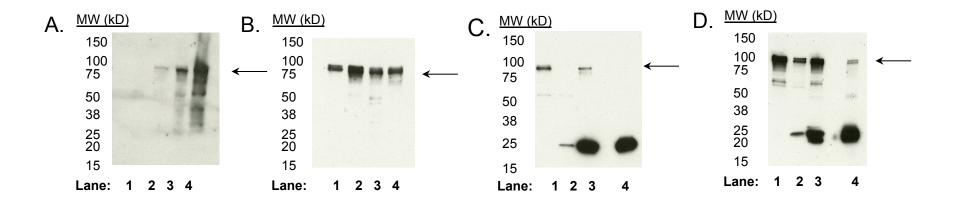
ENCODE DCC Antibody Validation Document

Date of Submission
Name: Email:
Lab
Lab
Antibody Name: Target:
Company/
Source:
Catalog Number, database ID, laboratory Lot Number
Antibody
Description:
Target
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Species Toyant Species Hest
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Validation Method #1 Validation Method #2
Purification Polyclonal/
Method Monoclonal
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Validation #1 Analysis					
Insert Validation Image (click here)					

Antibody sc13125 (UBF) and SAB1404509 (UBTF) Immunoblot/immunoprecipitation



A. Western blot using antibody sc13125 on nuclear lysates from cell lines GM12878 (Lane1), K562 (Lane2), HeLaS3 (Lane3), and HepG2 (Lane4). **B.** Western blot using antibody SAB1404509 on nuclear lysates from cell lines GM12878 (Lane1), K562 (Lane2), HeLaS3 (Lane3), and HepG2 (Lane4). **C.** Immunoprecipitation was performed on nuclear lysates from K562 cells using antibody sc13125. Lane1: Nuclear lysate. Lane 2: Unbound material from immunoprecipitation with sc13125. Lane 3: Bound material from immunoprecipitation was performed on nuclear lysates from K562 cells using antibody SAB1404509. Lane1: Nuclear lysate. Lane 2: Unbound material from immunoprecipitation with SAB1404509. Lane 3: Bound material from immunoprecipitation with SAB1404509. Lane 4: Bound material from control immunoprecipitation with mouse IgG. Arrow indicates band of expected size (97kD) that is highly enriched in the specifically immunoprecipitated fraction.

Validation #2 Analysis						
nsert Validation Image (Click here)						
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Validation 2: ChIPseq with alternate antibodies to the same factor

	UBTF sc-13125	UBTF SAB1404509
Total peaks	34443	66880
% Peak overlap	99.2	99.7

Antibodies/Immunogens:

sc-13125: raised against amino acids 1-220 of UBF of human origin

SAB1404509: Immunogen is UBTF (NP_055048, 551 a.a. ~ 651 a.a) partial recombinant protein with GST tag.

Comparison: K562 cells were used for ChIP-seq with antibody SAB1404509 or antibody sc-13125. 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 (

http://genome.ucsc.edu/ENCODE/protocols/dataStandards/ChIP_DNase_FAIRE_DNAme_v2_2011.pdf; 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.