ENCODE DCC Antibody Validation Document

Date of Submission
Name: Email:
Lab
Antibody Name: Target:
Company/
Source:
Catalag Nijumbay databasa ID labayataw
Catalog Number, database ID, laboratory Lot Number
Antibody Description:
Target
Description:
Species Target Species Host
Validation Method #1 Validation Method #2
Purification Polyclonal/
Method Monoclonal
V. 1. 1791
Vendor URL:
eference (PI/
ublication
nformation)
ease complete the following for antibodies to histone modifications:
your specifications are not listed in the drop-down box, ease write-in the appropriate information
tase mile in the appropriate information
istone Name AA modified AA Position Modification

Validation #1 Analysis		
Insert Validation II	mage (click here)	

Validation #2 Analysis			
		7	
Insert Validation Image (Click here)			

Identified Proteins (447)	Accession Number	Molecular Weight	Bowling FOSL1- 1
Fructose- bisphosphate aldolase A OS=Homo sapiens GN=ALDOA PE=1 SV=2	ALDOA_HUMAN	39 kDa	8
Poly(rC)-binding protein 1 OS=Homo sapiens GN=PCBP1 PE=1 SV=2	PCBP1_HUMAN	37 kDa	8
L-lactate dehydrogenase B chain OS=Homo sapiens GN=LDHB PE=1 SV=2	LDHB_HUMAN	37 kDa	6
60S acidic ribosomal protein P0 OS=Homo sapiens GN=RPLP0 PE=1 SV=1	RLA0_HUMAN	34 kDa	5
Alpha-enolase OS=Homo sapiens GN=ENO1 PE=1 SV=2	ENOA_HUMAN	47 kDa	5
Glyceraldehyde-3- phosphate dehydrogenase OS=Homo sapiens GN=GAPDH PE=1 SV=3	G3P_HUMAN	36 kDa	4
Aminoacyl tRNA synthase complex- interacting multifunctional protein 2 OS=Homo sapiens GN=AIMP2 PE=1	AIMP2_HUMAN	35 kDa	3

SV=2

Core histone macro-H2A.1 OS=Homo sapiens GN=H2AFY PE=1 SV=4	H2AY_HUMAN	40 kDa	3
Elongation factor 1-delta OS=Homo sapiens GN=EEF1D PE=1 SV=5	EF1D_HUMAN	31 kDa	3
Heat shock protein HSP 90- beta OS=Homo sapiens GN=HSP90AB1 PE=1 SV=4	HS90B_HUMAN	83 kDa	3
Mitochondrial import inner membrane translocase subunit TIM50 OS=Homo sapiens GN=TIMM50 PE=1 SV=2	TIM50_HUMAN	40 kDa	3
Sterol-4-alpha- carboxylate 3- dehydrogenase, decarboxylating OS=Homo sapiens GN=NSDHL PE=1 SV=2	NSDHL_HUMAN	42 kDa	3
Tubulin beta chain OS=Homo sapiens GN=TUBB PE=1 SV=2	TBB5_HUMAN	50 kDa	3
Twinfilin-2 OS=Homo sapiens GN=TWF2 PE=1 SV=2	TWF2_HUMAN	40 kDa	3
26S proteasome non-ATPase	PSD7_HUMAN	37 kDa	2

regulatory subunit 7 OS=Homo sapiens GN=PSMD7 PE=1 SV=2			
Biliverdin reductase A OS=Homo sapiens GN=BLVRA PE=1 SV=2	BIEA_HUMAN	33 kDa	2
Crk-like protein OS=Homo sapiens GN=CRKL PE=1 SV=1	CRKL_HUMAN	34 kDa	2
Cytosolic acyl coenzyme A thioester hydrolase OS=Homo sapiens GN=ACOT7 PE=1 SV=3	BACH_HUMAN	42 kDa	2
Dermcidin OS=Homo sapiens GN=DCD PE=1 SV=2	DCD_HUMAN	11 kDa	2
Elongation factor 1-alpha 1 OS=Homo sapiens GN=EEF1A1 PE=1 SV=1	EF1A1_HUMAN	50 kDa	2
Eukaryotic translation initiation factor 2 subunit 1 OS=Homo sapiens GN=EIF2S1 PE=1 SV=3	IF2A_HUMAN	36 kDa	2
Eukaryotic translation initiation factor 3 subunit I OS=Homo sapiens GN=EIF3I	EIF3I_HUMAN	37 kDa	2

PE=1 SV=1

Heterogeneous nuclear ribonucleoproteins A2/B1 OS=Homo sapiens GN=HNRNPA2B1 PE=1 SV=2	ROA2_HUMAN	37 kDa	2
Nucleophosmin OS=Homo sapiens GN=NPM1 PE=1 SV=2	NPM_HUMAN	33 kDa	2
Poly(rC)-binding protein 2 OS=Homo sapiens GN=PCBP2 PE=1 SV=1	PCBP2_HUMAN	39 kDa	2
Replication factor C subunit 2 OS=Homo sapiens GN=RFC2 PE=1 SV=3	RFC2_HUMAN	39 kDa	2
SUMO-activating enzyme subunit 1 OS=Homo sapiens GN=SAE1 PE=1 SV=1	SAE1_HUMAN	38 kDa	2
Transcriptional activator protein Pur-beta OS=Homo sapiens GN=PURB PE=1 SV=3	PURB_HUMAN	33 kDa	2
Tubulin alpha-1B chain OS=Homo sapiens GN=TUBA1B PE=1 SV=1	TBA1B_HUMAN	50 kDa	2
Vacuolar protein sorting-associated protein 26A OS=Homo sapiens GN=VPS26A	VP26A_HUMAN	38 kDa	2

PE=1 SV=2

Fos-related antigen 1 OS=Homo sapiens GN=FOSL1 PE=1

FOSL1_HUMAN 29 kDa SV=1

1

Heterogeneous nuclear ribonucleoproteins C1/C2 OS=Homo sapiens GN=HNRNPC

HNRPC_HUMAN 34 kDa PE=1 SV=4