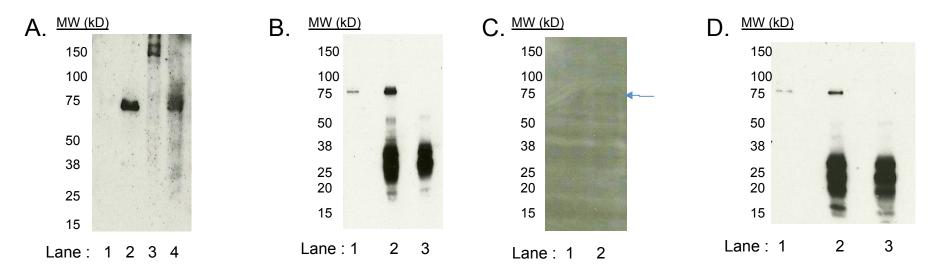
## **ENCODE DCC Antibody Validation Document**

Date of Submission
Name: Email:
Lab
Antibody Name: Target:
Company/ Source:
Catalog Number, database ID, laboratory
Antibody Description:
Target Description:
Species Target Species Host
Validation Method #1 Validation Method #2
Purification Method Monoclonal
Vendor URL:
Reference (PI/ Publication Information)
Please complete the following for antibodies to histone modifications: if your specifications are not listed in the drop-down box, please write-in the appropriate information
Histone Name AA modified AA Position Modification



Insert Validation Image (click here)

## ARID3A (NB100-279) & (sc-8821) Immunoblot / Immunoprecipitation



**A**. Western Blot using NB100-279 on nuclear lysates from cell lines GM12878 (Lane1), K562 (Lane2), HeLaS3 (Lane3), and HepG2 (Lane4). **B**. Immunoprecipitation was performed on nuclear lysates from K562 cells using antibody NB100-279. Lane1: Nuclear lysate. Lane 3: Bound material from control immunoprecipitation with rabbit IgG. . Lane 2: Bound material from immunoprecipitation with NB100-279. **C**. Western Blot using sc-8821 on nuclear lysates from cell lines GM12878 (Lane1), K562 (Lane2). **D**. Immunoprecipitation was performed on nuclear lysates from K562 cells using antibody sc-8821 and immunoblot with NB100-279. Lane1: Nuclear lysate. Lane 2: Bound material from immunoprecipitation with sc-8821. Lane 3: Bound material from control immunoprecipitation with Goat IgG. Arrow indicates band of expected size (~80kD) that is highly enriched in the specifically immunoprecipitated fraction.



Insert Validation Image (Click here)

## Validation 2: ChIPseq with alternate antibodies to the same factor

	ARID3A NB100-279	ARID3A sc-8821
Total peaks	122875	48018
% Peak overlap	86.8	86.5

## Antibodies/Immunogens:

**NB100-279 :** Immunogen: A synthetic peptide, which represented a portion of human Dead Ringer-Like 1 encoded within exon 8

sc-8821: epitope mapping at the N-terminus of ARID3A of human origin

**Comparison**: K562 cells were used for ChIP-seq with antibody sc-8821 or antibody NB100-279. Peaks were called from replicate experiments using PeakSeq with a .01 q-value cut-off. Comparisons between experiments were made using these peaks according to standard ENCODE replicate comparison parameters ( <u>http://genome.ucsc.edu/ENCODE/protocols/dataStandards/ChIP\_DNase\_FAIRE\_DNAme\_v2\_2011.pdf</u>; reported is the fraction of the top 40% of peaks in one list that are found in the full list of peaks obtained with the other antibody.