## Readers

TTD.HG1.0:UHRF1/UHRF2	ubiquitin,TTD,PHD,SAD_SRA	biquitin,TTD,PHD,SAD_SRA,zf-C3HC4			
BAH.HG1.9:BAHCC1/BAHD1/TNRC18	ВАН				
Bromodomain.HG1.49:BRD2/BRD3/BRD4/BRDT	Bromodomain,BET,BRD4_CDT	Bromodomain,BET			
PHD.HG12.1:MTF2/PHF1/PHF19	PHD	<b>_</b> Tudor,PHD	Tudor,PHD,Mtf2_C		
Chromo.HG1.9:CBX1/CBX3/CBX5	Chromo,Chromo_shadow	Chromo			
Chromo.HG1.44:CBX2/CBX4/CBX6/CBX7/CBX8	Chromo,CBX7_C	Chromo			
PWWP.HG1.18:ZMYND11	Bromodomain,PWWP	◆PHD,Bromodomain,PWWP			
PHD.HG14.1:UHRF1/UHRF2	• biquitin,TTD,PHD,SAD_SRA	ubiquitin,TTD,PHD,SAD_SRA,zf-C3HC4			
D.HG1.4:FBXL19/KDM2A/KDM2B/KDM7A/PHF2/PHF8	PHD,JmjC,JHD	₽HD	PHD,Cupin,JHD		
Bromodomain.HG8.0:ATAD2/ATAD2B	AAA,AAA_lid,Bromodomain				
PHD.HG4.8:SP140L/TRIM24/TRIM28/TRIM33/TRIM66	<b>2</b> f−B_box,PHD,Bromodomain	<b>●</b> HD,Bromodomain		PHD,Bromodomain,PWWP	
Chromo.HG5.6:CHD6/CHD7/CHD8/CHD9	Chromo,SNF2_N,Helicase_C	Chromo,SNF2_N,Helicase_C,BRK	©hromo,SNF2_N,Helicase_C,FYRN,FYRC		
Bromodomain.HG1.54:CREBBP/EP300	2f-TAZ,KIX,Bromodomain,DUF902,HAT_KAT11,ZZ				
PWWP.HG1.14:BRD1/BRPF1/BRPF3	EPL1,PHD,zf-HC5HC2H,Bromodomain,PWWP				
PHD.HG5.4:KMT2A/KMT2B/KMT2C/KMT2D	₽HD	<b>▼</b> f−HC5HC2H,PHD	▼f−HC5HC2H,PHD,FYRN,FYRC,SET		
BAH.HG3.0:MTA1/MTA2/MTA3/RERE	<b>B</b> AH	BAH,ELM2,Myb_DNA-binding,GATA,MTA_R1	BAH,PHD,zf-HC5HC2H		
PHD.HG7.0:PHF13/PHF23	PHD				
Chromo.HG4.1:CHD1/CHD2	Chromo,SNF2_N,Helicase_C,DUF4208	Chromo,SNF2_N,Helicase_C	Chromo,SNF2_N,Helicase_C,CDH1_2_SANT_HL	1,DCH748Q8	
Bromodomain.HG4.8:SMARCA2/SMARCA4	QLQ,HSA,BRK,SNF2_N,Helicase_C,SnAC,Bromodomain SNF2_N,Helicase_C,SnAC,Bromodomain SNF2				
Bromodomain.HG7.1:BRWD1/BRWD3/PHIP	WD40,Bromodomain	Bromodomain			
W4A/KDM4B/KDM4C/KDM5A/KDM5B/KDM5C/KDM5D	PHD	■mjN,ARID,PHD,JmjC,zf-C5HC2,PLU-1	JmjN,JmjC,PHD,zf−HC5HC2H,Tudor		
BAH.HG2.0:DNMT1	DNMT1-RFD,BAH,DNA_methylase	DNMT1-RFD,zf-CXXC,BAH,DNA_methylase	BAH,Chromo,DNA_methylase		
PHD.HG3.4:AC006064.6/CHD3/CHD4/CHD5	©HDNT,PHD,Chromo,SNF2_N,Helicase_C,DUF108 <b>7FD4DF,©6866</b> 660 <b>2</b> ,N,Helicase_C,DUF1087,DUF10 <b>88</b> +D,Chromo,SNF2_N,Helicase_C,DUF1087,DUF1086,CHDCT2				
YEATS.HG1.9:YEATS4	YEATS				
TUDOR.HG1.56:SND1	SNase,TUDOR				
Bromodomain.HG5.0:ZMYND8/ZMYND11	Bromodomain,PWWP	Bromodomain	PHD,Bromodomain,PWWP		
BAH.HG1.6:PBRM1	Bromodomain,BAH	■romodomain,BAH,HMG_box			
Bromodomain.HG6.0:TAF1/TAF1L	Bromodomain	■UF3591,Bromodomain	■BP-binding,DUF3591,zf-CCHC,Bromodomain	■UF3591,zf–CCHC,Bromodomain	
Bromodomain.HG4.5:ASH1L/PBRM1	<b>B</b> romodomain,BAH,HMG_box	Bromodomain	Bromodomain,BAH	♠WS,SET,Bromodomain,BAH	
Bromodomain.HG3.10:KAT2A/KAT2B	Acetyltransf, Bromodomain				
PHD.HG2.19:BRD1/BRPF1/BRPF3	■PL1,PHD,zf-HC5HC2H,Bromodomain	■PL1,PHD,zf-HC5HC2H,Bromodomain,PWWP			
3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2	MBT	MBT,SLED,SAM	MBT,zf-C2HC,SAM		
Bromodomain.HG2.3:BRD7/BRD9	Bromodomain	■romodomain,DUF3512			
Bromodomain.HG2.5:BRD1/BRPF1/BRPF3	<b>■ PL1</b> ,PHD,zf–HC5HC2H,Bromodomain	<b>■</b> PL1,PHD,zf–HC5HC2H,Bromodomain,PWWP			
MRG.HG1.1:MORF4L1/MORF4L2/MSL3	Judor-knot,MRG	MRG			
ING.HG1.2:ING1/ING2/ING4/ING5	ING	●NG,PHD			
Chromo.HG4.2:AC006064.6/CHD3/CHD4/CHD5	©HDNT,PHD,Chromo,SNF2_N,Helicase_C,DUF108 <b>7中地</b> 508666000000000000000000000000000000000				
PHD.HG1.1:CXXC1/DIDO1/PHF3	<b>₽</b> HD	PHD,zf-CXXC,zf-CpG_bind_C			
Bromodomain.HG5.2:BAZ1A/BAZ1B/BAZ2A/BAZ2B	WAC_Acf1_DNA_bd,DDT,WHIM1,WSD,PHD,Bromodomain				
PHD.HG4.5:BAZ1A/BAZ1B/BAZ2A/BAZ2B	PHD	WAC_Acf1_DNA_bd,DDT,WHIM1,WSD,PHD,Bro	modomain		
1/PWWP2A/PWWP2B/PWWP3B/ZCWPW1/ZCWPW2	PWWP	<b> ▼ f − CW ,PWWP</b>			