FIGURE S11.—Real and false induced mate-pair CNV candidates. Depictions of array data and anomalous mate-pairs are similar to Figures 1 and 2. An additional track shows traces correlating to the coverage density of ~3 kb mate-pair mappings for the parental (blue) and aphidicolin-treated (magenta) samples. (A) A validated induced 88 kb deletion on Chr11. (B) A false 61 kb candidate deletion on Chr20. Note the lack of change in either the array intensity or 3 kb fragment coverage as observed in (A).