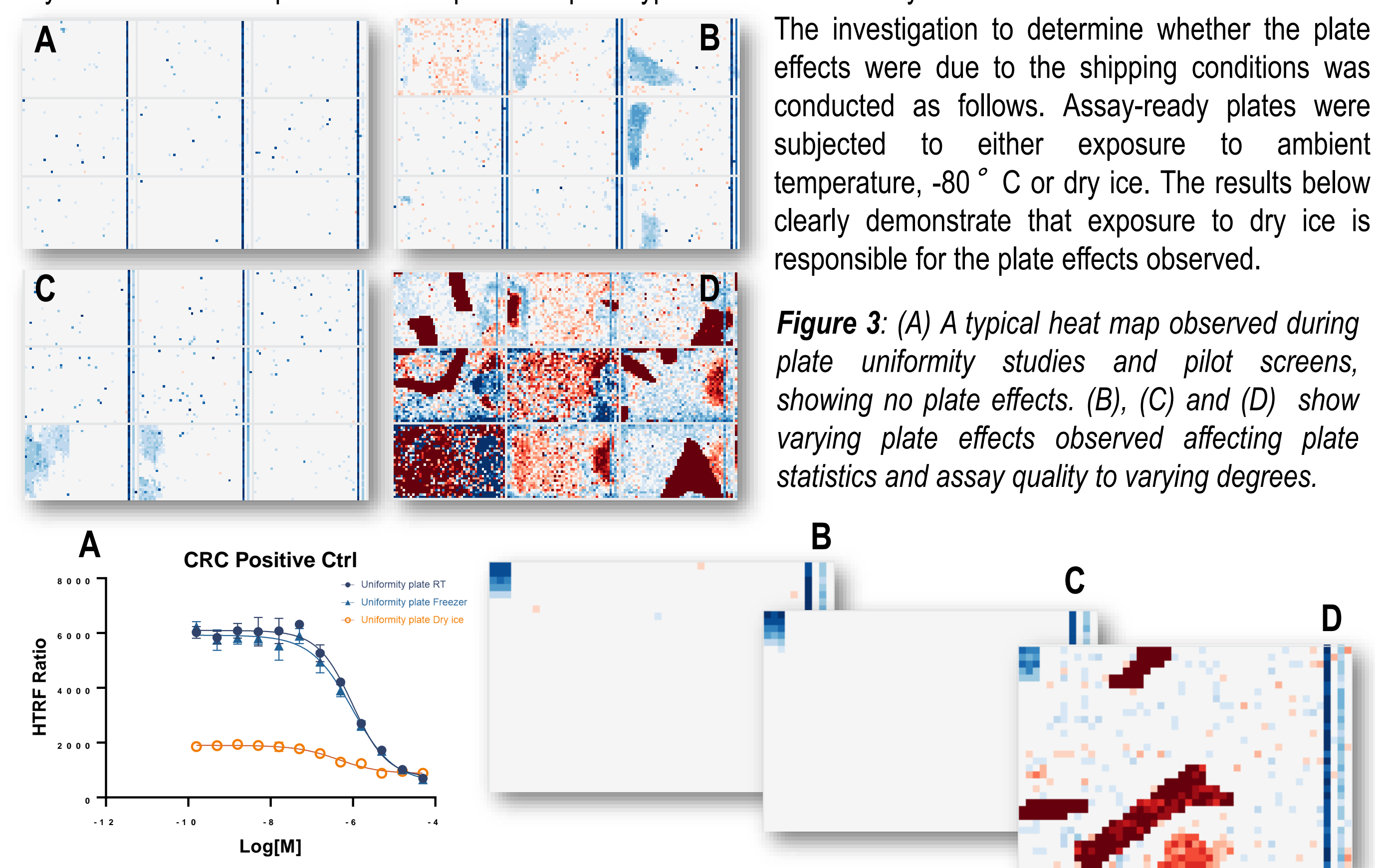


Figure 2: (A) 370,000 compounds were tested at single concentration (20 μM) for each target. (B) After filtering out compounds interfering with the read-out, frequent hitters or high Lilly demerit score* a cluster analysis was performed showing a Percentage inhibition of ≥ 25 using Tanimoto similarity of 0,6. (C) compounds selected for hit confirmation were subjected to re-test in duplicate showing a good correlation. (D) pIC50 correlation of potency determination of confirmed hits.

During the collaboration and screening activities an unusual plate effect which affected assay quality of all the targets was observed on assay-ready plates. An investigation to the cause of this was subsequently attributed to the dry ice used in the shipment with the particular plate type utilised in the assays.



4 CONCLUSION

Through the collaboration with our client, we have successfully developed high-throughput TR-FRET assays for a range of targets. The example given here highlights the HTS capabilities at Charles River across multiple sites not only utilising Charles River's Discovery UK compound libraries but also our clients or 3rd party compound collections to aid our partners hit finding campaigns and increasing chances of success. Our highly experienced HTS scientists and medicinal chemists work in close collaboration with our client's counterparts to deliver high quality assays and hits to progress the client's drug discovery process. We also highlight the importance of the shipping conditions used when transferring assay ready plates between facilities, we demonstrated that dry ice shipments can have a detrimental effect on the assay quality. Following the hit identification by the medicinal chemists, the client is now progressing compounds through the screening cascade and is currently confirming activity and mechanism of action through SPR studies at Charles River and in-house. Collaborating with a knowledgeable partner like Charles River can help our partners develop a good hit-finding approach, avoid pitfalls and ensure that the hits they identify are of high quality. Charles River's drug discovery experts support the hit identification process and the subsequent steps in the drug discovery workflow.