

Supplementary Data

The Biological Significance of Targeting Acetylation-mediated Gene Regulation for Designing New Mechanistic Tools and Potential Therapeutics

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The EMBL sequence analysis was then used to perform multiple sequence alignment using the Clustal Omega program tool' is missing the word tool after sequence analysis.

1cm0_b/1-163 KVIEFHVGN SLNQKPN -
 2rc4_b/1-264 RCPSVIEFG KYE IHTWYSSPYPQEYSRLPKLYLCEFCLKYMKSRTILQQHMKKGWFHPP
 3biy_b/1-317 KFSAKRLPST -
 3toa_b/1-266 KYVDKIHIGNYEIDAWYFSPFPEDYGKQPKLWLCEYCLKYMKYEKSYRFHLGQCQWRQPP
 4bhw_b/1-448 GYCCGRKLEFSPQTLCCYGKQLCTIPRDATYYSYQNRYHFCEKCFNEIQGESVSLGDDPS
 5u7g_b/1-410 NDTLDPEPFVDCKECGRKMHQICVLHYDIIWPSGFVCDNCLKKTGRPRKENKFSAKRLQT
 6pgu_b/1-313 SKFSAKRLPST -
 6v8n_b/1-327 FSAKRLPSTR -

Conservation



Quality



Consensus

KY+SK+E+GN+E I++WY+SPQPC+Y+RDPKL Y+CE+CLKYMKK+++RQH+K+++WR+PP

Occupancy



70 80 90 100 110

1cm0_b/1-163 - - - - -

2rc4_b/1-264 A N E I Y R K N N I S V F E V D G N V S T I Y - - - - -

3biy_b/1-317 - - - - -

3toa_b/1-266 G K E I Y R K S N I S V Y E V D G K D H K I Y C Q - - - - -

4bhw_b/1-448 Q P Q T T I N K E Q F S K R K N D T L D P E L F V E C T E C G R K M H Q I C V L H H E I I W P A G F V C D G C L K K S A - - - - -

5u7g_b/1-410 T - - - - -

6pgu_b/1-313 - - - - -

6v8n_b/1-327 - - - - -

Conservation



Quality

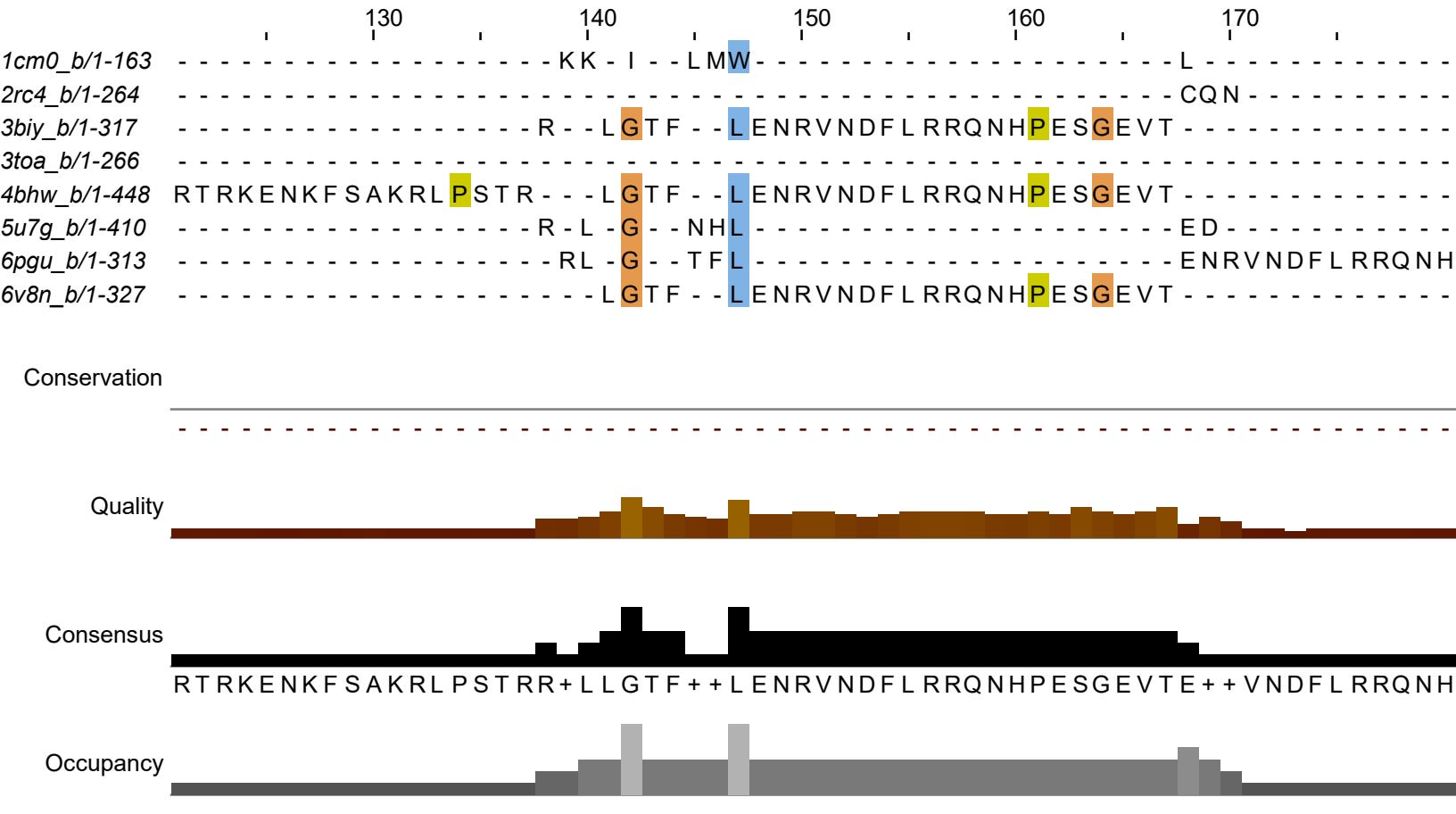


Consensus

+ + E I Y R K + N I S V + E V D G + + + + I Y + + E C T E C G R K M H Q I C V L H H E I I W P A G F V C D G C L K K S A

Occupancy





190 200 210 220 230

<i>1cm0_b/1-163</i>	- - - V G L - - -	Q N - V F - - -	S H Q L - - -	P R M - - -	P K E Y I T - - -	R L V F D
<i>2rc4_b/1-264</i>	- - - L - - -	C L - L A K L F - - -	L - D - H - - -	K T L - - -	Y Y - D V	
<i>3biy_b/1-317</i>	- - -	- - -	- - -	- - -	- - -	- - -
<i>3toa_b/1-266</i>	- - - N - L - - -	C L L - A - - -	K L F L D H - - -	? K T - - -	L Y F D - - -	V
<i>4bhw_b/1-448</i>	- - -	- - -	- - -	- - -	- - -	- - -
<i>5u7g_b/1-410</i>	- - - R V N K F L R R Q N H P E A G E V - - -	- - -	- - -	- - -	- - -	- - -
<i>6pgu_b/1-313</i>	P E S G E V T - - -	- - -	- - -	- - -	- - -	- - -
<i>6v8n_b/1-327</i>	- - -	- - -	- - -	- - -	- - -	- - -

Conservation



Quality



Consensus

P E S G E V T + G L R V N K F L R R Q N H P E A G E V C L L + A K L F + + + L D H + R + ? K T + + + Y + + F D + + V + V

Occupancy



250 260 270 280 290

1cm0_b/1-163 PKH - KT ----- L AL I K D G R V I

2rc4_b/1-264 E P F - L F ----- Y V L T Q N D V K G C H L V

3biy_b/1-317 - - - V R V V H A S D K T V E V K P G M K A R F V D S G E M A E S F P Y R T K A L F A F E - - - E I D G V D L C F

3toa_b/1-266 E P F - V F ----- Y I L T E V D R Q G A H I V

4bhw_b/1-448 - - - V R V V H A S D K T V E V K P G M K A R F V D S G E M A E S F P Y R T K A L F A F E - - - E I D G V D L C F

5u7g_b/1-410 - - - F V R V V V A S S D K T V E V K P G M K S R F V D S G E M S E S F P Y R T K A L F A F E - - - E I D G V D V C F

6pgu_b/1-313 - - - V R V V H A S D K T V E V K P G M K A R F V D S G E M A E S F P Y R T K A L F A F E - - - E I D G V D L C F

6v8n_b/1-327 - - - V R V V H A S D K T V E V ? P G M K A R F V D S G E M A E S F P Y R T K A L F A F E - - - E I D G V D L C F

Conservation



Quality

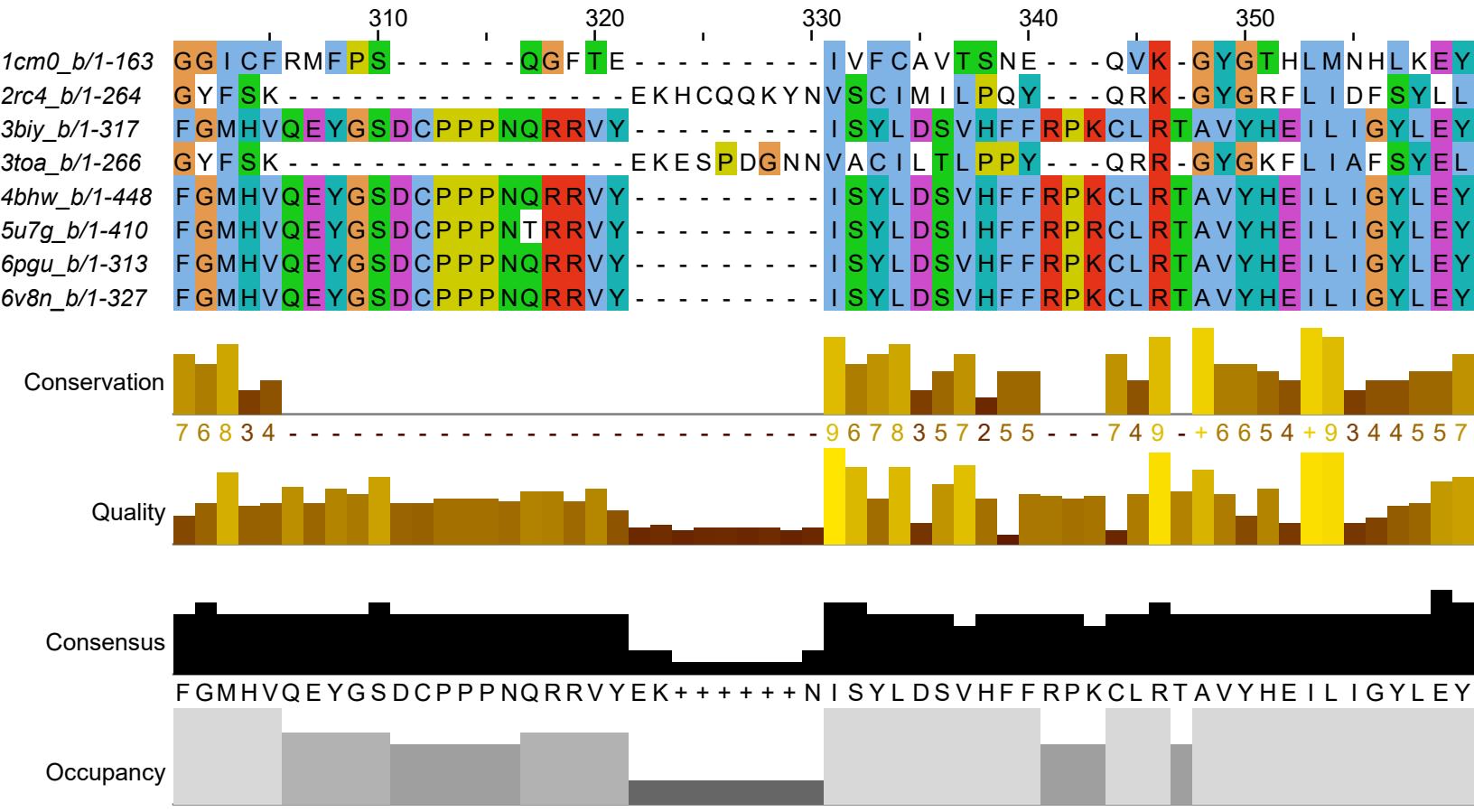


Consensus

E P F F V R V V H A S D K T V E V K P G M K A R F V D S G E M A E S F P Y R T K A L F A F E V D R Q G E I D G V D L C F

Occupancy





370 380 390 400 410
 1cm0_b/1-163 H I - - K H - - -
 2rc4_b/1-264 S K - - R E G Q A G S P E K P L S D L G R L S Y M A Y W K S V I L E C L Y H Q N D K Q I S I K K L S K L T G I C P Q D I
 3biy_b/1-317 V K K L G Y - - -
 3toa_b/1-266 S K - - L E - - -
 4bhw_b/1-448 V K K L G Y - - -
 5u7g_b/1-410 V K - - K L - - -
 6pgu_b/1-313 V K K L G Y - - -
 6v8n_b/1-327 V K K L G Y - - -



V K K L G Y G Q A G S P E K P L S D L G R L S Y M A Y W K S V I L E C L Y H Q N D K Q I S I K K L S K L T G I C P Q D I



430 440 450 460 470

1cm0_b/1-163 - - - D I - - -

2rc4_b/1-264 T STL HHL RML DRREKL I QDHMAKLQL NL RPVDVDPECL RWTPV - - -

3biy_b/1-317 - - - TT - - -

3toa_b/1-266 - - - ST VGSPEKPLSDLGKLS

4bhw_b/1-448 - - - TT - - -

5u7g_b/1-410 - - - GY - - -

6pgu_b/1-313 - - - TT - - -

6v8n_b/1-327 - - - TT - - -

Conservation



1 1

Quality



Consensus

T STL HHL RML DRREKL I QDHMAKLQL NL RPVDVDPECL RWTPVTTVGSPEKPLSDLGKLS

Occupancy



490 500 510 520 530

1cm0_b/1-163 - - - - -
2rc4_b/1-264 - - - - -
3biy_b/1-317 - - - - -
3toa_b/1-266 Y R S Y W S W V L L E I L R D G T L S I K D L S Q M T S I T Q N D I I S T L Q S L N M V K Y W K G Q H V I C V T P K L V
4bhw_b/1-448 - - - - -
5u7g_b/1-410 - - - - -
6pgu_b/1-313 - - - - -
6v8n_b/1-327 - - - - -

Conservation



Quality



Consensus

Y R S Y W S W V L L E I L R D G T L S I K D L S Q M T S I T Q N D I I S T L Q S L N M V K Y W K G Q H V I C V T P K L V

Occupancy



550 560 570 580 590

<i>1cm0_b/1-163</i>	- - - - -	L - NFLTYAD - - - - -
<i>2rc4_b/1-264</i>	- - - - -	- - - - -
<i>3biy_b/1-317</i>	- - - - -	G - H I W A C P P S E G D D Y I F H C H P P D Q K I P K P K R L Q E W Y K K M
<i>3toa_b/1-266</i>	E E H L K S A P P I T V D S V C L K W A P - - - - -	
<i>4bhw_b/1-448</i>	- - - - -	G - H I W A C P P - - - - -
<i>5u7g_b/1-410</i>	- - - - -	V T G H I W A C P P S E G D D Y I F H C H P P D Q K I P K P K R L Q E W Y K K M
<i>6pgu_b/1-313</i>	- - - - -	G - H I W A C P P S E G D D Y I F H C H P P D Q K I P K P K R L Q E W F K K M
<i>6v8n_b/1-327</i>	- - - - -	G - H I W A C P P S E G D D Y I F H C H P P D Q K I P K P K R L Q E W Y K K M

Conservation



Quality



Consensus

EEHLKSAPPITVDSVCLKWAPGTHIWACPPSEGDDYIFHCHPPDQKIPKPKRLQEWYKKM

Occupancy



610 620 630 640 650

1cm0_b/1-163 - - - - -
2rc4_b/1-264 - - - - -
3biy_b/1-317 L D K A V S E R I V H D Y K D I F K Q A T E D R L T S A K E L P Y F E G D F W P N V L E E S I K E S Q K L Y A T M E K H
3toa_b/1-266 - - - - -
4bhw_b/1-448 - - - - -
5u7g_b/1-410 M L D K A F A E R I I I N D Y K D I F K Q A N E D R L T S A K E L P Y F E G D F W P N V L E E S I K E S G K L Y A T M E K
6pgu_b/1-313 L D K A V S E R I V H D Y K D I F K Q A T E D R L T S A K E L P Y F E G D F W P N V L E E S I Q K L Y A T M E K H K E V
6v8n_b/1-327 L D K A V S E R I V H D Y K D I F K Q A T E D R L T S A K E L P Y F E G D F W P N V L E E S I K E L E Q K T S K N K S S

Conservation



Quality



Consensus

L D K A V S E R I V H D Y K D I F K Q A T E D R L T S A K E L P Y F E G D F W P N V L E E S I K + L + + K + + K + K E +

Occupancy



This figure displays a sequence alignment and quality analysis for a protein region spanning amino acid positions 670 to 710 across several homologs. The top part shows the sequence alignment with color-coded conservation and highlighted regions. The bottom part provides detailed quality metrics for each position.

Sequence Alignment:

- Conservation:** A horizontal dashed line indicates conservation levels across the sequence.
- Quality:** A horizontal bar chart where darker shades represent higher quality.
- Consensus:** A horizontal bar chart showing the most frequent amino acid at each position.
- Occupancy:** A horizontal bar chart showing the occupancy of each amino acid at each position.

Sequence Data (Amino Acid Positions 670-710):

Position	1cm0_b/1-163	2rc4_b/1-264	3biy_b/1-317	3toa_b/1-266	4bhw_b/1-448	5u7g_b/1-410	6pgu_b/1-313	6v8n_b/1-327
670	-	-	K	-	-	H	F	N
671	-	-	E	-	-	K	V	S
672	-	-	V	-	-	F	I	Q
673	-	-	F	-	-	Y	A	T
674	-	-	I	-	-	H	G	M
675	-	-	+ +	-	-	Q	P	S
676	-	-	F	-	-	+	+	+
677	-	-	+	-	-	+	+	+
678	-	-	+	-	-	+	+	+
679	-	-	+	-	-	+	+	+
680	-	-	+	-	-	+	+	+
681	-	-	+	-	-	+	+	+
682	-	-	+	-	-	+	+	+
683	-	-	+	-	-	+	+	+
684	-	-	+	-	-	+	+	+
685	-	-	+	-	-	+	+	+
686	-	-	+	-	-	+	+	+
687	-	-	+	-	-	+	+	+
688	-	-	+	-	-	+	+	+
689	-	-	+	-	-	+	+	+
690	-	-	+	-	-	+	+	+
691	-	-	+	-	-	+	+	+
692	-	-	+	-	-	+	+	+
693	-	-	+	-	-	+	+	+
694	-	-	+	-	-	+	+	+
695	-	-	+	-	-	+	+	+
696	-	-	+	-	-	+	+	+
697	-	-	+	-	-	+	+	+
698	-	-	+	-	-	+	+	+
699	-	-	+	-	-	+	+	+
700	-	-	+	-	-	+	+	+
701	-	-	+	-	-	+	+	+
702	-	-	+	-	-	+	+	+
703	-	-	+	-	-	+	+	+
704	-	-	+	-	-	+	+	+
705	-	-	+	-	-	+	+	+
706	-	-	+	-	-	+	+	+
707	-	-	+	-	-	+	+	+
708	-	-	+	-	-	+	+	+
709	-	-	+	-	-	+	+	+
710	-	-	+	-	-	+	+	+

730 740 750 760 770

1cm0_b/1-163 - - - - - E Y A I

2rc4_b/1-264 - - - - -

3biy_b/1-317 G C M L V E L H T Q S Q D - - - - -

3toa_b/1-266 - - - - -

4bhw_b/1-448 - - - - - S E G D

5u7g_b/1-410 L H T Q G Q D R F V Y T C N E C K H H V E T R W H C T V C E D Y D L C I N C Y N T K S H T H K M V K W G L G L D - - -

6pgu_b/1-313 L V E L H T Q S - - - - -

6v8n_b/1-327 H L E F S S L R R A Q W S T G C M L V E L H T Q - - - - -

Conservation



Quality



Consensus



L + E L + + L R + + + + + + + C + + + + + + + H C T V C E D Y D L C I N C Y N T K S H T H K M V K W G L G L D + + + +

Occupancy



850 860 870 880 890

1cm0_b/1-163 - - - - -
2rc4_b/1-264 - - - - -
3biy_b/1-317 - - - - -
3toa_b/1-266 - - - - -
4bhw_b/1-448 EGDFWPNVLEESIKESGGGSGSQKLYATMEKHKEVFFVIRLIAGPAANSLPPIVDPDPLP
5u7g_b/1-410 - - - - -
6pgu_b/1-313 - - - - -
6v8n_b/1-327 - - - - -

Conservation



Quality



Consensus

EGDFWPNVLEESIKESGGGSGSQKLYATMEKHKEVFFVIRLIAGPAANSLPPIVDPDPLP

Occupancy



910 920 930 940

1cm0_b/1-163 - - - -
2rc4_b/1-264 - - - -
3biy_b/1-317 - - - -
3toa_b/1-266 - - - -
4bhw_b/1-448 CDLMDG**R**D~~A~~F~~L~~T~~L~~A~~R~~D~~K~~H~~L~~E~~F~~S~~S~~L~~R~~R~~A~~Q~~W~~S~~T~~M~~C~~M~~L~~V~~E~~L~~H~~T~~Q~~S~~Q~~D
5u7g_b/1-410 - - - -
6pgu_b/1-313 - - - -
6v8n_b/1-327 - - - -

Conservation



Quality



Consensus

CDLMDG**R**D~~A~~F~~L~~T~~L~~A~~R~~D~~K~~H~~L~~E~~F~~S~~S~~L~~R~~R~~A~~Q~~W~~S~~T~~M~~C~~M~~L~~V~~E~~L~~H~~T~~Q~~S~~Q~~D

Occupancy

