# Automated Solutions for Cellular Screening and Characterization of Therapeutic Antibodies for Antibody-Dependent Cellular Cytotoxicity Utility

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### Introduction

BioTek

### **Target Receptor Binding Affinity**

Automated Assay Workflow

Since the end of the 1990's, the pharmaceutical industry has seen an increased interest in biologics, especially in the therapeutic areas of oncology and inflammation. The selection of potent and selective Mabs (monoclonal antibodies) for specific target receptors, such as Receptor Tyrosine Kinase (RTK) and G protein-coupled receptors (GPCRs), is the first step in successfully developing an antibody based drug. In addition to potent binding, some antibodies have the ability to recruit the immune system effector cells, a process known as ADCC (Antibody Dependent Cell Cytotoxicity). The ability to promote ADCC is an important attribute of successful candidates and for that reason should be sought in the final drug design.

Here we present the automation of two assays for the characterization and selection of potent antibody drug candidates. Both assays rely on HTRF® detection. The first assay guantifies the binding affinity of antibodies to their target antigen, on live cells. The second assay measures the affinity of the antibody Fc portion to the CD16 receptor also on live cells. As Fc-CD16 binding affinity correlates well with effector function, this method is a simple, yet efficient and precise way of assessing biological activity through ADCC.

Each assay procedure has been automated through the incorporation of easy-to-use, robust instrumentation. An 8-channel liquid handler is used for antibody titration and transfer to the lowvolume 384-well assay plates. Cell and reagent dispensing are then carried out using a non-contact liquid dispenser. The small footprint of the instrument, and ability to autoclave the dispensing pathway, allow for sterile manipulations of each component. Binding, as well as antibody competition experiments were performed to validate the automated assay procedure. Results demonstrate that the combination of assay and instrumentation create a powerful, easy to use and efficient method to screen and characterize antibodies being considered for use in ADCC applications.

### **BioTek Instrumentation**



Figure 1 – Precision<sup>™</sup> Microplate Pipetting System combines an 8-channel pipetting head and an 8-channel bulk reagent dispenser in one instrument. The instrument was used to perform the serial dilutions of test antibodies and binding proteins, and transfer aliquots into the LV384-well assay plates.



Figure 2 – MultiFlo<sup>™</sup> Microplate Dispenser offers fast, accurate plate dispensing capabilities through its two peristaltic and two syringe pumps, with volumes ranging from 1-3000  $\mu$ L. The instrument was used to dispense cryopreserved cells, media, EC<sub>80</sub> concentrations of test antibodies, as well as labeled acceptor antibodies.

Figure 3 – Synergy™ H4 Hybrid Multi-Mode Microplate Reader combines a filter-based and monochromator-based detection system in one unit. The HTRF<sup>®</sup> certified reader uses the filterbased system and Xenon flash lamp to detect the 665 nm and 620 nm fluorescent emissions from this chemistry using Ex: 330/80 nm; Em: 620/10 and 665/8 nm filters, and a 365 nm cutoff dichroic mirror.

The assays described here monitor the ability of test antibodies to bind to a target antigen on the receptor of choice, such as a Receptor Tyrosine Kinase or GPCR, which has been cloned into a HEK293 cell line, expressed, and labeled with SNAP-Tb. Two specific receptors were used in this project; EGFR-An ErbB family receptor targeted by immunotherapeutics such as Cetuximab, and CXCR4-A GPCR belonging to the chemokine family, and also a known anti-cancer therapy target.

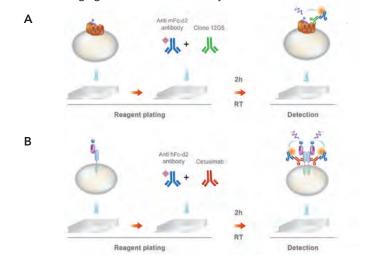




Figure 4 - Indirect Binding Measurement. Binding of antibodies to the target antigen is measured through the use of an anti-species specific Fc-d2 labeled antibody. The target antibody binds to the SNAP-Tag labeled receptor antigen (A.CXCR4; B.EGFR). Upon addition, the secondary labeled antibody binds to the primary, unlabelled antibody, causing an increase in HTRF signal.

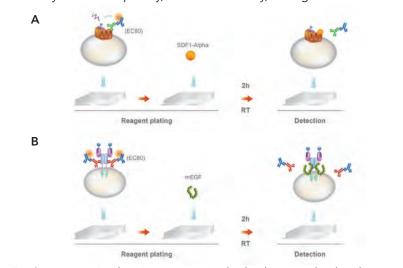


Figure 5 - Competitive Binding Assessment. Antibodies known to bind to the target receptor are added at an EC<sub>80</sub> concentration. Other binding proteins to the target receptor (A.SDF1-Alpha:CXCR4; B.EGF:EGFR) are then added. As the proteins bind to the receptor the primary:secondary antibody complex is dissociated, causing a decrease in HTRF signal.

# Fc Fragment: CD16a Binding Affinity

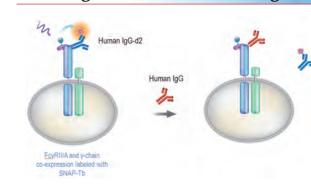
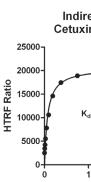
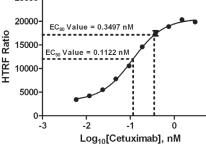


Figure 6 - Fc Binding on CD16a. Human IgG labeled with the d2 acceptor binds to HEK293 cells expressing the YRIIIA and gamma chain labeled with SNAP-Tb, generating a specific HTRF signal. Unlabeled IgG test antibodies can then be assessed for their ability to bind to the FcyRIIIA and gamma chain, which subsequently displaces the labeled IgG antibody, causing a decrease in HTRF signal





Cetuximab showed high affinity with a  $K_d$  in the subnanomolar range, which agrees with previously generated values of 0.03 nM, and those described in the literature<sup>1</sup>

# 12G5: CXCR4 Binding Assessment

# **CD16a Affinity Determination**



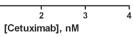
Figure 7 – Titration and transfer of antibody or competitive binding proteins initially accomplished. Subsequent antibody, cell, and media additions can be accomplished on the MultiFlo using a single or multiple addition methods.

# **Cetuximab: EGFR Binding Assessment**

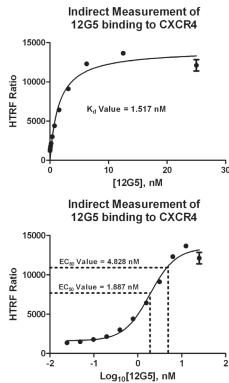
To gain insight into the binding characteristics of Cetuximab to EGFR, a binding experiment was performed in an indirect format through the use of a secondary anti-human labeled antibody.

> Indirect Measurement of Cetuximab binding to EGFR

> > K₄ Value = 0.071 nM



Indirect Measurement of Cetuximab binding to EGFR



**Competitive Binding Evaluation** 

Competitive Binding of Cetuximab and EGF

Log<sub>10</sub>[EGF], M

Competitive Binding of 12G5 and SDF1

to CXCR4

2G51 = 4 828e-9 M

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CXCR4 receptors, respectively.

15000-

10000-

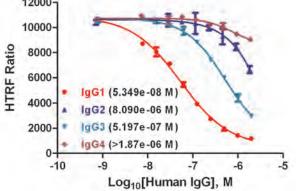
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performed once again using an indirect format through the use of a secondary labeled mouse anti-CXCR4 antibody. Figure 10 - 12G5 Equilibrium Binding Constant K, Determination. A 12G5

The binding characteristics of 12G5 to CXCR4 were also assessed. Initial binding of 12G5 was

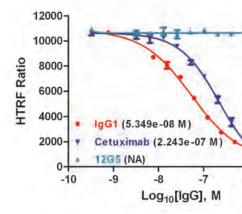
dose response curve was created using a 1:2 dilution scheme, starting at 25 nM. Cryopreserved cells were resuspended at a concentration of 8x10<sup>5</sup> cells/mL. [Antimouse Fc-d2 Ab] equaled 100 nM. The  $K_d$ value was determined using the Nonlinear Regression Michaelis-Menten curve fit.

Figure 11 – 12G5 Binding EC Calculation. The EC<sub>80</sub> value was determined from the  $EC_{50}$  and hill slope values, using a Nonlinear Regression Sigmoidal Dose-Response (variable slope) curve fit.



By comparing the IC<sub>50</sub> values found here to those generated previously using the manual method; 6.738e-08 M (lgG1), 4.225e-05 M (lgG2), 9.127e-07 M (lgG3), and 8.008e-05 M (lgG4), it is apparent that the rank order of IgG binding is accurately calculated, and also agrees with that reported in the literature.

The epidermal growth factor receptor (EGFR) inhibitor, Cetuximab, was also tested to ensure that the binding and effector cell recruitment of newly developed therapeutic antibodies could be accurately assessed with the automated assay procedure.



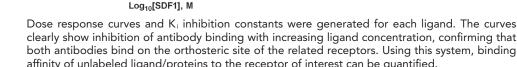
The IC<sub>50</sub> value for Cetuximab compared well to the manually generated value of 2.275e-07 M. This result, along with that from other human IgG antibodies, demonstrates that the automated assay is able to accurately assess binding of human IgGs to CD16a. As would be expected, the mouse 12G5 antibody does not dissociate the IgG-d2 antibody from the CD16a Fc receptor.

#### Conclusions

- competitive binding of unlabeled proteins or ligands, to target receptors
- 2. The CD16a assay is able to accurately detect the binding of test human IgG antibodies to CD16a Ec receptors
- 3. The Precision is capable of accurately creating and transferring dose response curves of antibodies and other competitive binding proteins, while the MultiFlo is able to rapidly dispense cells, media, and antibodies to LV384-well plates in a non-contact format
- 4. The HTRF® certified Synergy H4 is able to easily detect the fluorescent signals from each assay in a LV384-well format

Figure 8 – Cetuximab Equilibrium Binding Constant K<sub>d</sub> Determination. A Cetuximab dose response curve was created using a 1:2 dilution scheme, starting at 3 nM. Cryopreserved cells were resuspended at a concentration of 1x10<sup>6</sup> cells/mL. [Antihuman Fc-d2 Ab] equaled 30 nM. The  $K_d$ value was determined using the Nonlinear Regression Michaelis-Menten curve fit.

Figure 9 – Cetuximab Binding EC<sub>80</sub> Calculation. The  $EC_{80}$  value was determined from the EC<sub>50</sub> and hill slope values, using a Nonlinear Regression Sigmoidal Dose-Response (variable slope) curve fit.



K₁ Value = 9.217e-11 M

#### to EGER Cetuximab and EGF to EGFR. An EGF dose response curve was created using a 1:2 dilution scheme, starting at 1.67 $\mu$ M. Cetuximab was added at the EC<sub>80</sub> K, Value = 7.488e-10 M concentration previously determined. Cell and Fc-d2 antibody concentrations were

Competitive binding experiments were performed with the two systems previously described.

Known ligands EGF and SDF1 were used to illustrate competitive binding with the EGFR and

Figure 13 - Competitive Binding of 12G5 and SDF1 to CXCR4. An SDF1 dose response curve was created using a 1:2 dilution scheme, starting at 0.33  $\mu$ M. 12G5 was added at the EC<sub>80</sub> concentration previously determined. Cell and Fc-d2 antibody concentrations were as previously described.

Figure 12 - Competitive Binding of

as previously described.

clearly show inhibition of antibody binding with increasing ligand concentration, confirming that both antibodies bind on the orthosteric site of the related receptors. Using this system, binding affinity of unlabeled ligand/proteins to the receptor of interest can be quantified.



#### **IgG Binding Affinity**

The automated CD16a assay was validated using two different methods. The first included testing various IgG isotypes for their affinity to bind CD16a and recruit effector cells. Results were compared to values generated manually, as well as to those previously reported in the literature<sup>2</sup>.

> Figure 14 - Binding Affinity of IgG Antibodies to CD16a. Dose response curves were created using a 1:2 dilution scheme. starting at 1.98, 1.83, 2.00, and 1.87 µM for IgG1, 2, 3, and 4, respectively. Cryopreserved cells were resuspended at a concentration of 8x10<sup>5</sup> cells/mL [Human IgG-d2 Ab] equaled 50 nM

#### Therapeutic Ab Binding Affinity



Figure 15 - Therapeutic Antibody Binding Affinity to CD16a. Dose response curves were created using a 1:2 dilution scheme, starting at 1.98 µM for IgG1, and 0.83 µM for Cetuximab and 12G5. Cryopreserved cell and Human IgG-d2 Ab concentrations were as previously described.

1. The receptor binding assays are able to assess indirect binding of IgG antibodies, as well as

5. The combination of assay chemistry and instrumentation creates a perfect combination for highthroughput detection of therapeutic antibody binding in order to predict ADCC assay efficiency