

# PhosphoSens® Cell Lysate Activity Assay Format

# JNK Assay Validation Using the AQT1196 Selective Sensor Peptide Substrate

HGNC Name: MAPK8 (JNK1), MAPK9 (JNK2), and MAPK10 (JNK3)

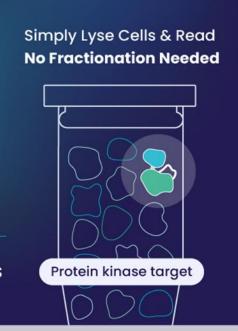
Long Names: c-Jun N-terminal Kinase, Mitogen-Activated Protein Kinase (MAPK)

# PhosphoSens-Lysate

PhosphoSens-Lysate assays are here!

Experience the Benefits of Running Kinase Activity Assays in Complex Samples

Study Kinase Enzyme Activity Where Biology Happens



### **PhosphoSens-Lysate Assays**

transform kinase activity
measurement by enabling
continuous, real-time kinetic analysis
directly in cell and tissue lysates. This
innovative approach provides key
insights into your target's native
environment throughout the drug
discovery pipeline.

NEW PRODUCTS - Available Now for ERK1/2, AKT1/2/3, GSK3 $\alpha$ / $\beta$ , JNK1-3, and DNA-PK

COMING SOON - From our active R&D pipeline of selective sensor peptide substrates for:

p38A/B/G/D, CDK1-3/5, CDK4/6, MEK1/2, SGK1/2/3, and PIM1/3.

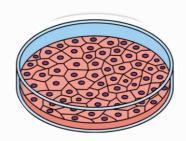
## PhosphoSens-Lysate Assays

### A Simple & Powerful Solution to a Complex Problem



## 1. Grow Cells (+/- Pathway Activation)

or Access Tissues



### 2. Harvest

ADD PhosphoPreserve

Lysis Buffer + Protease & Phosphatase Inhibitors





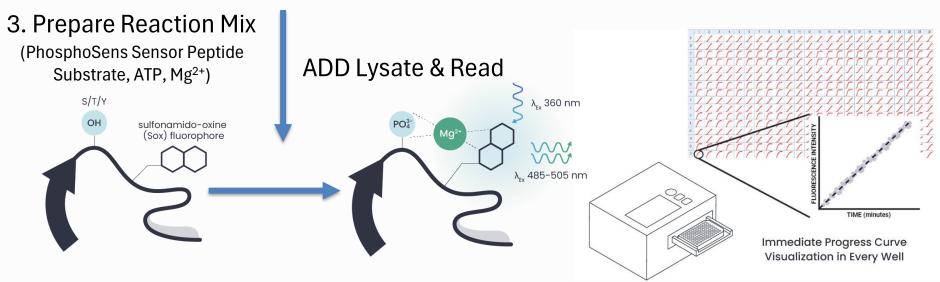
# Study Kinase Enzyme Activity Where Biology Happens

We have combined innovative PhosphoSens detection technology, invented at MIT, with high-throughput peptide synthesis methods to design sensor peptide substrates that are highly selective for the target of interest. This development integrates the advantages of the PhosphoSens platform (activity-based, direct, homogeneous, continuous/kinetic, quantitative) with the enabling capability of measuring endogenous target kinase activity in unfractionated cell or tissue lysates.



Easy-to-use Kits & Bulk Sensor Peptide

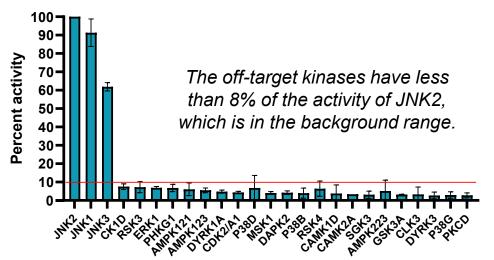
(Native substrates, stable, scalable, excellent lot consistency)



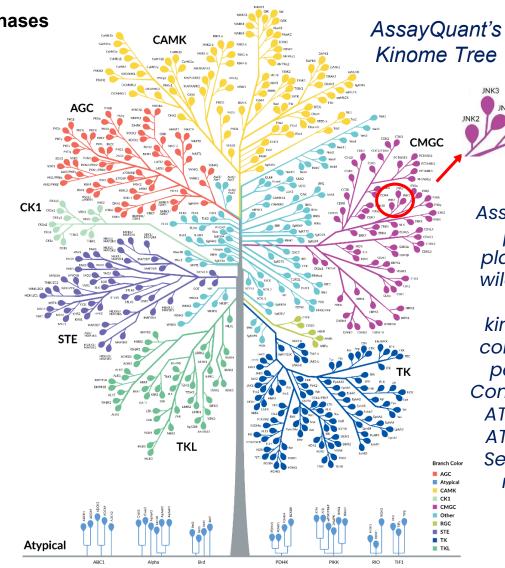
# Selectivity of AQT1196 for JNK1/2/3 Determined with AQT's Kinome Profiling Service



Top 25 Kinases from Kinome Profiling with 411 Wild-type Kinases



#	Kinase target	Kinase Group	AQT1196, μM	Enzyme, nM	RFU/pmol/min	% Activity
1	JNK2	CMGC	10.0	4.0	636.7	100.0
2	JNK1	CMGC	10.0	4.0	546.9	85.9
3	JNK3	CMGC	10.0	5.0	395.2	62.1
4	CK1D	CK1	10.0	1.0	48.5	7.6
5	RSK3	AGC	10.0	1.0	47.2	7.4
6	ERK1	CAMK	10.0	0.4	44.2	6.9
7	PHKG1	CAMK	10.0	1.7	42.4	6.7
8	AMPK121	CAMK	10.0	1.0	37.0	5.8
9	AMPK123	CAMK	10.0	0.2	34.4	5.4
10	DYRK1A	CMGC	10.0	2.0	30.9	4.9
11	CDK2/A1	CMGC	10.0	0.3	28.2	4.4
12	P38D	CMGC	10.0	0.4	26.5	4.2
13	MSK1	AGC	10.0	1.0	26.3	4.1
14	DAPK2	CAMK	10.0	2.4	26.2	4.1
15	P38B	CMGC	10.0	0.8	23.8	3.7



AssayQuant's Kinome profiling service platform features 411 wild-type kinases. All assays are run kinetically to assess compound or sensor peptide selectivity.

Compounds are run at ATP K<sub>m</sub> and/or 1 mM ATP (physiological).

Sensor peptides are run at 1 mM ATP.

## Outline for this Study



### PhosphoSens-Lysate Assay Validation Using the AQT1196 Sensor Peptide Substrate

### **Lysate Source:**

HEK293 cells (passage 17) +/- 3 μM Anisomycin (MedchemExpress, HY-18982; Resuspended in 100% DMSO)

### **Reference Compound Information:**

JNK-IN-8 (JNK Inhibitor XVI - Cayman Chemical, Item No. 18096)

### **Experimental Validation at AssayQuant:**

Lysate from HEK293 cells (+/-  $3 \mu M$  Anisomycin for 40 minutes, as a stress stimulus, to activate JNKs)

Phosphopeptide Control (AQT1232) and comparison to phosphorylation of AQT1196 sensor peptide

with recombinant JNK1 (Thermo Fisher, Part # PV3319)

AQT1196 sensor peptide substrate K<sub>m</sub> determination

**DMSO Tolerance Test** 

Reference Compound IC<sub>50</sub> Determinations with JNK-IN-8.

## Preparation of Cell Lysates from HEK293 Cells Treated +/- Anisomycin to Activate JNKs



- 1) HEK293 cells (passage 18) were plated in a 96-well tissue culture-treated plate and incubated for 48 hours at 37°C in EMEM medium supplemented with 10% FBS (ThermoFisher, A56708-01) and 1% Penicillin-Streptomycin (ThermoFisher, 15140122) in a 5% CO2 atmosphere.
- 2) Cells were washed with PBS, then incubated with culture medium containing 0.1% FBS (serum-starved to make the cells quiescent) and 1% Penicillin-Streptomycin for 24 hours. Subsequently, buffer (negative control) or Anisomycin (MeChemExpress, HY-18982; Resuspended in 100% DMSO) was added directly to the medium to a final concentration of 3 μM, and the cells were incubated for 40 minutes.
- 3) After the treatment, the culture medium was removed, the cells were washed with cold 1X PBS and then lysed with Cell Extraction Buffer (CEB) (see recipe below) for 15 minutes. The assay reaction mix (see next slide) and AQT1196 were added to lysates in each well to assess JNK activity.

### Cell Extraction Buffer (CEB) with Protease and Phosphatase Inhibitors:

- 50 mM HEPES, pH 7.4
- 150 mM NaCl
- 2 mM EGTA
- 1 mM DTT
- 1% Triton X-100

- PhosphoPreserve Phosphatase Inhibitor Cocktail 1, 100X
- PhosphoPreserve Phosphatase Inhibitor Cocktail 3, 60X
- PhosphoPreserve Protease Inhibitor Cocktail, 60X

## Titration of Cell Number and Lysate from HEK293 Cells +/-Anisomycin (3 µM) to Activate JNKs



Growth of HEK293 Cells, Treatment +/- Anisomycin, Cell Extraction, & Determination of JNK Activity with AQT1196 is Performed in the Same 96-well Plate

### Reaction Conditions and Set Up

#### **Reaction Conditions:**

54 mM HEPES, pH 7.5

1 mM ATP

1.2 mM DTT

0.012% Brij-35

1% glycerol

0.2 mg/ml BSA

0.54 mM EGTA

10 mM MgCl<sub>2</sub>

15 μM AQT1196 sensor peptide

#### JNK Enzyme:

- Cell Lysate from titration of HEK293 cells:
  - See below

### **Cell Plating and Protein Determination:**

HEK293 Cells were seeded in a 96-well plate (6 point, 2-fold serial dilution) at 2.5k, 5.0k, 10k, 20k, 40k, and 80k cells/well

Total protein was determined by the Bradford method: 0.1, 0.3, 2.3, 6.1, 16, and 26 µg of protein/well

### **Reaction Set Up:**

- •To **20 μL of Cell Lysate** in the 96-well plate
- •Add 78 µL Reaction Mix with AQT1196, ATP & DTT
- •Add **2 µL of 50X tool compound** in DMSO or DMSO control
- •To obtain 100 µL Final reaction volume

#### Notes:

Reaction was run at 30  $^{\circ}$ C for 240 minutes in Falcon $^{\otimes}$  96-well White Flat Bottom TC-treated Microplate (Corning Cat. #353296) at 100  $\mu$ L final well volume after sealing using optically-clear adhesive film (TopSealA-Plus plate seal, PerkinElmer [Cat. #6050185]) in a Biotek Synergy Neo 2 microplate reader with excitation (360 nm) and emission (485 nm) wavelengths.

1X Enzyme Dilution Buffer (EDB) is used to dilute enzyme and for the blank. Composition is 20 mM HEPES, pH 7.5, 0.01% Brij-35, 5% Glycerol, 0.5 mM EGTA, 1 mM DTT, 1 mg/ml Bovine Serum Albumin.

# Recombinant JNK1 Activity with AQT1196 and Comparison to AQT1232 Phospho-Control



Reaction Conditions and Set Up

### **Reaction Conditions:**

54 mM HEPES, pH 7.5

1 mM ATP

1.2 mM DTT

0.012% Brij-35

1% glycerol

0.2 mg/ml BSA

0.54 mM EGTA

10 mM MgCl<sub>2</sub>

#### 15 μM **Sensor peptides:**

- AQT1196 sensor peptide substrate
- AQT1232 phosphopeptide control

5 nM JNK1, full-length recombinant (Thermo Fisher, Part # PV3319)

### **Reaction Set Up:**

20  $\mu$ L Reaction Mix with AQT1196 or AQT1232, ATP, DTT 15 minutes incubation at 30°C (in the reader) 5  $\mu$ L of 5x enzyme or EDB with lysate buffer 25  $\mu$ L Final reaction volume

Reaction was run at 30 °C for 240 minutes in either Corning, low volume 384-well, white flat-bottom polystyrene NBS microplates (Cat. #3824) at 20 or 25  $\mu$ L final well volume or in in PerkinElmer, ProxiPlate-384 Plus, white shallow well microplates (Cat. #6008280) at 20  $\mu$ L final well volume after sealing using optically- clear adhesive film (TopSealA-Plus plate seal, PerkinElmer [Cat. #6050185]) in a Biotek Synergy Neo 2 microplate reader with excitation (360 nm) and emission (485 nm) wavelengths.

#### **Notes:**

Enzyme Dilution Buffer (EDB): 20 mM HEPES, pH 7.5, 0.01% Brij-35, 5% Glycerol, 0.5 mM EGTA, 1 mM DTT, 1 mg/ml Bovine Serum Albumin.

## Titration of Cell Number and Lysate from HEK293 Cells +Anisomycin (3 µM) to Activate JNKs

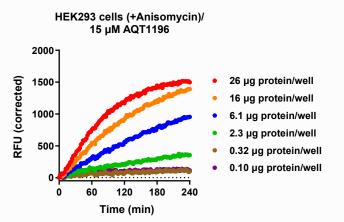


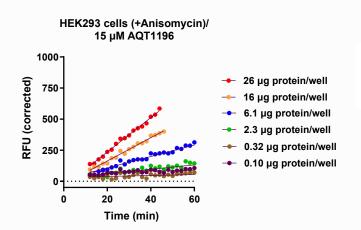
Progress Curves, Reaction Rates & Linearity for JNK Activity with AQT1196

### **Progress Curves**

Complete Time Course

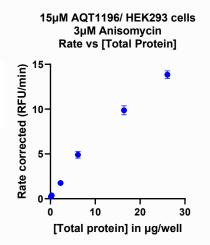
Linear Region



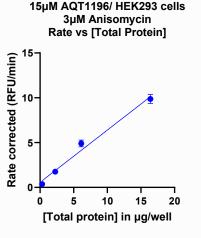


### <u>Assessment of Linearity</u>

All Data



**Linear Reaction** 



The JNK lysate activity assay is linear from 0.3 – 16 µg/well of crude protein (53-fold range), corresponding to 5k – 16k HEK293 cells seeded/well.

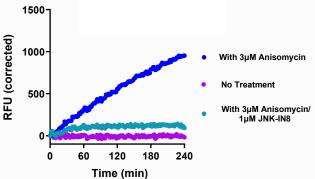
## JNK Lysate Activity Assay with AQT1196



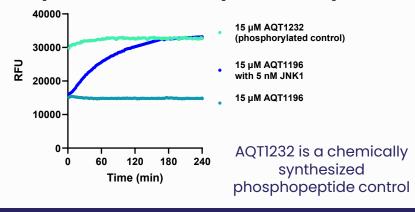


### A. Crude Lysate Samples: 20k HEK293 cells (or 6.1 µg protein)/well with 15 µM AQT1196

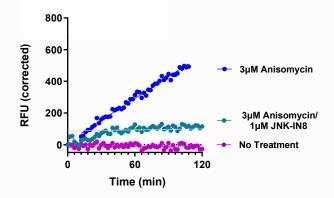
### 1) Full Time Course (0-240 min.)



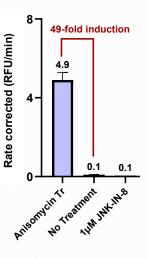
### **B. Purified JNK1 & AQT 1232 Control** 1) Full Time Course (0-240 min.)







### 3) Histogram



A. Crude lysate samples: The AQT1196 sensor peptide was used to generate RFU Corrected values (Total – Background) for 1) Full progress curve time course (0-240 min.), and 2) Linear range (4-120 min.), and to determine the slope for each condition, which is the Reaction rate (RFU Corrected/min. +/- standard deviations) shown as a histogram in 3), highlighting a 49-fold activation of JNK kinase activity in HEK293 cell lysates treated with 3µM Anisomycin for 40 minutes. The signal was eliminated by adding the selective JNK1/2/3 inhibitor JNK-IN-8 (1 μM). The amount of activation depends on several factors, including cell type, serum concentration, and duration of the pre-incubation to make cells quiescent, and the activating stimulus's nature, concentration, and duration. These conditions can be varied to optimize JNK activity. The total amount of JNK1/2/3 protein can be determined by Western Blotting or an ELISA; however, with the short stimulation times typically used, these levels are not expected to change.

B.1. Purified recombinant JNK1 enzyme & AQT1107 Control: JNK1 enzyme (5 nM) fully phosphorylated the AQT1196 sensor peptide substrate by 240 min., as shown by convergence with the signal obtained with the AQT1232 phosphopeptide positive control (a flat horizontal line defining the maximum RFU with this sensor peptide). The signal with JNK1 enzyme was eliminated by adding the JNK-IN-8 inhibitor (1 µM). The signal with AQT1232 is used to convert RFU (Corrected) values to nmoles of product.

# Sensor Peptide K<sub>m</sub> Determination for AQT1196 Using Lysate from HEK293 Cells (20k/well) + Anisomycin



### Reaction Conditions and Set Up

### **Reaction Conditions:**

54 mM HEPES, pH 7.5

1 mM ATP

1.2 mM DTT

0.012% Brij-35

1% glycerol

0.2 mg/ml BSA

0.54 mM EGTA

10 mM MgCl<sub>2</sub>

AQT1196 sensor peptide titration - 0, 0.20, 0.39, 0.78, 1.6, 3.1, 6.3, 13, 25, 50, and 100 μM.

### **Cell Plating and Protein Determination:**

HEK293 Cells were seeded at 20k cells/well in 24 wells (12 wells in duplicate) of the 96-well plate

Total protein was determined by the Bradford method: 6.1 µg of protein/well

### **Reaction Set Up:**

- To **20 μL Lysate** in the 96-well plate,
- Add 70 µL Reaction Mix with ATP, &DTT
- Add 10 μL of 10X AQT1196
- to obtain **100 μL** Final reaction volume

#### **Notes:**

Reaction was run at 30  $^{\circ}$ C for 240 minutes in Falcon $^{\otimes}$  96-well White Flat Bottom TC-treated Microplate (Corning Cat. #353296) at 100  $\mu$ L final well volume after sealing using optically-clear adhesive film (TopSealA-Plus plate seal, PerkinElmer [Cat. #6050185]) in a Biotek Synergy Neo 2 microplate reader with excitation (360 nm) and emission (485 nm) wavelengths.

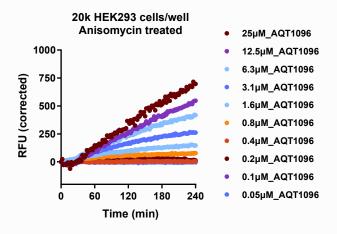
1X Enzyme Dilution Buffer (EDB) is used to dilute enzyme and for the blank. Composition is 20 mM HEPES, pH 7.5, 0.01% Brij-35, 5% Glycerol, 0.5 mM EGTA, 1 mM DTT, 1 mg/ml Bovine Serum Albumin.

# Sensor Peptide K<sub>m</sub> Determination with AQT1196 Using Lysate from HEK293 Cells (20k/well) + Anisomycin

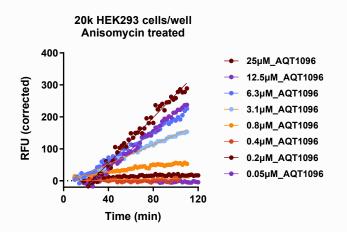


Titration Curves and K<sub>m</sub> Plot and Table

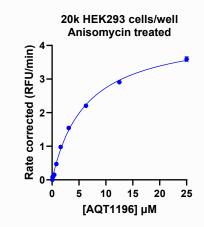
# Sensor Peptide progress Curves



### Sensor Peptide Linear range



### Sensor Peptide K<sub>m</sub> Plot



Michaelis-Menten	
Best-fit values	
Vmax	4.439
Km	6.206
Std. Error	
Vmax	0.1175
Km	0.4234
95% CI (asymptotic)	
Vmax	4.168 to 4.710
Km	5.229 to 7.182

The  $K_m$  value for AQT1196 is 6.2  $\mu$ M.

# DMSO Tolerance Test Using Lysate from HEK293 Cells (20k/well) + Anisomycin and AQT1196



#### **Reaction Conditions:**

54 mM HEPES, pH 7.5

1 mM ATP

1.2 mM DTT

0.012% Brij-35

1% glycerol

0.2 mg/ml BSA

0.54 mM EGTA

10 mM MgCl<sub>2</sub>

 $15 \mu M$  AQT1196 sensor peptide

### **Cell Plating and Protein Determination:**

HEK293 Cells were seeded at 20k cells/well in 24 wells (12 wells in duplicate) of the 96-well plate

Total protein was determined by the Bradford method: 6.1 µg of protein/well

### **Reaction Set Up:**

- To 20 μL Lysate in the 96-well plate,
- Add **70 μL Reaction Mix with AQT1196, ATP, &DTT**
- Add **10 μL of 10X DMSO**
- to obtain **100 μL** Final reaction volume

### Notes:

Reaction was run at 30  $^{\circ}$ C for 240 minutes in Falcon $^{\otimes}$  96-well White Flat Bottom TC-treated Microplate (Corning Cat. #353296) at 100  $\mu$ L final well volume after sealing using optically-clear adhesive film (TopSealA-Plus plate seal, PerkinElmer [Cat. #6050185]) in a Biotek Synergy Neo 2 microplate reader with excitation (360 nm) and emission (485 nm) wavelengths.

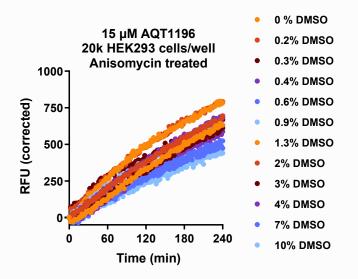
1X Enzyme Dilution Buffer (EDB) is used to dilute enzyme and for the blank. Composition is 20 mM HEPES, pH 7.5, 0.01% Brij-35, 5% Glycerol, 0.5 mM EGTA, 1 mM DTT, 1 mg/ml Bovine Serum Albumin.

# DMSO Tolerance Test Using Lysate from HEK293 Cells (20k/well) + Anisomycin and AQT1196

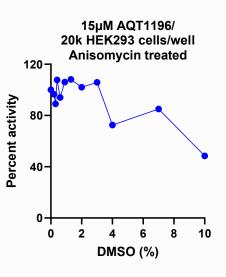


Titration Curves and Inhibition Plot

Complete Progress Curves



# Reaction Rate vs [DMSO] Plot



No significant loss in JNK activity was observed up to 3% DMSO. A final concentration of 2% DMSO will be used to assess inhibitor potency.

# IC<sub>50</sub> Determination using AQT1196 in 96-well plates with Lysates from HEK293 Cells (20k/well)



### Reaction Conditions and Set Up

### **Reaction Conditions:**

54 mM HEPES, pH 7.5

1 mM ATP

1.2 mM DTT

0.012% Brij-35

1% glycerol

0.2 mg/ml BSA

0.54 mM EGTA

10 mM MgCl<sub>2</sub>

15 μM AQT1196 sensor peptide

**Tool Compound** - JNK-IN-8 was titrated with 3-fold dilutions in 100% DMSO at 50X the final concentrations and then diluted 50-fold into the assay for final concentrations from 0-10  $\mu$ M in 2% DMSO.

### **Reaction Set Up:**

- To 20 μL Lysate in the 96-well plate,
- Add 78 μL Reaction Mix with AQT1196, ATP, &DTT
- Add 2 μL of 50X serially diluted JNKJ-IN-8 in 100% DMSO
- to obtain **100 μL** Final reaction volume

#### Notes:

Reaction was run at 30  $^{\circ}$ C for 240 minutes in Falcon $^{\otimes}$  96-well White Flat Bottom TC-treated Microplate (Corning Cat. #353296) at 100  $\mu$ L final well volume after sealing using optically-clear adhesive film (TopSealA-Plus plate seal, PerkinElmer [Cat. #6050185]) in a Biotek Synergy Neo 2 microplate reader with excitation (360 nm) and emission (485 nm) wavelengths.

1X Enzyme Dilution Buffer (EDB) is used to dilute enzyme and for the blank. Composition is 20 mM HEPES, pH 7.5, 0.01% Brij-35, 5% Glycerol, 0.5 mM EGTA, 1 mM DTT, 1 mg/ml Bovine Serum Albumin.

### **Cell Plating and Protein Determination:**

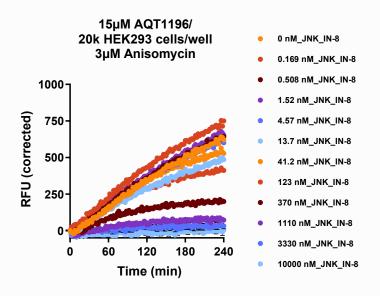
HEK293 Cells were seeded at 20k cells/well in 24 wells (12 wells in duplicate) of the 96-well plate

Total protein was determined by the Bradford method: 6.1 μg of protein/well

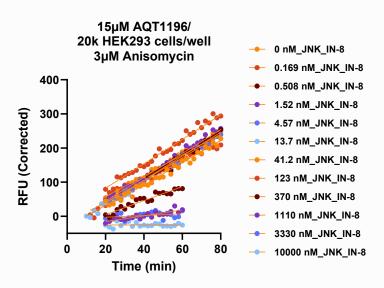
# IC<sub>50</sub> Determination using AQT1196 in 96-well plates with Lysates from HEK293 Cells (20k/well)



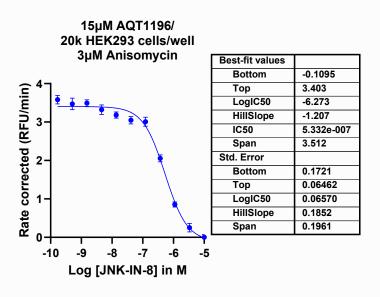
### **Full Progress Curves**



# Progress Curves (Linear Region)



### IC<sub>50</sub> Curve



The IC<sub>50</sub> value for JNK-IN-8 in lysate from anisomycin-treated HEK293 cells is 533 nM, which is close to the value of 290 nM reported by Peterson et al. (Biochemistry 2014, 53, 5771-5778) with HeLa cell lysates

## Summary



15µM AQT1196/

The PhosphoSens-Lysate Activity Assay for JNK1-3 using the selective sensor peptide AQT1196 demonstrates a robust and physiologically relevant assay that provides a functional assessment of endogenous JNK activity with all the cellular components and signaling complexes. This JNK activity assay is direct, highly quantitative, and in an easy-to-use format.

### Results include:

- Anisomycin treatment resulted in vigorous activation of JNK activity in HEK293 (49-fold) and HeLa (20-fold) cells.
- JNK activity with lysates from Anisomycin-treated HEK293 cells was linear from 0.3 to 16 μg protein/well, a 53-fold linear range.
- The Sensor peptide substrate AQT1196 has a  $K_m$  of 6.5  $\mu$ M when tested with Anisomycintreated HEK293 cell lysates.
- 97% of AQT1196 phosphorylation in lysates from Anisomycin-treated HEK293 cells is inhibited by the reference compound for JNK (JNK-IN-8, 1 μM).
- The IC<sub>50</sub> value for JNK-IN-8 in lysates from anisomycin-treated HEK293 cells with AQT1196 was 533 nM.

20k cells/well(6 µg protein/well) 20-fold induction RFU (Corrected)/min 0.1 0.1

AQT1196 enables selective and precise quantitation of JNK1/2/3 activity with different cell types (demonstrated for HEK293 and HeLa cells), providing a powerful tool for evaluating pathway activation and inhibition in complex samples from normal and disease states



### **Company & Technology Supporting Slides**



### **AQT Senior Scientific Team**

### **Experienced & Diverse Team from Leading Institutions**





Dr. Bill Radany CEO & Head of **Business Development** 



Dr. Erik Schaefer Co-Founder, CSO



Dr. Barbara **Imperiali** Co-Founder & CTO, MIT Faculty



Dr. Earl May Senior Director, Discovery Technologies (DT)



Dr. Eric Berg Director of Peptide Operations



Dr. Daniel Urul Senior Scientist, DT



Dr. Venky Nemmara Senior Scientist, DT

- We apply our >200 years of collective experience in kinase biology, drug or assay development, chemical-biology and automation to solve critical problems in target biology and drug development.
- AQT services allow us to evaluate drugs in diverse ways, delivering high-quality data combined with strong communication to enable discovery



Susan Cornell-Kennon Senior Scientist, DT



Dr. Satish **Pimrale** Business **Development Leader** 



















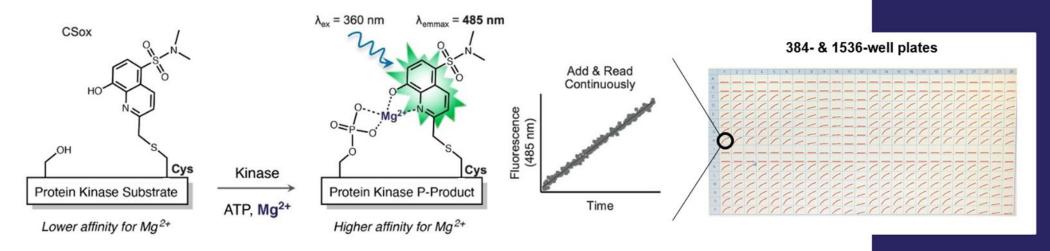






# Sensor Peptide Substrates for Continuous (Kinetic) Monitoring of Protein Kinase Activity





- Uses chelation-enhanced fluorescence via sulfonamido-oxine (Sox) chromophore, invented by Barbara Imperiali (MIT).
- 10 patents with 1 pending, exclusively licensed from MIT. Rich trade secret portfolio
- Sox is small, minimally hydrophobic, and neutral
- Assay is direct, homogeneous, and kinetic (continuous)
- Can use a wide range of ATP concentrations (Low μM, ATP K<sub>m</sub> or mM [physiological])
- Runs on commonly available readers with any plate type (96-, 384- or 1536-well)

Simple, Powerful & Flexible!

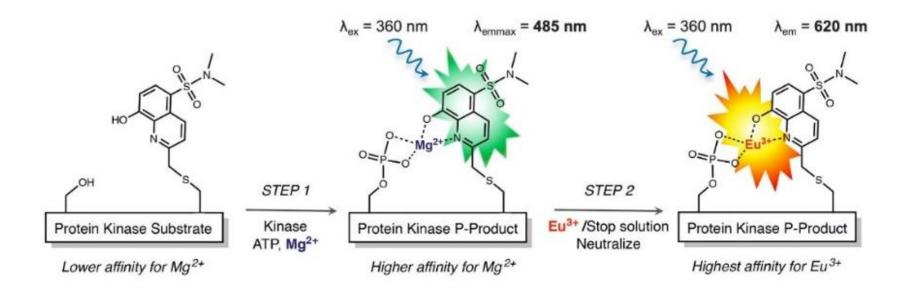
# a full progress curve in every well

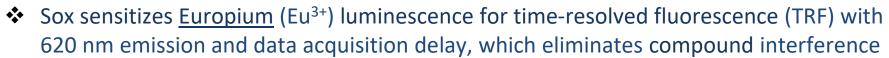
Ideal for quantitative assessments including Profiling, Potency & MOA



# Replacing Mg<sup>2+</sup> with Eu<sup>3+</sup> Create PhosphoSens®- Red







- Run as an Endpoint format; ideal for high-numbers of tests for HTS and SAR
- Patent approved in 2020

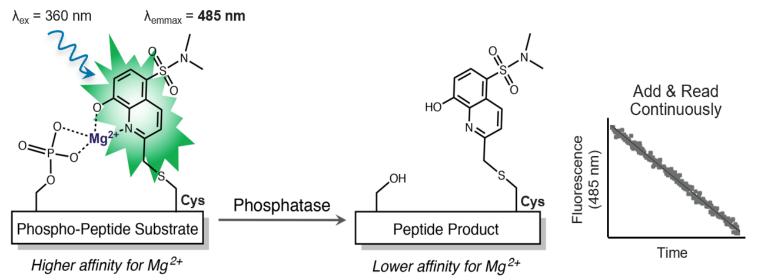
Eliminates autofluorescence while using the same small fluorophore



### **Ideal for:**

# Same Sox-based Detection Platform Enables Continuous Monitoring of **Protein**





- Same mechanism ChEF (Mg<sup>2+</sup>) for kinetic assays and Eu<sup>3+</sup>/TRF for Endpoint (Red), as for protein kinases
- Uses <u>CSox-based phosphopeptide substrates</u> derived from physiological targets to monitor loss of signal
- Covered by 10 patents (with 1 pending) and trade secret portfolio

*Increasing requests for tyrosine & serine/threonine phosphatases* 



Kinetic and Endpoint

One Technology,
Two Fit-for-Purpose
Formats

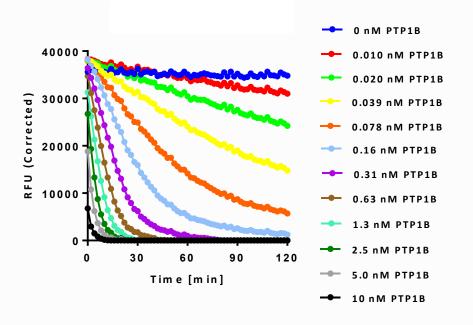
Expanding menu (currently 34 of the 190 Protein Phosphatases)

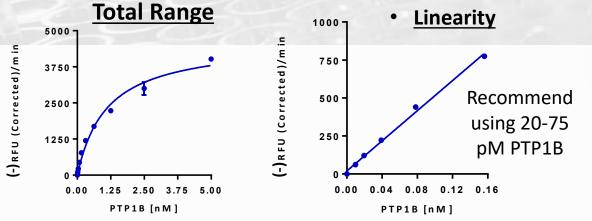
## Analysis of PTP1B & PTPN2 Tyrosine Phosphatases

Optimized *PhosphoSens* Phosphopeptide Substrates

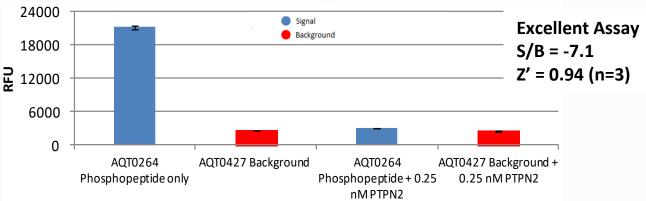


# Progress Curves & Linearity for PTP1B with AQT0266 Sensor Peptide, Net Signal





### Total & Background for PTPN2 with AQT0264 in PhosphoSens®-Red

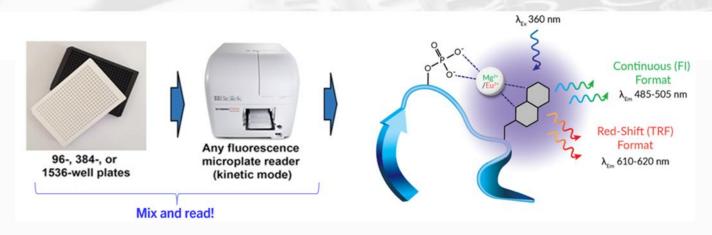


High-sensitivity & physiologically-relevant. Analysis of allosteric or substrate-competitive inhibitors

## PhosphoSens® Platform

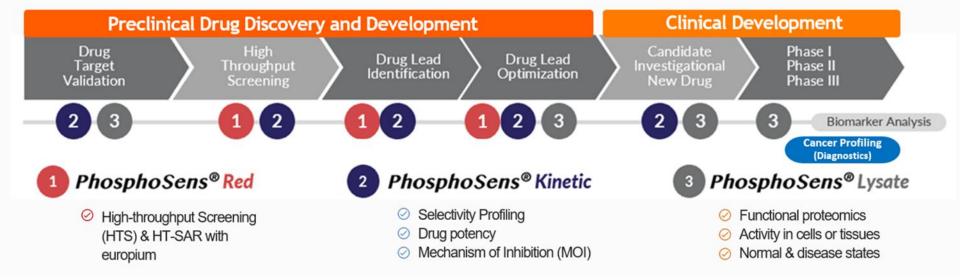
### Enabling Analysis Across the Drug Development Workflow





One Technology, Two Fit-for-Purpose Formats

Kinetic and Endpoint

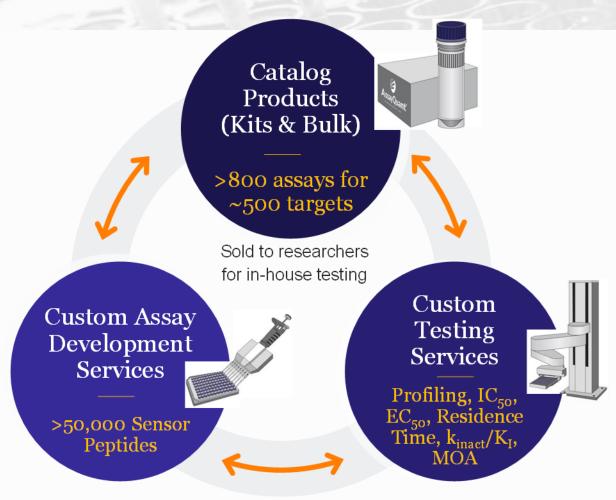


Accelerates progress and improves outcomes

### Integration of Catalog Products & Services

Multiple entry & transition points to address your needs





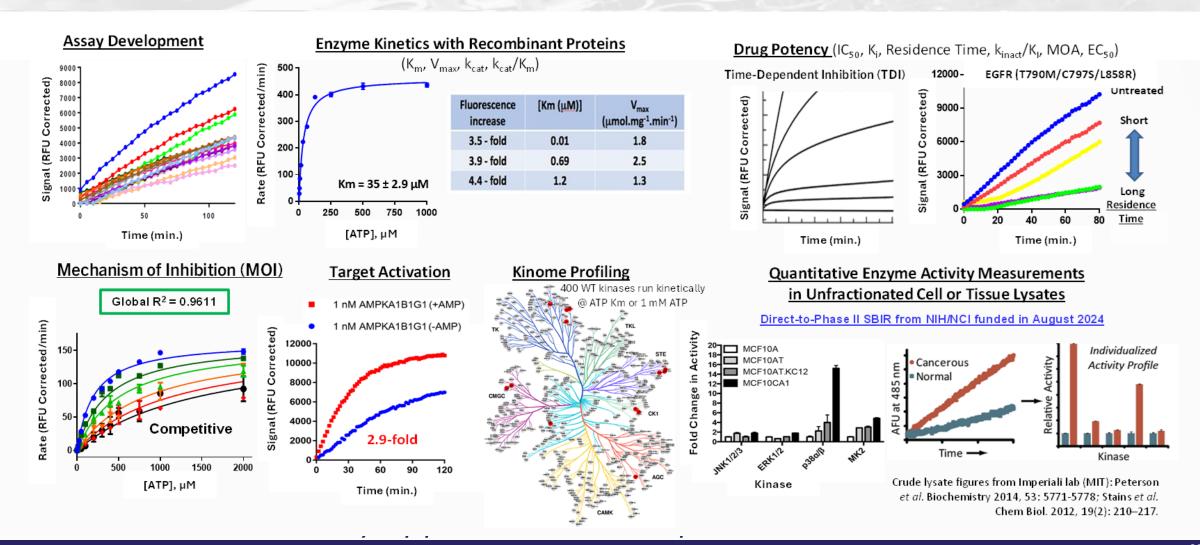


AQT creates custom PhosphoSens assays and AQT retains IP

Drug developers send compounds to AQT to test and tap our expertise

# *PhosphoSens®* Platform Continuous Workflow Applications





## PhosphoSens® Platform Advantages:



- Sensitive and continuous (kinetic) format based on Chelation-enhanced fluorescence.
- Homogeneous (add & read) and direct measure of enzyme activity resulting in a complete progress curve in every well for each condition.
- Sensor peptides or lipids based on <a href="mailto:physiologically-relevant">physiologically-relevant</a> substrates.
- Run under <u>optimal enzyme conditions</u>, with physiological  $Mg^{2+}$ ,  $Mn^{2+}$  &  $Ca^{2+}$  ions and <u>any ATP</u> concentration (1-2 mM = physiological; or at ATP  $K_m$ ) or sample type (lysates via selective sensor peptides or IP-kinase assays).
- Determination of <u>initial reaction rate</u> from <u>linear</u> portion of curve provides <u>high accuracy & precision</u> (Z'>0.7) and enables assessment of time-dependent inhibitor or activator potency ( $IC_{50}$ ,  $EC_{50}$ ,  $k_{inact}/K_{I}$ , residence time).
- \* Corrects for any compound autofluorescence Kinetic: Background signal doesn't change over time; Red: Uses Europium (Eu<sup>3+</sup>) for time-resolved fluorescence (TRF) with a 100 μsec delay & 620 nm emission.
- \* Rapid & predictable development of new assays is achieved even with difficult targets using >30,000 sensor peptide panel for protein kinases or phosphatases and any DAG species for DGK assays.
- ❖ Rigorous manufacturing process for PhosphoSens® sensors provides excellent lot to lot consistency.
- Works across entire drug development workflow Improving efficiency and performance

We apply our >200 years of collective experience in kinase biology, drug or assay development, and automation to address complex biology and evaluate drugs in diverse ways, delivering high-quality data to accelerate your programs.