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## **APPLICATION NOTE AND PROTOCOL**

Method	Enzyme Linked Immunosorbent Assay (ELISA)		
Detection Range	$5-640$ pM for $\gamma$ H2AX (pS139-H2AX) and		
	50 – 6400 pM for Total H2AX		
Sample Type	18-gauge needle biopsies, xenograft tumor quadrants, and lysates of cultured cells or PBMCs		
Sample Size	$25 \ \mu L$ lysate/75 $\mu L$ final well volume		

This Research Use Only Document has not been assessed via clinical assay performance standards and is intended for preclinical use. Related SOPs and documentation can be found on the DCTD Biomarkers site at: <a href="http://dctd.cancer.gov/ResearchResources/ResearchResources-biomarkers.htm">http://dctd.cancer.gov/ResearchResources/Res

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#### **1.0 INTENDED USE**

RESEARCH USE ONLY (RUO) Protocol. This SOP is supplied as an RUO protocol for the  $\gamma$ H2AX (gamma or phospho-Serine 139 H2AX) and Total H2AX assays for preclinical development laboratory use. Provided here are representative standard curves, performance characteristics, procedures for the assays, preparation of assay controls, detailed material/supply lists, and recommended assay quality control. This RUO SOP does not have an associated formal training program or qualified reagent supply and has not been assessed via clinical performance standards.

#### 2.0 INTRODUCTION

H2AX is a histone H2A variant that constitutes 2–25% of mammalian histone H2A depending on the organism and cell type. Like most other histone proteins, H2AX is composed of a central globular domain, flanked by N-terminal and C-terminal tails which possess sites for a variety of post-translational modifications such as acetylation, biotinylation, phosphorylation, methylation, and ubiquitination. H2AX is structurally similar to other H2A species except for the presence of a unique COOH terminal tail, containing a serine four residues from the C terminus<sup>1</sup>. Upon induction of a DNA double-strand break (DSB), the H2AX omega-4 serine residue becomes rapidly phosphorylated to form gamma-H2AX. Phosphorylated histone H2AX ( $\gamma$ H2AX) proves to be an early marker for DNA double-strand breaks, and signal levels directly correlate with the number of breaks formed. Therefore,  $\gamma$ H2AX can serve as a pharmacodynamic (PD) marker to measure the chemotoxic effect of potential DNA damaging agents. For this purpose, 96-well ELISAs for quantifying  $\gamma$ H2AX and Total H2AX have been developed and are described in this RUO SOP<sup>2-4</sup>.

For the  $\gamma$ H2AX ELISA,  $\gamma$ H2AX is captured from total cell extracts on plates coated with a  $\gamma$ H2AX capture monoclonal antibody. The captured protein is then detected using a H2AX polyclonal detection antibody followed by a HRP-conjugate to allow chemiluminescent readout and quantitation of  $\gamma$ H2AX levels. The assay for quantifying Total H2AX is based on the same principle using a monoclonal and polyclonal antibody to different epitopes for capture and detection. Using Total H2AX as denominator for  $\gamma$ H2AX reporting provides more reliable quantitation of the fraction that is phosphorylated H2AX ( $\gamma$ H2AX) and provides a more relevant PD readout for monitoring treatment induced DNA damage.

#### 3.0 CONSIDERATIONS

- 3.1 Unknown samples are processed following the validated PAR Immunoassay biopsy extraction SOP340520 (http://dctd.cancer.gov/ResearchResources/ResearchResources-biomarkers.htm)<sup>5</sup>.
- 3.2 These assays were developed with the following specific lot numbers for key reagents. Performance with new lots will need to be assessed by individual laboratories to meet expected performance criteria.

#### *YH2AX Assay*

- 3.2.1 Anti-phospho-H2AX (Ser139) mouse monoclonal antibody, clone JBW301(Millipore, Cat#: 05-636, Lot# DAM1567248) was used at 1:250 dilution (1 μg/μL) for assay development. Other lots used at 1:250 dilutions (1 μg/μL) during development were: Lot# JBC1367868 and Lot# DAM1493341.
- 3.2.2 Histone H2AX rabbit polyclonal antibody (Abcam, Cat# ab10475, Lot# 778559) was used at 1:850 dilution (1.7 μg/μL) for assay development.





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- 3.2.3 Goat anti-rabbit HRP-conjugated polyclonal antibody (KPL, Cat# 074-15-061, Lot# 101008) was used at 1:1000 dilution (1  $\mu$ g/ $\mu$ L) for assay development.
- 3.2.4  $\gamma$ H2AX peptide standard, a lyophilized powder (custom-preparation from Invitrogen, synthetic peptide: AVLLPKKTSATVGPKAPSGGKKATQA[PS]QEY) yH2AX JJ2, Lot# 113011 was used at a standard stock concentration of 20680 pM for assay development.

#### **Total H2AX Assay:**

- 3.2.5 HIST1H2AC monoclonal antibody 4F10 (Novus, Cat#: H00008334-M01, Lot# D8231-4F10) was used at 1:250 dilution (1  $\mu$ g/ $\mu$ L) for assay development.
- Histone H2AX rabbit polyclonal antibody (Abcam, Cat#: ab10475, Lot# 778559) was 3.2.6 used at 1:850 dilution (1.7  $\mu$ g/ $\mu$ L) for assay development.
- 3.2.7 Goat anti-rabbit HRP-conjugated polyclonal antibody (KPL, Cat#: 074-15-061, Lot# 120504) was used at 1:1000 dilution (1  $\mu$ g/ $\mu$ L) for assay development.
- 3.2.8 Total H2AX recombinant protein standard, a lyophilized powder (Axxora, Cat#: ALX-201-176-M005, Lot# L16677) was used at a standard stock concentration of 33000 pM for assay development.

#### ASSAY DEVELOPMENT DESCRIPTION 4.0

#### 4.1 yH2AX Standard Curve

Following the YH2AX assay protocol described in this RUO SOP, the RLU (relative light unit) values from the synthetic peptide standards were used to generate a standard curve with an Infinite M200 Pro Microplate Reader. An example of a typical yH2AX standard curve is shown below (Figure 1).







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**Figure 1**. Standard curve for the  $\gamma$ H2AX ELISA. This standard curve is for the **<u>purpose of</u> <u>illustration only</u>**, and should not be used to calculate unknowns. Each site should generate its own standard curve and data.

#### 4.2 Total H2AX Standard Curve

Following the Total H2AX assay protocol described in this RUO SOP, the RLU (relative light unit) values from the synthetic peptide standards were used to generate a standard curve with an Infinite M200 Pro Microplate Reader. An example of a typical total H2AX standard curve is shown below (Figure 2).



**Figure 2**. Standard curve for the Total H2AX ELISA. This standard curve is for the **<u>purpose of</u> <u>illustration only</u>**, and should not be used to calculate unknowns. Each site should generate its own standard curve and data.

#### 4.3 γH2AX Assay Performance

The inter-assay performances of the  $\gamma$ H2AX assay was evaluated using the defined critical reagent lot numbers in Section 3.2.

**Table 1**. Inter-assay Precision of the  $\gamma$ H2AX assay was measured by 3 runs of  $\gamma$ H2AX standards and assay controls. All 3 runs passed the inter-assay acceptance criteria (%CV <25%) with a range from 1 to 12% for the standards and 1.9 to 6.0% for the controls.





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	γH2AX Standar	d (pM)	Average RI	LU (n=3)	SD	%CV
	5		9926	67	122261	12
	10		20168	350	134112	7
	20		45986	500	277146	6
	40		10021	317	467893	5
	80		20811	817	1459080	7
	160		37371	983	930881	2
	320		58789	817	2240098	4
	640		79191	483	1063448	1
	Assay Contr	ol	Average p	M (n=3)	SD	%CV
	Low-C		79.9	9	1.5	1.9
	Mid-C		221.	.5	7.6	3.4
	High-C		558.	.3	33.4	6.0

#### 4.4 Total H2AX Assay Performance

The intra-assay performances of the Total H2AX were evaluated using the defined critical reagent lot numbers in Section 3.2.

**Table 2.** Inter-assay precision was measured by 3 runs of total H2AX standards and assay controls. All 3 runs passed the inter-assay acceptance criteria (%CV < 25%) with a range from 6 to 21% for the standards and 2.2 to 7.3% for the controls.

Total H2AX Standard (pM)	Average RLU (n=3)	SD	%CV
50	865720	184054	21
100	1063562	168332	16
200	1418546	218046	15
400	2298028	327995	14
800	4611161	370373	8
1600	10482498	851476	8
3200	26055096	1696172	7
6400	59158022	3389425	6





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Assay Control	Average pM (n=3)	SD	%CV
Low-C	216.1	15.8	7.3
Mid-C	844.8	49.3	5.8
High-C	2337.0	51.0	2.2

#### 4.5 Tumor lysate controls established during assay development

4.5.1 γH2AX Readout Ranges for the prepared lot of MCF7 Control Lysates (Low-C; Mid-C and High-C) used during assay development are provided as a reference below. These values are for the purpose of illustration only. Each site should generate its own lot-specific readout ranges.

Tumor Lysate Control Level	Example Acceptable Readout Value (pM)
Low-C	53-82
Mid-C	181-228
High-C	458-583

4.5.2 **Total H2AX Readout Ranges** for the prepared lot of tumor cell lines as Controls (Low-C; Mid-C and High-C) used during assay development are provided as a reference below. These values are for the **purpose of illustration only**. Each site should generate its own lot-specific readout ranges.

Tumor Lysate Control Level	Example Acceptable Readout Value (pM)
Low-C	154 - 275
Mid-C	602 - 1004
High-C	2075 - 2844





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#### 4.6 γH2AX /Total H2AX Preclinical Testing

The details of the preclinical testing results were previously presented and published <sup>2-4</sup>. The 96well plate-based ELISAs for quantifying  $\gamma$ H2AX and total H2AX levels have been assessed for in vitro and in vivo applications in irradiation exposure monitoring and in pharmacodynamic evaluation of anti-cancer agents.

In vitro, dose-dependent increases in the ratio of  $\gamma$ H2AX to total H2AX were detected after escalating ionizing radiation exposure and concentration-dependent increases after Topoisomerase 1 (Top1) inhibitor exposure. Treating with inhibitors of PARP or ATR alone did not significantly induce  $\gamma$ H2AX. Combinations of Top1 inhibitors with PARP or ATR inhibitors led to synergistic induction of DNA damage. Among five ATR inhibitors evaluated in combination with Top1 inhibitors, VE-822 and AZD-6738 were observed to have the highest synergy for  $\gamma$ H2AX induction, while NU-6027 showed none. Combinations of CPT-11 (Irinotecan, Camptosar, Campto) with ABT-888, AZD-2281 or MK-4827 showed synergistic induction of  $\gamma$ H2AX in A375 xenografts in vivo. Additional testing of human specimens including PBMCs, bone marrow, and tumor biopsies supports the assay's clinical suitability and potential advantages.

To conclude, the quantitative ELISA for measuring both  $\gamma$ H2AX and total H2AX is ready for Research Use for monitoring DNA damage induced by chemotherapeutic agents or irradiation.

#### 5.0 MATERIALS AND EQUIPMENT

#### 5.1 γH2AX Key Reagents:

- 5.1.1 γH2AX peptide standard, lyophilized powder, powder (custom-preparation from Invitrogen, synthetic peptide: AVLLPKKTSATVGPKAPSGGKKATQA[pS]QEY)
- 5.1.2 Anti-phospho-H2AX (Ser139) mouse monoclonal antibody, clone JBW301 (Millipore, Cat#: 05-636)
- 5.1.3 Histone H2AX rabbit polyclonal antibody (Abcam, Cat#: ab10475)

#### 5.2 Total H2AX Key Reagents:

- 5.2.1 Total H2AX recombinant protein standard, lyophilized powder (Axxora, Cat#: ALX-201-176-M005)
- 5.2.2 HIST1H2AC mouse monoclonal antibody, clone 4F10 (Novus, Cat#: H00008334-M01)
- 5.2.3 Histone H2AX rabbit polyclonal antibody (Abcam, Cat#: ab10475)
- 5.3 Tumor Lysate Control (custom preparation prepared to target low, mid and high γH2AX and H2AX ranges)
- 5.4 Goat anti-rabbit HRP-conjugated polyclonal antibody (KPL, Cat#: 074-15-061). Reconstitute to 1 mg/mL stock solution in HRP Stabilizer (KPL, Cat#: 54-15-01).
- 5.5 SuperSignal ELISA Pico Chemiluminescent Substrate (Thermo Scientific Pierce, Cat#: 37070)
- 5.6 Acetate plate sealers (Thermo Scientific Pierce, Cat #: 3501)
- 5.7 Reacti-Bind White Opaque 96-well Plate (Thermo Scientific Pierce, Cat#: 15042)
- 5.8 Carbonate-bicarbonate buffer capsules, pH 9.6 (e.g., Sigma-Aldrich, Cat#: C3041-50CAP)
- 5.9 Tween 20 nonionic, aqueous solution, 10% w/v (Roche Applied Science, Cat#: 11332465001)
- 5.10 20% sodium dodecyl sulfate (SDS; e.g., Sigma-Aldrich, Cat#: 05030-500ML-F)
- 5.11 10X Phosphate Buffered Saline, pH 7.2 (PBS; e.g., Invitrogen, Cat#: 70013-073)
- 5.12 SuperBlock (TBS) Blocking Buffer (Thermo Scientific Pierce, Cat#: 37535)





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- 5.13 Albumin, bovine serum (BSA; e.g., Sigma-Aldrich, Cat#: A7030)
- 5.14 Mouse serum (e.g., Sigma-Aldrich, Cat#: M5905)
- 5.15 Protamine sulfate salt from salmon (Sigma-Aldrich, Cat#: P4020-5G)
- 5.16 UltraPure DNase/RNase free distilled water (e.g., Invitrogen, Cat#: 10977-015) or Milli-Q water
- 5.17 Cell Extraction Buffer (CEB; Invitrogen, Cat#: FNN0011)
- 5.18 Protease Inhibitor Cocktail (Sigma-Aldrich, Cat#: P-2714 or Roche, Cat#: 11697498001)
- 5.19 PhosSTOP, phosphatase inhibitor cocktail tablets (Roche, Cat#: 04906837001)
- 5.20 Pipettors (200-1000 µL, 50-200 µL, and 2-20 µL) and tips
- 5.21 Multichannel pipettors (50-300 µL, 5-50 µL) and tips
- 5.22 Reagent reservoirs (Fisher Scientific, Cat#: 21-381-27C)
- 5.23 1.5-mL Sarstedt tubes (Sarstedt, Cat#: 72.692.005)
- 5.24 15-mL polypropylene tubes (e.g., Fisher Scientific, Cat#: 14-959-49B)
- 5.25 50-mL polypropylene tubes (e.g., Becton Dickinson, Cat#: 352098)
- 5.26 Ice bucket
- 5.27 Sorvall Fresco microcentrifuge (Fisher Scientific)
- 5.28 Vortex Genie 2 (Daigger, Cat#: 3030A)
- 5.29 Dry, heated incubator able to maintain  $37^{\circ}C \pm 3^{\circ}C$
- 5.30 Dry, heated incubator able to maintain  $25^{\circ}C \pm 3^{\circ}C$
- 5.31 Infinite® 200 or M200Pro Microplate Reader (Tecan US)
- 5.32 BioTek ELx405 Select Microplate Washer (BioTek Instruments)
- 5.33 -80°C freezer
- 5.34 4°C refrigerator





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#### 6.0 **EXAMPLE ELISA PLATE CONFIGURATION**

#### 6.1 Plate map for γH2AX IA

	1	2	3	4	5	6	7	8	9	10	11	12
Α	1X PBS-2% BSA (Assay Buffer) Only				5 p	5 pM 1X PBS-2% BSA (Assay Buff			er) Only			
B	High C					10	pМ					Low C
С	nigii-C	<b>S</b> 1	<b>S</b> 3	<b>S</b> 5	<b>S</b> 7	20	pМ	<b>S</b> 9	S11	S13	S15	Low-C
D	Mid C					40	pМ					Mid C
E	MIId-C					80	pМ					Mid-C
F	Low C	S2	<b>S</b> 4	<b>S</b> 6	<b>S</b> 8	160	pМ	S10	S12	S14	S16	High C
G	Low-C					320	pМ					nigii-C
Η	1X PBS	-2% BSA	A (Assay	Buffer)	Only	640	pМ	1X PBS-2% BSA (Assay Buffer) Only			er) Only	
	Control Samples	Unkno	own Samples, Triplicate		γH2 Pept Stand Dupl	AX tide lards, icate	Unknown Samples, Triplicate			plicate	Control Samples	

S1 through S16 are unknown sample (S) wells in triplicate.

#### Plate map for total H2AX IA 6.2

	1	2	3	4	5	6	7	8	9	10	11	12	
Α	1X PBS-2% BSA (Assay Buffer) Only		50 pM		1X PBS-2% BSA (Assay Buffer) Only								
B	High C					100	pМ					Low C	
С	nigii-C	<b>S</b> 1	<b>S</b> 3	<b>S</b> 5	<b>S</b> 7	200	pМ	<b>S</b> 9	S11	S13	S15	Low-C	
D	Mid C	Mid-C				400	pМ					Mid C	
E	Mid-C					800	pМ					Milu-C	
F	Low C	S2	<b>S</b> 4	<b>S</b> 6	<b>S</b> 8	1600	) pM	S10	S12	S14	S16	High C	
G	LOW-C						3200	) pM					High-C
Η	1X PBS	-2% BSA	A (Assay	Buffer)	Only	6400	) pM	1X	1X PBS-2% BSA (Assay Buffer) Only				
	Control Samples	ontrol mples Unknown Samples, Triplicate			Total I Pep Stand Dupl	H2AX tide lards, icate	Unknown Samples, Triplicate			Control Samples			

S1 through S16 are unknown sample (S) wells in triplicate.





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#### 6.3 Optional plate map for running both total H2AX and $\gamma$ H2AX on the same plate.

		Total H	2AX IA					γH2A	X IA				
	1 2	3	4	5	6	7	8	9	10	11	12		
A	50 pM	1X PBS-	2% BSA (4	BSA (Assay Buffer) Only			рΜ	1X PBS-	1X PBS-2% BSA (Assay Buffer)				
B	100 pM					10	pМ						
С	200 pM	<b>S</b> 1	<b>S</b> 3	S5	S5	S5	Mid-	20	pМ	S1a	S3a	S5a	Mid-C
D	400 pM				č	40	pМ						
E	800 pM			T		80	pМ			Ŧ			
F	1600 pM	S2	S4	Low-	High-	160	pМ	S2a	S4a	C Low-	High-		
G	3200 pM			C	C	320	pМ				C		
Η	6400 pM	1X PBS-	2% BSA (4	Assay Buff	er) Only	640	pМ	1X PBS-	2% BSA (	Assay Buf	fer) Only		
	Total H2AX Peptide Standards, Duplicate	Unkno	Inknown and Control Samples, Triplicate		γH2AX Stand Dup	Peptide lards, licate	Unknown and Control Samples, Triplicate		mples,				

for Total H2AX analysis.

S1 to S5 are unknown sample wells in triplicate S1a to S5a are unknown sample wells in triplicate for yH2AX analysis.

The plate map above can be used for running both total H2AX and  $\gamma$ H2AX assays on the same plate if a few samples are used.

#### 7.0 SAMPLE LYSATE PREPARATION

Unknown samples (solid tissue and cells, respectively) should be processed following SOP340520-Biopsy Specimen Processing and SOP340506-PBMC Protein Extraction for the PAR Immunoassay with the following modifications<sup>5</sup>:

- Landing Page: http://dctd.cancer.gov/ResearchResources/ResearchResources-biomarkers.htm •
- **SOP340520 LINK:** • http://dctd.cancer.gov/ResearchResources/biomarkers/docs/par/SOP340520\_Biopsy\_Tissue.pdf

#### **SOP340506 LINK:** • https://dctd.cancer.gov/ResearchResources/biomarkers/docs/par/SOP340506 PBMC Extraction. pdf

- 7.1 SOP340520 sample lysis references use of an 18-g needle tumor biopsy. For preclinical sample processing, a similar tissue piece would weigh approximately 10 mg.
- 7.2 In SOP340520 (tumor tissue) and SOP340506 (PBMC/Cells), add 1 tablet PhosSTOP (Roche Applied Science, Cat#: 04906837001) per 10 mL Cell Extraction Buffer (CEB) for all steps requiring "CEB with Protease Inhibitors (PIs)."





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#### 8.0 γH2AX ELISA PLATE SET-UP

#### 8.1 Key Reagents/Supplies

It is recommended to record the lot numbers, stock reagent concentration, and manufacturer's expiration dates for the Key Reagents in the Assay Record (Appendix 1, Section 1). It is advised to follow manufacturer's recommendations for the handling and storage of the reagents/supplies.

- 8.1.1  $\gamma$ H2AX Peptide Standard: Prepared as a 20680 pM stock solution. Aliquot in sufficient volumes for one 96-well plate. NCI has stored the standard successfully at 80°C for > 1 y.
- 8.1.2 **Tumor Lysate Control**: Lysates prepared from tumor cell lines to target Low, Medium and High  $\gamma$ H2AX ranges. NCI has stored the control lysates successfully at -80°C for > 1 y.

See Appendix 3 for details on preparation of tumor lysate controls.

- 8.1.3 **γH2AX mAb**: Stock solution qualified from the manufacturer. Dilutions for assay performance for specific lot numbers of antibody will need to be determined by the assay site and should be matched to the pAb. Aliquot in sufficient volumes for one 96-well plate. NCI has stored the γH2AX mAb (glycerol containing buffer) successfully at -20°C for > 1 y.
- 8.1.4 **H2AX pAb**: Stock solution qualified from the manufacturer. Dilutions for assay performance for specific lot numbers of antibody will need to be determined by the assay site and should be matched to the mAb. NCI has stored the H2AX pAb successfully at -80°C for > 1 y.
- 8.1.5 **Goat Anti-Rabbit HRP-Conjugated pAb**: Prepare a 1 mg/mL stock solution in HRP Stabilizer. Aliquot in sufficient volumes for three 96-well plates. NCI has stored the HRP-Conjugated pAb successfully at 2°C to 8°C for up to 1 y.
- 8.1.6 **Chemiluminescent Substrate Solutions**: Stock solutions (Peroxide and Pico Luminol/Enhancer Solutions) qualified from the manufacturer. Protect from light during storage and use. NCI has stored the substrate solutions successfully at room temperature at  $25^{\circ}C \pm 3^{\circ}C$  for up to 1 y.
- 8.1.7 **Reacti-Bind White Opaque 96-well Plate**: Store at  $25^{\circ}C \pm 3^{\circ}C$  away from volatile chemicals.

#### 8.2 **Plate Map and Buffer Preparation**

- 8.2.1 Based on the number of unknown samples to be analyzed, generate a Plate Map (example, Section 6.1) to define the location and replicates of unknown samples, tumor lysate controls, and γH2AX peptide standards. Samples from a single experiment should be analyzed on one 96-well plate, not split over two plates, to ensure consistent sample handling.
- 8.2.2 Once the number of wells is known, determine the amount of reagents required for the assay using the Assay Record in Appendix 1. Prepare Wash Buffer and Assay Buffer according to Appendix 1, Section 2A.

#### IMPORTANT: For all wash and aspiration steps, do not let the wells dry out.





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Note that it is recommended that both 37°C and 25°C incubation steps for the assay be carried out in fixed-temperature incubators with each 96-well plate placed on a CoolSink thermoconductive plate (pre-warmed for at least 1 hr) during these incubation steps. These thermoconductive plates should be placed horizontally inside the incubator in direct contact with the incubator bottom or shelf and should not be stacked. The assay plate should be placed and carefully centered onto a prewarmed thermoconductive plate inside the incubator for each incubation step.

#### 8.3 **Plate Preparation**

- 8.3.1 Use the calculations in the Assay Record (Appendix 1, Section 3A) to prepare 11 mL γH2AX mAb Coating Solution for the assay. This is sufficient for one 96-well plate (preparing enough for 110 wells). Thaw coating antibody immediately prior to dilution; do not allow to sit for extended periods upon thawing.
- 8.3.2 If more than one 96-well plate is to be coated, pool the aliquots of coating antibody and then dilute appropriately. This will ensure that all plates are exposed to identical coating antibody. Discard excess diluted antibody.
- 8.3.3 Add 100  $\mu$ L of the  $\gamma$ H2AX mAb Coating Solution per well using a multichannel pipettor, cover the plate with an acetate sheet, and incubate at 37°C for 2 h. Record the coating antibody incubation conditions in the Assay Record (Appendix1, Section 3B).
- 8.3.4 Alternatively, the plate can be incubated overnight at 2°C to 8°C.
- 8.3.5 Following incubation with the  $\gamma$ H2AX mAb Coating Solution, aspirate the plate using a plate washer (for the BioTek Plate Washer, use the *Aspirate* program). After aspiration, tap the plate on paper towels to remove any residual liquid.
- 8.3.6 Add 250 μL of SuperBlock to each well. Cover the plate with an acetate sheet and incubate at 37°C for 1 to 1.5 h. Record the incubation conditions in the Assay Record (Appendix 1, Section 4).
- 8.3.7 After blocking, move plate to  $25^{\circ}C \pm 3^{\circ}C$  until washing step (Step 8.7.1).
- 8.3.8 Alternative, once coated plates have been blocked, they can be stored at 2°C to 8°C for up to one week. It is important to not let wells dry out.

#### 8.4 Unknown Sample Lysate Preparation

- 8.4.1 Tumor/tissue stock lysates to be analyzed in the assay are initially normalized to 0.2  $\mu g/\mu L$  in  $\gamma$ H2AX sample diluent (CEB Complete). Tumor stock lysates with a total protein concentration of < 0.2  $\mu g/\mu L$  should not be used in the  $\gamma$ H2AX Immunoassay.
  - 8.4.1.1 Place unknown samples to be assayed on ice. Record the sample name/IDs and starting lysate concentration in the Assay Record (Appendix 1, Section 5) for each sample to be used.
  - 8.4.1.2 For unknown stock lysates with stock protein concentrations  $\geq 0.2 \ \mu g/\mu L$ , calculate the volume of stock lysate required to prepare 85  $\mu L$  of a 0.2  $\mu g/\mu L$  as follows:





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			<u>0.2 μg/μL</u> <u>XX</u> μg/μL	* 85 Stock Lysate	$5 \mu L = XX \mu L V$	ol. Stock Lysate							
		8.4.1.3	Record the volume concentration of w 1, Section 5).	es stock lysate orking dilutio	e, $\gamma$ H2AX sample diluer ons of the lysate in the A	nt (CEB Complete) and Assay record (Appendix							
		8.4.1.4	Do not pipette less yield volumes of s to 1:5 pre-dilution	than 5 μL of tock lysate lea of the lysate	the stock lysate. If the ss than 5 $\mu$ L, prepare subefore proceeding.	calculations above fficient volume of a 1:2							
		8.4.1.5	In a labeled 1.5 ml Complete) to the c volume to 85 $\mu$ L.	L tube, add su alculated volu Keep the wor	fficient γH2AX sample ume of stock lysate need king lysate on ice.	diluent (CEB led to bring total							
		8.4.1.6	Flash freeze remaining stock lysate in liquid nitrogen or dry/ice ethanor return to -80°C freezer. As with all lysates, freeze/thaw cycles should l minimized.										
	8.4.2	From the prepared (CEB Co	From the normalized working tumor/tissue lysates of 0.2 $\mu$ g/ $\mu$ L, three final dilutions are prepared in order to analyze the lysates at 2, 1 and 0.5 $\mu$ g/well in $\gamma$ H2AX sample diluent (CEB Complete).										
		8.4.2.1	Perform the follow needed to prepare	ving calculation 3 different lys	on to calculate the volur sate dilutions (2, 1, or 0	ne of working lysate 5 µg/well).							
			<u>2, 1 or 0.5 μg/well</u> 0.2 μg/μL Workin	l * g Lysate	$4 \text{ wells} = 40, 20 \text{ or } 10^{-10}$	0 μL Working Lysate							
		8.4.2.2	For each Diluted I used to bring the to dilution in triplicat	_ysate, γH2A2 otal volume to te (plus 1 extr	X sample diluent (CEB 100 μL. This is suffic a well).	Complete) should be ient volume to run each							
		8.4.2.3	Record the Sample (CEB Complete) u Record (Appendix	e ID, volume used to prepar 1, Section 6A	Working Lysate and γH e each Diluted Tumor L A).	2AX sample diluent sysate in the Assay							
		8.4.2.4	Clearly label 1.5 n sufficient volume volume of working	nL tubes with γH2AX samp g lysate to bri	the sample number (e.g le diluent (CEB Comple ng the total volume to 1	s., S1, S2), add ete) to the calculated 00 μL.							
		8.4.2.5	Keep the Diluted I Lysate.	Lysates on ice	until use. Discard rem	aining unused Working							
	8.4.3	For prepa	For preparation of PBMC/Cell Lysates samples:										
		8.4.3.1	Stock lysates for F Section 7.0 and A <sub>1</sub>	PBMCs/Cells ppendix 3.	$(1 \times 10^7 \text{ cells/mL})$ are p	repared according to							
_		8.4.3.2	Place 100 µL of th number (e.g., S1, S for triplicate well j	e stock lysate S2). No other preparation (+	into a 1.5 mL tube labe sample preparation is r 1 well extra).	eled with the sample necessary; this is enough							





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- 8.4.3.3 Record the stock lysate concentration in cells/mL, the volume of stock PBMC/Cell lysate set aside for each sample in the Assay Record (Appendix 1, Section 6A).
- 8.4.3.4 Lysates will be diluted an additional 3-fold once loaded into the 96-well plate yielding a relative load of  $2.5 \times 10^5$  cells/well. This relative load will provide quantitative values in the range of the assay for most PBMC and cellular lysates. It may be necessary to pre-dilute lysates prepared from certain cell lines with high levels of  $\gamma$ H2AX. In the event that a cell line provides quantitative values above the range of the assay (>ULQ) the samples should be prediluted in CEB Complete prior to analysis.
- 8.4.3.5 Keep the PBMC/Cell lysates aliquoted for use in the assay on ice until use. Flash freeze the remaining stock lysate in liquid nitrogen or dry ice/ethanol bath and return to -80°C freezer. As with all lysates, freeze/thaw cycles should be minimized.

#### 8.5 Preparation of γH2AX peptide standards (run in duplicate)

- 8.5.1 For one 96-well plate, retrieve a  $\gamma$ H2AX peptide standard stock tube (20680 pM) from the -80°C freezer and thaw on ice. Vortex and mix by inverting 5-8 times before use. Label eight 1.5-mL tubes, numbered 1 through 8, for the  $\gamma$ H2AX peptide standards. Prepare a 9th tube as the blank.
- 8.5.2 Prepare the γH2AX peptide standards by serial dilution as outlined in the Assay Record (Appendix 1, Section 6B) with final concentrations ranging from 1920 to 15 pM in Assay Buffer (1X PBS-2% BSA).
- 8.5.3 Standards will be diluted an additional 3-fold when added to the 96-well plate to generate a reference curve ranging from 640 to 5 pM γH2AX peptide standard. Keep standards on ice until use. Only make enough standards for the assay and discard any excess.

#### 8.6 **Control Lysates** (run in duplicate)

- 8.6.1 For one 96-well plate, retrieve one of each High-C, Mid-C and Low-C tumor lysate control vials from the -80°C freezer and thaw on ice. Controls are prepared at a concentration ready for use in the assay and no further dilution is required. Vortex and mix by inverting 5-8 times before use. See Appendix 3 for more information on preparation of controls.
- 8.6.2 Keep controls on ice until use. Controls will be diluted 3-fold with Assay Buffer (1X PBS-2% BSA) once loaded into the 96-well plate. Only thaw enough of the controls for the assay and discard any excess.
- 8.6.3 Note the Control Lot Numbers in Appendix 1, Section 6C.





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#### 8.7 γH2AX Protein Capture

8.7.1 Following incubation with SuperBlock (Step 8.3.6), the plates are aspirated and washed once with 350 μL of Wash Buffer (1X PBS-0.1% Tween) using a plate washer.

For the BioTek Microplate Washer, the settings are:

METHOD	
Number of Cycles:	1
Soak/Shake:	Yes
Soak Time:	5 Sec
Shake before soak:	No
Prime after soak:	No
DISPENSE	
Dispense Volume:	350 μL/well
Dispense Flow Rate:	06
Dispense Height:	120 (15.240 mm)
Horizontal DISP POS:	00 (0.000 mm)
Bottom Wash First:	No
Prime Before Start:	No
ASPIRATE	
Aspirate Height:	031 (3.937 mm)*
Horizontal ASPR POS:	-20 (-0.914 mm)*
Aspiration Rate:	05 (6.4 mm/sec)
Aspirate Delay:	1000 MSec
Crosswise ASPIR:	No
Final Aspiration:	Yes
Final Aspirate Delay:	1000 MSec

\*Recommended initial setting, optimize Aspirate Height and Horizontal ASPR POS to allow for complete aspiration of an individual unit following manufacturer's recommendations.

- 8.7.2 After the wash, tap the plate on paper towels to remove residual buffer. Proceed immediately to the next step; do not allow the plate to dry out.
- 8.7.3 Immediately, add 50 μL of Assay Buffer (1X PBS-2% BSA) to each well using a multichannel pipettor. Each well will hold a final volume of 75 μL after sample addition.





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8.7.4 Use the Plate Map Design (Section 6.1) and the Sample Calculation Table (Appendix 1, Section 6A) as a guide to set up the 96-well plate for incubation with unknown samples, γH2AX peptide standards (Appendix 1, Section 6B), and tumor cell controls (Appendix 1, Section 6C and Appendix 3). Pipette reagents in the following order; do not deviate from order of addition:

Order	Sample/Reagent and Volume
1	$25 \ \mu L$ of specified concentrations of $\gamma$ H2AX peptide standards into designated duplicate wells. Load the lowest concentration first.
2	$25 \ \mu L$ of each unknown sample into designated triplicate wells.
3	25 μL each of assay controls (Low-C, Mid-C, and High-C) into both sets of designated duplicate wells.
4	$25 \ \mu L$ of additional Assay Buffer (1X PBS-2% BSA) into each of the Background wells.

8.7.5 Cover the plate with an acetate sheet and incubate at  $2^{\circ}$ C to  $8^{\circ}$ C for  $18 \pm 2$  h. Record the date, start time, and incubation temperature in the Assay Record (Appendix 1, Section 7).

#### 8.8 γH2AX Detection (next day)

Go to Section 10.0 for the Detection Method which is the same for both the  $\gamma$ H2AX and Total H2AX assays.

#### 9.0 Total H2AX ELISA PLATE SET UP

#### 9.1 Key Reagents/Supplies

It is recommended to record the lot numbers, stock reagent concentration, and manufacturer's expiration dates for the Key Reagents in the Assay Record (Appendix 2, Section 1). It is advised to follow manufacturer's recommendations for the handling and storage of the reagents/supplies.

- 9.1.1 **Total H2AX Peptide Standard**: Prepared as a 33000 pM stock solution. Aliquot in sufficient volumes for one 96-well plate. NCI has stored successfully at -80°C for > 1 y.
- 9.1.2 **Tumor Lysate Control**: Lysates from tumor cell lines prepared to target Low, Medium and High Total H2AX ranges. NCI has stored successfully at -80°C for > 1 y.

See Appendix 3 for details on preparation of tumor cell line lysate controls.

- 9.1.3 **Total H2AX mAb**: Stock solution qualified from the manufacturer. Dilutions needed for assay performance of specific lot numbers of antibody will need to be determined by the assay site and should be matched to the pAb. NCI has stored successfully at -80°C for > 1 y.
- 9.1.4 **H2AX pAb**: Stock solution qualified from the manufacturer. Dilutions needed for assay performance of specific lot numbers of antibody will need to be determined by the assay site and should be matched to the mAb. Aliquot in sufficient volumes for one 96-well plate. NCI has stored successfully at -80°C for > 1 y.





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- 9.1.5 **Goat Anti-Rabbit HRP-Conjugated pAb**: Prepare a 1 mg/mL stock solution in HRP Stabilizer. Aliquot in sufficient volumes for three 96-well plates. NCI has stored successfully at 2°C to 8°C for up to 1 y.
- 9.1.6 **Chemiluminescent Substrate Solutions**: Stock solutions (Peroxide and Pico Luminol/Enhancer Solutions) qualified from the manufacturer. Protect from light during storage and use. NCI has stored successfully at 25°C ± 3°C for up to 1 y.
- 9.1.7 **Reacti-Bind White Opaque 96-well Plate**: Store at  $25^{\circ}C \pm 3^{\circ}C$  away from volatile chemicals.

### 9.2 Plate Map and Buffer Preparation

- 9.2.1 Based on the number of unknown samples to be analyzed, generate a Plate Map (Section 6.2) to define the location and replicates of unknown samples, tumor lysate controls, and Total H2AX peptide standards. Samples from a single experiment should be analyzed on one 96-well plate, not split over two, to ensure consistent sample handling.
- 9.2.2 Once the number of wells is known, determine the amount of reagents required for the assay using the Assay Record in Appendix 2.

### **IMPORTANT:** For all wash and aspiration steps, do not let the wells dry out.

 Note that it is recommended that both 37°C and 25°C incubation steps for the assay be carried out in fixed-temperature incubators with each 96-well plate placed on a CoolSink thermoconductive plate (pre-warmed for at least 1 hr) during these incubation steps. These thermoconductive plates should be placed horizontally inside the incubator in direct contact with the incubator bottom or shelf and should not be stacked. The assay plate should be placed and carefully centered onto a prewarmed thermoconductive plate inside the incubator for each incubation step.

#### 9.3 **Plate Preparation**

- 9.3.1 Use the calculations in the Assay Record (Appendix 2, Section 3A) to prepare 11 mL Total H2AX mAb Coating Solution for the assay. This is sufficient for one 96-well plate (preparing enough for 110 wells). Thaw coating antibody immediately prior to dilution; do not allow to sit for extended periods upon thawing.
- 9.3.2 If more than one 96-well plate is to be coated, pool the coating antibody aliquots and then dilute appropriately. This will ensure that all plates are exposed to identical coating antibody. Discard excess diluted antibody.
- 9.3.3 Add 100 μL of the Total H2AX mAb Coating Solution per well using a multichannel pipettor, cover the plate with an acetate sheet, and incubate at 37°C for 2 h. Record the coating antibody incubation conditions in the Assay Record (Appendix 2, Section 3B).
- 9.3.4 Alternatively, the plate can be incubated overnight at 2°C to 8°C.
- 9.3.5 Following incubation with the Total H2AX mAb Coating Solution, aspirate the plate using a plate washer (for the BioTek Plate Washer, use the *Aspirate* program). After aspiration, tap the plate on paper towels to remove any residual liquid.





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- 9.3.6 Add 250 μL of SuperBlock to each well. Cover the plate with an acetate sheet and incubate at 37°C for 1 to 1.5 h. Record the incubation conditions in the Assay Record (Appendix 2, Section 4).
- 9.3.7 After blocking, move plate to  $25^{\circ}C \pm 3^{\circ}C$  until washing step (Step 9.5.1).
- 9.3.8 Alternatively, once coated plates have been blocked they can be stored at 2°C to 8°C for up to one week. It is important to not let wells dry out.

### 9.4 Unknown Sample Lysate Preparation

- 9.4.1 Tumor/tissue stock lysates with stock protein concentrations > 0.75  $\mu g/\mu L$  are initially prediluted to 0.75  $\mu g/\mu L$ . For stock lysates with protein concentrations between 0.2 and 0.75  $\mu g/\mu L$ , no predilution is required and the stock lysates are used directly to prepare the final assay lysate preparations as described below. Unknown sample lysates with a total protein concentration of < 0.2  $\mu g/\mu L$  should <u>not</u> be used in the Total H2AX Immunoassay.
- 9.4.2 For unknown stock tumor/tissue lysates with stock protein concentrations  $> 0.75 \ \mu g/\mu L$ :
  - 9.4.2.1 Place unknown samples to be assayed on ice. Record the sample name/IDs and starting lysate concentration in the Assay Record (Appendix 2, Section 5) for each sample to be used.
  - 9.4.2.2 Calculate the volume of stock lysate required to prepare 10  $\mu$ L of a 0.75  $\mu$ g/ $\mu$ L Working Lysate as follows:

 $0.75 \,\mu g/\mu L \quad * \quad 10 \,\mu L = \quad XX \,\mu L \text{ Vol. Stock Lysate}$ 

XX µg/µL Stock Lysate

- 9.4.2.3 Record the stock lysate concentration, working lysate concentration, volume stock lysate and volume H2AX sample diluent (CEB Complete) in the Assay Record (Appendix 2, Section 5).
- 9.4.2.4 Do not pipette less than 5  $\mu$ L of the stock lysate. If the calculations above yield volumes of stock lysate less than 5  $\mu$ L, prepare sufficient volume of a 1:2 to 1:5 pre-dilution of the lysate before proceeding.
- 9.4.2.5 In a labeled 1.5 mL tube, add sufficient Total H2AX Sample Diluent (CEB Complete) to the calculated volume of stock lysate needed to bring the total volume to 10 μL.
- 9.4.2.6 Add 65  $\mu$ L of Assay Buffer (1X PBS-2% BSA) to generate 75  $\mu$ L of 0.1  $\mu$ g/ $\mu$ L Diluted Working Lysate. Record the volume prediluted lysate (10  $\mu$ L), diluted working lysate concentration (0.1  $\mu$ g/ $\mu$ L) and volume Assay Buffer (65  $\mu$ L) in the Assay Record (Appendix 2, Section 5).
- 9.4.2.7 Keep the diluted working lysate on ice. Flash freeze remaining stock lysate in liquid nitrogen or dry ice/ethanol and return to -80°C freezer. As with all lysates, freeze/thaw cycles should be minimized.





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- 9.4.3 For unknown stock lysates with stock protein concentrations <u>between</u> 0.2 and 0.75 μg/μL, no normalization is needed and stock lysate are directly diluted with Assay Buffer (1X PBS-2% BSA) for the Diluted Working Lysate of 0.1 μg/μL.
  - 9.4.3.1 Place unknown samples to be assayed on ice. Record the sample name/IDs and starting lysate concentration in the Assay Record (Appendix 2, Section 5) for each sample to be used.
  - 9.4.3.2 Calculate the volume of stock lysate required to prepare 75  $\mu$ L of a 0.1  $\mu$ g/ $\mu$ L Working Lysate as follows:

 $\frac{0.1 \,\mu g/\mu L}{XX} \,\mu g/\mu L \text{ Stock Lysate} = \frac{XX}{\mu} \mu L \text{ Vol. Stock Lysate}$ 

- 9.4.3.3 Record the stock lysate concentration, working lysate concentration, volume stock lysate and volume Assay Buffer (1X PBS-2% BSA) in the Assay Record (Appendix 2, Section 5).
- 9.4.3.4 In a labeled 1.5 mL tube, add sufficient volume Assay Buffer (1X PBS—2% BSA) to the calculated volume of stock lysate needed to bring the total volume to 75  $\mu$ L.
- 9.4.3.5 Keep the diluted working lysate on ice. Flash freeze remaining stock lysate in liquid nitrogen or dry ice/ethanol and return to -80°C freezer. As with all lysates, freeze/thaw cycles should be minimized.
- 9.4.4 From the normalized working tumor/tissue lysate of  $0.1 \,\mu g/\mu L$ , three final dilutions are prepared in order to analyze the lysates at 0.5, 0.375 and 0.25  $\mu g/well$ .
  - 9.4.4.1 Perform the following calculation to calculate the volume of normalized working lysate needed to analyze the lysates at 0.5, 0.375 and 0.25 µg/well.

<u>0.5, 0.375 or 0.25  $\mu$ g/well</u> \* 4 wells = 20, 15 or 10  $\mu$ L Working Lysate

0.1 µg/µL Working Lysate

- 9.4.4.2 For each diluted working lysate, Assay Buffer (1X PBS-2% BSA) should be used to bring the total volume to 100 μL. This is sufficient volume to run each dilution in triplicate (+1 well extra).
- 9.4.4.3 Record the Specimen ID, Working Lysate Concentration, Volume Working Lysate and Assay Buffer used to prepare each final diluted lysate in the Assay Record (Appendix 2, Section 6A).
- 9.4.4.4 Clearly label 1.5 mL tubes with the sample number (e.g., S1, S2), add sufficient volume Assay Buffer to the calculated volume of Working Lysate to bring the total volume to 100 μL.
- 9.4.4.5 Keep the Diluted Lysates on ice until use. Discard remaining unused Working Lysate.
- 9.4.5 For preparation of PBMC/Cell Lysate samples:





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	9.4	.5.1 Stoc Sect	k lysates for F ion 7.0 and Aj	PBMCs/Cells ( ppendix 3.	1 x 10 <sup>7</sup> cells/mL) are pr	repared according to		
	9.4	.5.2 PBM into x 10 info	PBMC/Cell diluted lysates are prepared by adding 7.5 $\mu$ L of the stock lysate into 117.5 $\mu$ L of Assay Buffer (1X PBS-2% BSA) to a final concentration of 6 x 10 <sup>5</sup> cells/mL in 125 $\mu$ L for the Total H2AX assay. Record the dilution information in the Sample Calculation Table (Appendix 2, Section 6A).					
	9.4	.5.3 Clea the s	Clearly label all tubes with the sample number (e.g., S7, S8) and add 7.5 $\mu$ L of the stock lysate into 117.5 $\mu$ L of 1X PBS-2% BSA for each sample.					
	9.4	.5.4 Lysa with	Lysates will be diluted an additional 3-fold once loaded into the 96-well plate with various assay reagents yielding a relative load of $1.5 \times 10^4$ cells/well.					
	9.4	.5.5 Kee rema 80°C	Keep the diluted PBMC/Cell lysates on ice until use. Flash freeze the remaining stock lysate in liquid nitrogen or dry ice/ethanol bath and return to - 80°C freezer. As with all lysates, freeze/thaw cycles should be minimized.					
	9.4.6 <b>P</b> ı	reparation	of Total H2A	X Recombina	nt Standards (run in e	luplicate)		
	9.4	.6.1 For stoc Vort num 9th t well	one 96-well pl k solution tube ex and mix by bered 1 throug ube of only A s.	late, retrieve on e (33000 pM) f y inverting 5-8 gh 8, for the To ssay Buffer (1	times before use. Labert table H2AX recombinant times before use. Labert tal H2AX recombinant X PBS-2% BSA) to loa	inant standard working and thaw on ice. el eight 1.5 mL tubes, t standards. Prepare a d into the background		
	9.4	.6.2 Prep the 1 1920	are the Total Assay Record 00 to 150 pM	H2AX recomb (Appendix 2, 9 in Assay Buffe	inant standards by seria Section 6B) with concer r (1X PBS-2% BSA).	l dilution as outlined in ntrations ranging from		
9.4.6.3 Standards will be diluted an additional 3-fold when added to the 96-w to generate a reference curve ranging from 6400 to 50 pM Total H2A2 recombinant standard.					ed to the 96-well plate M Total H2AX			
	9.4	.6.4 Kee disc	p standards on ard any excess	i ice until use.	Only make enough star	ndards for the assay and		
	9.4.7 <b>P</b> i	reparation	of Tumor Ce	ll Lysate Cont	rols (run in duplicate	)		
	9.4	.7.1 For lysa prov requ Sect	one 96-well pl te control vial ided at a conc ired. Vortex a ion 6C). See A	late, retrieve or from the -80°( centration ready nd mix by invo Appendix 3 for	ne of each High-C, Mid C freezer and thaw on id y for use in the assay ar erting 5-8 times before more information on p	-C and Low-C tumor ce. Controls are id no further dilution is use (Appendix 2, reparation of controls.		
	9.4	.7.2 Kee Buff	p controls on i fer (1X PBS-2	ice until use. C % BSA) once	Controls will be diluted loaded into the 96-well	3-fold with Assay plate.		
	9.4	.7.3 Note	e the Control I	Lot Numbers o	n the Assay Record (Aj	opendix 2, Section 6C).		







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#### 9.5 **Total H2AX Protein Capture**

9.5.1 Following incubation with SuperBlock (SOP Step 9.3.6), the plates are aspirated and washed once with 350 μL of Wash Buffer (1X PBS-0.1% Tween) using a plate washer.

For the BioTek Microplate Washer, the settings are:

METHOD	
Number of Cycles:	1
Soak/Shake:	Yes
Soak Time:	5 Sec
Shake before soak:	No
Prime after soak:	No
DISPENSE	
Dispense Volume:	350 μL/well
Dispense Flow Rate:	06
Dispense Height:	120 (15.240 mm)
Horizontal DISP POS:	00 (0.000 mm)
Bottom Wash First:	No
Prime Before Start:	No
ASPIRATE	
Aspirate Height:	031 (3.937 mm)*
Horizontal ASPR POS:	-20 (-0.914 mm)*
Aspiration Rate:	05 (6.4 mm/sec)
Aspirate Delay:	1000 MSec
Crosswise ASPIR:	No
Final Aspiration:	Yes
Final Aspirate Delay:	1000 MSec

\*Recommended initial setting, optimize Aspirate Height and Horizontal ASPR POS to optimize complete aspiration for an individual unit following manufacturer's recommendations.

- 9.5.2 After the wash, tap the plate on paper towels to remove residual buffer. Proceed immediately to the next step; do not allow the plate to dry out.
- 9.5.3 Immediately, add 50  $\mu$ L of Assay Buffer (1X PBS-2% BSA) containing 100 ng/well of Protamine to each well using a multichannel pipettor. Each well will hold a final volume of 75  $\mu$ L after sample addition.
- 9.5.4 Use the Plate Map Design (Section 6.2) and the Sample Calculation Table (Appendix 2, Section 6A) as a guide to set up the 96-well plate for incubation with samples, Total H2AX standards (Appendix 2, Section 6B), and assay controls (Appendix 2, Section 6C and Appendix 3). Pipette reagents/lysates in the following order; do not deviate from order of addition:





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Order	Sample/Reagent and Volume
1	25 µL of specified concentrations of Total H2AX standards into designated
	duplicate wells. Load the lowest concentration first.
2	25 μL of each unknown sample into designated triplicate wells.
3	25 µL each of assay control (Low-C, Mid-C, and High-C) into both sets of
	designated duplicate wells.
4	$25 \ \mu L$ of additional 1X PBS-2% BSA into each of the Background wells.

9.5.5 Cover the plate with an acetate sheet and incubate at  $2^{\circ}$ C to  $8^{\circ}$ C for  $18 \pm 2$  h. Record the date, start time, and incubation temperature in the Assay Record (Appendix 2, Section 7).

#### 10.0 DETECTION FOR BOTH γH2AX AND TOTAL H2AX ASSAYS (NEXT DAY)

#### 10.1 **Detection Antibody Incubation**

- 10.1.1 Prepare a sufficient amount of the rabbit detection pAb 15 min before washing the plate(s) after the capture incubation for the  $\gamma$ H2AX and/or Total H2AX assays.
- 10.1.2 Using the calculations in Appendices 1 and/or 2, Sections 8A, to prepare the rabbit detection pAb working solution in Assay Buffer (1X PBS-2% BSA). Be sure to record the lot number of mouse serum used.
- 10.1.3 Allow the prepared rabbit detection pAb to incubate for 15 min at  $25^{\circ}C \pm 3^{\circ}C$  and record this incubation in Appendices 1 and/or 2, Section 8A.
- 10.1.4 After the 18-h incubation of the coated plates with samples is complete, aspirate and wash the wells 4 times with 350 μL of Wash Buffer (1X PBS-0.1% Tween) using the same wash program as Step 8.7.1, except run for **4 cycles**. Record the date and time samples were removed from the wells in the Assay Record (Appendices 1 and/or 2, Section 7).
- 10.1.5 After the wash, tap the plate on paper towels to remove residual Wash Buffer. Proceed immediately to the next step; do not allow the plate to dry out.
- 10.1.6 Add 100  $\mu$ L of the rabbit detection pAb working solution per well using a multichannel pipettor, cover the plate with an acetate sheet, and incubate for 2 to 2.5 h at 25°C ± 3°C. Discard residual working solution and record the incubation conditions in the Assay Record (Appendices 1 and/or 2, Section 8B).
- 10.1.7 15 min before the incubation with the rabbit detection pAb is complete, prepare a sufficient amount of HRP conjugate for the assay. Using the calculations in Appendices 1 and/or 2, Sections 9A, prepare the HRP conjugate working solution in Assay Buffer.
- 10.1.8 Allow the prepared HRP conjugate to incubate in the dark at  $25^{\circ}C \pm 3^{\circ}C$  for 15 min and record the incubation conditions in the Assay Record (Appendices 1 and/or 2, Section 9A).
- 10.1.9 After the 2 to 2.5 h incubation with the rabbit detection pAb is complete, aspirate and wash the wells 4 times with 350  $\mu$ L of Wash Buffer (1X PBS-0.1% Tween) using the same wash program as SOP Step 8.7.1, except run for **4 cycles**. Tap plate on paper towels to remove residual liquid and proceed immediately to the next step.





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10.1.10 Add 100  $\mu$ L of the HRP conjugate working solution per well using a multichannel pipettor. Cover the plate with an acetate sheet and incubate in the dark for 1 to 1.5 h at 25°C ± 3°C. Discard residual working solution and record the incubation conditions in the Assay Record (Appendices 1 and/or 2, Section 9B).

#### 10.2 Signal Detection

10.2.1 Turn on the Tecan Infinite Plate Reader at least 30 min before use. For luminescence optical density readings, the plate reader should be set to the following reading parameters:

Shaking duration:	5 sec
Mode:	linear
Amplitude:	1 mm
Attenuation:	OD1
Integration Time:	100 ms

- 10.2.2 Just before the HRP conjugate incubation is finished, prepare SuperSignal ELISA Pico Chemiluminescent Substrate Solution as outlined in Appendices 1 and/or 2, Section 10A, being sure to note the time of preparation. This must be made up immediately before use, kept in the dark, and at a sufficient volume for the assay.
- 10.2.3 After the 1 to 1.5 h HRP conjugate incubation is complete, aspirate and wash the wells 4 times with 350  $\mu$ L of Wash Buffer (1X PBS-0.1% Tween) using the same wash program as SOP Step 8.7.1, except run for **4 cycles**. Tap plate on a paper towel to remove excess buffer and proceed immediately to the next step.
- 10.2.4 Add 100 µL of the freshly made Substrate Solution per well with a multichannel pipettor and avoid bright light. Record the time of addition to wells (Appendices 1 and/or 2, Section 10B).
- 10.2.5 The first chemiluminescence reading should be within 2 min of substrate addition. Record the time of the initial relative light unit (RLU) reading in the Assay Record (Appendices 1 and/or 2, Section 10B).
  - 10.2.5.1 Refer to Sections 4.3 or 4.4 for examples of expected RLU readout ranges for standards and tumor cell lysate controls for the  $\gamma$ H2AX and Total H2AX assays, respectively.
  - 10.2.5.2 If the signal is too high from the initial reading, wait 5 min and read the plate again at the same instrument setting. Continue reading until the RLU signal is on scale.
  - 10.2.5.3 Record time the final RLU reading is taken in Appendices 1 and/or 2, Section 10B.
- 10.2.6 Save the resulting readings.
- 10.2.7 The equation for the line (y=mx+b) of the standard curve can be used to convert RLU readings of the unknowns to  $\gamma$ H2AX and Total H2AX readings in pM.
- 10.3 Review and finalize the Assay Records (Appendices 1 and/or 2). Document ANY and ALL deviations from this SOP in the Assay Record (Appendices 1 and/or 2, Section 11).





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#### 11.0 QUALITY CONTROL RECOMMENDATIONS FOR YH2AX/TOTAL H2AX ASSAYS

#### 11.1 Background Well QC

- 11.1.1 From the "Plate Map Design" identify the background wells that are to be used in QC determination. A total of 14 wells are used for background determination; the 4 corner wells and 2 adjacent to the high standard are not used for background level calculation.
- 11.1.2 A  $\pm$  2 SD criterion is applied to the initial 14-well dataset to identify outliers.
- 11.1.3 If a background well RLU value is  $\geq$  2 SD from the mean, delete that value from the background data set.
- 11.1.4 Once all wells that were  $\geq 2$  SD from the initial background data set mean have been deleted, the %CV for the background wells must be < 20%.
- 11.1.5 If the %CV for the background wells is < 20%, the assay passes QC, proceed to Step 11.2.
- 11.1.6 If the %CV is  $\geq$  20%, the Assay Fails QC, do not continue with the analysis. State in the Assay Record (Appendices 1 and/or 2, Section 11) the reason for assay failure. Rerun the assay with fresh reagents.

#### 11.2 Standard Curve QC

11.2.1 Low Standard QC and LLQ Assignment for yH2AX Assay:

- 11.2.1.1 In order to use the 5 to 10 pM range of the standard curve, the mean RLU readout of the 5 pM standard must be  $\geq$  3 SD above the mean RLU readout of the background; this value is referred to as the LLQ-RLU.
- 11.2.1.2 If the 5 pM standard fails, then the mean RLU readout of the 10 pM standard must be  $\geq$  3 SD above the mean RLU readout of the background.
- 11.2.1.3 If the 10 pM standard also fails, the Assay Fails QC.
- 11.2.1.4 The lowest passing standard is assigned as the LLQ (pM) for the assay.
- 11.2.2 Signal-to-background (S/B) ratio QC and ULQ Assignment, yH2AX:
  - 11.2.2.1 The ratio for the lowest passing standard (5 or 10 pM) RLU readout to the mean RLU readout of the background must be  $\geq$  1.1. If not, the Assay Fails QC.
  - 11.2.2.2 The ratio of the highest standard RLU readout (640 pM) to the mean RLU of the background must be  $\geq$  15. If not, the Assay Fails QC.
  - 11.2.2.3 If the high standard passes QC, it is assigned as the ULQ (pM) for the assay.
- 11.2.3 Low Standard QC and LLQ Assignment, Total H2AX Assay:
  - 11.2.3.1 In order to use the 50 to 100 pM range of the standard curve, the mean RLU readout of the 50 pM standard must be  $\geq$  3 SD above the mean RLU readout of the background; this value is referred to as the LLQ-RLU.
  - 11.2.3.2 If the 50 pM standard fails, then the mean RLU readout of the 100 pM standard must be  $\geq$  3 SD above the mean RLU readout of the background.
  - 11.2.3.3 If the 100 pM standard also fails, the Assay Fails QC.





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- 11.2.3.4 The lowest passing standard is assigned as the LLQ (pM) for the assay.
- 11.2.4 Signal-to-background (S/B) ratio QC and ULQ Assignment, Total H2AX Assay:
  - 11.2.4.1 The ratio for the lowest passing standard (50 or 100 pM) RLU readout to the mean RLU readout of the background must be  $\geq$  1.1. If not, the Assay Fails QC.
  - 11.2.4.2 The ratio of the highest standard RLU readout (6400 pM) to the mean RLU of the background must be  $\geq$  15. If not, the Assay Fails QC.
  - 11.2.4.3 If the high standard passes QC, it is assigned as the ULQ (pM) for the assay.

#### 11.3 Control Samples

- 11.3.1 The QC determination for the control samples should have the following criteria:
  - 11.3.1.1 At least one control at each level (Low-, Mid-, and High-C) must have a CV of < 20% for the replicate wells.
  - 11.3.1.2 At least one control at each level and at least 4 of 6 controls overall must fall within the defined  $\gamma$ H2AX or Total H2AX pM range determined for the specific lot of critical reagent.
- 11.3.2 If any of these criteria are not met, the Assay Fails QC. State in the Assay Record (Appendices 1 and/or 2, Section 11) the reason for assay failure. Rerun the assay with fresh reagents.

#### 11.4 Unknown Sample Replicate QC and LLQ/ULQ QC

- 11.4.1 Triplicate repeats for each sample must have a CV < 20%.
- 11.4.2 Review the average  $\gamma$ H2AX or Total H2AX levels and identify any values that are < LLQ or > ULQ.
  - 11.4.2.1 If a sample is > ULQ and there is sufficient sample volume, it can be re-run with fresh reagents at a lower protein or cell number load/well. If a sample is < LLQ and there is sufficient sample volume, it can be re-run at a higher protein or cell number load/well.





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#### 12.0 REFERENCES

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- 3. Ji J, Zhang Y, Redon C, Chen A, Holbeck S, Pommier Y, Parchment R, Hollingshead M, Rubinstein L, Tomaszewski J, Doroshow J, Bonner, W. Gamma-H2AX and H2AX Quantitative ELISA for Monitoring DNA Damage Induced by Chemotherapeutic Agents or Irradiation. (2015) J Clin Oncol 33 (suppl; abstr 2559).
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- 5. SOP documents can be found on the DCTD Biomarkers website:
  - Landing Page: <u>http://dctd.cancer.gov/ResearchResources/ResearchResources-biomarkers.htm</u>
  - SOP340520 LINK: http://dctd.cancer.gov/ResearchResources/biomarkers/docs/par/SOP340520\_Biopsy\_Tissue.pdf
  - SOP340506 LINK:

https://dctd.cancer.gov/ResearchResources/biomarkers/docs/par/SOP340506\_PBMC\_Extraction. pdf





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## **APPENDIX 1: γH2AX ASSAY RECORD**

NOTE:

Record times using **military** time (24-h designation); for example, specify 16:15 to indicate 4:15 PM.

Assay Technician:

Date: \_\_\_\_\_\_
Plate ID (optional): \_\_\_\_\_

#### 1. Key Reagents/Supplies for γH2AX Assay

The reagents listed below are considered key to the success of a reproducible assay. Tracking performance of lots used in this RUO protocol and qualifying new reagent lots against the previous lots (suggested performance criteria  $\pm 25\%$  agreement) is recommended.

Reagent Name	Lot Number	Stock Concentration	Expiration Date
γH2AX Peptide Standard			
Tumor Lysate Controls: • High-C • Mid-C • Low-C			
γH2AX Mouse mAb			
H2AX Rabbit pAb			
Goat Anti-Rabbit HRP Conjugate			
Chemiluminescent Substrate: Pico Stable Peroxide and Luminol/Enhancer Solutions		N/A	
Reacti-Bind White Opaque 96-well Plate		N/A	N/A

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### 2. Preparation of Reagents for *γ*H2AX Assay

### A. <u>Reagents</u>

Buffers should be prepared based on volumes needed to complete all the steps with the number of 96-well plates in the experimental run. Always prepare at least 10% excess volume of buffer to ensure adequate volume to complete the study (scale-up appropriately for additional plates).

- i. <u>Coating Buffer</u>: Dissolve one capsule of Carbonate-Bicarbonate Buffer in 50 mL of deionized water yielding 0.1 M Carbonate-Bicarbonate Buffer, pH 9.6 final. For each 96-well plate (prepare enough for 110 wells), 11 mL coating buffer will be needed. Keep at 2°C to 8°C. Discard unused buffer at end of the experimental run.
- ii. <u>SuperBlock</u>: For one 96-well plate (preparing for 110 wells), pipette 40 mL SuperBlock into a 50 mL tube. Keep at 2°C to 8°C. Discard unused buffer at end of assay run.
- iii. <u>Wash Buffer (1X PBS-0.1% Tween)</u>: To prepare 1 L of buffer pipette 100 mL 10X PBS and 10 mL of 10% Tween 20 into 890 mL deionized water. Keep at  $25^{\circ}C \pm 3^{\circ}C$  for up to 1 wk.
- iv. <u>Assay Buffer/Diluent (1X PBS-2% BSA)</u>: To prepare 400 mL of assay diluent add 8 g BSA and 40 mL 10X PBS to 360 mL deionized water. Keep at 2°C to 8°C for up to 2 wks.
- v. <u>γH2AX Sample Diluent (CEB Complete)</u>: To prepare 10 mL CEB Complete add 500 μL of 20% SDS, 100 μL of 100 mM PMSF, 400 μL of 25X PI, one tablet of PhosSTOP to 9.0 mL CEB. Keep at 2°C to 8°C. Discard unused buffer at end of assay run.

### **3.** Capture Antibody Preparation for γH2AX Assay:

A. <u>Preparation of γH2AX mAb Coating Solution</u>

Remove antibody from -20  $^\circ C$  freezer and place on ice.

For one 96-well plate, prepare 110 wells:  $(100 \,\mu\text{L/well*110})/(1000 \,\mu\text{L/mL}) = 11 \,\text{mL}$ . Prepare  $\gamma$ H2AX mAb Coating Buffer using the following calculations:

i. Dilution of γH2AX mAb **STOCK** = 1:

e.g., γH2AX mAb STOCK recommended dilution for Lot# DAM1567248 is 1:250.

 $\frac{11 \text{ mL}}{\text{Recommended dilution of}} * 1000 \text{ } \mu\text{L/mL} = \underline{XX} \text{ } \mu\text{L} \text{ } \gamma\text{H2AX mAb} \text{ } \textbf{STOCK}$  $\frac{11 \text{ mL}}{2} \text{ } \gamma\text{H2AX mAb} \text{ } \textbf{STOCK}$ 

11 mL		
	* 1000 $\mu$ L/mL = $\mu$ L $\gamma$ H2AX mAb <b>STOCK</b>	
(dilution factor)		

- ii. Place the following in a 15-mL polypropylene tube and mix by inversion 5 to 8 times.
  - 11 mL Coating Buffer
  - μL γH2AX mAb **STOCK**

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B.	Incubation Conditions	s for Coating Plate	<u>2</u>					
	Add 100 $\mu$ L $\gamma$ H2AX mAb Coating Solution to each well, and incubate at 37°C for 2 h. Plates can also be coated overnight at 2°C to 8°C.							
	Date:	Start Time:	: Incub	ation Temp:	°C			
	Date:	Stop Time:	:					
4. Bloo Afte	<b>ck Step for γH2AX Assa</b> er the capture antibody inc bate at 37°C for 1 to 1.5 h	y pubation, aspirate t n (move to 25°C ±	he wells and add 3°C if blocking	l 250 μL SuperBlo longer).	ock to each well and			
Incu	bation conditions for bloc	king plate:						
Da	te: Start Ti	me: :	Incubation Te	emp:	°C			
Da	te: Stop Ti	me: :	_					
5. Pre	paration of Working Dil	utions of Unknov	vn Sample Lysa	tes for γH2AX As	ssay			
Nor	malize unknown lysates to	ο 0.2 μg/μL worki	ng dilution prior	to preparation of s	samples for the			
imm	iunoassay.	Stock Lysate	Working Lysa	te Vol Stock				
Sample	<u>SampleID</u>	<u>Conc.</u>	<u>Conc.</u>	Lysate	Vol. CEB Complete			
<u>110.</u>		xx μg/μL	0.2 μg/μL	(μL)	Lysate used)			
<b>S1</b>								
S2								
<b>S</b> 3								
<b>S4</b>								
<b>S</b> 5								
<b>S</b> 6								

ASSAY RECORD:

**S7** 

**S8** 

**S9** 

**S10** 

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#### 6. Preparation of Unknown Samples (A) and yH2AX Peptide Standards (B) for yH2AX Assay

#### A. <u>Unknown Sample Calculation Table</u>

Unknown samples are run in triplicate, 25  $\mu$ L sample/well (preparing 1 well extra). Sample numbers correspond to those on the Plate Map Design (Section 6.1). \*It may be necessary to further dilute certain cell line lysates with high  $\gamma$ H2AX in order to obtain a quantitative value in the range of the assay. In the event that a cell line provides values above the range of the assay (>ULQ), appropriately pre-dilute with CEB Complete and re-run.

All	All Samples Tumor/Tissue Samples			PBMC/Cell Samples			
		Working Lysate Conc.	Diluted Lys	sate (2, 1, and 0.5 µg	g/well)	Stock Cell Conc.	Stock Lysate Vol. (µL)
Sample Sample No. Name/ID	Sample Name/ID	0.2 μg/μL	Vol. Working Lysate (µL)	Vol. CEB Complete (100 μL - Vol. Lysate)	Final conc. (µg/well)	*1 x 10 <sup>7</sup> cells/mL	100 µL
<b>S1</b>		μg/μL			µg/well	cells/mL	
<b>S2</b>		μg/μL			µg/well	cells/mL	
<b>S</b> 3		μg/μL			µg/well	cells/mL	
<b>S4</b>		μg/μL			µg/well	cells/mL	
<b>S</b> 5		μg/μL			µg/well	cells/mL	
<b>S6</b>		μg/μL			µg/well	cells/mL	
<b>S7</b>		μg/μL			µg/well	cells/mL	
<b>S8</b>		μg/μL			µg/well	cells/mL	
<b>S</b> 9		μg/μL			µg/well	cells/mL	
S10		μg/μL			µg/well	cells/mL	
S11		μg/μL			µg/well	cells/mL	
S12		μg/μL			µg/well	cells/mL	
S13		μg/μL			µg/well	cells/mL	
S14		μg/μL			µg/well	cells/mL	
S15		μg/μL			µg/well	cells/mL	
S16		μg/μL			µg/well	cells/mL	

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#### B. <u>γH2AX Peptide Standards</u>

Calculations for preparation of the 1920 pM  $\gamma$ H2AX standard in tube #1.

Supplied  $\gamma$ H2AX peptide standard = \_\_\_\_\_ pM

e.g.,  $\gamma$ H2AX peptide standard **Stock**  $\gamma$ H2AX JJ2, Lot# 113011 is supplied at 20680 pM (64 ng/ml).



Serial dilutions of the  $\gamma$ H2AX peptide standards are used to prepare the remaining tubes with concentrations ranging from 960 to 15 pM in Assay Buffer (1X PBS-2% BSA). 25 µL of each diluted standard will be added to the standard wells in the 96-well plate containing 50 µL of Assay Buffer (1X PBS-2% BSA) to give a 3-fold dilution which generates a reference standard curve ranging from 640 to 5 pM  $\gamma$ H2AX standards. Label tubes with final concentration of standard.

Tube # (Plate Row)	Vol. and Source of Concentrated Standard	Vol. 1X PBS- 2% BSA	Resulting Conc. of Diluted Standard	Conc. of Standard in Plate (1:3 Dilution)
1 (H)	μL of γH2AX Standard Stock Solution	μL (bring to 200 μL)	1920 pM	640 pM
2 (G)	100 $\mu$ L of tube #1	100 µL	960 pM	320 pM
3 (F)	100 $\mu$ L of tube #2	100 µL	480 pM	160 pM
4 (E)	100 $\mu$ L of tube #3	100 µL	240 pM	80 pM
5 (D)	100 $\mu$ L of tube #4	100 µL	120 pM	40 pM
6 (C)	100 $\mu$ L of tube #5	100 µL	60 pM	20 pM
7 (B)	100 $\mu$ L of tube #6	100 µL	30 pM	10 pM
8 (A)	100 µL of tube #7	100 µL	15 pM	5 pM
9 (Blank)	0 μL	600 μL	0 pM	0 pM (0 pM)

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#### C. <u>Controls (see Appendix 3 for preparation)</u>

The High-, Mid- and Low-C tumor lysate controls for the  $\gamma$ H2AX assay are prepared from 1 x 10<sup>7</sup> cells/mL stock lysates of irradiated MCF7 and untreated MCF7 cultured tumor cell lines. Controls will be diluted an additional 3-fold when added to the 96-well plate.

High-C: irradiated MCF7 lysate at  $1 \times 10^7$  cells/mL.

Mid-C: mix 30% High-C and 70% Low-C.

Low-C: untreated MCF7 lysate 1 x  $10^7$  cells/mL.

Controls are aliquoted for single use, thaw only what is needed for the run.

Note the Control Numbers below:

High-C:			
Mid-C:			
Low-C:			

#### 7. Plate Incubation for γH2AX Assay

Add 25  $\mu$ L unknown samples, tumor controls, and  $\gamma$ H2AX peptide standards to the 96-well plate (wells contain 50  $\mu$ L 1X PBS-2% BSA), cover plate, and incubate at 2°C to 8°C for 18  $\pm$  2 h.

 Date:
 Start Time:
 :
 Incubation Temp:
 °C

 Date:
 Stop Time:
 :
 :

#### 8. Detection Antibody: H2AX Rabbit pAb

#### A. <u>Preparation of H2AX Rabbit pAb Working Solution (100 µL/well)</u>

Remove antibody from -20°C freezer and place on ice.

For one 96-well plate, prepare 110 wells:  $(100 \,\mu\text{L/well*110})/(1000 \,\mu\text{L/mL}) = 11 \,\text{mL}$ . Prepare **H2AX Rabbit pAb Working Solution** using the following calculations:

- i. Dilution of H2AX Rabbit pAb **STOCK** = 1: \_\_\_\_\_
- ii. e.g., H2AX pAb **STOCK** recommended dilution for Lot# 778559 is 1:850

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	11 Recommend H2AX R ST	mL ed dilution of abbit pAb OCK	<sup></sup> * 1000 μL	$/mL = XX \mu L H2AX$	K Rabbit pAb <b>STOCK</b>	
	(dilutio	mL on factor)	* 1000 μL	$mL = - \mu L H2$	2AX Rabbit pAb	
	iii. Place the follow	wing in a 15-r	nL polypropyl	ene tube:		
	11 mL 11 μL μL	1X PBS-2% Mouse serui H2AX Rabb	BSA m (1:1000) L pit pAb <b>STOCK</b>	Lot #:		
	iv. Mix by inversi	on 5 to 8 time	es, and let stand	d at $25^{\circ}C \pm 3^{\circ}C$ for 15 i	nin before use.	
	Start Time:	:	Stop Time:	: Incubation	Temp: <u>°C</u>	
<ul> <li>B. <u>Addition of Prepared H2AX Rabbit pAb Working Solution</u></li> <li>Add 100 µL of the H2AX <b>Rabbit pAb Working Solution</b> to each well and incubate for 2 to</li> </ul>						
	2.5 h at $25^{\circ}C \pm 3^{\circ}C$ .	<b>F</b>	~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~			
	Start Time: :	Stop Ti	me: :	Incubation Temp:	°C	

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#### 9. Reporter: HRP Conjugate for yH2AX Assay

A. <u>Preparation of HRP Conjugate Working Solution (100 µL/well)</u>

For one 96-well plate, prepare 110 wells:  $(100 \ \mu L/well*110)/(1000 \ \mu L/mL) = 11 \ mL$ . Prepare **HRP Conjugate Working Solution** using the following calculations:

- i. Recommended dilution of Goat Anti-Rabbit HRP Conjugate **STOCK** = 1: \_\_\_\_\_
- ii. e.g., HRP Conjugate **STOCK** recommended dilution for Lot# 101008 is 1:1000

 $\frac{11 \text{ mL}}{\text{Recommended dilution of}} * 1000 \text{ }\mu\text{L/mL} = \underline{XX} \text{ }\mu\text{L} \text{ HRP Conjugate STOCK}$   $\frac{11 \text{ mL}}{\text{HRP Conjugate STOCK}} = \frac{1000 \text{ }\mu\text{L/mL}}{1000 \text{ }\mu\text{L/mL}} = \frac{1000 \text{ }\mu\text{L}}{1000 \text{ }\mu\text{L}}$ 

11 mL				
	*	$1000 \ \mu L/mL$	=	 _ μL HRP Conjugate STOCK
(dilution factor)				

iii. Place the following in a 15-mL polypropylene tube:

11 mL	1X PBS-2% BSA		
11 µL	Mouse serum (1:1000)	Lot #:	
μL	HRP Conjugate STOCK		

iv. Mix by inversion 5 to 8 times, and let stand at  $25^{\circ}C \pm 3^{\circ}C$  for 15 min before use.

Start Time: : Stop Time: : Incubation Temp: °C

B. Addition of HRP Conjugate Working Solution

Add 100  $\mu$ L of the **HRP Conjugate Working Solution** to each of the washed wells and incubate in the dark for 1 to 1.5 h at 25°C ± 3°C.

Start Time: : Stop Time: : Incubation Temp: °C

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## **10.** Chemiluminescent Substrate for γH2AX Assay

#### A. <u>Preparation of Substrate Solution (100 µL/well)</u>

Calculate volume of substrate required for the experimental run. For one 96-well plate, prepare 110 wells:  $(100 \,\mu\text{L/well*110})/(1000 \,\mu\text{L/mL}) = 11 \,\text{mL}$ . Immediately before washing the plate, prepare the following in a 15-mL polypropylene tube wrapped with aluminum foil. Mix by inversion 5 to 8 times and keep at  $25^{\circ}\text{C} \pm 3^{\circ}\text{C}$  in the dark until use.

5.5 mL Pico Stable Peroxide	(50 µL/well*110)/ (1000 µL/mL)
5.5 mL Pico Luminol/Enhancer	(50 µL/well*110)/ (1000 µL/mL)

Time of Substrate Preparation:

## :

#### B. <u>Substrate Solution Incubation and RLU Reading Times</u>

Time of Substrate Addition to Wells:	:
Time Initial RLU Reading is Captured:	:
Time Final RLU Reading is Captured (opt):	:

#### 11. Notes, including any deviations from the SOP:

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## **APPENDIX 2: TOTAL H2AX ASSAY RECORD**

NOTE:

Record times using military time (24-h designation); for example, specify 16:15 to indicate 4:15 PM.

Assay Technician:

Date:

Plate ID (optional):

#### 1. Key Reagents/Supplies for Total H2AX Assay

The reagents listed below are considered key to the success of a reproducible assay. Tracking performance of lots used in this RUO protocol and qualifying new reagent lots against the previous lots (suggested performance criteria  $\pm 25\%$  agreement) is recommended.

Reagent Name	Lot Number	Stock Concentration	Expiration Date
Total H2AX Recombinant Standard			
Tumor Lysate Control:			
High-C			
Mid-C			
Low-C			
Total H2AX Mouse mAb			
H2AX Rabbit pAb			
Goat Anti-Rabbit HRP Conjugate			
Chemiluminescent Substrate: Pico Stable Peroxide and Luminol/Enhancer Solutions		N/A	
Reacti-Bind White Opaque 96-well Plate		N/A	N/A

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#### 2. Preparation of Reagents for Total H2AX Assay

#### A. <u>Reagents</u>

Buffers should be prepared based on volumes needed to complete all the steps with the number of 96-well plates in the experimental run. Always prepare at least 10% excess volume of buffer to ensure adequate volume to complete the study (scale-up appropriately for additional plates).

- i. <u>Coating Buffer</u>: Dissolve one capsule of Carbonate-Bicarbonate Buffer in 50 mL of deionized water yielding 0.1 M Carbonate-Bicarbonate Buffer, pH 9.6 final. For each 96-well plate (prepare enough for 110 wells), 11 mL coating buffer will be needed. Keep at 2°C to 8°C. Discard unused buffer at end of the experimental run.
- ii. <u>SuperBlock</u>: For one 96-well plate (preparing for 110 wells), pipette 40 mL SuperBlock into a 50-mL tube. Keep at 2°C to 8°C. Discard unused buffer at end of assay run.
- iii. <u>Wash Buffer 1X PBS- 0.1% Tween</u>: To prepare 1 L of buffer pipette 100 mL 10X PBS and 10 mL of 10% Tween 20 into 890 mL deionized water. Keep at  $25^{\circ}C \pm 3^{\circ}C$  for up to 1 wk.
- iv. <u>Assay Buffer 1X PBS-2% BSA</u>: To prepare 400 mL of buffer add 8 g BSA and 40 mL 10X PBS to 360 mL deionized water. Keep at 2°C to 8°C for up to 2 wks.
- v. <u>Sample Diluent Buffer–CEB Complete:</u> To prepare 10 mL CEB Complete add 500 μL of 20% SDS, 100 μL of 100 mM PMSF, 400 μL of 25X PI, one tablet of PhosSTOP to 9.0 mL CEB. Keep at 2°C to 8°C. Discard unused buffer at end of assay run.
- vi. <u>Assay Buffer (1X PBS-2% BSA) containing 2 μg/mL (100 ng/well) of Protamine</u>: To prepare 6 mL of buffer add 20 μL of 600 μg/mL Protamine working stock and 5980 μL of 1X PBS-2% BSA. Keep at 2°C to 8°C. The final concentration used in the assay is 100 ng/well. Discard unused buffer at end of the experimental run.

Note: To make 600  $\mu$ g/mL Protamine stock: dilute 120 mg of Protamine powder in 2 mL of dH<sub>2</sub>O, this gives 60 mg/mL stock. Make two additional 10X serial dilution (1:100 dilution) with dH<sub>2</sub>O to generate 600  $\mu$ g/mL working stock.

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#### 3. Capture Antibody: Total H2AX mAb

#### A. <u>Preparation of Total H2AX mAb Coating Solution</u>

Remove antibody from -20°C freezer and place on ice.

For one 96-well plate, prepare 110 wells:  $(100 \,\mu\text{L/well*110})/(1000 \,\mu\text{L/mL}) = 11 \,\text{mL}$ . Prepare **Total H2AX mAb Coating Solution** using the following calculations:

#### i. Dilution of Total H2AX mAb **STOCK** = 1:

e.g., Total H2AX mAb  ${\bf STOCK}$  recommended dilution for Lot# 09281-4F10 (1mg/mL) is 1:250.

 $\frac{11 \text{ mL}}{\text{Recommended dilution of}} * 1000 \text{ }\mu\text{L/mL} = \underline{XX} \text{ }\mu\text{L} \text{ Total H2AX mAb } \text{STOCK}$ 

 $\frac{11 \text{ mL}}{(\text{dilution factor})} * 1000 \ \mu\text{L/mL} = \___ \mu\text{L} \text{ Total H2AX mAb STOCK}$ 

ii. Place the following in a 15-mL polypropylene tube and mix by inversion 5 to 8 times.

11 mL	Coating Buffer
μL	Total H2AX mAb STOCK

#### B. Incubation Conditions for Coating Plate

Add 100  $\mu$ L **Total H2AX mAb Coating Solution** to each well, and incubate at 37°C for 2 h. Plates can also be coated overnight at 2°C to 8°C.

 Date:
 Start Time:
 Incubation Temp:
 °C

Date: \_\_\_\_\_ Stop Time: \_\_\_\_\_

#### 4. Block Step for Total H2AX Assay

After the capture antibody incubation, aspirate wells, and place 250  $\mu$ L SuperBlock in each well and incubate at 37°C for 1 to 1.5 h (move to 25°C ± 3°C if blocking longer).

Incubation conditions for blocking plate:

 Date:
 Start Time:
 :
 Incubation Temp:
 °C

 Date:
 Stop Time:
 :

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#### 5. Preparation of Working Dilutions of Unknown Tumor/Tissue Lysates for Total H2AX Assay

Normalize unknown biopsy lysates to 0.75  $\mu$ g/ $\mu$ L working dilution with CEB Complete if the unknown stock lysate concentrations are  $\geq$  0.75  $\mu$ g/ $\mu$ L prior to preparation of samples for the immunoassay. Then dilute the 0.75  $\mu$ g/ $\mu$ L normalized working lysate or 0.2 to 0.75  $\mu$ g/ $\mu$ L stock lysates to 0.1  $\mu$ g/ $\mu$ L with 1X PBS-2% BSA.

<u>Sample</u> <u>No.</u>	Sample ID	<u>Stock Lysate</u> <u>Conc.</u>	$\frac{Prediluted}{Lysate Conc.}$ (Only for $\geq$ 0.75 <u>µg/µL</u> )	$\frac{Vol. Stock}{Lysate (Only for}$ $\frac{for \ge 0.75}{\mu g/\mu L)}$	$\frac{\text{Vol. CEB}}{\text{Complete (Only}}$ $\frac{\text{for} \ge 0.75}{\mu g/\mu L)}$	<u>Vol. 0.2 μg/μL to</u> <u>0.75 μg/mL</u> <u>Stock or</u> <u>Prediluted</u> <u>Lysate</u>	<u>Diluted</u> <u>Working Lysate</u> <u>Conc.</u>	<u>Vol. 1X PBS-2%</u> <u>BSA</u>
		xx μg/μL	0.75 μg/μL or N/A	(µL)	(10 µL - Vol. Stock Lysate)	(10 μL or XX μL)	0.1 µg/µL	(75 μL - Vol. 0.1 μg/μL Lysate)
<b>S1</b>								
S2								
<b>S</b> 3								
<b>S4</b>								
<b>S</b> 5								
<b>S</b> 6								
<b>S7</b>								
<b>S8</b>								
<b>S9</b>								
S10								

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#### 6. Preparation of Unknown Samples (A), Total H2AX Recombinant Standards (B) for Total H2AX Assay

#### A. <u>Unknown Sample Calculation Table:</u>

Unknown samples are run in triplicate, 25 µL sample/well (preparing 1 well extra). Sample numbers correspond to those on the Plate Map.

All Samples			Tumor/	<b>Fissue Samples</b>	PBMC/Cell Samples			
Sample		Working Lysate Conc.	Diluted L	ysate (0.5, 0.375 and 0.25	µg/well)	Stock Cell Conc.	Stock Lysate Vol. (µL)	Vol. 1X PBS- 2% BSA (µL)
No.	Sample ID	0.1 μg/μL	Vol. Working Lysate (µL)	<b>Vol. 1X PBS-2% BSA</b> (100 μL - Vol. Lysate)	Final conc. (µg/well)	1 x 10 <sup>7</sup> cells/mL	7.5 μL	117.5 μL
<b>S1</b>		μg/μL			µg/wel	cells/mL		
<b>S2</b>		μg/µL			µg/wel	cells/mL		
<b>S</b> 3		μg/μL			µg/wel	cells/mL		
<b>S4</b>		μg/μL			µg/wel	cells/mL		
<b>S</b> 5		μg/μL			µg/wel	cells/mL		
<b>S6</b>		μg/μL			µg/wel	cells/mL		
<b>S7</b>		μg/μL			µg/wel	cells/mL		
<b>S8</b>		μg/μL			µg/wel	cells/mL		
<b>S9</b>		μg/μL			µg/wel	cells/mL		
S10		μg/μL			µg/wel	cells/mL		
S11		μg/μL			µg/wel	cells/mL		
S12		μg/μL			µg/wel	cells/mL		
S13		μg/μL			µg/wel	cells/mL		
S14		μg/μL			µg/wel	cells/mL		
S15		μg/μL			µg/wel	cells/mL		
S16		μg/μL			µg/wel	cells/mL		

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#### B. <u>Total H2AX Recombinant Protein Standards</u>

Calculations for preparation of the 19200 pM Total H2AX standard in tube #1.

Supplied Total H2AX recombinant protein standard = \_\_\_\_\_ pM

e.g., Total H2AX standard Stock Lot# L16677 is supplied at 33000 pM (500 ng/ml)

19200 pM					
Conc, of Total H2AX standard Stock (pM)	)	*	200 µL	=	<u>XX</u> μL Total H2AX standard Stock solution in 200 μL final

$$\left(\begin{array}{c} \underline{19200 \text{ pM}} \\ \underline{\phantom{0}} \end{array}\right) * 200 \,\mu\text{L} = \underline{\phantom{0}} \mu\text{L} \text{ total H2AX standard Stock} \\ \text{solution in } \underline{200 \,\mu\text{L}} \text{ final} \end{array}$$

Serial dilutions of the Total H2AX recombinant standards are used to prepare the remaining tubes with concentrations ranging from 9600 to 150 pM in 1X PBS-2% BSA. 25  $\mu$ L of each diluted standard will be added to the standard wells in the 96-well plate containing 50  $\mu$ L of Assay Buffer (1X PBS-2% BSA) with 2  $\mu$ g/mL Protamine (100 ng/well of Protamine at 50  $\mu$ L) which equals a 3-fold dilution of standards to generate a reference standard curve ranging from 6400 to 50 pM Total H2AX standards. Label tubes with final concentration of standard.

Tube # (Plate Row)	Vol. and Source of Concentrated Standard	Vol. 1X PBS-2% BSA	Resulting Conc. of Diluted Standard	Conc. of Standard in Plate (1:3 Dilution)
1 (H)	µL of Stock Solution	μL	19200 pM	6400 pM
2 (G)	100 $\mu$ L of tube #1	100 µL	9600 pM	3200 pM
3 (F)	100 $\mu$ L of tube #2	100 µL	4800 pM	1600 pM
4 (E)	100 µL of tube #3	100 µL	2400 pM	800 pM
5 (D)	100 $\mu$ L of tube #4	100 µL	1200 pM	400 pM
6 (C)	100 $\mu$ L of tube #5	100 µL	600 pM	200 pM
7 (B)	100 $\mu$ L of tube #6	100 µL	300 pM	100 pM
8 (A)	100 µL of tube #7	100 µL	150 pM	50 pM
9 (Blank)	0 µL	600 µL	0 pM	0 pM

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#### C. <u>Controls (see Appendix 3 for preparation)</u>

Controls for the Total H2AX are prepared as specified in the table below and are aliquoted for single use. Thaw control amounts as needed for the run.

Control Tube Dilution	Vol. and Source of Control Lysate	Vol. 1X PBS-2% BSA	Conc. of Control in Plate (1:3 Dilution)
High-C (1:16.7) *1 x 10 <sup>7</sup> cells/mL stock lysate	7.5 μL of SN12 C Tumor Cell Stock	117.5µL	High-C (1:50)
Mid-C (1:16.7) *1 x 10 <sup>7</sup> cells/mL stock lysate	$\underline{9}$ µL of MCF7 Tumor Cell Stock	141 μL	Mid-C (1:50)
Low-C (1:66.7)	<u><b>30</b></u> µL of Mid-C (MCF7) (1:50)	90 µL	Low-C (1:200)

Note the Control Lot Numbers below:

#### 7. Plate Incubation for Total H2AX Assay

Add 25  $\mu$ L unknown samples, tumor controls, and Total H2AX standards to the 96-well plate to wells containing 50  $\mu$ L 1X PBS-2% BSA - 2  $\mu$ g/mL Protamine (100 ng/well of Protamine), cover plate, and incubate at 2°C to 8°C for 18  $\pm$  2 h.

 Date:
 Start Time:
 Incubation Temp:
 °C

 Date:
 Stop Time:
 :

#### 8. Detection Antibody: H2AX Rabbit pAb

A. <u>Preparation of H2AX Rabbit pAb Working Solution (100 µL/well)</u>

Remove antibody from -20°C freezer and thaw on ice.

For one 96-well plate, prepare 110 wells:  $(100 \,\mu\text{L/well*110})/(1000 \,\mu\text{L/mL}) = 11 \,\text{mL}$ . Prepare **H2AX Rabbit pAb Working Solution** using the following calculations:

- i. Dilution of H2AX Rabbit pAb **STOCK** = 1: \_\_\_\_\_
- ii. e.g., total H2AX pAb **STOCK** recommended dilution for Lot# 778559 is 1:850.

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	11 Recommend H2AX R STO	mL ed dilution of abbit pAb DCK	· * 1000 μL	$mL = XX \mu L H2AY$	K Rabbit pAb <b>STOCK</b>		
	(dilutio	mL n factor)	- * 1000 μL	$mL = \frac{\mu L H^2}{STOCK}$	2AX Rabbit pAb		
	iii. Place the follow	ving in a 15-r	nL polypropyl	ene tube:			
	11 mL 11 μL μL	1X PBS-2% Mouse serui H2AX Rabb	BSA m (1:1000) L bit pAb <b>STOCK</b>	Lot #:			
	iv. Mix by inversion 5 to 8 times, and let stand at $25^{\circ}C \pm 3^{\circ}C$ for 15 min before use.						
	Start Time:	:	Stop Time:	: Incubation	Temp: <u>°C</u>		
В.	Addition of Prepared H Add 100 $\mu$ L of the H2A 2.5 h at 25°C ± 3°C.	2AX Rabbit <u>-</u> AX <b>Rabbit p</b> A	pAb Working ( Ab Working S	Solution Solution to each well an	id incubate for 2 to		
	Start Time: :	Stop Ti	me: :	Incubation Temp:	<u>°C</u>		

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#### 9. Reporter: HRP Conjugate for Total H2AX Assay

A. <u>Preparation of HRP Conjugate Working Solution (100 µL/well)</u>

For one 96-well plate, prepare 110 wells:  $(100 \,\mu\text{L/well*110})/(1000 \,\mu\text{L/mL}) = 11 \,\text{mL}$ . Prepare **HRP Conjugate Working Solution** using the following calculations:

- i. Recommended dilution of Goat Anti-Rabbit HRP Conjugate **STOCK** = 1: \_\_\_\_\_
- ii. e.g., HRP Conjugate **STOCK** recommended dilution for Lot# 120504 is 1:1000

11 mL Recommended dilution of \* 1000  $\mu$ L/mL = <u>XX</u>  $\mu$ L HRP Conjugate **STOCK** HRP Conjugate STOCK 11 mL \* 1000  $\mu$ L/mL = \_\_\_\_  $\mu$ L HRP Conjugate **STOCK** (dilution factor) iii. Place the following in a 15-mL polypropylene tube: 1X PBS-2% BSA 11 mL 11 µL Mouse serum (1:1000) Lot #: HRP Conjugate STOCK \_\_\_\_μL Mix by inversion 5 to 8 times, and let stand at  $25^{\circ}C \pm 3^{\circ}C$  for 15 min before use. iv. Start Time: : Stop Time: : Incubation Temp: °C Addition of HRP Conjugate Working Solution B. Add 100 µL of the HRP Conjugate Working Solution to each of the washed wells and incubate in the dark for 1 to 1.5 h at  $25^{\circ}C \pm 3^{\circ}C$ . : Stop Time: : Incubation Temp: Start Time: °C

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### 10. Chemiluminescent Substrate Total H2AX Assay

#### A. <u>Preparation of Substrate Solution (100 µL/well)</u>

Calculate volume of substrate required for the experimental run. For one 96-well plate, prepare 110 wells:  $(100 \ \mu L/well*110)/(1000 \ \mu L/mL) = 11 \ mL$ . Immediately before washing the plate, prepare the following in a 15-mL polypropylene tube wrapped with aluminum foil. Mix by inversion 5 to 8 times and keep at  $25^{\circ}C \pm 3^{\circ}C$  in the dark until use.

5.5 mL Pico Stable Peroxide	(50 μL/well*110)/ (1000 μL/mL)
5.5 mL Pico Luminol/Enhancer	(50 µL/well*110)/ (1000 µL/mL)

Time of Substrate Preparation:

# 

#### B. <u>Substrate Solution Incubation and RLU Reading Times</u>

Time of Substrate Addition to Wells:	:
Time Initial RLU Reading is Captured:	<u> </u>
Time Final RLU Reading is Captured (opt):	:

#### 11. Notes, including any deviations from the SOP:

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## **APPENDIX 3: PREPARATION OF TUMOR LYSATE CONTROL SAMPLES**

### 1.0 MATERIALS AND REAGENTS

- 1.1 Pipettors (200-1000 µL) and tips
- 1.2 Automatic pipettor
- 1.3 Vacuum filter/storage bottle system, 0.22-µm pore, 500 mL (e.g., Corning, Cat#: 430758).
- 1.4 1-, 5-, 10-, and 25-mL pipettes, sterile, individually wrapped (e.g., Fisher Scientific, Cat#: 13-675-15C, 13-675-22, 13-675-20, and 13-668-2)
- 1.5 1.5-mL Sarstedt tubes (e.g., Sarstedt, Cat#: 72.692.005)
- 1.6 15-mL polypropylene tubes (e.g., Fisher Scientific, Cat#: 14-959-49B or Becton Dickinson, Cat#: 352097)
- 1.7 50-mL polypropylene tubes (e.g., Becton Dickinson, Cat#: 352098)
- 1.8 Cell culture flask, 75 cm<sup>2</sup>, vent cap (T75; e.g., Corning, Cat#: 3290)
- 1.9 Ice bucket
- 1.10 100% Ethanol
- 1.11 MCF7 human breast adenocarcinoma (ATCC, Cat#: HTB-22)
- 1.12 SN12C human renal cancer cell (NCI)
- 1.13 10X phosphate buffered saline (PBS; e.g., Invitrogen, Cat#: 70013-073) [Dilute 1:10 in DI water to prepare 1X PBS for use in assay]
- 1.14 RPMI-1640 medium, 500-mL bottles (e.g., Invitrogen, Cat#: 22400089)
- 1.15 L-Glutamine, 200 mM (e.g., Invitrogen, Cat#: 25030164)
- 1.16 Fetal bovine serum, 500-mL bottles (FBS; Gemini Bio-Products, Cat#: 100-106) [Store at -20°C as 50-mL aliquots in 50-mL polypropylene tubes]
- 1.17 Trypsin Solution, 0.25%, 1X with EDTA (e.g., Invitrogen, Cat#: 25200056)
- 1.18 Cell Extraction Buffer (CEB; Invitrogen, Cat#: FNN0011)
- 1.19 20% sodium dodecyl sulfate (SDS; e.g., Sigma-Aldrich, Cat#: 05030-500ML-F)
- 1.20 Phenylmethanesulfonyl fluoride (PMSF) (Sigma-Aldrich, Cat#: 93482-50ML-F)
- 1.21 Protease Inhibitor (PI) Cocktail (Sigma-Aldrich, Cat#: P-2714 or Roche, Cat#: 11 697 498 001)
- 1.22 PhosSTOP, phosphatase inhibitor cocktail tablets (Roche, Cat#: 04906837001)
- 1.23 Liquid nitrogen or dry ice/ethanol bath
- 1.24 Hemocytometer and cover slips or an automated cell counter (e.g., Vi-Cell)
- 1.25 100°C heat block or boiling water bath
- 1.26 Vortex Genie 2 (Daigger, Cat#: 3030A)
- 1.27 Table-top centrifuge with a swing-bucket rotor, refrigerated (e.g., Sorvall Legend RT centrifuge (Fisher Scientific) with a Swing Bucket Rotor (Fisher Scientific, Cat#: 75-006-434) and manufacturer-recommended tube adaptors
- 1.28 Sorvall Fresco microcentrifuge (Fisher Scientific)
- 1.29 Class II Type A2 biosafety cabinet/tissue culture laminar flow hood
- 1.30 37°C tissue culture incubator, humidified, 5% CO<sub>2</sub>,
- 1.31 37°C water bath
- 1.32 -20°C and -80°C freezers





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#### 2.0 PROTOCOL

2.1 <u>Prepare Tissue Culture Medium</u>

*Note: Growth medium can be prepared ahead of time and stored at* 4°*C for up to 3 weeks.* 

- 2.1.1 Using sterile technique, prepare complete RPMI-1640 growth medium (RPMI-1640 + 10% FBS + 1% L-glutamine).
- 2.1.2 For each 500-mL bottle of RPMI-1640 to be prepared, thaw a 50-mL aliquot of FBS and a 5-mL aliquot of 200 mM L-glutamine in a 37°C water bath and then move to a laminar flow hood. Clean the outside of tubes with 70% ethanol.
- 2.1.3 Using a disposable filter unit and sterile technique, filter 445 mL RPMI-1640, 50 mL FBS, and 5 mL of 200 mM L-glutamine.
- 2.1.4 Label bottle with lot number and expiration date of medium components (if applicable) and date of preparation. Store at 4°C until use.

#### 2.2 <u>Thawing Frozen MCF7 or SN12C Cells</u>

- 2.2.1 Pre-warm growth medium in a 37°C water bath for 15 to 20 min.
- 2.2.2 Frozen cell stocks (~1 mL) should be quickly thawed in a 37°C water bath until just a little bit of ice remains. Immediately transfer the vial to a laminar flow hood, wipe the exterior with ethanol, and transfer into a 15 mL polypropylene tube containing 9 mL growth medium.
- 2.2.3 Gently pellet cells by centrifugation at 200 x g in a swing-bucket rotor at 4°C for 10 min. Aspirate and discard the supernatant.
- 2.2.4 Gently resuspend the cell pellet in 10 mL growth medium and transfer into a T75 flask containing an additional 15 mL growth medium (25 mL total volume).
- 2.2.5 Label the flask with the date and passage number of the cells and place in 37°C, 5% CO<sub>2</sub>, humidified incubator.
- 2.2.6 Cultured cells should become confluent with 4 to 5 days.

### 2.3 <u>Subculture MCF7 or SN12C Cells</u>

Note: Cells maintained in tissue culture should be regularly split to maintain overall cell line health. Cells should be cultured for no longer than <u>30 passages</u> before preparing a new culture with a fresh vial of cells.

- 2.3.1 Pre-warm growth medium in a 37°C water bath and trypsin at RT for 15 to 20 min.
- 2.3.2 In a laminar flow hood, aspirate old medium and add 5 mL RT trypsin solution. Gently rock the container to get complete coverage of the cells. Leave at RT for 10 to 30 sec.
- 2.3.3 Carefully, aspirate and discard the trypsin without disturbing the cells and place flask in a  $37^{\circ}$ C, 5% CO<sub>2</sub>, humidified incubator for 30 to 60 sec.
- 2.3.4 Tap flask by hand to release cells from the bottom and then add 10 mL of growth medium to the flask. Pipette up and down 2 to 3 times to disperse cell clumps.





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	2.3.5	2.3.5 Prepare 5 new T75 flasks to split the current MCF7 cell culture by adding 23 mL gromedium to each flask and labeling the flask with the date and the new passage number the cells. Add 2 mL of cell suspension from previous step into each flask.				
	2.3.6	Incubate flasks with 4 to 5 day	s in a 37°C, 59 ys.	% CO <sub>2</sub> , humid	ified incubator. Cells sl	hould become confluent
	2.3.7	2.3.7 For γH2AX induction to create the High γH2AX Control (High-C), MCF-7 cells are irradiated at 10 Gy and returned to the 37°C, 5% CO <sub>2</sub> incubator for 30 min prior to harvest.				
2.4	Harves	t Cultured MCF	7 or SN12C 0	Cells		
	2.4.1	On the day of o SOP340024, fi	cell harvest fo ve T75 flasks	or preparation of cells should	of the Tumor Cell Contr d be 80% to 90% conflu	ol Lysates for ient.
	2.4.2	Pre-warm grow 20 min.	wth medium in	n a 37°C water	bath and trypsin and 12	X PBS at RT for 15 to
	2.4.3	In a laminar flo monolayer sur	ow hood, aspi face once with	rate old mediu h 10 mL RT 12	m from all five T75 flas X PBS.	sks and wash the cell
	2.4.4	Aspirate PBS, complete cove	add 5 mL RT rage of the ce	trypsin to each lls. Leave at F	h flask. Gently rock the T for 2 to 3 min.	e container to get
	2.4.5	Carefully, aspi incubate for an to detach from	rate and disca additional 2 flask bottom.	rd the trypsin to 3 min at RT	without disturbing the c . Observe the cells, the	ells and let flask monolayer will begin
	2.4.6	Add 10 mL RT cell clumps. T	Γ 1X PBS to e ransfer the ce	each flask and j ells from each f	pipette up and down sev lask into 15-mL polypr	veral times to disperse opylene tubes.
	2.4.7	Gently pellet c Aspirate and d	ells by centrif iscard the sup	fugation at 200 ernatant.	x g in a swing-bucket	rotor at 4°C for 10 min.
	2.4.8	Wash the cell j disperse cell cl polypropylene	pellets with 10 umps. Pool a tube, respecti	0 mL 1X RT P all MCF7 or SN ively, so cells a	BS and pipette up and c N 12C cell suspensions are evenly dispersed.	lown several times to into a 50-mL
	2.4.9	Using either a viable cell cou	hemocytomet nt in the pool	er or automation ed cell suspens	c cell counter (e.g., Vi- ion.	Cell) determine the total
		<ul> <li>MCF7 or S preparation culture of</li> </ul>	SN12C_cells n n. If the cells cells will need	nust be $\ge 90\%$ are $< 90\%$ via 1 to be used for	viable to be used for co ble, they should be disc r control lysate preparat	ntrol lysate arded and a fresh ion.
	2.4.10	Adjust the tota volumes into 1	l viable cell c .5 mL Sarstee	oncentration to lt tubes.	$3 \times 10^6$ cells/mL and t	hen aliquot 1 mL
	2.4.11	Centrifuge alice and discard the	uots at 12,00 e supernatant	0 x g in a Sorv and place tube	all microcentrifuge at 4 s on ice.	°C for 5 min. Aspirate
	2.4.12	If not used imr bath. Store the	nediately, sna e frozen cell p	p-freeze the co ellets in a free	ell pellets in liquid nitro zer box at -80°C.	gen or a dry ice/ethanol





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2.5	MCF7	or SN12C Cell I	<u>Lysis</u>			
	2.5.1	Add 300 µL Ce and 1 mM PMS relative cell cor	Il Extraction F per 3 x 10 <sup>6</sup> Incentration of	Buffer (CEB) cells to fresh $7 \sim 1 \times 10^7$ cells	containing 1X protease or frozen cell pellets, th /mL.	inhibitor (PI) cocktail is should yield a
	2.5.2 Vortex tube for 3 to 5 sec at medium speed (setting 5-6 on Vortex Genie 2). Ensu cell pellet is dislodged and mixing gently in the CEB.			Genie 2). Ensure the		
	2.5.3 Place tube on ice and incubate the cells in the CEB for 30 min with 3 to 5 sec of vortexing at 10-min intervals.			h 3 to 5 sec of		
	2.5.4	2.5.4 Move samples to room temperature and add 20% SDS to a final concentration of 1% SDS (e.g., 15 μL 20% SDS to 300 μL lysate).			oncentration of 1%	
	2.5.5	2.5.5 Vortex tube for 3 to 5 sec at medium speed to distribute the SDS in the buffer.			n the buffer.	
	2.5.6	Boil the cell ex on ice.	tract for 5 mi	n in a 100°C h	eat block or boiling wat	ter bath, and cool down
	2.5.7	Clarify the extr 10 min at 4°C. the original tub	act by centrif Transfer sup e with any pr	ugation in a So ernatant into a ecipitated mate	orvall Fresco microcent new Sarstedt tube and erial in the appropriate	rifuge at 12,000 x g for hold on ice. Discard waste container.
	2.5.8	Label each Sars freezing).	stedt tube wit	h a Lot Identif	ication Code (date of ce	ell harvesting, lysis, and
	2.5.9	If not used imm	nediately for t	he immunoass	ay, snap-freeze the prot	tein extract in liquid

- nitrogen or a dry ice/ethanol bath. Store the frozen lysates in a freezer box at -80°C.
- 2.5.10 Designation of High, Mid, and Low Controls
- 2.6 yH2AX and H2AX Assay Controls

Note: NCI has used the following strategy for creation of the controls but each laboratory should qualify their own batch of controls.

- 2.6.1 yH2AX Assay Controls
  - 2.6.1.1 The High-, Mid- and Low-C tumor lysate controls are prepared from a  $1 \times 10^7$ cells/mL stock lysates of irradiated MCF7 and untreated MCF7 cultured tumor cell lines. Controls will be diluted an additional 3-fold when added to the 96well plate.
    - High-C: irradiated MCF7 lysate at  $1 \times 10^7$  cells/mL.
    - Mid-C: mix 30% High-C and 70% Low-C.
    - Low-C: untreated MCF7 lysate 1 x  $10^7$  cells/mL.





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- 2.6.2 Total H2AX Assay Controls
  - 2.6.2.1 The High-, Mid- and Low-C tumor lysate controls are dilutions prepared from a  $1 \times 10^7$  cells/mL stock lysates of SN12C and MCF7 cultured tumor cell lines. Controls will be diluted an additional 3-fold when added to the 96-well plate.
  - 2.6.2.2 SN12C lysate (~1 x:  $10^7$  cells/mL) diluted at 1:50 will be used as High-C. MCF7 lysate (~1 x  $10^7$  cells/mL) diluted at 1:50 will be used as MID-C and MCF7 lysate diluted at 1:200 will be used as Low-C. A representative dilution scheme is shown in the table below.

Control Tube Dilution	Vol. and Source of Control Lysate	Vol. 1X PBS-2% BSA	Conc. of Control in Plate (1:3 Dilution)
High-C (1:16.7) *1 x 10 <sup>7</sup> cells/mL stock lysate	<u>7.5</u> μL of SN12 C Tumor Cell Stock	117.5μL	High-C (1:50)
Mid-C (1:16.7) *1 x 10 <sup>7</sup> cells/mL stock lysate	<b>9</b> μL of MCF7 Tumor Cell Stock	141 μL	Mid-C (1:50)
Low-C (1:66.7)	<u><b>30</b></u> µL of Mid-C (MCF7) (1:50)	90 μL	Low-C (1:200)

- 2.6.3 New lots of tumor control lysates should be assayed in parallel with previously qualified controls using qualified assay reagents. The new tumor control lysates should provide control readouts with  $\pm 25\%$  agreement to the previous lot. Use of the Westgard Multirule System and a control grid are recommended to identify changes in performance of the control samples across assay runs within a given laboratory. Controls are aliquoted for single use at 120-125 µL per vial.
- 2.6.4 Representative assay readout ranges for the γH2AX and Total H2AX Control Lysates are outlined in the Steps 4.3 and 4.4.





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## **APPENDIX 4: ABBREVIATIONS**

BSA	=	Bovine Serum Albumin
С	=	Control
Cat#	=	Catalog Number
CEB	=	Cell Extraction Buffer
CV	=	Coefficient of Variation
ELISA	=	Enzyme-Linked ImmunoSorbent Assay
Н	=	Hour
γH2AX	=	Histone H2AX Phosphorylated at Serine 139 (designed as Gamma)
HRP	=	Horse Radish Peroxidase
IA	=	Immunoassay
LHTP	=	Laboratory of Human Toxicology and Pharmacology
LLD	=	Lower Limit of Detection
LLQ	=	Lower Limit of Quantitation
LLQ-c	=	Lower Limit of Quantitative Concentration
min	=	Minute
mm	=	Milli-meter
MSec	=	Milli-second
mAb	=	Monoclonal Antibody
pAb	=	Polyclonal Antibody
PAR	=	Poly(ADP-ribose)
PBS	=	Phosphate Buffered Saline
PI	=	Protease Inhibitors
PMSF	=	Phenylmethanesulfonyl Fluoride
PD	=	Pharmacodynamic
RLU	=	Relative Light Units
RT	=	Room Temperature
RUO	=	Research Use Only
SD	=	Standard Deviation
SDS	=	Sodium Dodecyl Sulfate
SOP	=	Standard Operating Procedure
Temp	=	Temperature
UD	=	Undetectable
UQ	=	Unquantifiable
μL or μg	=	Microliter or Microgram
Y	=	Year





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## **RUO PROTOCOL APPROVAL**

Developed at:	National Clinical Target Validation Laboratory (NCTVL)				
	Applied Developmental Directorate				
	Frederick National Laboratory for Cancer Research				
NCTVL Approval:	Jiuping Ji	Date: 4-25-17			
LHTP Approval:	Ralph Parchment	Date: 28 Gpul 2017			
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## **CHANGE HISTORY**

Revision	Approval Date	Description	Originator	Approval
	4/25/2017	New Document	YZ, KG, KFG	JJ



