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Kinetic measurements of intracellular ATP levels in co-culture models using live-cell analysis

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Summary & Impact

- Differential metabolic requirements of tumor cells have gained recognition as attractive therapeutic targets (e.g. glutaminase1 (GLS1) inhibitor CB-839 in clinical trials for a variety of indications, including triple-negative breast cancer (TNBC)).

for the duration of the 3-day time course. In contrast, most receptor-positive cell lines displayed little to no effect of GLS1 inhibition.

– Co-culture with CCD-1086SK fibroblasts attenuated the effect of CB-839 in TNBC cell lines, an effect which was not due to drug buffering.

Incucyte[®] System for continuous live-cell analysis: Methodology



IncuCyte[®] ATP Software Module

Fast, flexible and powerful control hub for

- Graphing of normalized kinetic data

NEW: Incucyte[®] ATP Analysis

Software Module

continuous live-cell analysis comprising image

acquisition, processing and data visualization.

- Color scaled visualization of cytosolic ATP levels

- Standard approaches to monitoring drug induced metabolic perturbations are limited to endpoint assays that lack cell-specific data in complex co-culture models and provide limited kinetic information.
- The Incucyte[®] ATP Assay provides a live-cell approach to image and analyze cytosolic ATP levels of tumor cells in mono- or co-culture with stromal cells over time.
- Across all TNBC lines tested, CB-839 treatment resulted in a reduction in ATP that was sustained
- Visualization tools included in the Incucyte[®] ATP Analysis Software Module provide a quick, qualitative assessment of data across the assay plate and a view into the heterogeneity of treatment responses.
- Measurements of proliferation, cell death, and mitochondrial membrane potential enabled by the Incucyte[®] Metabolism Optical Module and associated reagents can provide additional insight and supporting data.

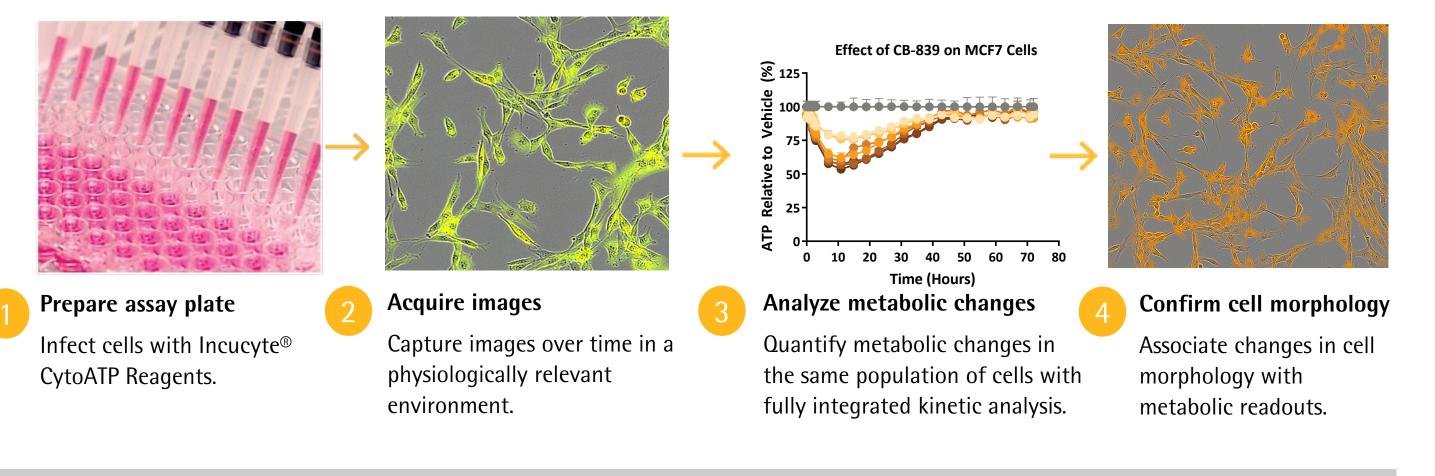
Incucyte[®] Metabolism Optical Module A single, flexible platform designed to combine the analysis of metabolism with additional phenotypic assays

 End-to-end solution: reagents, protocols, plus built-in acquisition and analysis software modules. – Uninterrupted incubation to reduce biological artifacts.

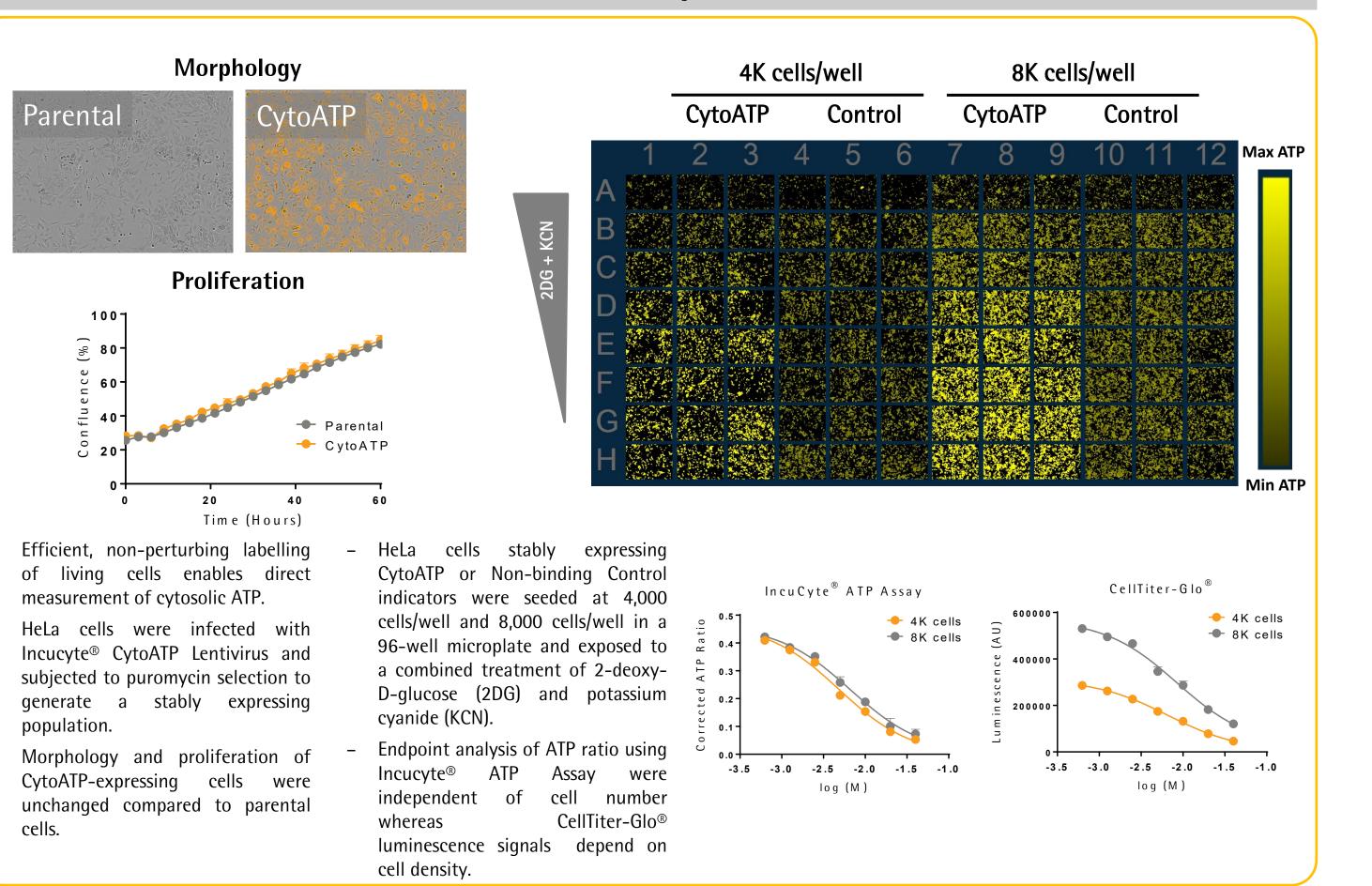
NEW: Metabolism Optical Module

- Incucyte[®] Metabolism Reagents A suite of non-perturbing cell labelling and reporter
- reagents. - CytoATP, CytoATP Non-binding Control, and
- NucLight Orange Reagents
- Mitochondrial Membrane Potential (MMP) Kit
- Annexin V and Caspase 3/7 Cell Health Reagents

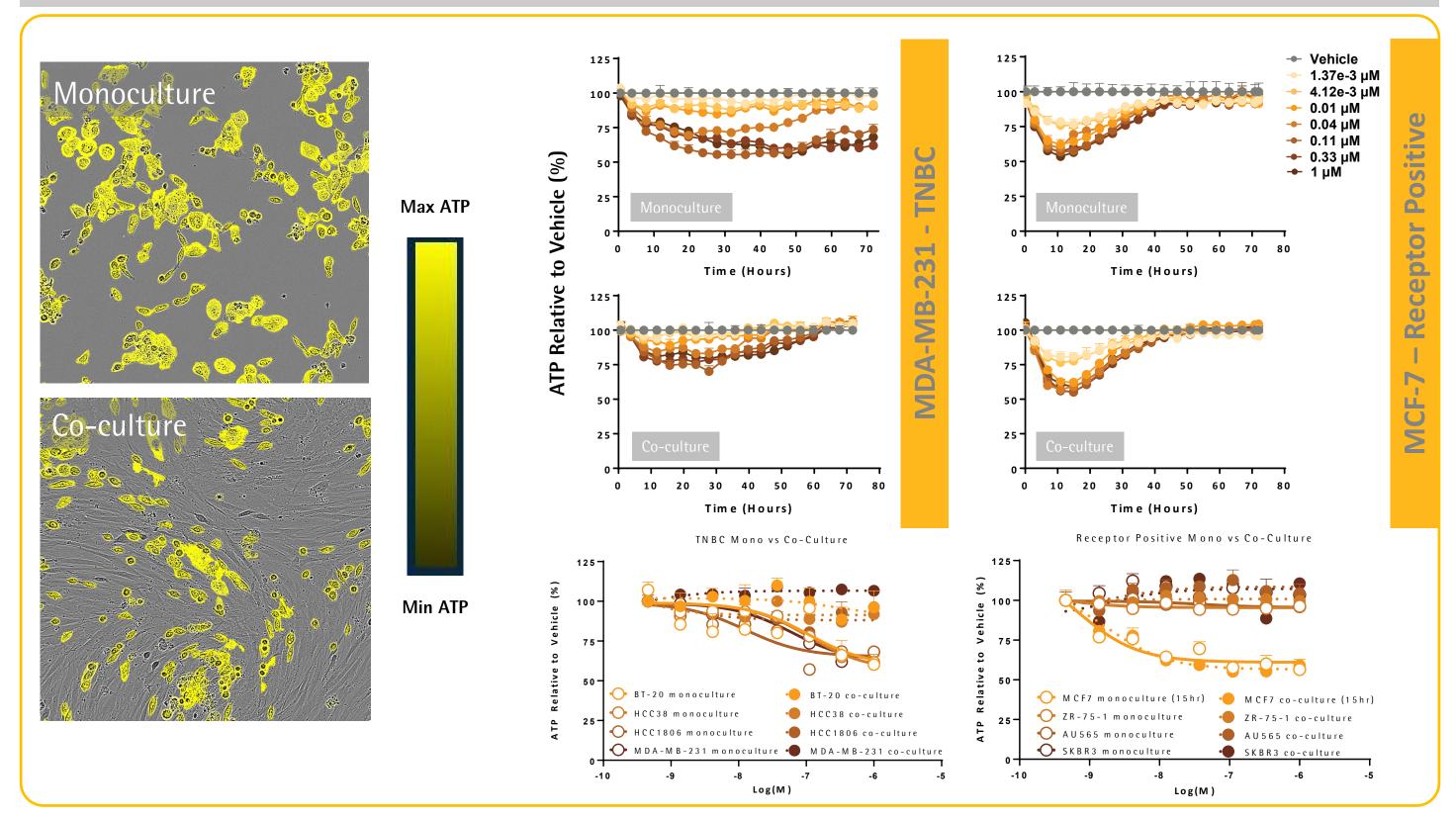
NEW: Incucyte[®] CytoATP and CytoATP Non-binding Control Lentivirus Reagents, Incucyte[®] MMP Orange Kit



Automated measurement of ATP independent of cell number



Cell type-specific ATP measurements in co-culture

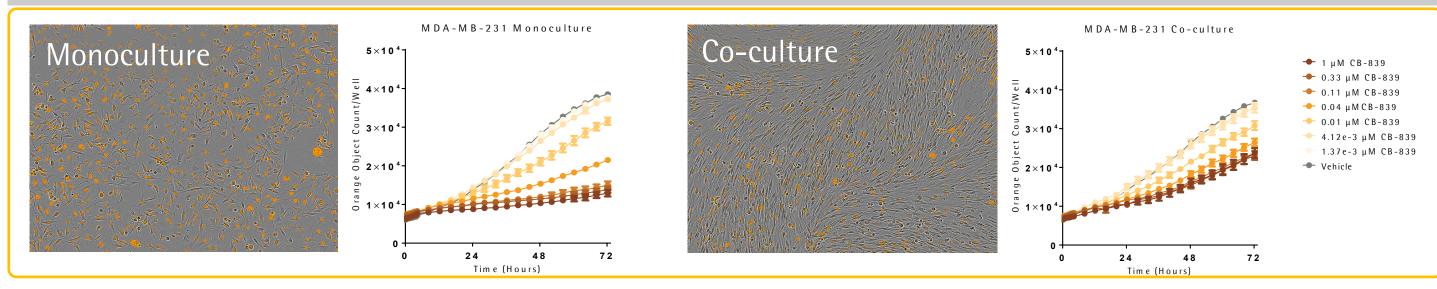


- TNBC or receptor-positive breast cancer cell lines stably expressing CytoATP or Non-binding Control indicators were seeded in the presence and absence of CCD14086SK fibroblasts (8K cells/well) and treated with CB-839 (1.37 nM-1 μM).
- Identification of cellular changes in ATP (color scaled masking) is performed using the integrated Incucyte[®] ATP Analysis Software Module.
- Masked images provide visualization of lower ATP in HCC1806 cells treated with 330 nM CB-839 in monoculture compared to those in coculture.
- Kinetic data shows a sustained depletion of CytoATP in TNBC lines grown in monoculture following CB-839 treatment, whereas receptor-positive

cell lines displayed little to no effect or, in the case of MCF7 cells, a transient decrease followed by full recovery.

- Co-culture with stromal cells mediated resistance to CB-839 in a panel of TNBC cells, while effects on receptor-positive cell lines were unaltered between the two conditions (72 h time points shown except where indicated).
- This effect was not due to drug buffering, as the receptor-positive cell line MCF7 showed overlapping concentration-response curves in both conditions (15 h time point shown to capture transient ATP depletion).

Cell type–specific measurement of proliferation in co-culture



- MDA-MB-231 cells were infected with Incucyte[®] NucLight Orange (NLO) Lentivirus and subjected to puromycin selection to generate a stably expressing population.
- MDA-MB-231 (TNBC) cell line displayed a sustained depletion of ATP (see

ATP

above figure) and robust inhibition of proliferation as measured by orange nuclear count following CB-839 treatment.

- Co-culture of MDA-MB-231 cells with CCD14086SK fibroblasts mitigated

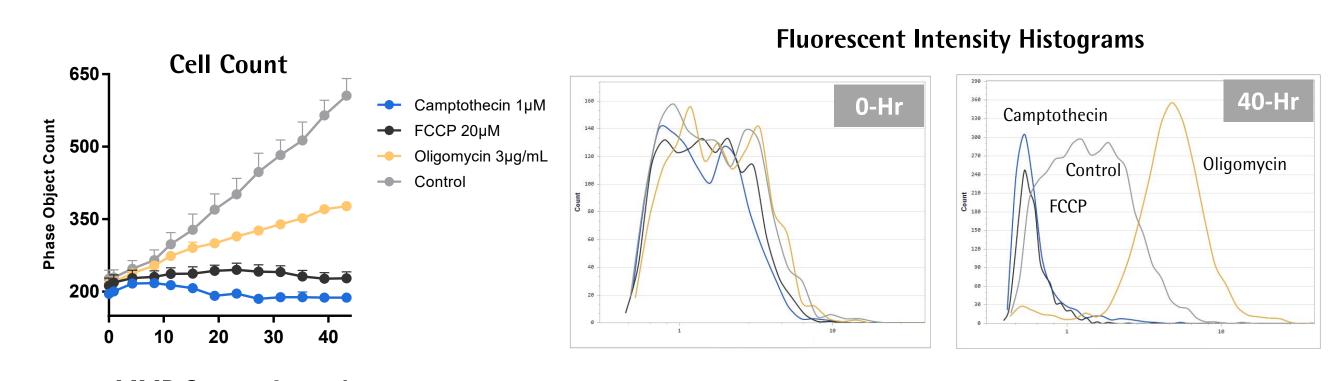
Caspase

ATP depletion and reduced the anti-proliferative effect of CB-839.

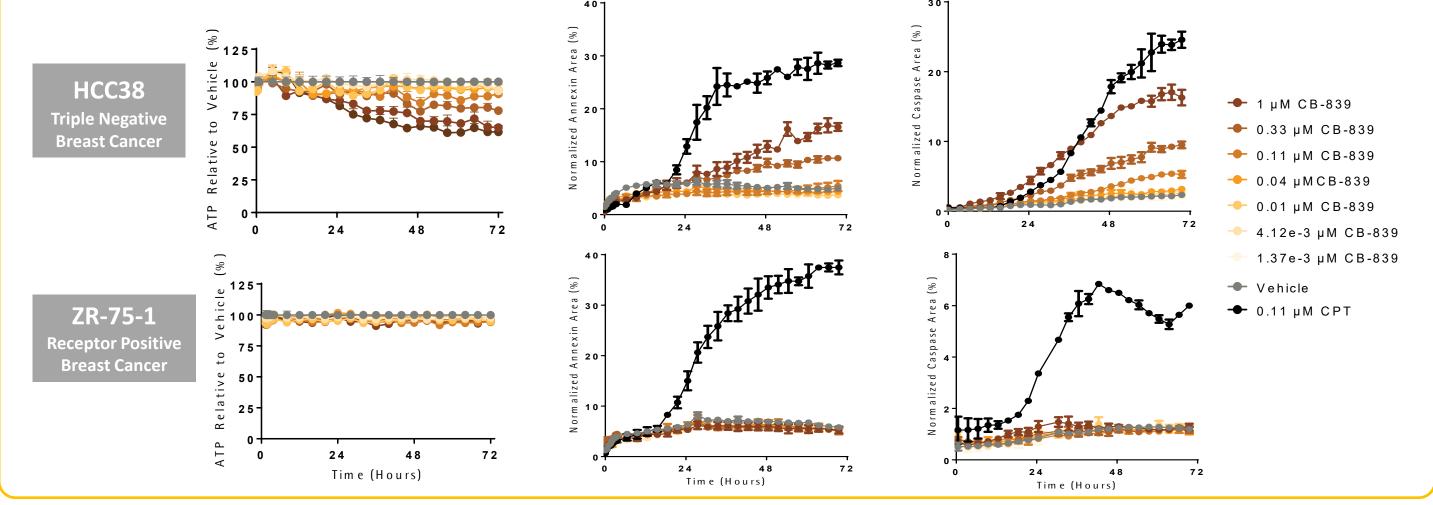
Measure apoptosis with Annexin V and Caspase 3/7 Orange Reagents

Annexin

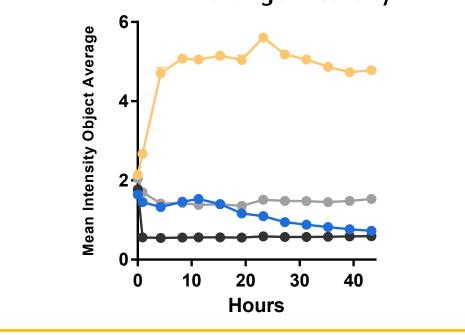
Evaluation of mitochondrial membrane potential using live-cell analysis

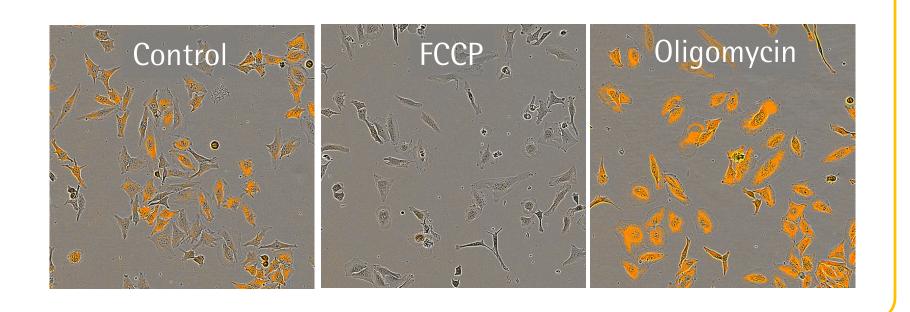


MMP Orange Intensity



- TNBC (HCC38) and receptor-positive (ZR-75-1) breast cancer cell lines stably expressing CytoATP or Non-binding Control indicators show expected ATP response when treated with CB-839 (1.37 nM-1 μ M). Specifically, CB-839 selectively depletes ATP in HCC38 cells (TNBC).
- IncuCyte[®] Annexin V Orange and IncuCyte[®] Caspase-3/7 Orange reagents
- are non-perturbing, mix and read apoptosis reagents that provide supporting cell health data.
- Apoptosis was observed over time selectively in the TNBC cell line treated with CB-839, while only the positive control (Camptothecin, CPT) generated a response in the receptor-positive cell line.





- HeLa cells were treated with the mix-and-read Incucyte[®] MMP Orange Reagent to detect live-cell changes in mitochondrial membrane potential.
- Histograms show mean fluorescent intensity for control and drugtreated wells at 0 and 40hrs. Oligomycin increases fluorescent intensity

(i.e. MMP) while FCCP and Camptothecin show a reduction.

- Proliferation is also measured in parallel by using the Cell-by-Cell Analysis software (i.e. Phase Object Count).
- Images show the MMP Orange Reagent used with control compounds, Oligomycin and FCCP.

Acknowledgments

We thank Hinnah Campwala for providing the NLO figures and John Rauch for generating TMRM data.