Table 1. Cardiac phenotype	s associated with	mutations in PcG of	or TrxG genes
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	Complex	Enzymatic activity	Subunits (core & accessory) <sup>c</sup>	Cardiac phenotypes in mutant or transgenic mice	Additional information	
PcGª	DBC2	methyltransferase specific for H3K27 (Czermin <i>et al.</i> 2002, Muller <i>et al.</i> 2002)	Ezh1/2	<i>Ezh2<sup>NK</sup></i> : hypoplasia in complex myocardium, excessive trabeculation, spetal defects, dilation in RV.	Ezh2 methylates GATA4 and inhibits GATA4 activity	
				<i>Ezh2</i> <sup>SHF</sup> : cardiac hypertrophy and fibrosis in RV during adulthoood		
	11(02		Eed	<i>Eed</i> <sup><i>TnT</i></sup> : hypoplasia in complex myocardium, excessive trabeculation		
			Suz12	n/d	]	
			Jmj	<i>Jmj</i> <sup>-/-</sup> : excessive trabeculation, DORV, VSD, ventricular wall noncompaction		
	PRC1	chromatin compaction (Fischle <i>et al.</i> 2003, Francis <i>et al.</i> 2004), ubiquitin ligase specific for H2AK119 (Wang <i>et al.</i> 2004)	Phc1/2/3	<i>Phc1<sup>-/-</sup></i> : looping defect, VSD, pulmonary stenosis, aortic stenosis, tetralogy of Fallot	n/a	
				<i>6 MHC-Pch1</i> : reduced contractility, sarcomere disorganization, cardiomyocyte apoptosis		
			Cbx2/4/8	n/d	]	
			Bmi1	n/d		
			Ring1	n/d		
			Rnf2	n/d		
	PR-DUB	deubiquitinase specific for H2AK119 (Scheuermann <i>et</i> <i>al.</i> 2010)	Bap1	n/d	n/a	
			Asxl1/2	n/d		
TrxG⁵		chromatin remodeling (Kwon <i>et al.</i> 1994, Phelan <i>et al.</i> 2000, Wang <i>et al.</i> 1996) E	Brm/Brg1	<i>Tie2Cre;Brg1<sup>F/F</sup></i> : sparse trabeculation	<i>Brg1</i> genetically interacts with <i>Nkx2.5</i> , <i>Tbx5</i> , <i>GATA4</i> , <i>Tbx20</i> ;	
				<i>Sm22a-cre;Brg1</i> <sup>f/f</sup> : thin compact myocardium and failure to form IVS, both due to proliferation defect; early differentiation of cardiomyocyte		
				<i>Brg1</i> inactivation in adult heart ( <i>Tnnt2-rtTA;Tre-cre;Brg1</i> <sup>f/f</sup> ): reduced hypertrophic response to TAC		
	BAF/BRM		BAF60a/b/c	<i>siBaf60c</i> : trabeculation defect, impaired development of SHF	BAF60c together with GATA4 induce cardiac fate in non-cardiogenic mesoderm; BAF60c enables GATA4 binding to target promoters; BAF60c potentiates GATA4, Nkx2.5, Tbx5 activity in reporter assays by recruiting BAF	
			BAF250	n/d		
			BAF170	n/d	]	
			BAF47	n/d		
	MLL	methyltransferase specific for H3K4 (Milne <i>et al.</i> 2002)	MLL1/2/3	n/d	Ash2I interacts with Tbx1 in a cancer cell line	
			Ash2l	n/d	and promotes Tbx1 activity in luciferase assays	
			Wdr5	n/d		

References
Delgado-Olguín <i>et al.</i> 2012, He <i>et al.</i> 2012a, He <i>et al.</i> 2012b, Jung <i>et al.</i> 2005, Kim <i>et al.</i> 2004, Lee <i>et al.</i> 2000, Mysliwiec <i>et al.</i> 2011, Nakajima <i>et al.</i> 2011, Shirato <i>et al.</i> 2009, Takeuchi <i>et al.</i> 1999, Toyoda <i>et al.</i> 2003
Koga <i>et al.</i> 2002, Shirai <i>et al.</i> 2002, Takihara <i>et al.</i> 1997
n/a

Hang *et al.* 2010, Licket *et al.* 2004, Stankunas *et al.* 2008, Takeuchi and Bruneau 2009, Takeuchi *et al*. 2011

Stoller et al. 2010

<sup>a</sup>Drosophila contains a fourth PcG complex, PhoRC, which is thought to recruit other PcG complexes to special regulatory elements known as Polycomb Response Elements (PRE). However, the mechanism of PcG recruitment in mammals appears to be distinct from that in Drosophila, and a mammalian complex that is functionally equivalent to PhoRC has not been identified.

<sup>b</sup>A "supercomplex" containing components of BRM/BAF, MLL and several non-TrxG complexes has also been identified in mammalian cells (Nakamura et al. 2002).

<sup>c</sup>Subunits that are essential for the complex's biochemical properties are considered core subunits. In some cases, a core component can be encoded by multiple homologs that are expressed in a cell type-specific manner. Subunits that are not essential for the complex's activity but are part of the complex under specific circumstances are considered accessory subunits. n/d: not determined

n/a: not available