

Figure 9. Nuclear organization in S. pombe. (A) Electron microscopy analysis of an S. pombe nucleus. (Top) Micrograph of a cross-section through a high-pressure fixed and Lowicryl-embedded interphase S. pombe cell. The cellular structures are indicated: cell wall, nuclear envelope, nucleolus, heterochromatin region, and SPB. (Bottom) A higher magnification of the same nucleus. The nuclear structures indicated are SPB, γ-tubulin region, anchor structure, and the centromeric heterochromatin. (B) Two interphase nuclei with heterochromatin (centromeres, telomeres, and the silent mat2-mat3 loci) decorated by red fluorescent immunolocalization of Swi6, and kinetochore chromatin (centromeres only) decorated by green fluorescent immunolocalization of CENP-A^{Cnp1}. The red signals, not in close proximity to green, represent telomeres or the mat2-mat3 loci. All centromeres are clustered at the nuclear periphery adjacent to the SPB. (C) A model for chromatin organization at the fission yeast nuclear periphery. (Top) Genes with low expression levels tend to associate with the nuclear periphery, whereas highly expressed genes tend to reside in the nuclear interior. (Top) Localization of divergent intergenic regions and H2A.Z at the nuclear envelope may present a mechanism for anchoring the promoters of convergent gene pairs at the periphery. (Bottom left) Differential localization of Ima1, nuclear pores, and Man1. The inner membrane proteins Ima1 and Man1 are not equally distributed at the nuclear periphery, but rather occupy distinct areas that interact with different chromosomal regions. The subtelomeric chromatin is associated with Man1-rich peripheral regions in which Swi6 is also located. Ima1 is colocalized with Dcr1 and Rdp1 at nuclear pores. (Bottom right) Organization of centromeric DNA at the SPB. Central domain cnt and imr regions are localized closer to the SPB than the heterochromatic dg and dh repeats. The two centromeric domains are shaded in colors symbolizing the different IF localization of Swi6 (dg-dh repeats) and CENP-A^{Cnp1} (imr/cnt regions). (A, Reprinted from Kniola et al. 2001; C, bottom right, Adapted from Takahashi et al. 1992.)