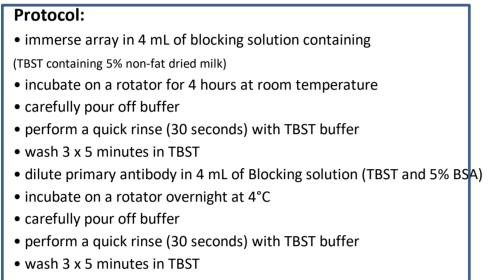


There is no graph for H3K36



2-27-2020

H3K36me3 Cell Signaling

4909

1:10000

O/N at 4C

Abcam

ab6721

1:2500

1 hr@ RT

GR119224-3

5

RI

- dilute secondary antibody in 4 mL of Blocking solution
- incubate on a rotator for 1 hour at room temperature
- carefully pour off buffer

Test Date:

Antibody:

Part number:

Lot number:

Vendor:

Dilution Incubation:

Vendor:

Part number:

Lot number:

Dilution: Incubation:

Primary antibody:

Secondary antibody:

User:

- perform a quick rinse (30 seconds) with TBST buffer
- wash 3 x 5 minutes in TBST
- detect using Chemi-Doc XR

Gridded image of the array

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Annotated Array PchAb 1419_Exposure_4 Thursday, February 27, 2020 5:31:36 PM	H3 K36me3 H3 K36me3, H4 R3me2s H4 R3me2s H3 K36me3, H4 K5ac H4 K5ac H4 R3me2s, H4 K5ac H3 K36me3, H4 R3me2s, H4 K5ac
Show Assigned Spots Map Both Sides of Array Show Most Specific H4 K5ac Reset	

H3K36me3_Cell Signaling_4909_5_DB _PchAb 1419

Reactivity of test antibody with control peptides and singly modified peptides Graph is scaled so that the intended primary reactivity is set to 1

Select Modification For Scaling: H3 K36me3 V					
Single Modification	on Peptide Reactivity (Scaled)				
	0 1419_Exposure_4				
Thursday, Fel	oruary 27, 2020 5:31:36 PM				
13 K30me1 H3 K9me2 H3 K9me2 H3 K9me2 H3 K9me2 H3 K17me2 H3 K17me2 H3 K17me2 H3 K17me2 H3 K27me2 H3 K3	H3 K36me3 H3 K36me3 H4 K336ac H4 K336ac H4 K336ac H4 K12ac H4 K12ac H4 K17me2s H4 K17me2s H4 K17me2s H4 K17me2s H4 K17me2s H4 K17me2s H4 K17me2s H4 K17me2s H4 K17me2s H2 K17me2				
H3 K3me2 H3 K9me2 H3 K9me2 H3 K9me2 H3 K9me2 H3 K17me2 H3 R17me2 H3 R170 H3 R170 H3 R170 H3 R26me2 H3 K27me2 H3 K27me2 H3 K27me2 H3 K27me2 H3 K27me2 H3 K27me2	H3 K36me3- H3 K36me3- H4 K3me2s- H4 K3me2s- H4 K13me2s- H4 K13me2s- H4 K17me2s- H4 K17me2s- H4 K17me2s- H4 K17me2s- H4 K17me2s- H4 K17me2s- H4 K17me2s- H4 K20me1- H4 K20me2- H4 K20me2- H4 K20me2- H4 K20me2- H2 K16me2- H2 K15me2- H2 H1-19 unmod- H2 H1-19 unmod- H2 H1-19 unmod-				
	H3 K36me3				

Cross inhibition of reactivity by other modifications of the same peptide Graph is scaled so that saturation of signal = 1

6me3 since there is only one spont on the array that contains this, namely, the singly modified peptide				