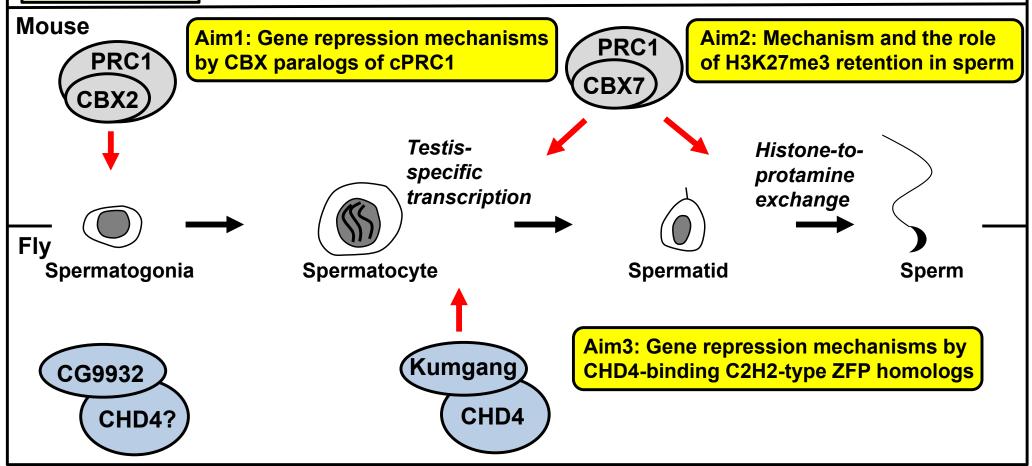
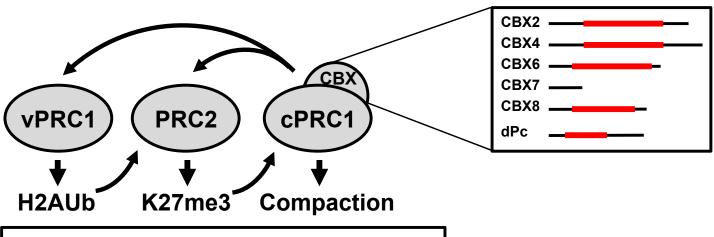


How to block aberrant transcription?

Choosing right promoters by cell type-specific chromatin modifying complexes



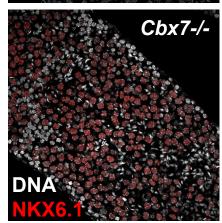
Aim1. Gene repression mechanism by CBX paralogs of cPRC1



Cbx7+/+

DNA
NKX6.1

- Q) Determinants of gene repression?
- Q) Why CBX7 but not other CBX?



Wild type

Cbx7-/-





vPRC1-cUbMut

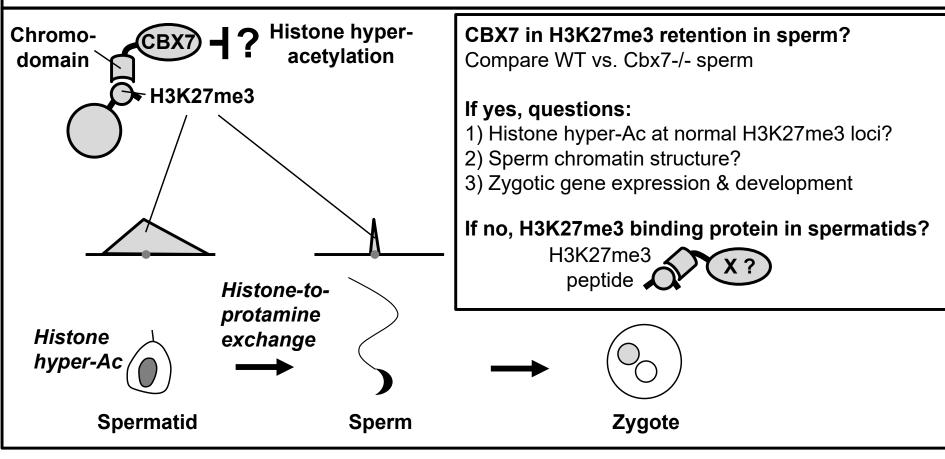
Cbx7-/-; rescue-CBX7-dTAG

Cbx7-/-; rescue-CBX2-dTAG

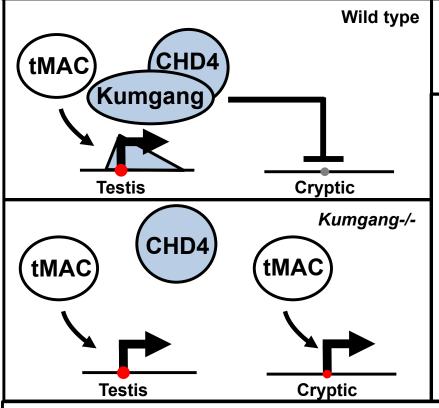
Measure genome-wide: vPRC1, PRC2, cPRC1, H2AUb, H3K27me3, Compaction

Analyze cellular phenotypes

Aim2. Mechanism and the role of H3K27me3 retention in sperm



Aim3. Gene repression mechanisms by CHD4-binding ZFP homologs



Hypothesis

Global disruption of chromatin structure, such as yeast Spt6 mutant (elongation factors)

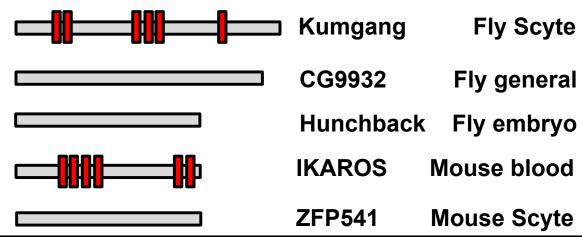
Approach

Acute-depletion of Kumgang

DD-DD-Kumgang flies (depletion of Kumgang by TMP withdrawal)

Measure time course

- Gene expression
- Genome-wide CHD4, tMAC binding
- Nucleosome positioning by MNase-seq



- **Q) Role of ZF in Kumgang** Individual Zf mutant rescue
- Q) Role of CHD4-binding ZFPs

Kumgang homolog rescue

Future: isolation of dominant suppressor EMS mutants reverting fertility of kumgang