

Due Diligence Report Lead optimization ADC-1013 Type: Version: Date: Page: Document ID: Report V2 2015-04-09 2 of 2 L0Y0T104C503

Aim

To develop an second antibody to be used in local immunotherapy of cancer based on a fully human antibody by increasing the affinity/potency from 1.7 nM to below 0.1 nM.

1 Summary

A **Construction** high affinity antibody for local immunotherapy of cancer was developed **Construction** a **Second Construction** antibody originating from a human antibody library

Five molecular libraries were designed based on the sequence and structure **descent**. These libraries contained restricted variability at specific positions or random variability covering the entire variable sequence. From these libraries, an initial selection round using phage display was made to enrich **descent** binders with moderate and high affinity.

The enriched molecular libraries were then recombined using FIND[™]. The resulting FINDlibrary contained recombined sequences from **Sequences** originating from all the starting libraries. This FIND-library was subjected to several rounds of phage selection with increasing stringency. From these selection rounds approximately 10,000 individual clones were screened in a concentration independent affinity ELISA. The top 100 clones were selected and screened in Biacore and in an in vitro functional assay and ranked based on protein behavior. The top 10 clones went further to hit-to-lead, where protein behavior, affinity, in vitro efficacy and potency were analyzed in more detail. ADC-1013 was chosen based on the results from this process. Finally, different Fc-format was evaluated, and it was decided to use the common gamma heavy chain for ADC-1013.

In conclusion the lead optimization using FIND resulted in a drug candidate exceeding the established goals for the project

This report was sent to **Constant of the Due Diligence** process

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