Snapshot: SMC Protein Complexes I

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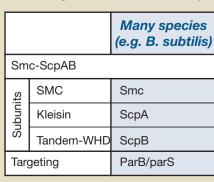
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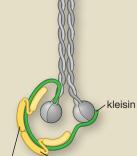
Eukaryotic SMC Complexes

			S. cerevisiae (Budding Yeasts)	<i>S. pombe</i> (Fission Yeasts)	<i>A. thaliana</i> (Plants)	<i>C. elegans</i> (Nematodes)	<i>D. melanogaster</i> (Insects)	<i>H. sapiens</i> (Vertebrates)
Со	hesi	n						
Subunits	-S	MC	Smc1	Psm1 (Smc1)	SMC1 (TTN8)	HIM-1 (SMC-1)	SMC1	SMC1α, SMC1β*
	-S	MC	Smc3	Psm3 (Smc3)	SMC3	SMC-3	SMC3	SMC3
	-k	leisin	Scc1 (Mcd1), Rec8 *	Rad21, Rec8*	SYN2-4, SYN1*	SCC-1 (COH-2), COH- REC-8*, COH-3/4*	Rad21, C(2)M?*	SCC1, REC8*, RAD2
	HEAT-A		Pds5	Pds5	PDS5	EVL-14	Pds5	PDS5A, PDS5B
	HEAT-B		Scc3 (Irr1)	Psc3, Rec11*	SCC3	SCC-3	SA, SA-2 (SNM)*	SA1, SA2, SA3*
Regulators	Kollerin Ioading complex		Scc2	Mis4	SCC2	PQN-85 (SCC-2)	Nipped-B	NIPBL (SCC2)
			Scc4	Ssl3	-	MAU-2	Mau-2	MAU2 (SCC4)
	Acetyltransferase		Eco1 (Ctf7)	Eso1	ECO1 (CTF7)	F08F8.4	Eco (Deco) + San	ESCO1, ESCO2
	Dea	acetylase	Hos1	-	-	-	-	HDAC8
	Stabilizer		-	-	-	-	Dalmatian	Sororin (CDCA5)
	Destabilizer		Wpl1 (Rad61)	Wpl1	WAPL	WAPL-1	Wapl	WAPL
	Separase		Esp1	Cut1	ESP	SEP-1	Sse + Thr	ESPL1
	Shugoshin-		Sgo1	Sgo1	SGO1	-	MEI-S332	SGOL1
	phosphatase complex		PP2A	PP2A	-	-	-	PP2A
Cor	nden	sin						
Subunits	-SMC		Smc4	Cut3 (Smc4)	SMC4A	SMC-4, DPY-27**	Smc4 (Gluon)	SMC4
	-SMC		Smc2	Cut14 (Smc2)	SMC2A/B	MIX-1 (SMC-2)	Smc2	SMC2
	in (I)	-kleisin	Brn1	Cnd2	CAP-H	DPY-26	Barren (Cap-H)	CAP-H
	Condensin (I)	HEAT-IA	Ycs4	Cnd1	CAP-D2	DPY-28	Cap-D2	CAP-D2
	Con	HEAT-IB	Ycg1	Cnd3	CAP-G	CAPG-1	Cap-G	CAP-G
	in II	-kleisin	-	-	CAP-H2	KLE-2	Cap-H2	CAP-H2
	Condensin	HEAT-IIA	-	-	CAP-D3	HCP-6	Cap-D3	CAP-D3
	Con	HEAT-IIB	-	-	CAP-G2	CAPG-2	-	CAP-G2
Regulators	Cyclin-dependent kinase		Cdc28	Cdc2	-	-	-	CDK1
	Aurora B kinase		lpl1	Ark1	-	AIR-2	Aurora B	Aurora B (AURKI
	Polo-like kinase		Cdc5	-	-	-	-	PLK1
Sm	c5/6							
Subunits	-SMC		Smc5	Smc5 (Spr18)	SMC5	SMC-5	Smc5	SMC5
	-SMC		Smc6 (Rhc18)	Smc6 (Rad18)	SMC6A/B	SMC-6	Smc6 (Jnj)	SMC6
	Kleisin		Nse4 (Qri2)	Nse4 (Rad62)	NSE4A/B	-	Nse4	NSE4A, NSE4B
	Tandem-WHD E3 ligase		Nse1	Nse1	NSE1	-	Nse1	NSE1
	Tandem-WHD		Nse3	Nse3	NSE3	-	Mage (Nse3)	MAGE-G1 (NSE
	SUMO ligase		Mms21 (Nse2)	Nse2 (Pli2)	NSE2	-	Quijote, Cervante	NSE2
ors			Nse5	Nse5	-	-	-	_
Co-factors			Kre29 (Nse6)	Nse6	-	-	-	SLF2
Ś	BR	CT domain	Rtt107?	Brc1?	-	_	-	SLF1

*Meiosis-specific subunit **Specific subunit of the dosage compensation complex

Prokaryotic SMC Complexes

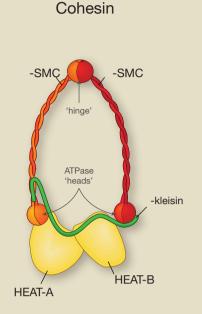


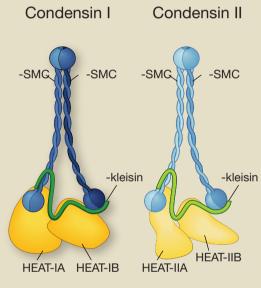


SMG SMC

Tandem-WHD Tandem-WHD

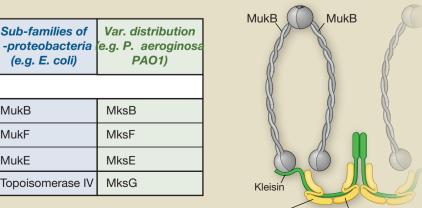






-SMC -SMC SUMO Ligase Kleisin Tandem-WHD

Smc5/6



Sub-families of Var. distribution

MksB

MksF

MksE

MksG

PAO1)

(e.g. E. coli)

Topoisomerase IV

MukB

MukF

MukE

MukBEF/MksBEF

SMC

Kleisin

Co-factor

Tandem-WHD

Subunits

Tandem-WHD Tandem-WHD

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This two-part SnapShot depicts the composition and architecture of SMC protein complexes and their regulators (in part I) and their roles at different stages of the cell cycle (in part II).

Part I: The architecture of SMC-kleisin complexes

The core of any Structural Maintenance of Chromosomes (SMC) protein complex is formed by a homoor heterodimer of SMC proteins. A single kleisin subunit binds via its N- and C-terminal interfaces to the neck domain of a v-SMC subunit and to the head domain of a κ -SMC subunit to create a tripartite ringlike architecture. The activities of the core complexes are regulated by diverse peripheral subunits, many of which are either composed of α -helical HEAT-repeats or Tandem Winged-Helix Domains (WHD) and associate via the kleisin protein.

In eukaryotic cells, at least three distinct SMC complexes (Cohesin, Condensin and Smc5/6) are needed for the correct segregation of chromosomes during mitotic and meiotic cell division cycles. During gametogenesis, several subunits of the Cohesin complex are substituted by paralogous proteins with specialized meiotic functions (marked by a single asterisk). In addition, specialized forms of some SMC complexes exist in most metazoans, such as Condensin I and Condensin II, or a *C. elegans*-specific Condensin I variant with roles in dosage compensation (marked by a double asterisk). Furthermore, The functions of SMC protein complexes are controlled by other regulatory proteins and through posttranslational modifications. These modifications include phosphorylation by several kinases (Polo-Like Kinase 1, Dbf4-dependent Kinase and Casein Kinase 1 for Cohesin; Cyclin-Dependent Kinase, Aurora B Kinase, Polo-Like Kinase 1, and Casein Kinase 2 for Condensin), acetylation, sumoylation, or proteolytic cleavage.

Most prokaryotic cells express a single SMC-ScpAB complex. MukBEF and MksBEF complexes found in certain bacteria feature an architecture that apparently deviates from the canonical SMC–kleisin organization. The N-terminal domain of kleisin MukF (and MksF) forms dimers and might thus be able to create structures with four or more SMC–kleisin subunits, such as tetrapartite rings or dimers of dimer (see cartoon).

ABBREVIATIONS

CDK1, Cyclin-Dependent Kinase 1; HEAT, Huntingtin, Elongation factor 3, protein phosphatase 2A, Tor1 kinase; PLK1, Polo-Like Kinase 1; SLF, Smc5/6 Localization Factor; SMC, Structural Maintenance of Chromosomes; WHD Winged-Helix Domain

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