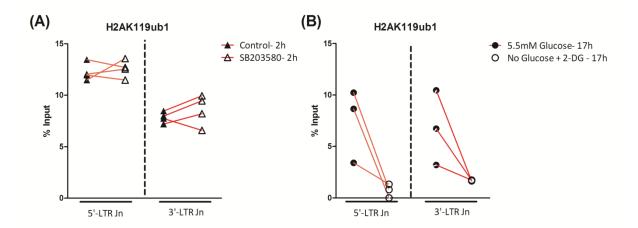


Supplementary figure 1. Transcriptional and epigenetic impact of PRC1 modulators. (A) Cryopreserved PBMCs isolated from HTLV-1-infected individuals (N=5) were cultured for 2 hr in the presence of either control (DMSO) or the H2A-ubiquitylation inhibitor PRT4165 (50  $\mu$ M). (B) Cryopreserved PBMCs isolated from HTLV-1-infected individuals (N=9) were cultured for 17 hr in the presence of either control (DMSO); the deubiquitinase (DUB) inhibitor PR619 (50  $\mu$ M); or the H2A-ubiquitylation inhibitor PRT4165 (50  $\mu$ M). Cells were harvested for RNA and subjected to qRT-PCR with primers specific for tax mRNA (plus-strand) and sHBZ mRNA (minus-strand). Error bars represent the SEM. (C) Cryopreserved HTLV-1-infected PBMCs (N=4) were fixed after 17 hr of culture in the presence of control (DMSO) or deubiquitinase inhibitor PR619 (50  $\mu$ M). Fixed cells were subjected to ChIP-qPCR, using antibodies directed against H2AK119ub1, H3K27me3, H3K4me3, H3K9/14Ac and IgG, and primers specific respectively for the 5′-LTR and 3′-LTR junctions of the HTLV-1 provirus. Enrichment is expressed as percent input DNA. Statistical significance was calculated using the two-tailed Student's t test (\*p < 0.05, \*\*p < 0.005, \*\*\*p<0.0005). N=5 for (A); N=9 for (B) and N=4 for (C).



Supplementary figure 2. Effect of p38-MAPK inhibition and glucose levels on H2AK119ub1 at the HTLV-1 provirus. Cryopreserved HTLV-1-infected PBMCs were fixed: (A) after 2 hr of culture in the presence of a control (DMSO) or p38-MAPK inhibitor (SB203580), and (B) after 17 hr of culture in medium containing 5.5mM glucose (5.5mM glucose-17h) or in glucose-free medium containing glycolysis inhibitor 2-deoxy-D-glucose (no glucose + 2-DG-17h). Fixed cells were subjected to ChIP-qPCR, using antibodies directed against H2AK119ub1 or IgG, and primers specific respectively for the 5'-LTR and 3'-LTR junctions of the HTLV-1 provirus. Enrichment is expressed as percent input DNA. N=4 for (A); N=3 for (B).

## Supplementary Table S1. Patient samples used in the study. (Table relates to Figures 1-4; Supplementary Figures 1 and 2)

	Figure 1						Figure 4A- H3K4me3					Suppl. Fig. 1A			
Sr. No.	Patient Code	Age	Sex	Clinical status	S	Sr. No.	Patient Code	Age	Sex	Clinical status	Sr. No.	Patient Code	Age	Sex	Clinical status
1	TDZ	61.0	F	HAM	1	1	TCQ	78.0	М	HAM	1	TFF	55.1	F	HAM
2	TW	52.7	F	HAM	2	2	TCX	50.7	М	HAM	2	TEJ	52.5	F	HAM
3	TEB	61.6	М	HAM	3	3	TED	59.0	F	HAM	3	TCG	58.7	F	HAM
4	HEZ	61.7	F	AC	4	1	TCJ	54.6	F	HAM	4	TCK	69.6	F	HAM
5	HBX	64.6	F	AC							5	TEF	53.3	М	HAM
6	TCD	32.2	F	HAM			Figure 4A- H3K9/14Ac								
7	TED	59.0	F	HAM	S	Sr. No.	Patient Code	Age	Sex	Clinical status		Suppl. Fig. 1B			
8	TEF	53.3	M	HAM	1	l	TBC	76.5	F	HAM	Sr. No.	Patient Code	Age	Sex	Clinical status
					2	2	TCX	50.7	М	HAM	1	TCJ	54.6	F	HAM
	Figure 2A				3	3	TEB	61.6	М	HAM	2	TCK	69.6	F	HAM
Sr. No.	Patient Code	Age	Sex	Clinical status	4	1	TED	59.0	F	HAM	3	TEF	53.3	М	HAM
1	TEB	61.6	М	HAM							4	TEK	43.5	F	HAM
2	TET	54.3	F	HAM			Figure 4B- H3K4me3				5	TFF	55.1	F	HAM
3	TEP	50.8	F	HAM	S	Sr. No.	Patient Code	Age	Sex	Clinical status	6	TEJ	52.5	F	HAM
4	TDZ	61.0	F	HAM	1	1	TEB	61.6	М	HAM	7	TCQ	78.0	М	HAM
					2	2	TED	59.0	F	HAM	8	TCR	59.2	М	HAM
	Figure 2B				3	3	TCX	50.7	М	HAM	9	TCG	58.7	F	HAM
Sr. No.	Patient Code	Age	Sex	Clinical status	4	1	TCJ	54.6	F	HAM					
1	TEB	61.6	М	HAM								Suppl. Fig. 1C			
2	TED	59.0	F	HAM			Figure 4B- H3K9/14Ac				Sr. No.	Patient Code	Age	Sex	Clinical status
3	TBC	76.5	F	HAM	S	Sr. No.	Patient Code	Age	Sex	Clinical status	1	TEK	43.5	F	HAM
4	TCX	50.7	M	HAM	1	l	TEB	61.6	М	HAM	2	TCK	69.6	F	HAM
					2	2	TCY	60.0	F	HAM	3	TCG	58.7	F	HAM
					3	3	TCR	59.2	М	HAM	4	TEF	53.3	М	HAM
	Figure 3A				4	1	TCX	50.7	F	HAM					
Sr. No.	Patient Code	Age	Sex	Clinical status	5	5	TED	59.0	F	HAM		Suppl. Fig. 2A			
1	TCK	69.6	F	HAM	6	5	TCQ	78.0	М	HAM	Sr. No.	Patient Code	Age	Sex	Clinical status
2	TCJ	54.6	F	HAM							1	TCX	50.7	М	HAM
3	TEJ	52.5	F	HAM			Figure 4C				2	TCR	59.2	М	HAM
4	TBP	79.5	M	HAM	S	Sr. No.	Patient Code	Age	Sex	Clinical status	3	TEB	61.6	М	HAM
5	TDZ	61.0	F	HAM	1	1	TCJ	54.6	F	HAM	4	TED	59.0	F	HAM
					2	2	TCK	69.6	F	HAM					
					3	3	TAD	57.8	F	HAM		Suppl. Fig. 2B			
	Figure 3B										Sr. No.	Patient Code	Age	Sex	Clinical status
Sr. No.	Patient Code	Age	Sex	Clinical status							1	TCJ	54.6	F	HAM
1	TCR	59.2	М	HAM	1	AC : A:	symptomatic carrier		-		2	TCK	69.6	F	HAM
2	TEJ	52.5	F	HAM			HTLV-1 Associated M	velona	thy		3	TAD	57.8	F	HAM
3	TCJ	54.6	F	HAM	'	17-XIVI - I	THEY I ASSOCIATED IVI	устора	city						

## Supplementary Table S2. Chemical compounds and supplier list. (Table relates to Figures 1-4; Supplementary Figures 1 and 2)

<u>Sr.</u>				<u>Final</u>		
No.	<u>Name</u>	<u>Supplier</u>	<u>Solvent</u>	conc.	Mechanism of action	<u>Reference</u>
					Broad spectrum 2-OG oxygenase inhibitor	
1	DMOG	Cayman Chem.	DMSO	0.5mM	(and isocitrate dehydrogenase inhibitor)	(24)
		Cell Guidance				
2	SB203580	Systems	DMSO	10μΜ	Selective p38-MAPK inhibitor	(21)
		Cell Guidance				
3	BIRB-796	Systems	DMSO	5μΜ	Selective p38-MAPK inhibitor	(21)
		Cell Guidance				
4	PD184352	Systems	DMSO	10μΜ	Selective ERK-MAPK inhibitor	(21)
5	2-Deoxy-D-Glucose	Sigma	Water	10mM	Glycolysis inhibitor	(30)
		Cell Guidance				
6	U0126	Systems	DMSO	10μΜ	Selective ERK-MAPK inhibitor	(21)
7	Anisomycin	Cayman Chem.	DMSO	5μΜ	MAPK Inducer, Protein synthesis inhibitor	(26)
8	PR619	Cayman Chem.	DMSO	50μΜ	Broad-spectrum Deubiquitinase Inhibitor	(28)
9	PRT4165	Cayman Chem.	DMSO	50μΜ	H2A ubiquitiylation inhibitor	(29)

## Supplementary Table S3. Primers and probes used in the study. (Table relates to Figures 1-4; Supplementary Figures 1 and 2)

qRT-PCR				
<u>Sr. No</u>	Gene		<u>Sequence</u>	Reference
1	Tax	F	5'-CCGGCGCTGCTCTCATCCCGGT-3'	
1	Tax	R	5'-GGCCGAACATAGTCCCCCAGAG-3'	
2	sHBZ	F	5'-GGACGCAGTTCAGGAGGCAC-3'	(22)
2	SUPT	R	5'-CCTCCAAGGATAATAGCCCG-3'	(22)
3	18S rRNA	F	5'-GTAACCCGTTGAACCCCATT-3'	
3	IOSTRINA	R	5'-CCATCCAATCGGTAGTAGCG-3'	
4	c-fos	F	5'-CGGGCTTCAACGCAGACTA-3'	(25)
	C-103	R	5'-GGTCCGTGCAGAAGTCCTG-3'	(23)
		F	5'-TGATGCAACGCTCTCCAAGC-3'	(32)
5	ATF3	R	5'-TTAGCTCTGCAATGTTCCTTC-3'	(32)
		F	5'-CGGAGTCAACGGATTTGGTC-3'	
6	GAPDH	R	5'-AAGCTTCCCGTTCTCAGCC-3'	
			1	

ChIP-qPCR								
1	5'-LTR	F	5'-GACAGCCCATCCTATAGCACTC-3'					
_	Junction	R	5'-CTAGCGCTACGGGAAAAGATT-3'	(22)				
2	3'-LTR	F	5'-AATACACCAACATCCCCATTTC-3'					
_	Junction	R	5'-GTTTTCACTGGGAGGCTCTAA-3'					
3	ATF3	F	5'-CGAACTTGCATCACCAGTGC-3'	(32)				
	promoter	R	5'-GGTCGTTTACTCCGTGTTGC-3'	. (/				
qPCR Probe	<u>s</u>		,					
1	5'-LTR Junction		FAM -TCGTCCGGGATACGAGCGCC- TAM					
2	3'-LTR Junction		FAM -AGAGGCAGATGACAATGACCATGAGCC- TAM					
3	ATF3 promoter		FAM-AGCATTACGTCAGCCTGGGACTG -TAM					
4	ATF3 mRNA		FAM-CACTGCACAGCTCTCTCTCTCGC-TAM					
5	5 GAPDH mRNA		FAM- CATTGACCTCAACTACATGGTTTACATGTTCCAATATGATTC- TAM					