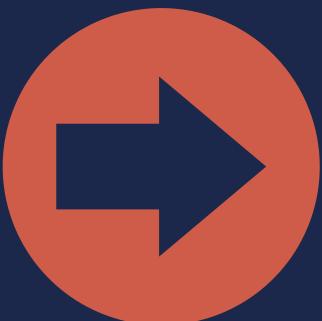


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# GPCR ASSAY TECHNOLOGIES GUIDE



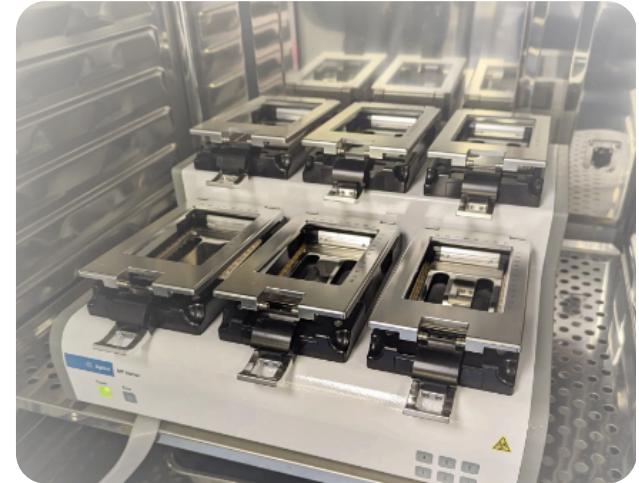
# CELLULAR IMPEDANCE

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## HOW IT WORKS

Microelectronic biosensor system that monitors whole-cell responses without labels

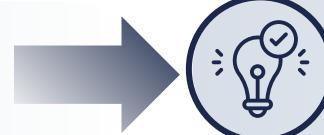


## TECHNOLOGY

Agilent xCELLigence cell analysis system



## KEY FEATURES



**Label-free** functional assay

**Broad applicability** across various cell types: endogenous GPCRs in primary cells, over-expressing cell lines, stem cells, or disease relevant cell lines

Simultaneously screen GPCR function across **all coupling classes**: Gas, Gaq, as well as Gai and Ga12/13

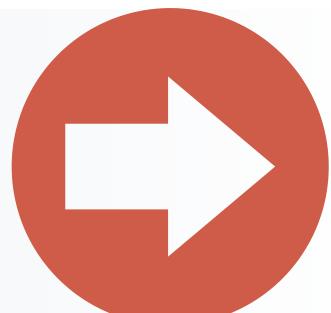
## INSIGHTS GENERATED

**Real-time, whole-cell kinetic data** enables deep, mechanistic insight into GPCR signalling

Interrogation of **complex signalling pathways** with high resolution, especially useful for receptors with **unknown signalling profiles** (eg. Orphan receptors)

Gain a **differentiated** view of GPCR pharmacology that helps uncover **nuanced signalling signatures and identify functional selectivity**

**YOUR PARTNER FOR SUCCESS IN GPCR DRUG DISCOVERY**



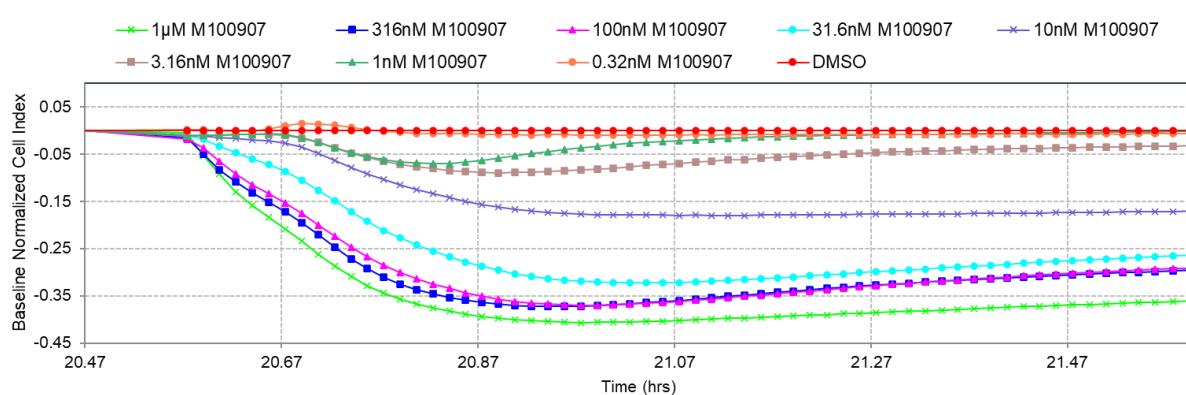


## CASE STUDY

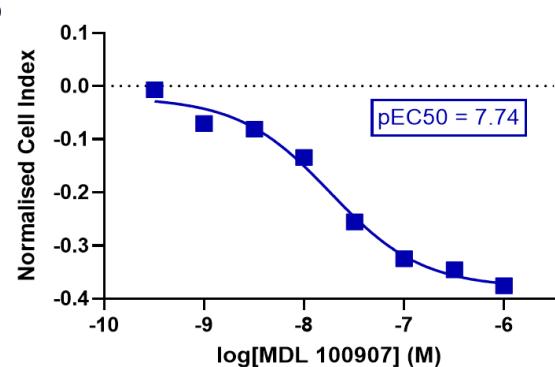
5-HT<sub>2A</sub> is a GPCR which is abundantly expressed in the CNS and an attractive clinical target for anxiety, psychosis and OCD.

MDL 100907 is characterised in the literature as a selective 5-HT<sub>2A</sub> antagonist. Profiling of MDL 100907 in the xCELLigence assay revealed that this ligand also displays inverse agonist activity at the 5-HT<sub>2A</sub> receptor. This result highlights the importance of incorporating orthogonal assay approaches into pre-clinical drug safety and efficacy profiling.

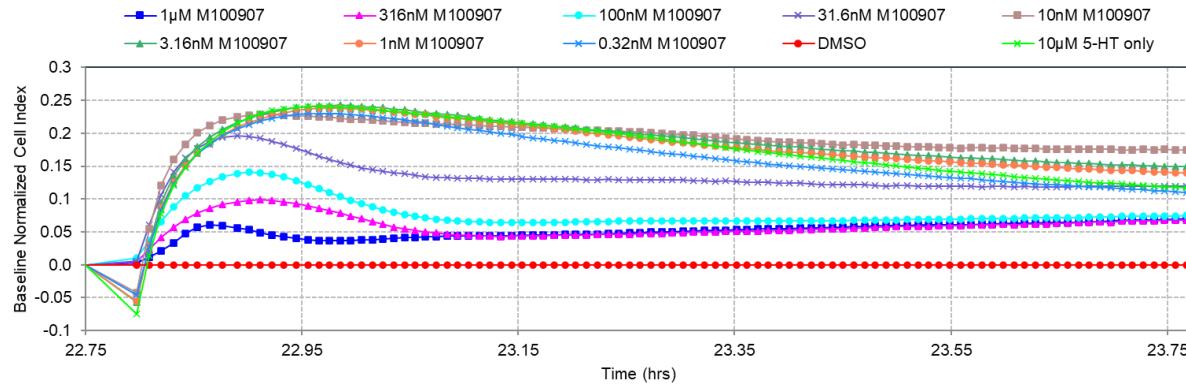
A



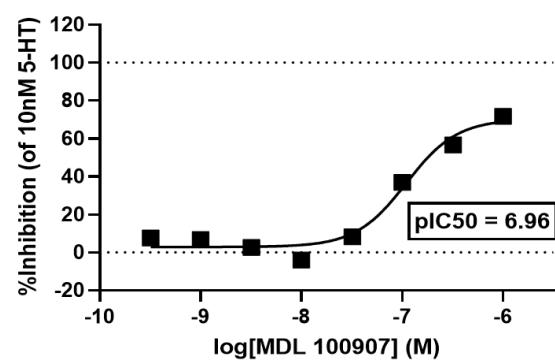
B



C



D



**Figure 1.** MDL 100907 shows antagonist and inverse agonist activity at the 5-HT<sub>2A</sub> receptor in the label-free xCELLigence assay. Baseline normalised cell index trace for MDL 100907 alone (A) or 10nM 5-HT (EC<sub>80</sub>) following pre-treatment with the indicated concentrations of MDL 100907 (C). Normalised cell index values plotted as a dose-response curve for MDL 100907 alone (B) or dose-inhibition curve for MDL 100907 pre-treatment, plotted at the peak agonist response (D).



# $\beta$ -ARRESTIN RECRUITMENT bio:ascent



## HOW IT WORKS

A live cell, luminescence based protein-protein interaction assay measuring  $\beta$ -arrestin recruitment to GPCRs in real time

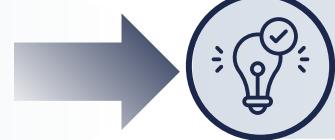


## TECHNOLOGY

Promega NanoBiT Arrestin



## KEY FEATURES



Generic method for all GPCRs;  
**useful for non G protein pathways**

Useful for **biased agonism** studies

Useful for orphan receptors where endogenous ligands/  
**coupling pathways are unknown**

Readily **scaled** for HTS

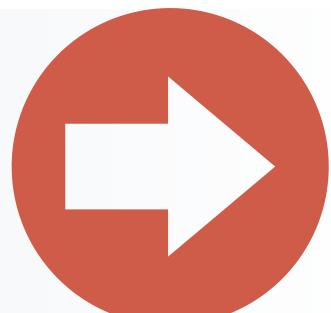
## INSIGHTS GENERATED

Uncovers GPCR pharmacology **beyond traditional G-protein signalling**

Quantifies **ligand bias**. Helps determine whether compounds favour G-protein signalling,  $\beta$ -arrestin recruitment, or demonstrate a balanced profile

Supports the development of **safer, more targeted therapeutics** by enabling **precise control** over downstream signalling pathways

**YOUR PARTNER FOR SUCCESS IN GPCR DRUG DISCOVERY**





## HOW IT WORKS

Fluorescent dye and high speed plate imaging system detects intracellular  $\text{Ca}^{2+}$  changes

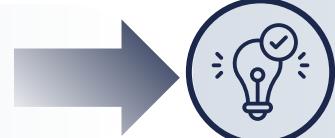


## TECHNOLOGY

FLIPR Penta



## KEY FEATURES



Pathway specific for **G<sub>aq</sub> signalling**

Can be used with **G<sub>a16</sub>-engineered** systems, enabling receptors to be “forced” through the calcium pathway for rapid functional assessment

Highly **sensitive** to early signalling

FLIPR Penta - extremely **high-throughput**

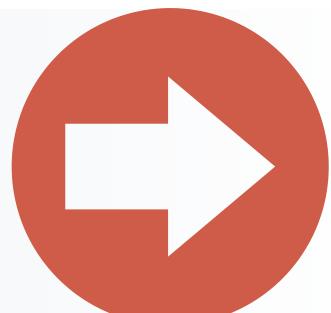
## INSIGHTS GENERATED

Captures **full real-time** fluorescence traces, going far **beyond simple max-min endpoints**

Detailed **interrogation of signalling behaviour**, including onset time, peak amplitude, and the rate of signal decay as cells return to baseline

**Rich kinetic insight** supports deeper mechanistic understanding, ligand differentiation, and confident decision-making

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# IP<sub>1</sub> ACCUMULATION

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## HOW IT WORKS

Measures accumulation of IP<sub>1</sub> caused by Gaq activation using TR-FRET

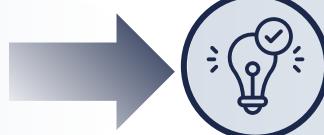


## TECHNOLOGY

HTRF IP-One assay



## KEY FEATURES



Suitable for **Gaq-coupled GPCRs**

Functional assay for live cells which works with **many cell systems**

Robust assay suitable for **miniaturisation and HTS** cascades



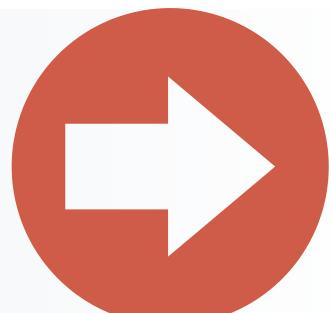
## INSIGHTS GENERATED

Directly **quantifies Gaq-coupled GPCR signalling**

Unlike rapid and transient calcium flux, IP<sub>1</sub> accumulation provides a **longer, more forgiving measurement window**, delivering a stable and **highly reproducible** readout

Ideally suited for HTS applications - this assay is an **excellent choice for reliable compound screening**

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# cAMP MEASUREMENT

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## HOW IT WORKS

Measures intracellular cAMP levels as a readout of Gas or Gai/o signalling

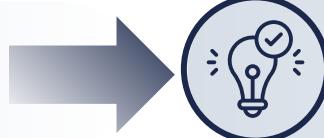


## TECHNOLOGY

Promega Glosensor cAMP  
Bellbrooks transcreener cAMP  
Revvity HTRF cAMP



## KEY FEATURES



Highly **sensitive**, quantitative readout

Versatile - suitable for **both Gas and Gai** receptors

**High-throughput** format ideally suited to HTS

**Flexible** - performs equally well in endogenously expressing cell lines as in stably or transiently transfected systems

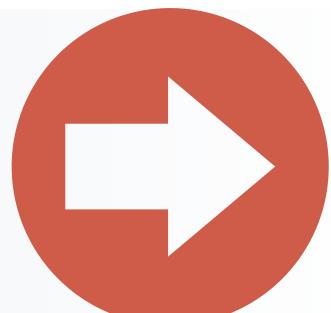
## INSIGHTS GENERATED

**Reliable insight** into compound behaviour and receptor signalling - kinetic insight empowers deeper mechanistic understanding and more confident decision-making

Excellent for **SAR generation and medicinal chemistry** due to quantitative data

High sensitivity enables detection of **weak agonists, partial agonists, and inverse agonists**

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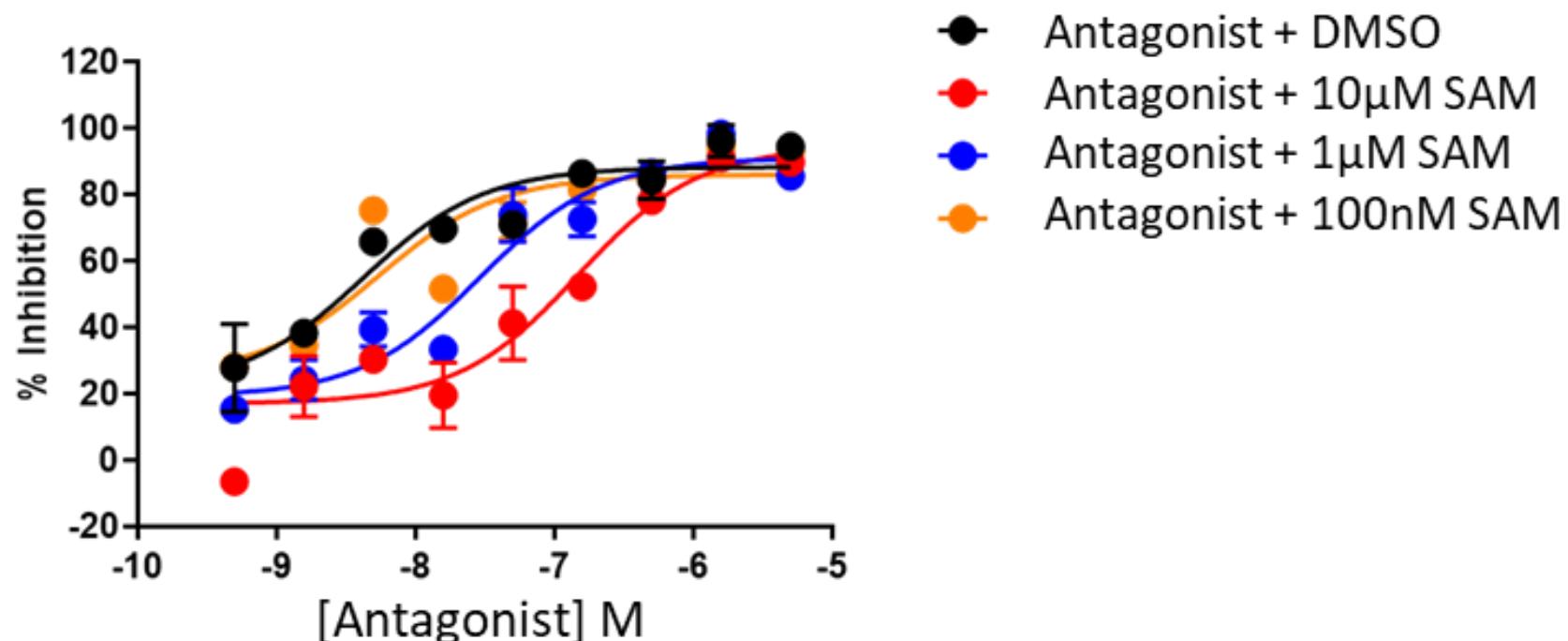


## CASE STUDY

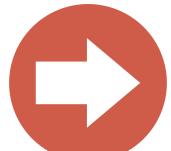
During a long running Lead Optimisation project on a GPCR target, we observed some very steep SAR, which was unexpected.

The possibility of silent allosteric modulators (SAMs) was investigated using the functional cAMP primary assay. We pre-incubated  $\pm$  the suspected SAM with a known allosteric antagonist before the addition of the agonist.

Dis-inhibition of the antagonist with no agonist activity on its own indicates that the compound is indeed binding at the same site but doing so without causing a functional effect.



**Figure 2.** Allosteric antagonist dose response curve  $\pm$  increasing concentrations of potential silent allosteric modulator in the presence of agonist EC<sub>80</sub>. Data are expressed as percent inhibition of the agonist EC<sub>80</sub> response and are the mean  $\pm$  SEM of a minimum of 4 replicates.





## HOW IT WORKS

Measures phosphorylation of ERK1/2, a key downstream signaling output of many GPCRs



## TECHNOLOGY

Alphascreen Phospho-ERK assay

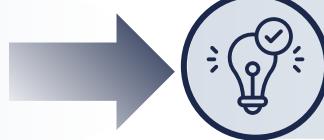


## KEY FEATURES

**Universal** readout - suitable for Gaq-, Gai/o-, and some Gas-coupled receptors

Sensitive technique which can detect **weak or partial agonists**

Captures **both** G-protein and  $\beta$ -arrestin mechanisms



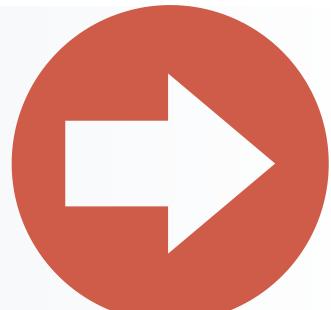
## INSIGHTS GENERATED

Provides a readout of integrated GPCR signalling - especially valuable for receptors that are **difficult to monitor using traditional second-messenger assays**

Captures ERK as a convergence point for both G-protein dependent and  $\beta$ -arrestin mediated pathways, offering a **broad, biologically meaningful view** of receptor activity

Enables **reliable detection** of ligand-driven signalling events

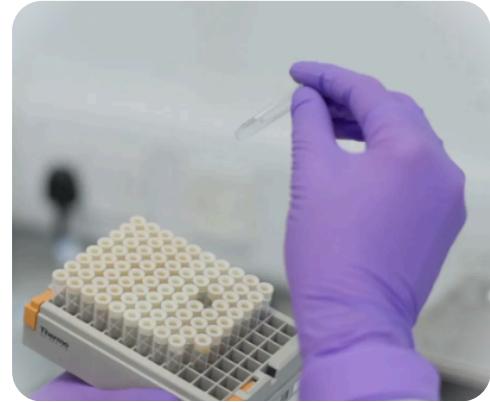
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## HOW IT WORKS

Directly monitors G-protein activation by tracking the separation of tagged Ga and Gy subunits



## TECHNOLOGY

Bioluminescence Resonance Energy Transfer (BRET)



## KEY FEATURES

Reports proximal, immediate G-protein engagement, letting you see exactly **which G-protein family a compound activates**

Delivers a **depth of mechanistic pharmacology** that traditional assay formats simply cannot achieve

**Optimised** constructs available for nearly every Ga, G $\beta$ , and Gy subtype



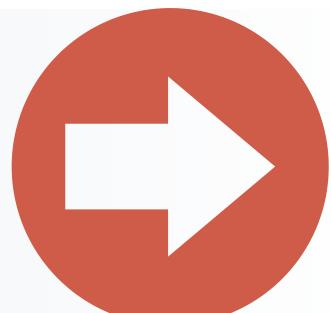
## INSIGHTS GENERATED

Enables **precise mapping** of GPCR G-protein coupling and signaling bias - reveals exactly which G-proteins a receptor couples to, and the strength of those interactions

Enables the discovery and optimisation of ligands with **finely tuned, G-protein-specific bias**

Uncovers **unique signalling signatures** - particularly valuable when working with first-in-class compounds or orphan receptor programmes

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READY TO MOVE YOUR  
GPCR PROJECT  
FORWARD?

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