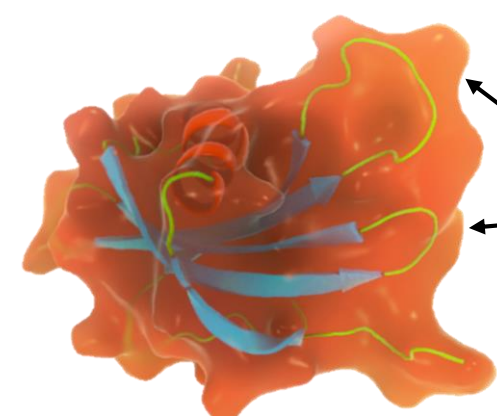


## Introduction

### Affimer Technology

- The Affimer biotherapeutic protein scaffold is based on human Stefin A
- Two surface loops engineered into the scaffold backbone



Binding loops: Two randomised 9 amino acid loop regions

- Phage display compatible - Large Affimer phage libraries (1x10<sup>11</sup>)

### Benefits of Affimer Therapeutics

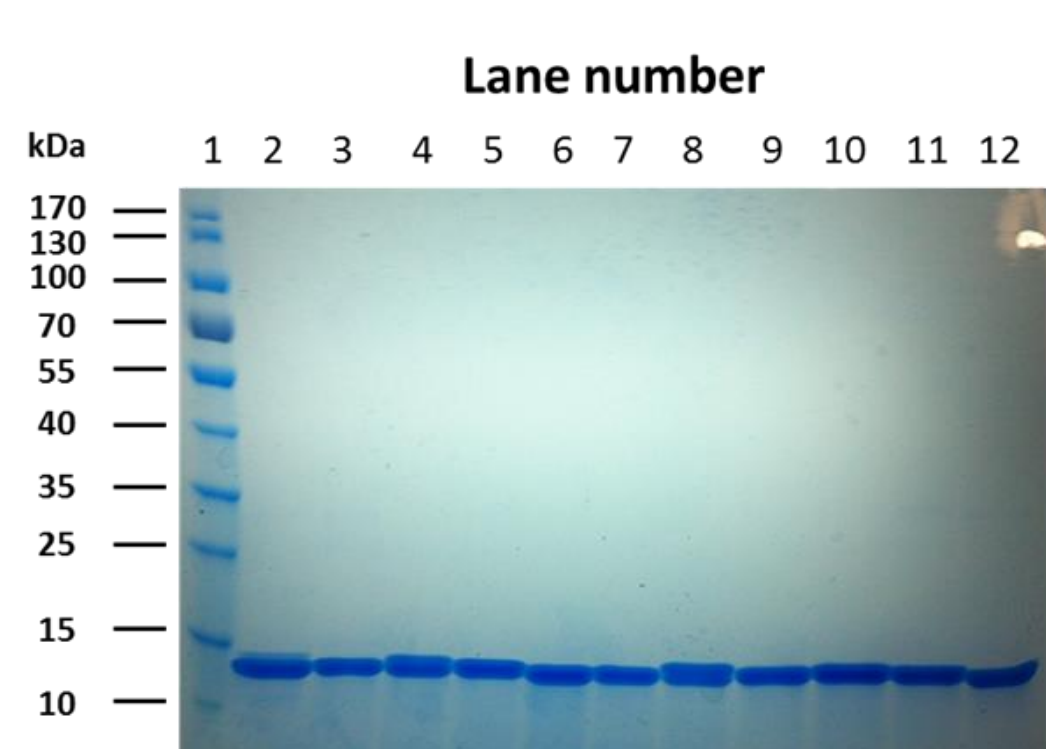
- Small size:** 14 kDa, 1/10<sup>th</sup> the size of an antibody
- High expression:** 100's mg/L in flasks (*E. coli*)
- No post translational modifications:** ease of manufacturing and improved stability
- Improved tissue penetration:** small size gives greater potential for increased efficacy
- Ease of formatting:** Fc format and in-line fusions, potential to generate multi-specific drugs to blockade multiple disease pathways

## Targeting the Immune Checkpoint PD-L1

- Programmed death-ligand 1 (PD-L1) is clinically validated in oncology and shown to play an important role in down regulating the immune system allowing tumour cells to evade detection and metastasize
- Affimer inhibitors with low single digit nM affinity were identified to either human or mouse PD-L1 (AVA04-182) using phage display.

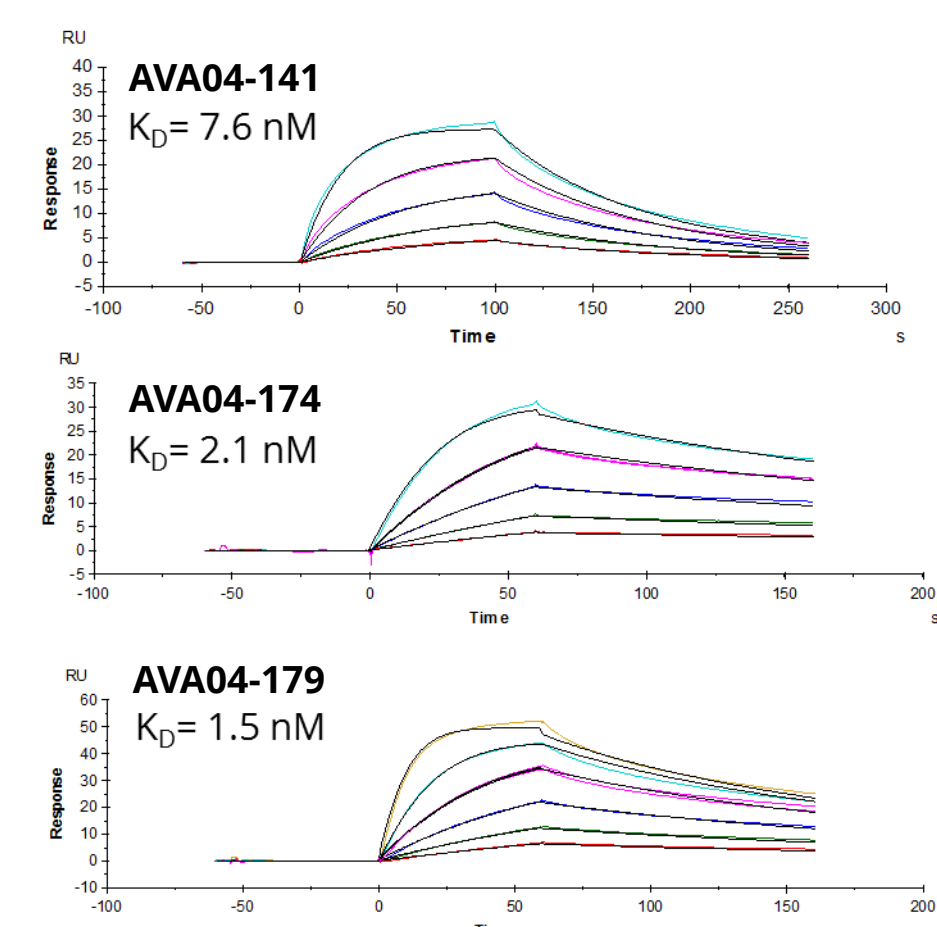
## Anti-PD-L1 Affimer Programme (AVA04): Lead Identification

AVA04 Affimer SDS-PAGE Analysis



Lane	Sample	Purified Yield (mg/L culture)
1	Markers	na
2	AVA04-141	300
3	AVA04-142	265
4	AVA04-144	340
5	AVA04-145	323
6	AVA04-146	295
7	AVA04-147	264
8	AVA04-148	231
9	AVA04-149	220
10	AVA04-150	239
11	AVA04-151	44
12	SQT (2 ug)	370

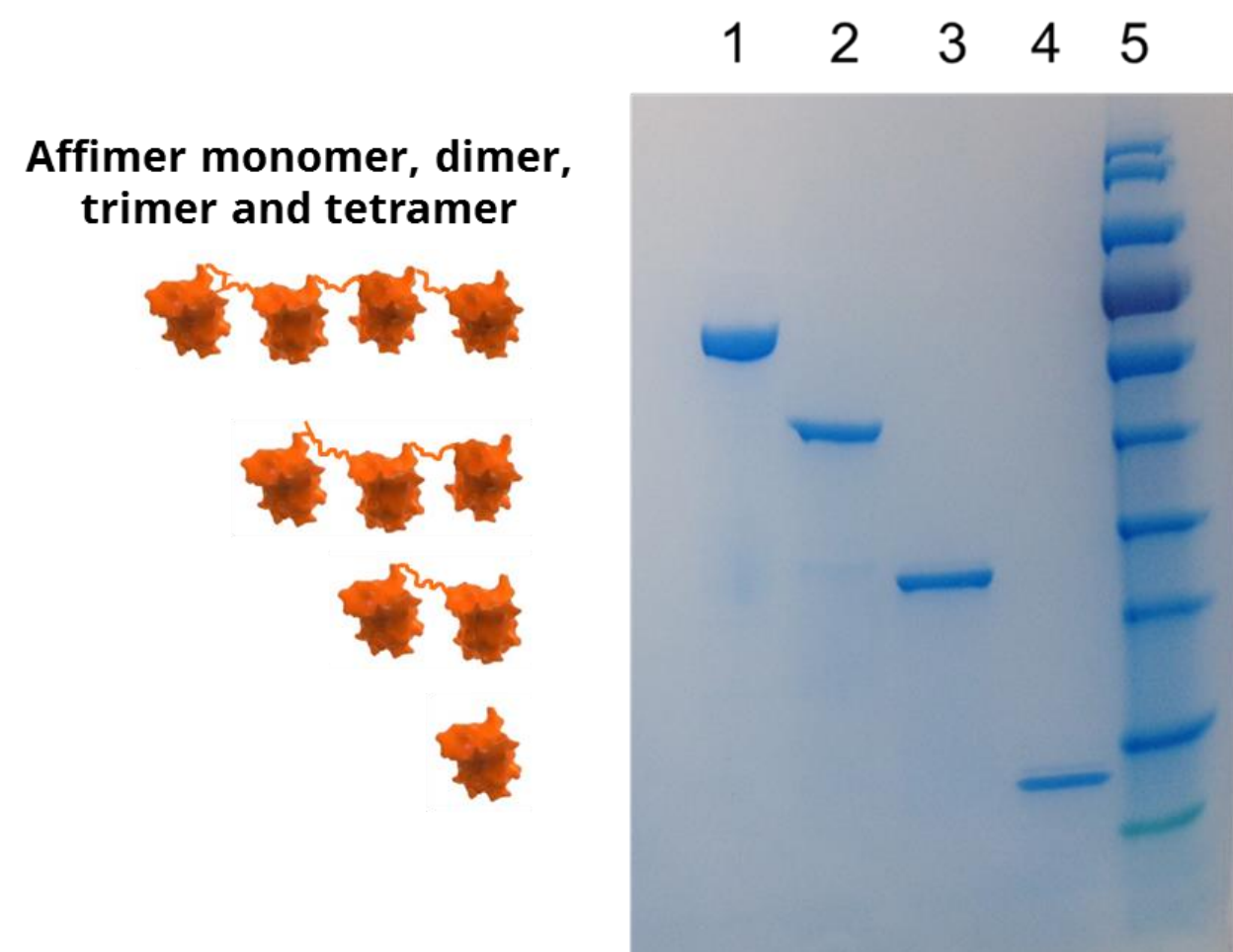
AVA04 Affimer Biacore



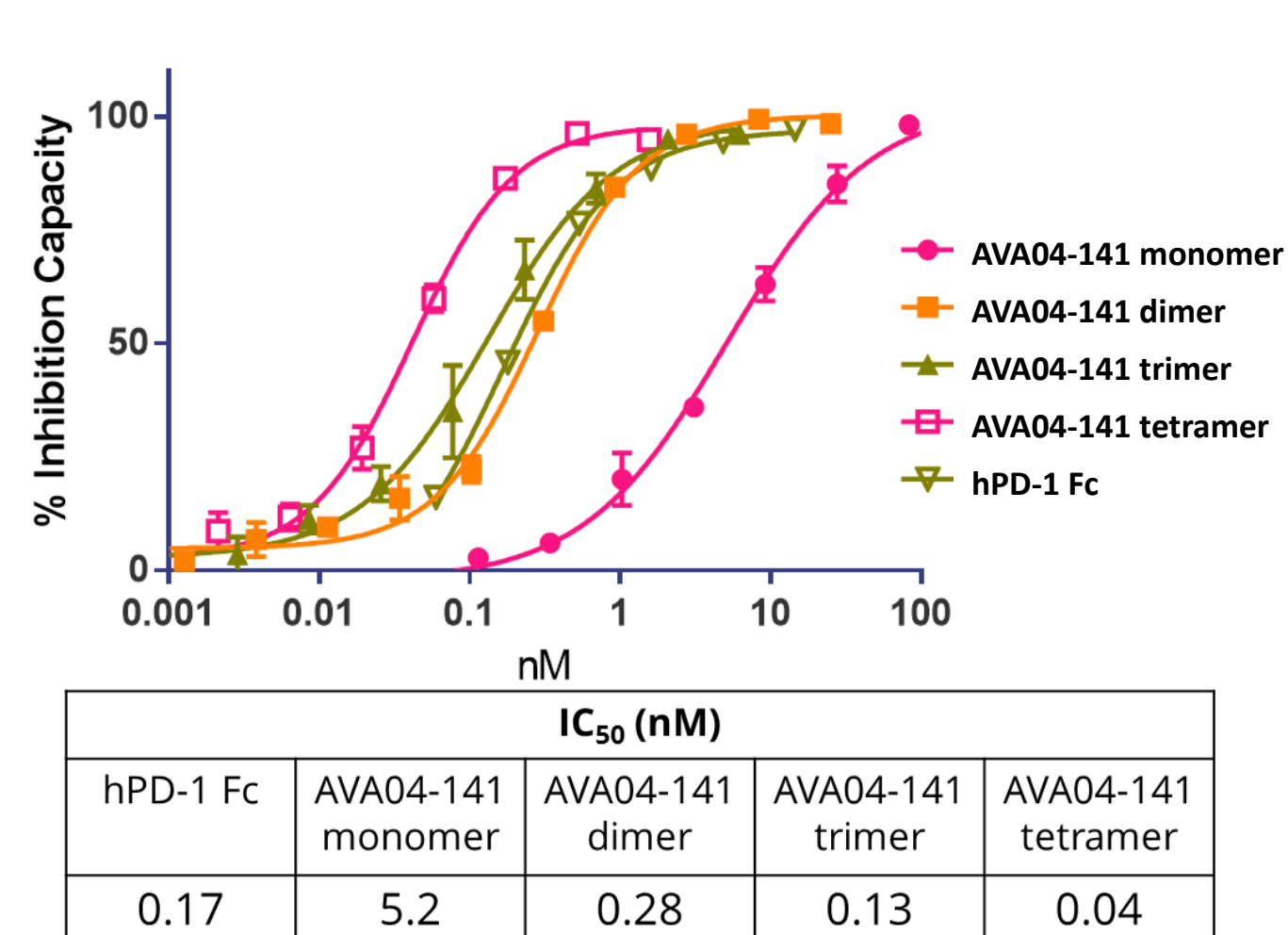
Lead Affimer proteins against human PD-L1 were identified by phage display and expressed, purified from *E. coli*. Over 60 unique Affimer sequences identified with high affinity to PD-L1 as determined by Biacore.

## Affimer Formatting : Multimers

AVA04 Affimer Multimer SDS-PAGE Analysis



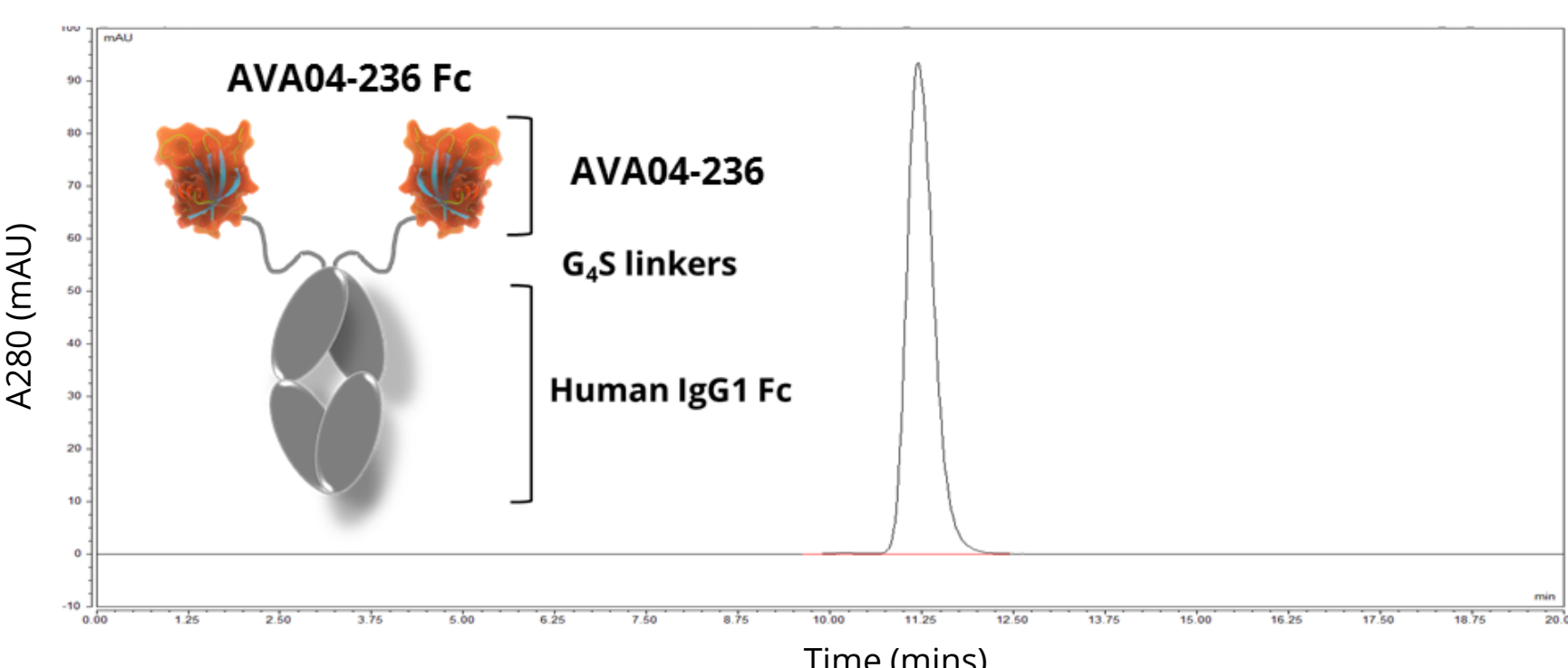
AVA04-141 Multimers Competition ELISA



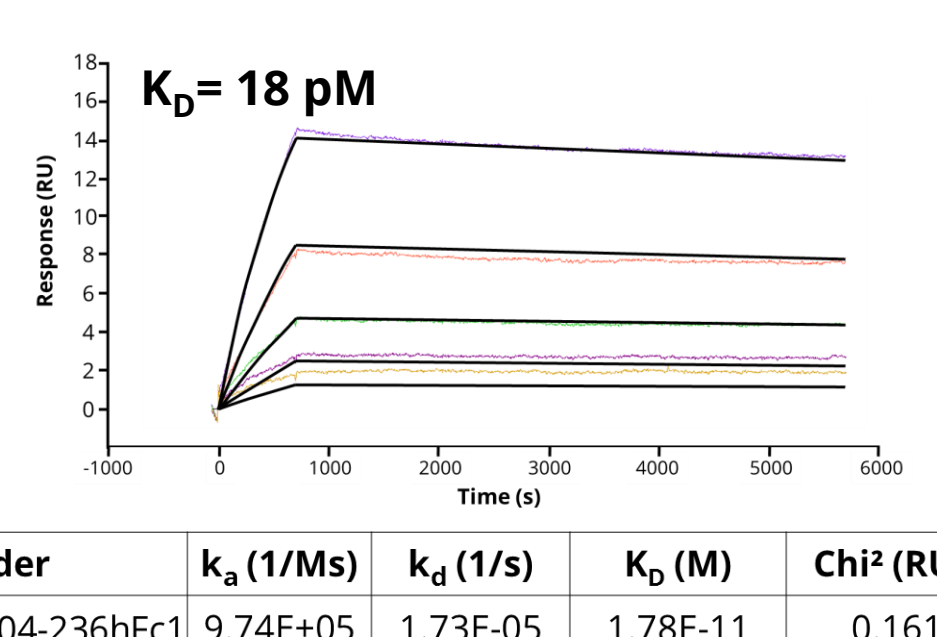
Affimer proteins can be formatted as multimers and expressed in *E. coli* to generate highly potent molecules. Unoptimised yields of purified Affimer multimers from *E. coli* >100mg/L in shake flasks.

## Affimer Formatting : Fc Fusions

AVA04-236hFc1 SEC-HPLC

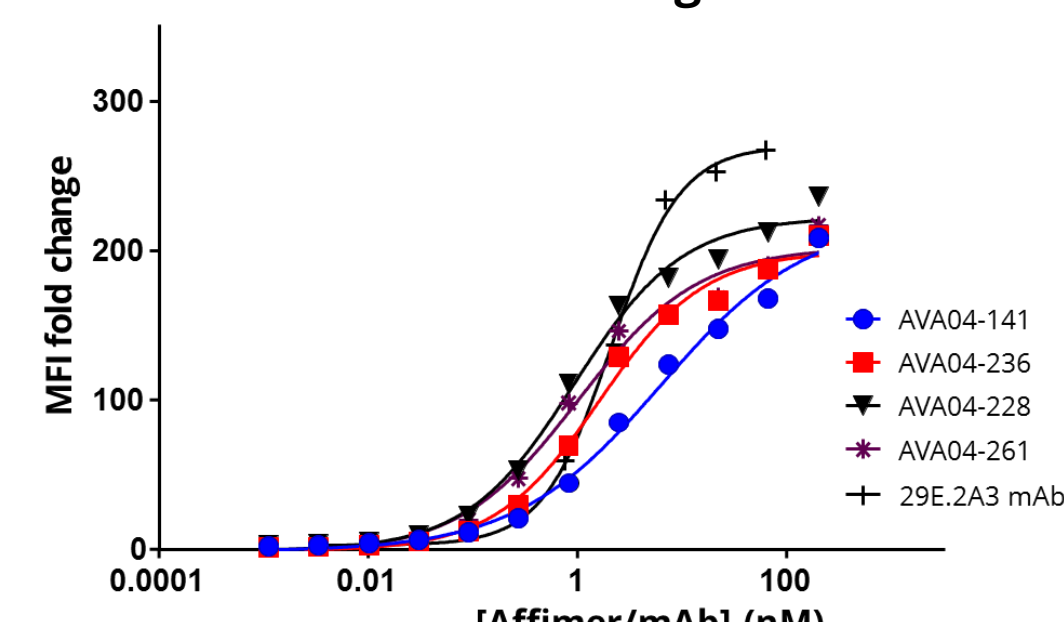


AVA04-236hFc1 Biacore Kinetics

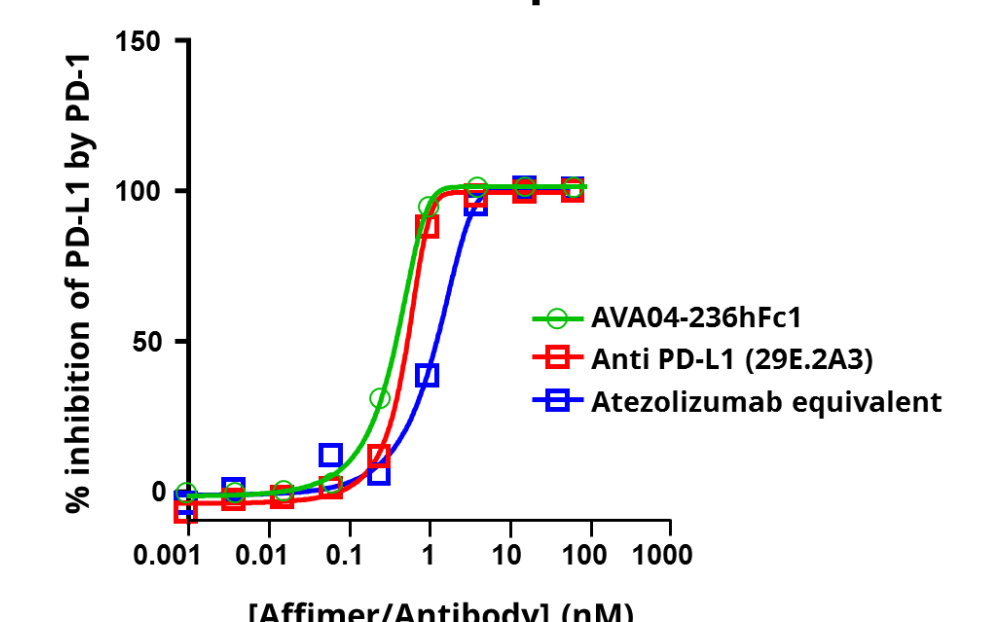


Affimer proteins can be formatted as Fc fusions to add effector function, half-life extension and enhanced affinity. AVA04-236hFc1 gives purified yields of ~200mg/L from transient Expi293F cells and shows similar affinity in the PD-1/PD-L1 competition ELISA to an Atezolizumab surrogate

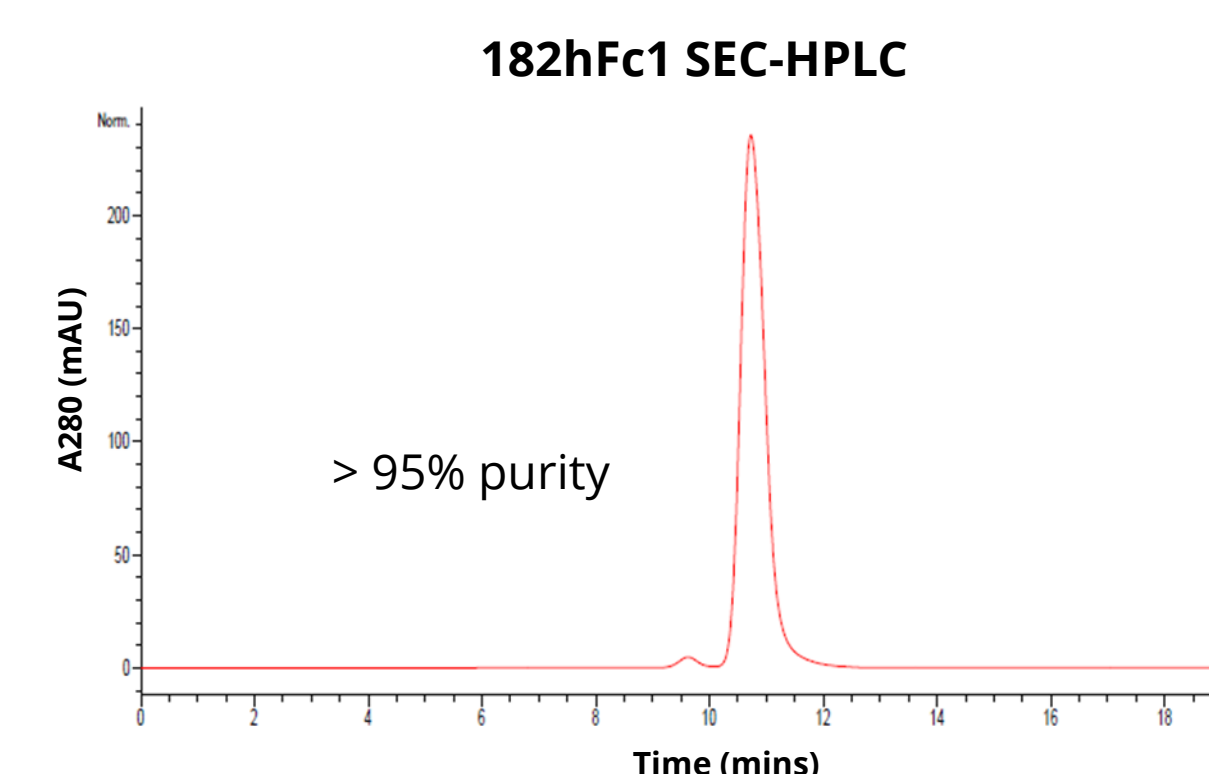
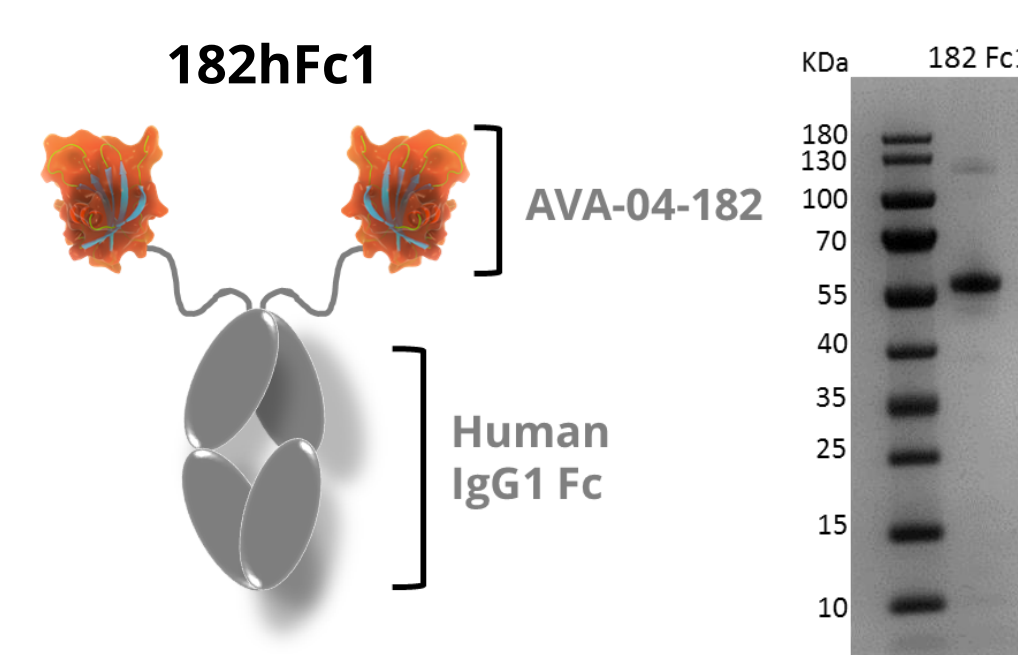
Affimer Monomer FACS Binding to CHO PD-L1 Cells



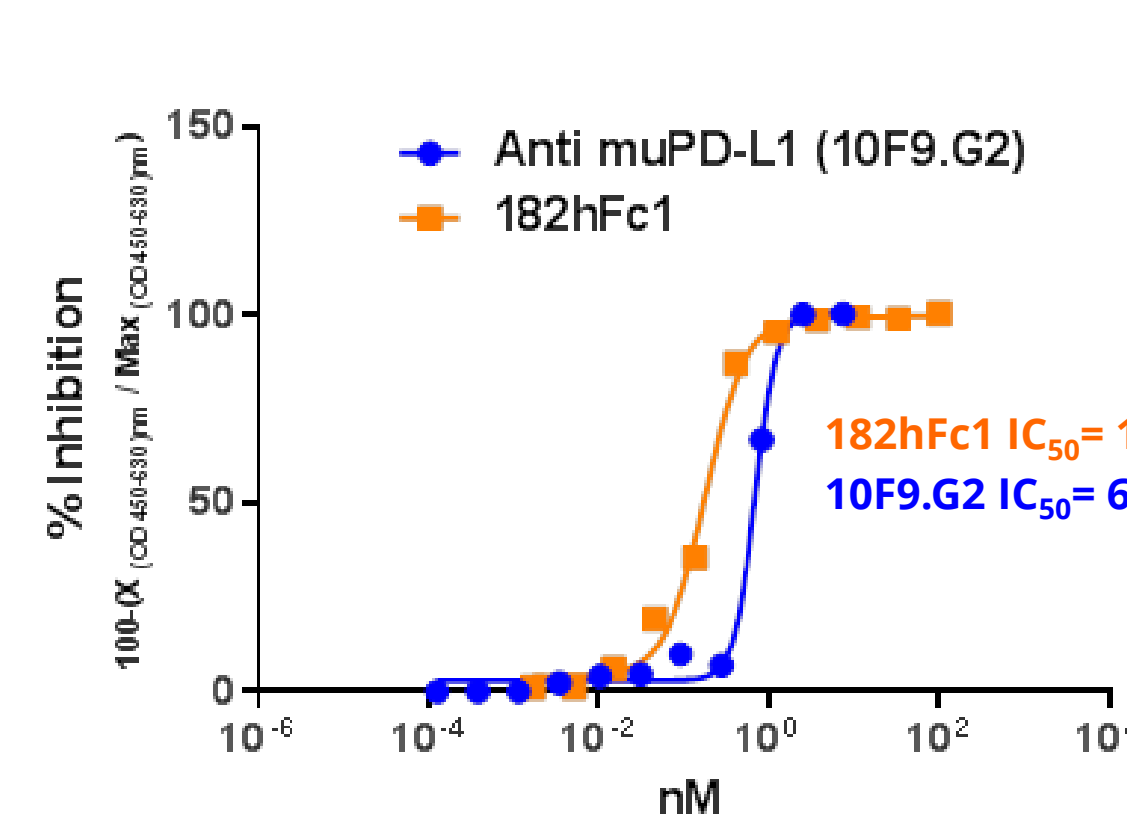
PD-1/PD-L1 Competition ELISA



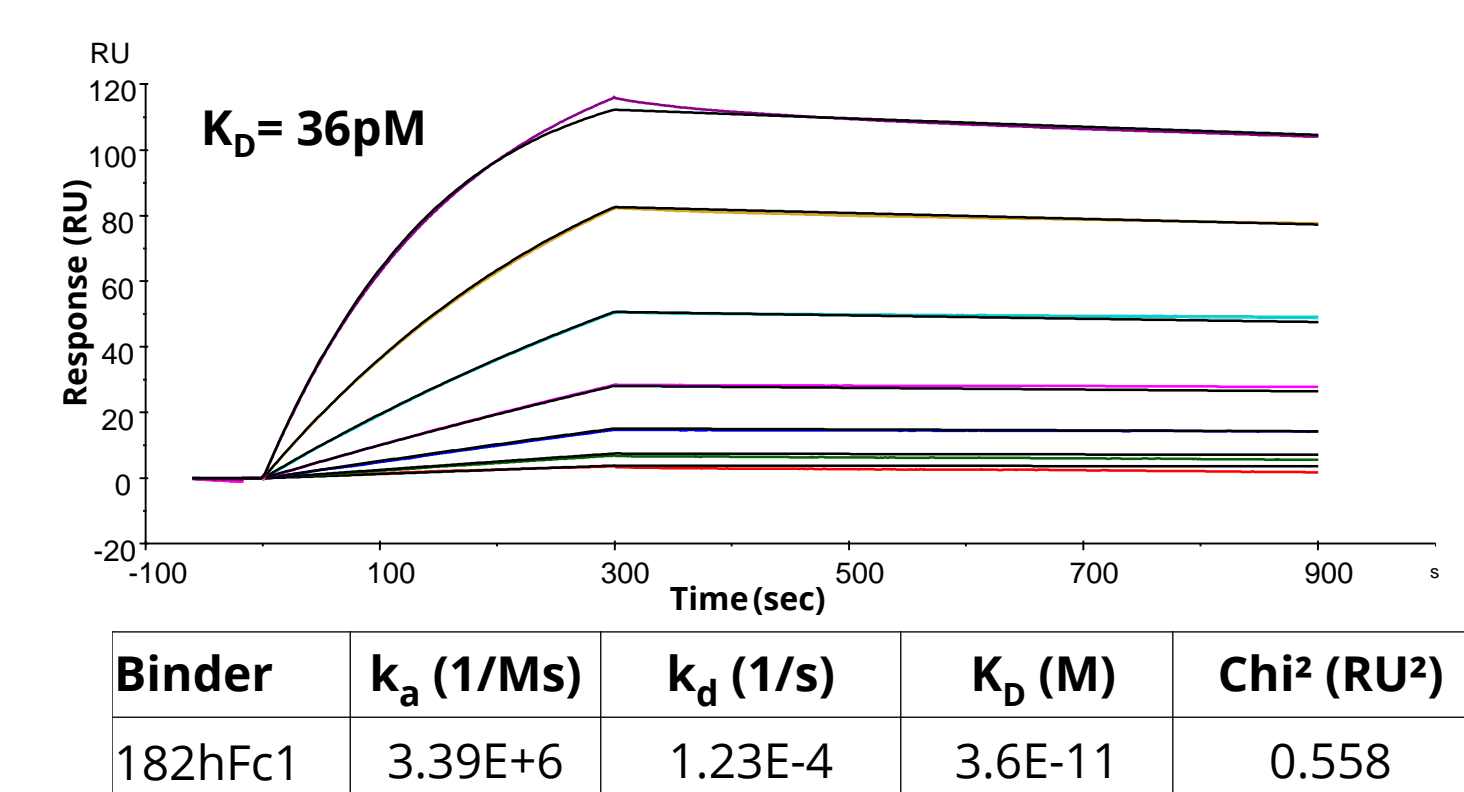
## AVA04-182 Formatting and Characterisation



182hFc1 PD-1/PD-L1 competitive ELISA

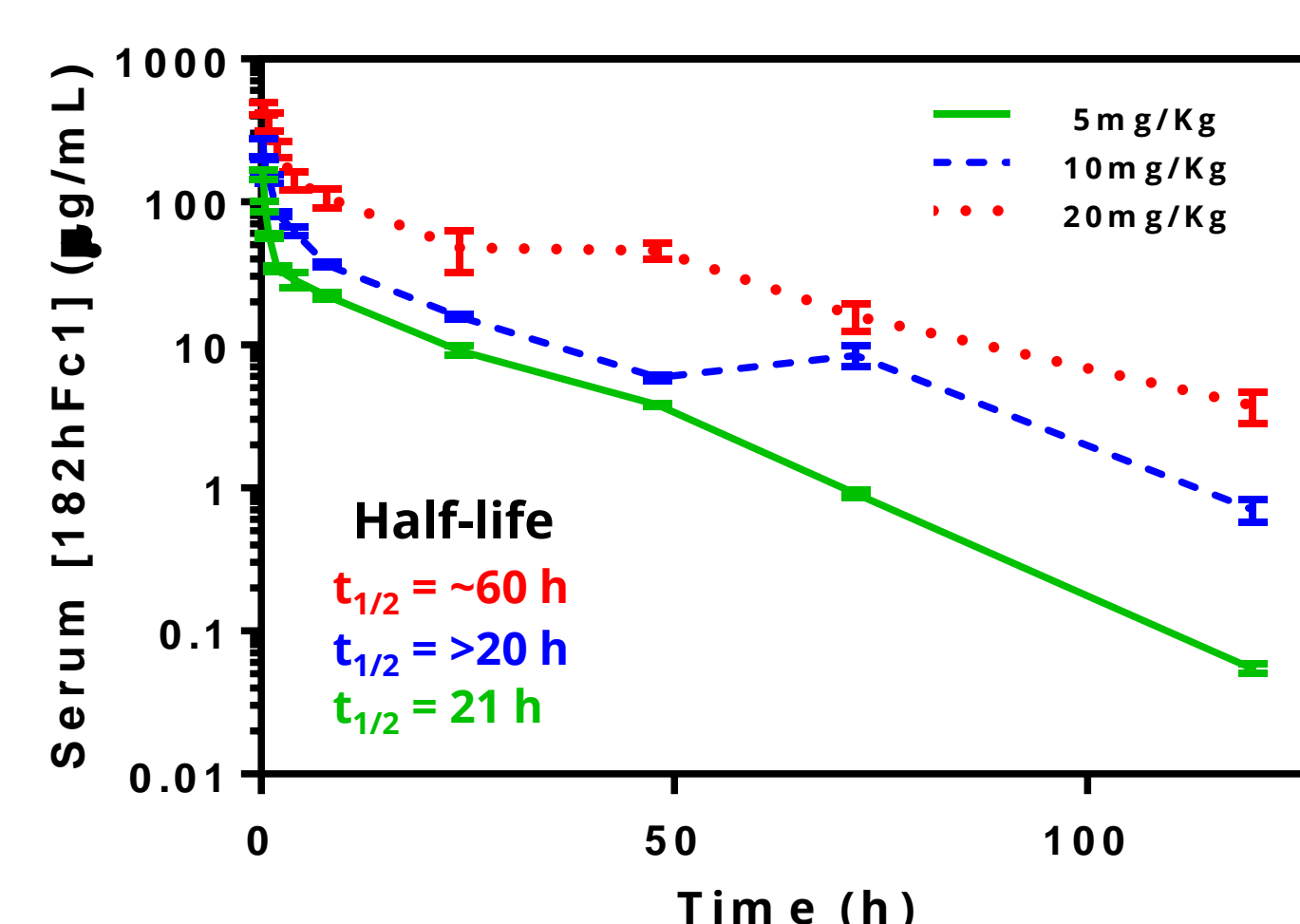


182hFc1 Biacore Kinetics



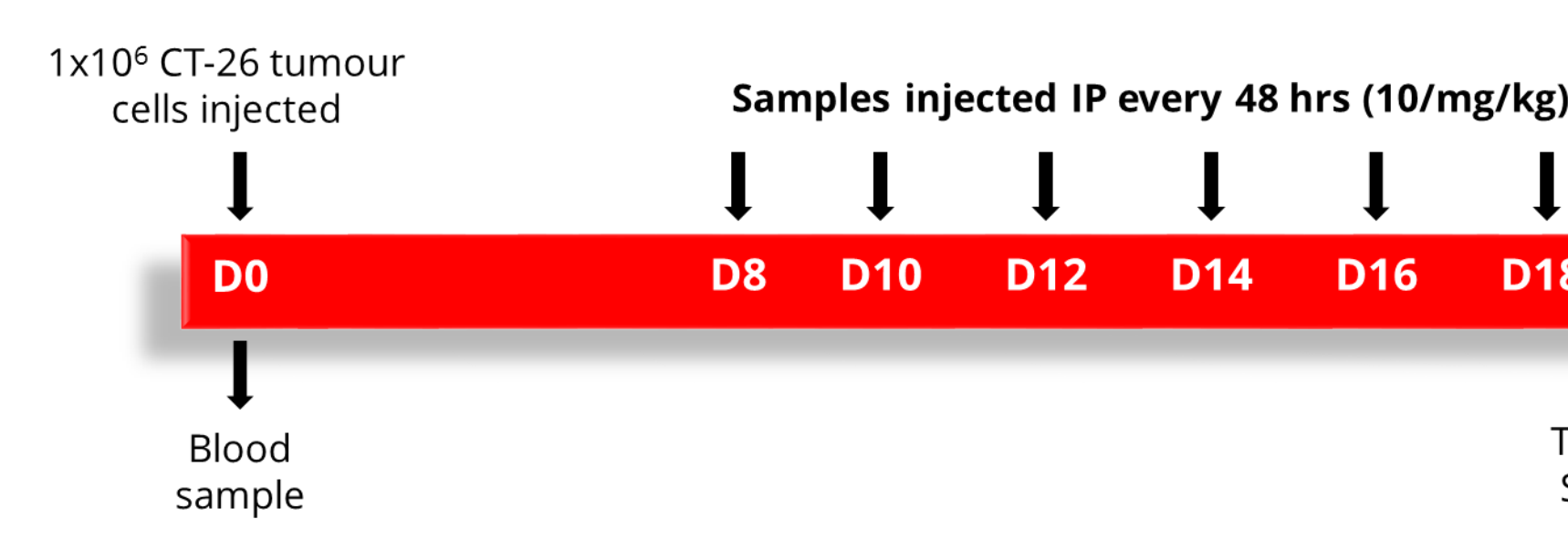
AVA04-182 only binds mouse PD-L1 and was formatted as an Fc fusion (182hFc1) for PK and efficacy models. Expression from Expi293 gave purified yields >100mg/L. 182hFc1 competes against PD-1 for binding to mouse PD-L1 (IC<sub>50</sub> = 178 pM) and has a K<sub>D</sub> of ~40 pM as determined by Biacore.

## Pharmacokinetics of 182hFc1 in Mice

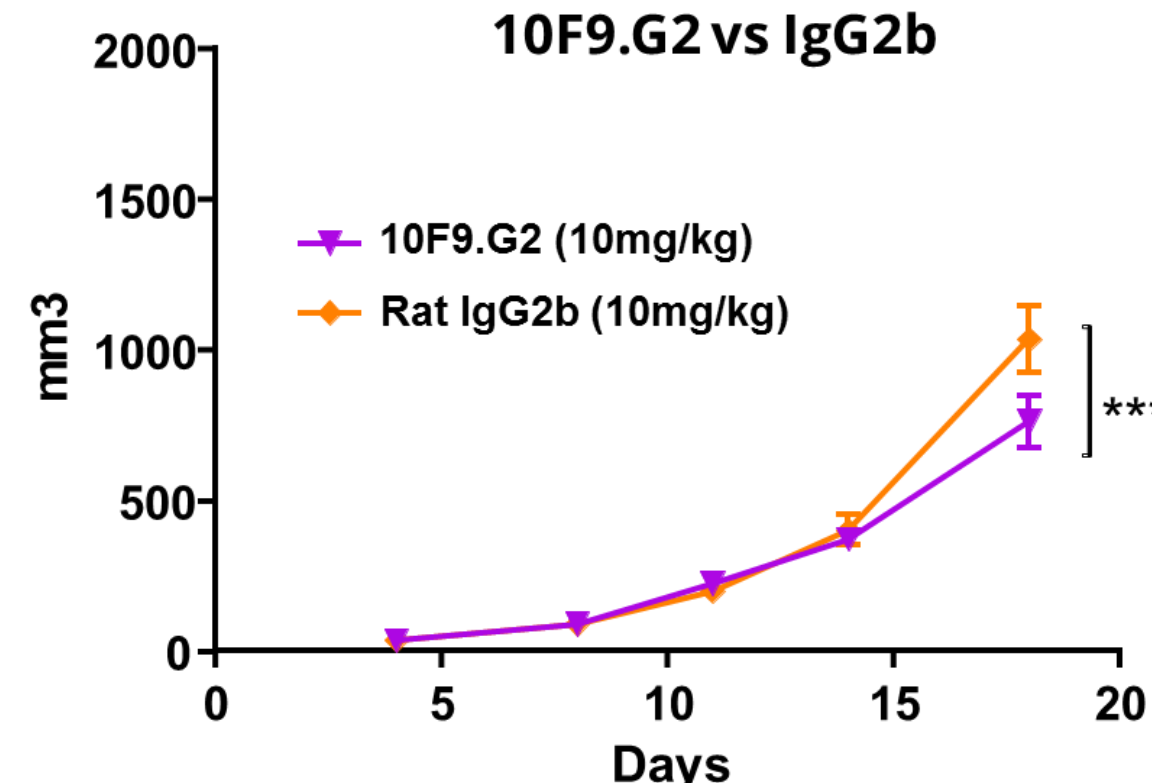


182hFc1 dosed as a bolus IV injection in 3 mice/time point. PK followed for 7 days and serum levels of 182hFc1 determined by binding ELISA. Affimer protein was well tolerated *in vivo* at all doses.

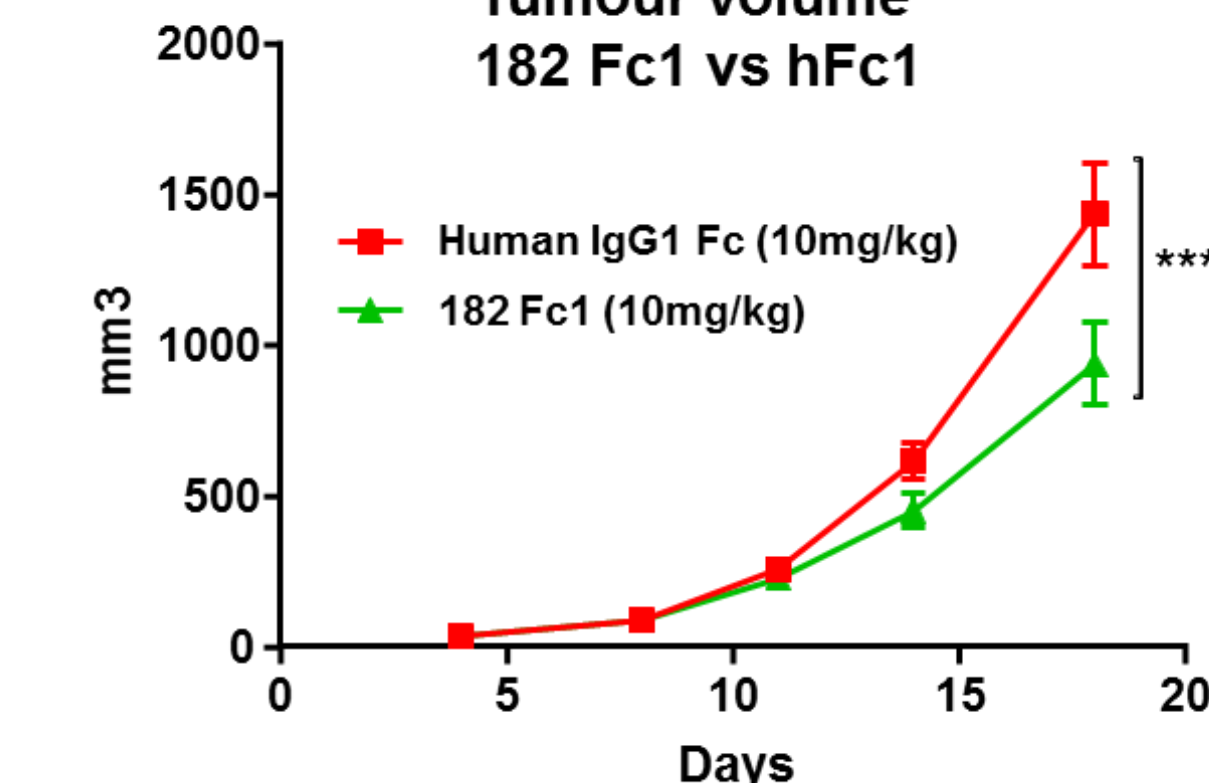
## Efficacy of 182hFc1 in a CT26 Mouse Syngeneic Model



Tumour Volume 10F9.G2 vs IgG2b



Tumour Volume 182 Fc1 vs hFc1



Balb/c mice received CT-26 tumour cells (1x10<sup>6</sup>) subcutaneously at D0. D8 to D18 182hFc1, mAb 10F9.G2 (rat anti-mouse PD-L1), hIgG1 Fc control and rat IgG2b isotype control were injected i.p. at 10 mg/kg every other day. 182hFc1 showed statistically significant inhibition of tumour growth compared to the hIgG1 Fc (hFc1) control.

## Conclusions

- Affimer biotherapeutics can be identified using phage display and easily produced at high levels in bacterial and mammalian expression systems
- The human PD-L1 Affimer antagonists can be formatted in a variety of ways to generate high affinity molecules as determined by Biacore and PD-1/PD-L1 competitive ELISA
- 182hFc1 was shown to be well tolerated in mice, even with repeat dosing at 10 mg/kg in the syngeneic model
- 182hFc1 inhibited tumour growth in the tumour of a CT-26 syngeneic model
- This work demonstrates that the Affimer technology has the necessary properties for a therapeutic platform: generation of high affinity binding proteins that can be formatted to extend the serum half-life and blockade a biologically relevant disease pathway *in vivo*

Acknowledgments : This work was part supported by an Innovate UK grant. The CT-26 syngeneic mouse model was run by Oncodesign (Nugues A. and Fancon M.)

