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Supplemental Figure S2. Characteristics of PacBio sequencing. (A) Length distribution of CCS, CCS-FL and Iso-Seq reads at the 6 different time points. (B) Scatter plots of log2 transformed CCS-FL read counts from replicates at 6 time points with cutoff of 5. Pearson correlation was shown in each plot. (C) Scatter plot showing a strong Pearson correlation between PacBio CCS-FL read counts and RPKMs from Illumina-based short read RNA-seq (Duncan et al. 2014) at 0 h of meiosis.