Identification of G9a inhibitors by AlphaLisaTM technology and hit confirmation using MT-GloTM

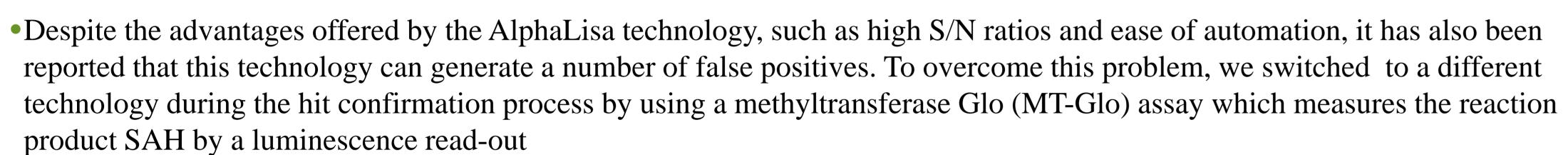
Abstract # 99

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BACKGROUND

- •G9a (EHMT2) is a HKMT that contains a SET domain and localizes in euchromatin regions where it catalyzes the mono- and dimethylation of H3K9
- •Overexpressed in several tumors such as lung, CRC, HCC, and bladder
- •It represses tumor suppressor gene expression (p21, RunX3), and silencing of G9a leads to decreased proliferation of colon and prostate cancer cell lines
- •To identify novel G9a inhibitors, we screened Inventiva's proprietary compound library (IVALib) using an AlphaLisa assay. In order to optimally screen the full library, the AlphaLisa assay, utilizing H3-derived peptides and a specific antibody against H3K9me2, was developed in a 384 well format with robust Z-factor (0.84) and signal to noise ratios of around 800



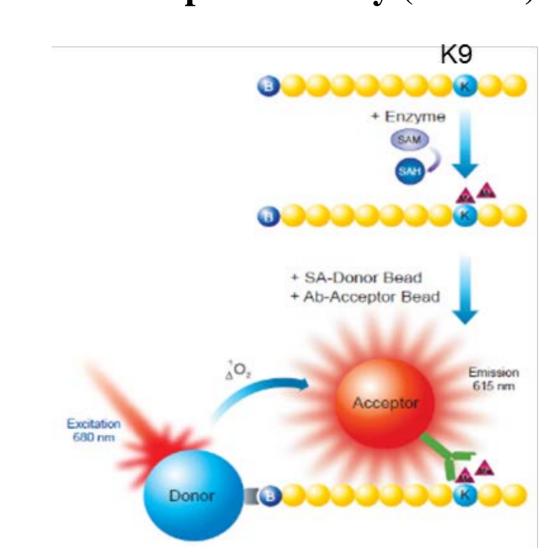
IVALib

- 240,000 Compounds
- IVALib has been designed and optimized over years for drug discovery programs
- More than 70% of the compounds are original when compared to Zinc library
- Compounds are available as liquid solutions and 70% as powders, and all the library is stored in controlled environment
- Regular quality controls are performed and a collection enrichment to maintain diversity and originality is in place
- Good hit rate on internal screening programs achieved
- Library available for external drug discovery partnerships

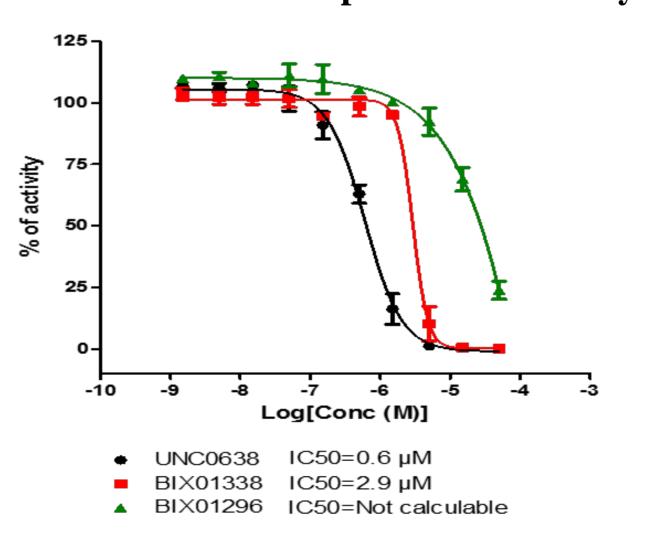
PRIMARY SCREENING ASSAY

We internally optimized an AlphaLisa® assay using G9a enzyme (BPS Bioscience), H3 biotinylated-peptide (Anaspec) and a specific Ab against H3K9me2 (Perkin-Elmer)

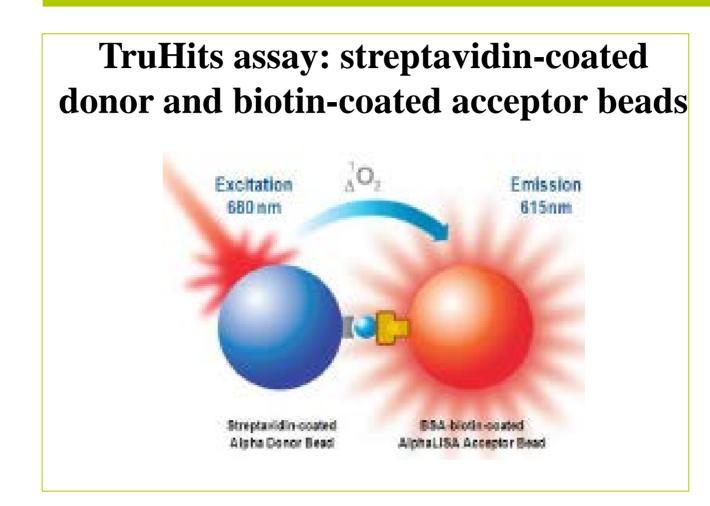
The assay gives excellent reproducibility (Z=0.84) in 384w format

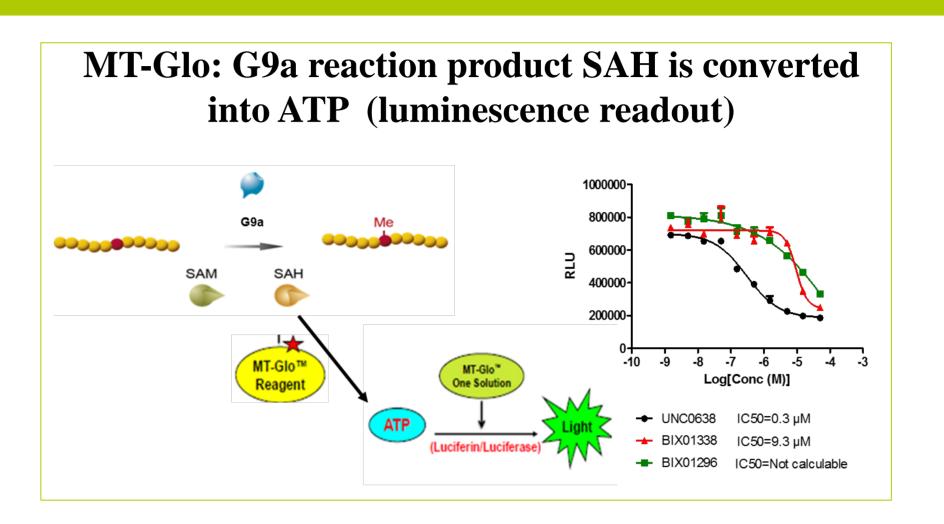


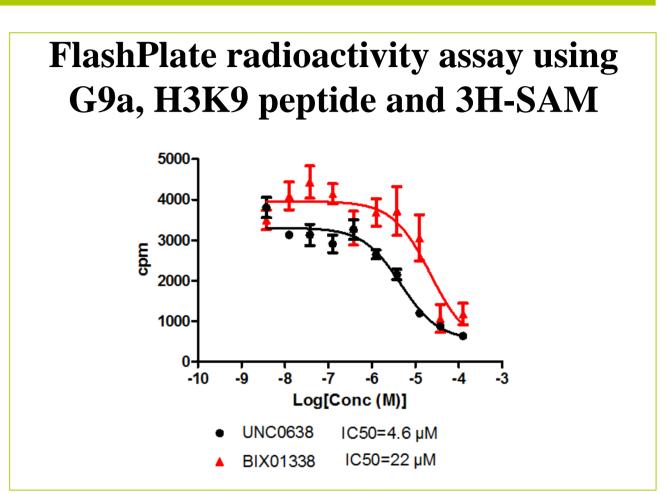
Activities of reference compounds were assayed



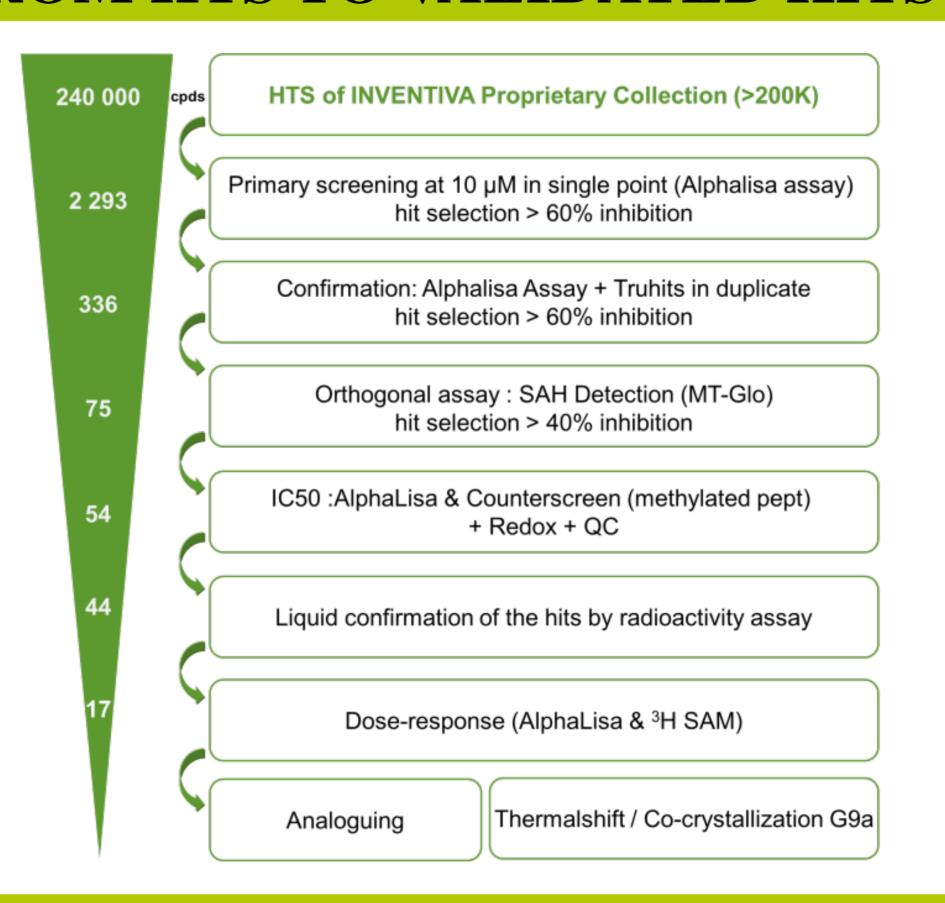
COUNTER SCREENING



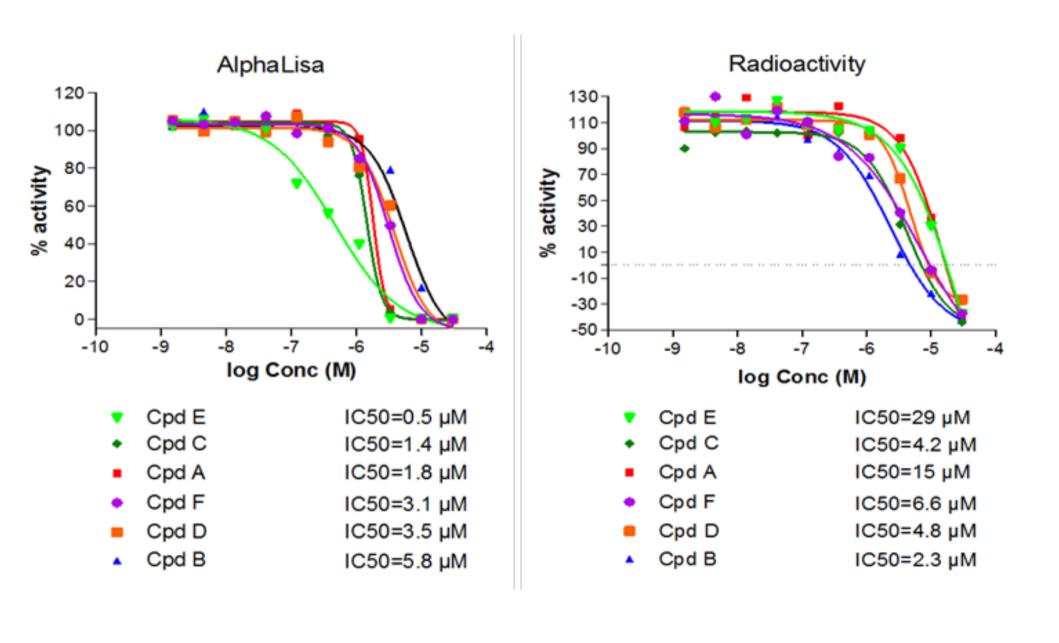




FROM HTS TO VALIDATED HITS



Example of hits showing activity in the µM range



Thermal Shift Assay

• Selected series exhibit $\Delta T_{G9a} = 1.6-2.4$ °C

CONCLUSIONS

- •Inventiva's proprietary compound collection screen allowed the identification of G9a hit inhibitor compounds
 - False positive compounds were rejected using orthogonal counter-screens assays
 - The biomolecular interaction of our hits with G9a was confirmed using a thermal shift assay
- G9a crystallography is ongoing
- We have identified chemical matters to enter into H2L phase to develop selective inhibitors of G9a
- G9a program is available for setting-up a drug discovery partnership.
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