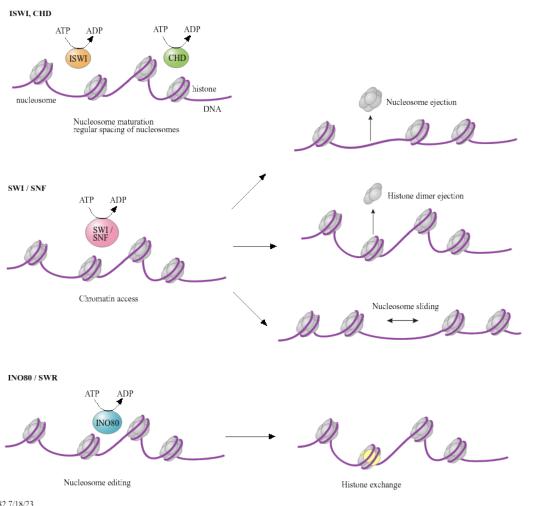


ATP - DEPENDENT CHROMATIN REMODELING



Chromatin remodeling complexes (CRCs)

ISWI family CHD family NURD complex SMARC CHD3 CHD4 HDAC ACF A5 BAZ1A MBD2 MBD3 MTA WICH A5 BAZ1B GATAD2 RBBP4 RBBP7 NORC A5 PHF6 CDK2AP BAZ2A BRF A5 BAZ2B CHRAC A5 BAZIA CHRACI POLE3 RSF A5 RSF1 NURF BPTF RBBP4 RBBP7 BAP18 NURF38 CERF A1 CECR2

SWI / SNF family

BAF complex

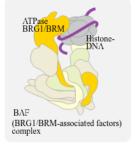
BRG1/BRM

DIKUI/DIKWI								
MARC	A2/4 B1		C	D	E1			
	ACTB	ACTL6	BCL7		2210	1		
		SS18						
	1.70.770.4	TO TOTAL IO	TO TOTAL					

BAF ARID1 DPF1/3 DPF2

PBAF ARID2 PBRM1 BRD7 PHF10

ncBAF BRD9 BICRA BICRAL



INO80 / SWR family

INO80 complex

	INO80	INC©0B	INC&0C	INC80D	INC80E	TEFT	NFRKB	MCRS1
	UCHL5	RUVBL	ACTL6	ACTR5	ACTR8	YY1		
yeast	IES1	IES2	IES3	IES4	IES5	TAF14	NHP10	ARP4

SRCAP complex

SRCAP RUVBL ACTL6 ACTB DMAP1 YEATS4 VPS72 H2A.Z ACTR6 ZNHIT1

p400 (TIP60) complex

p400	RUVBL	ACTL6	ACTB	DMAP1	YEATS4	VPS72	H2A.Z
TRRAP	MORF4L	MRGBP	BRD8		EPC	EAF6	TIP60

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