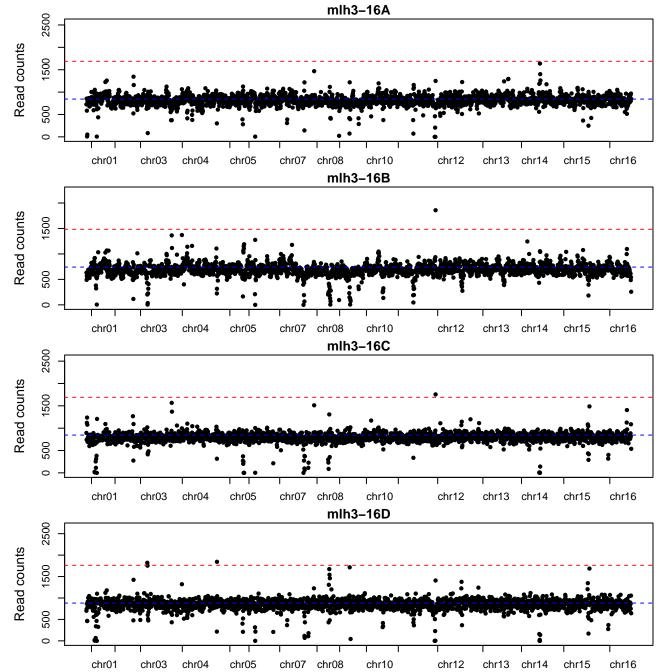
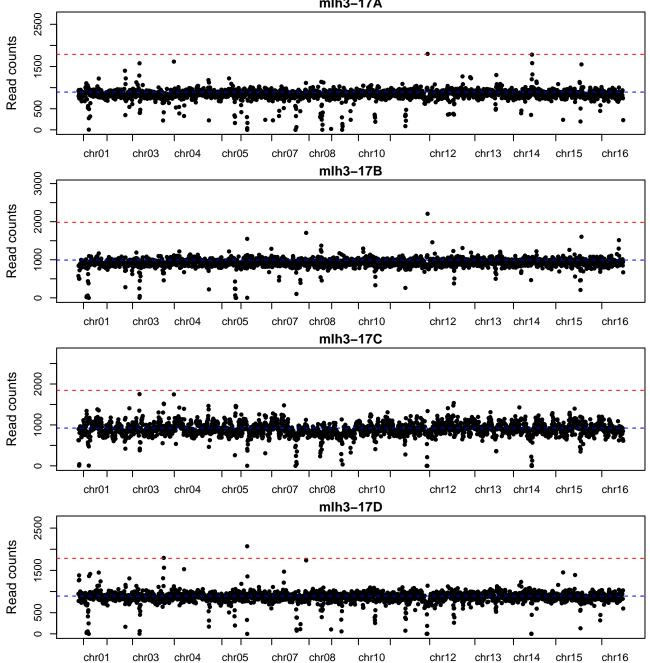
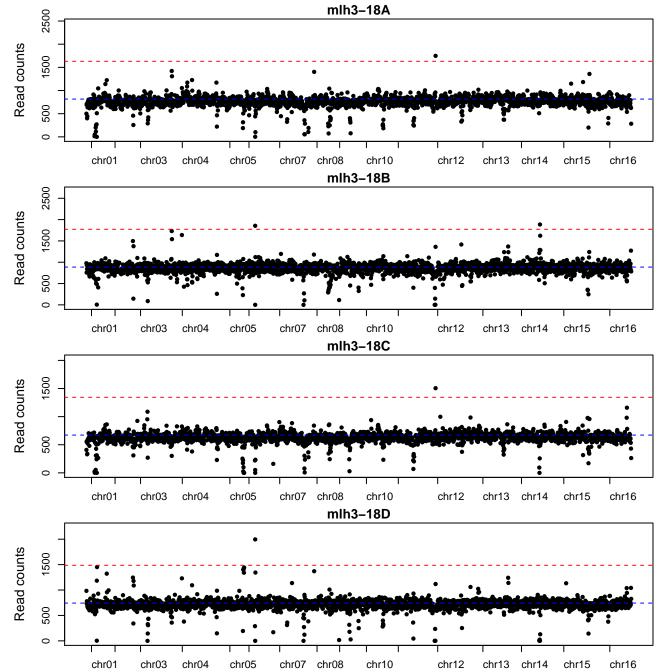


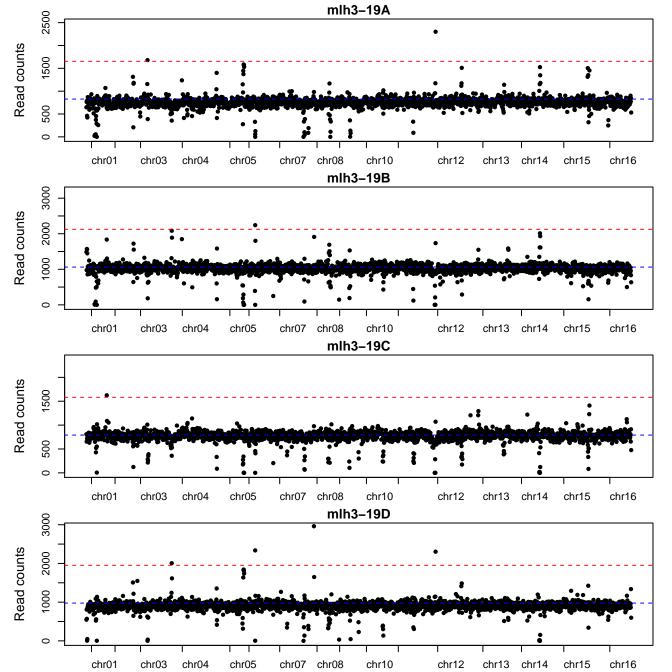
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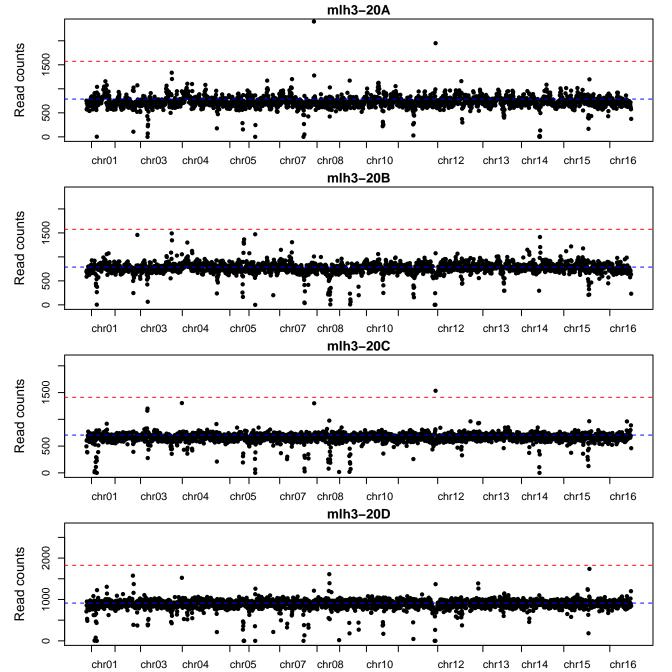


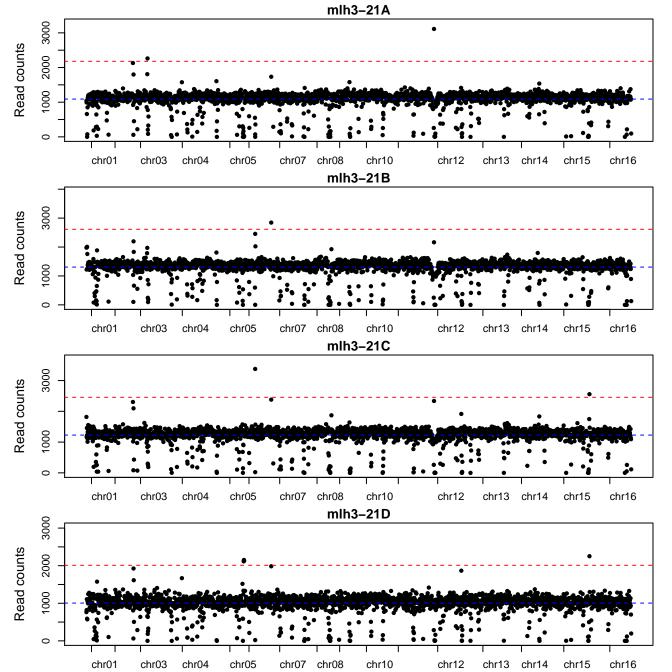


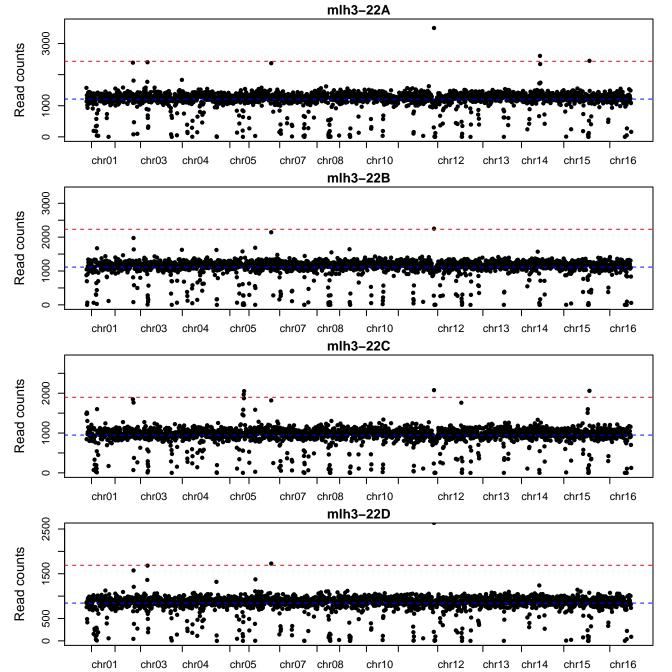
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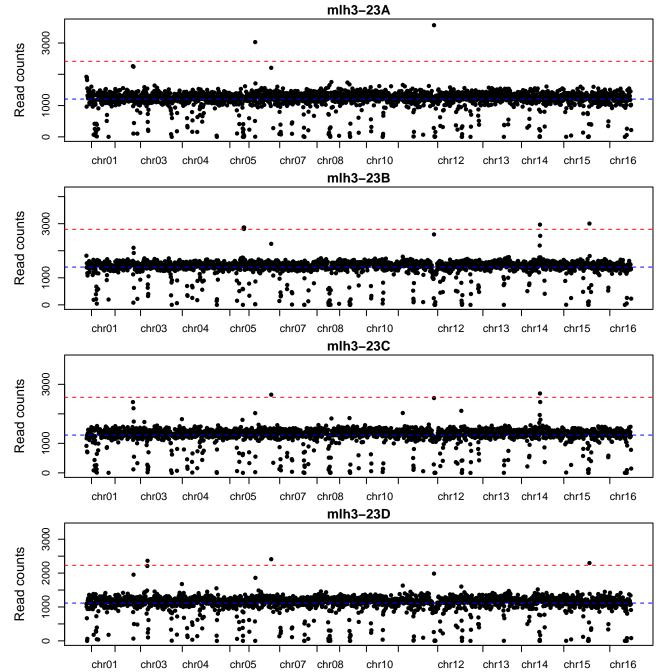


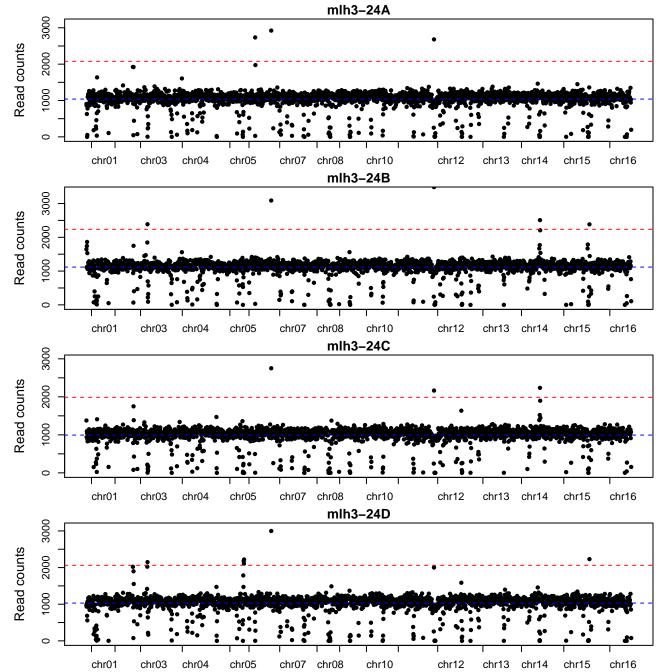


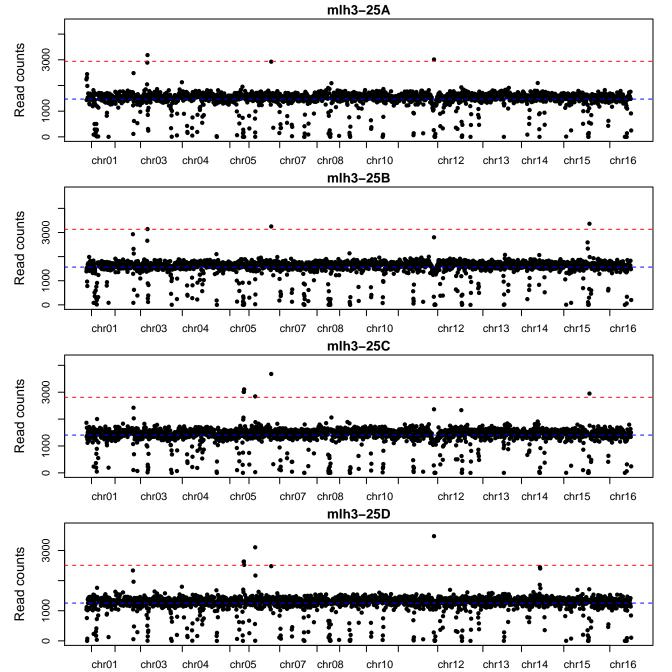


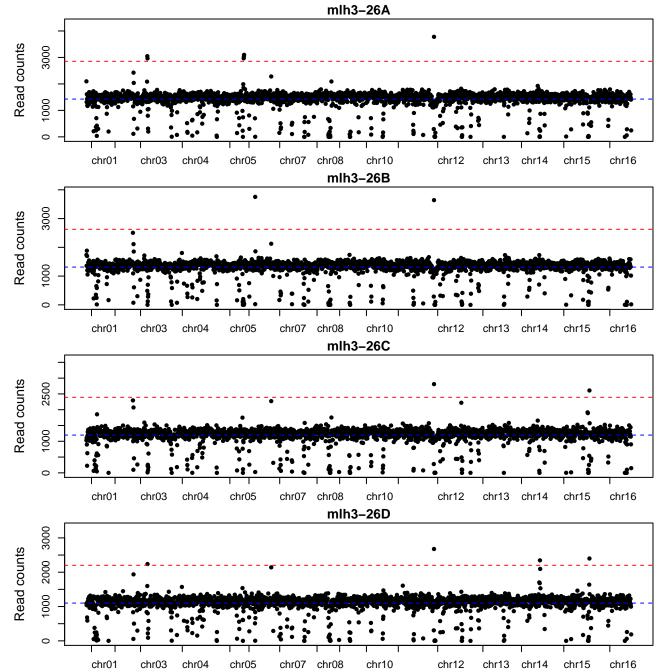


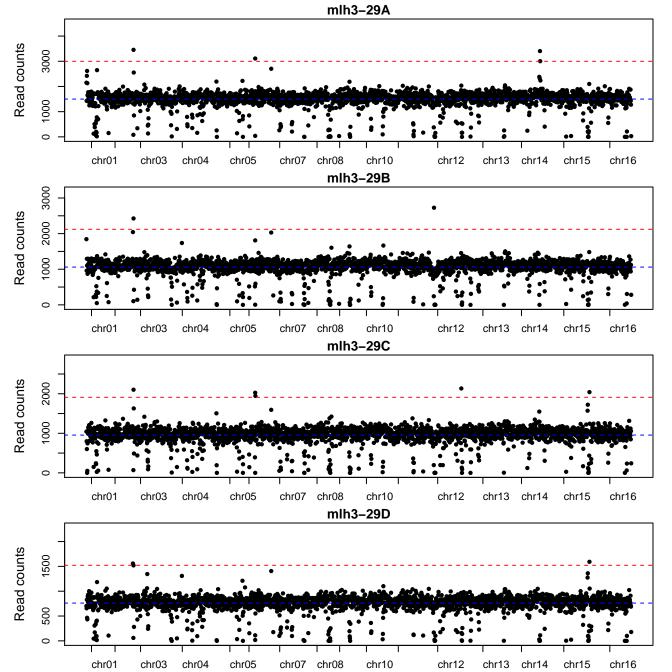


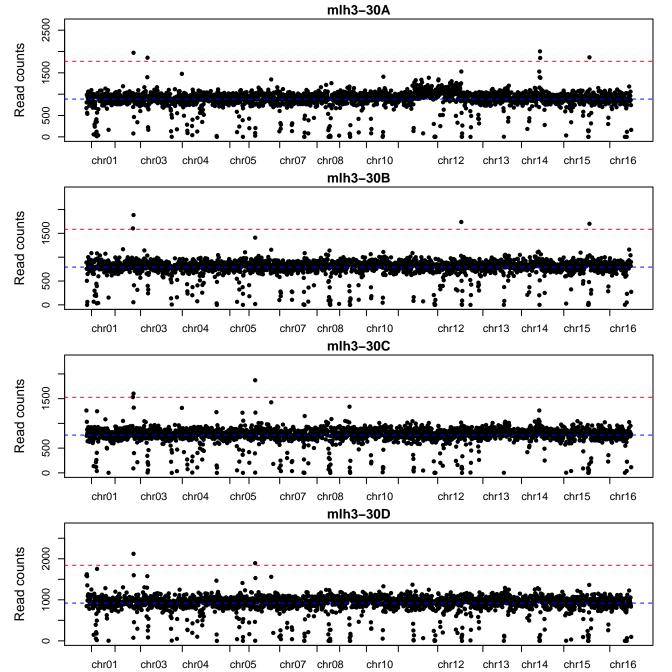


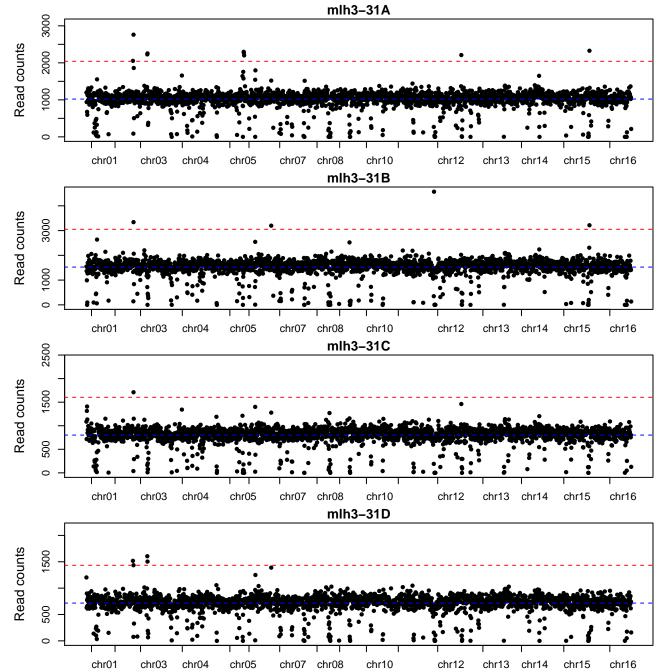


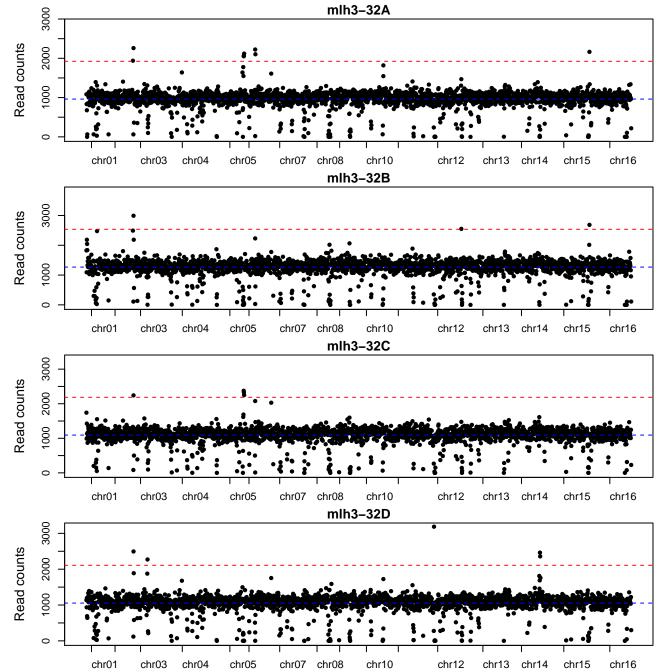


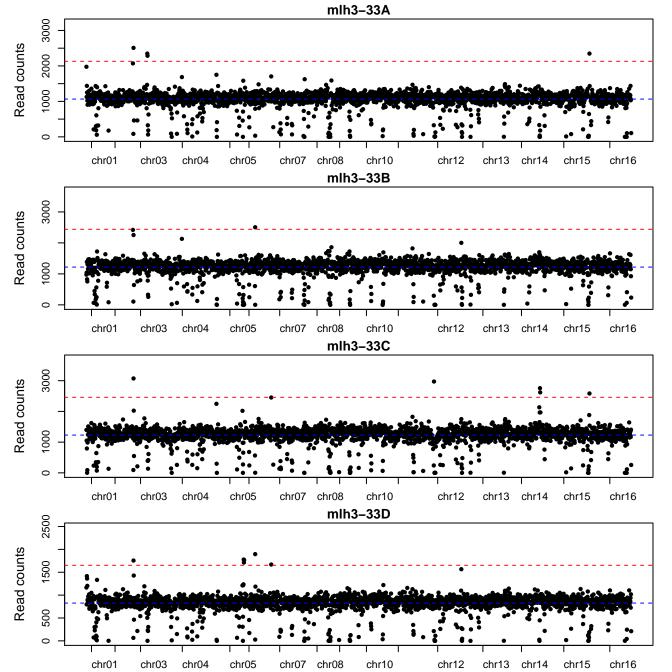


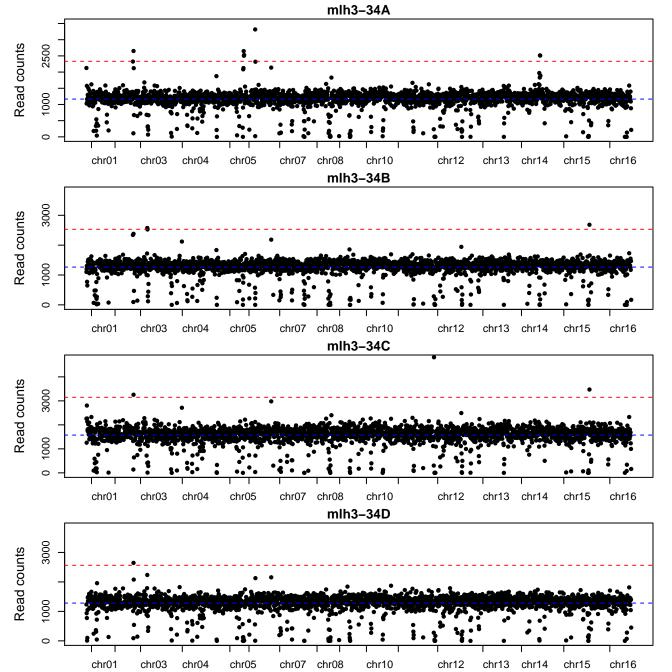


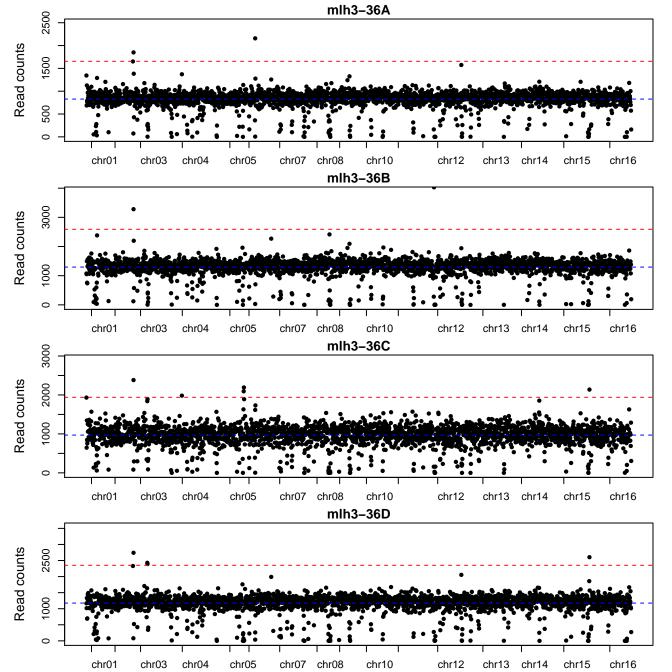


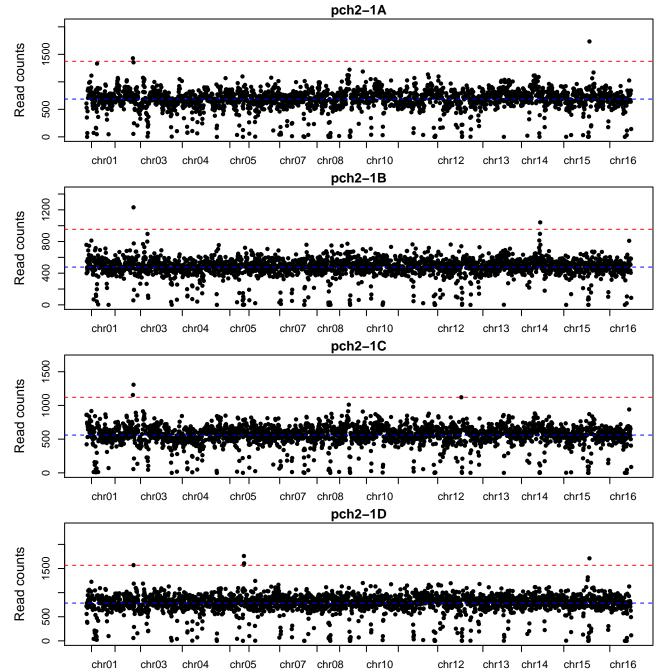


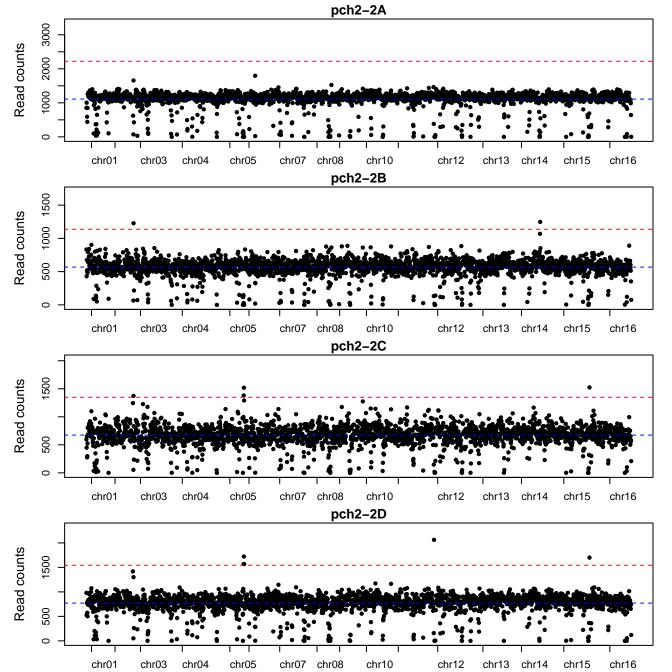


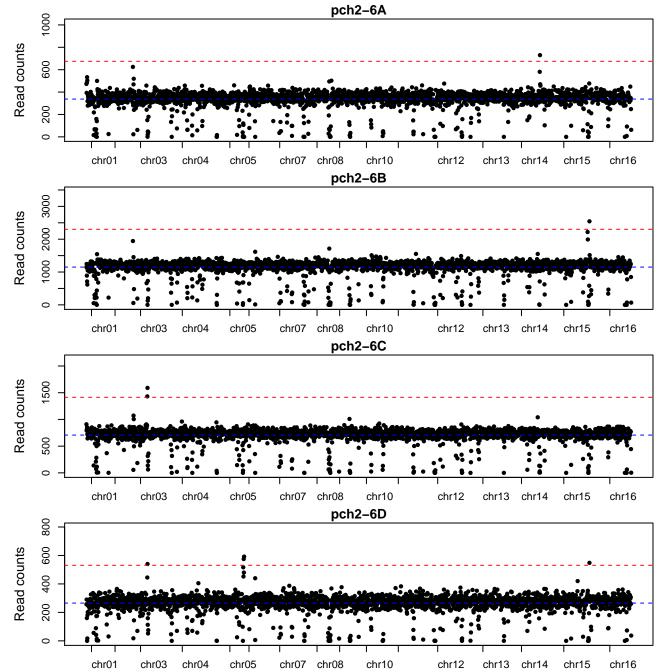


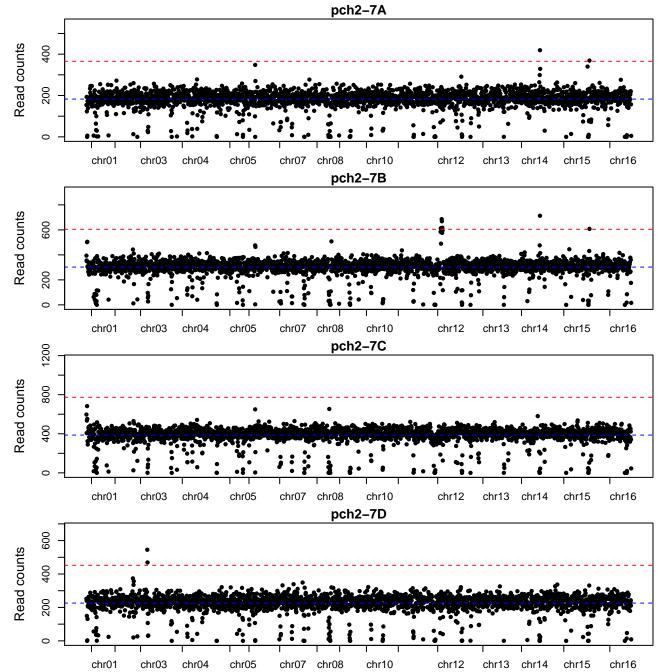


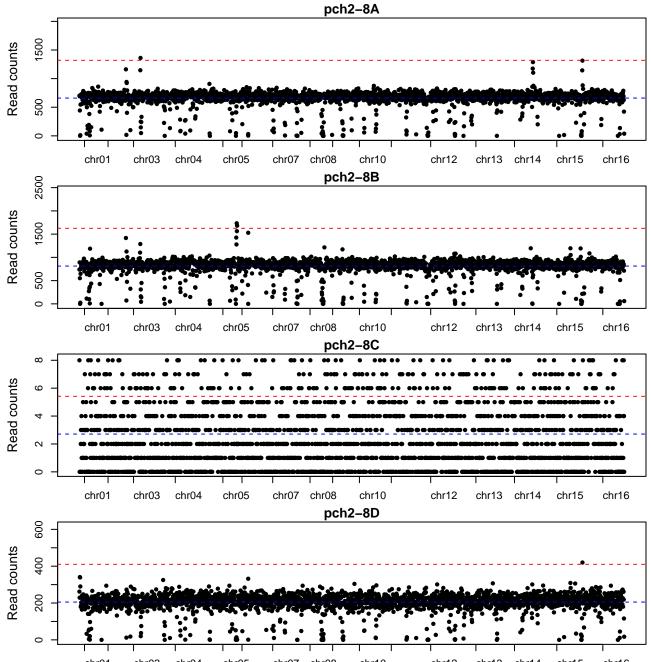




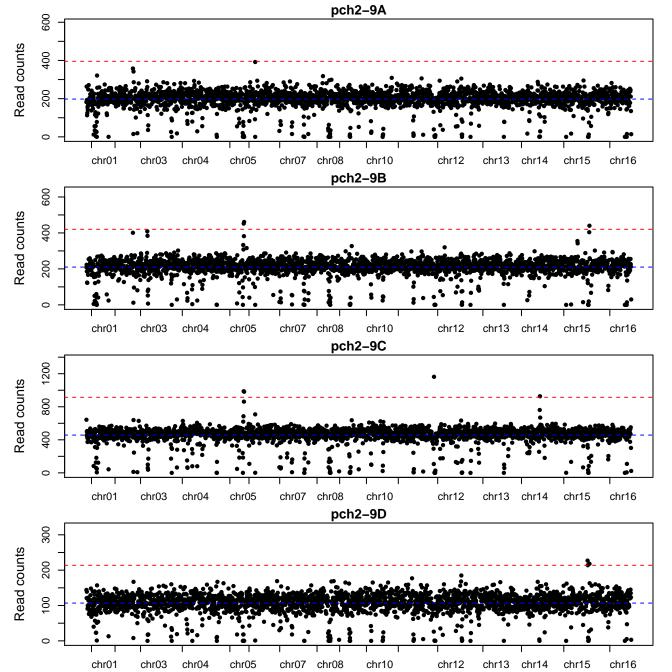


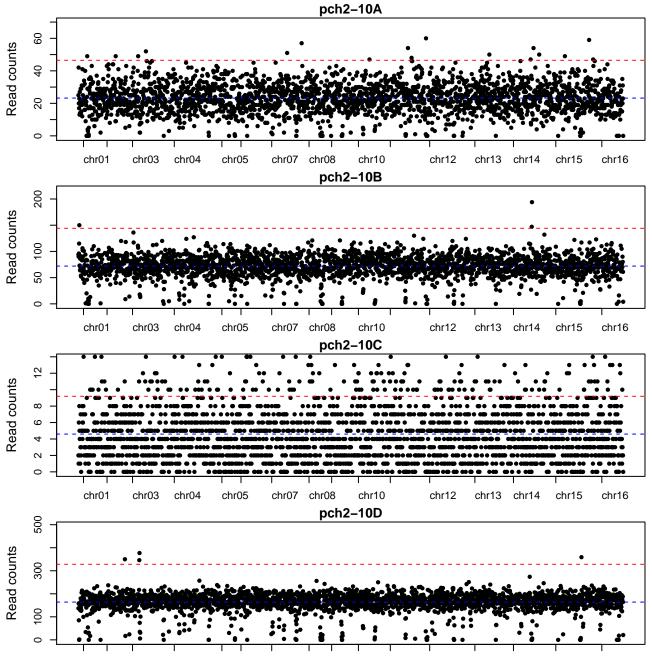




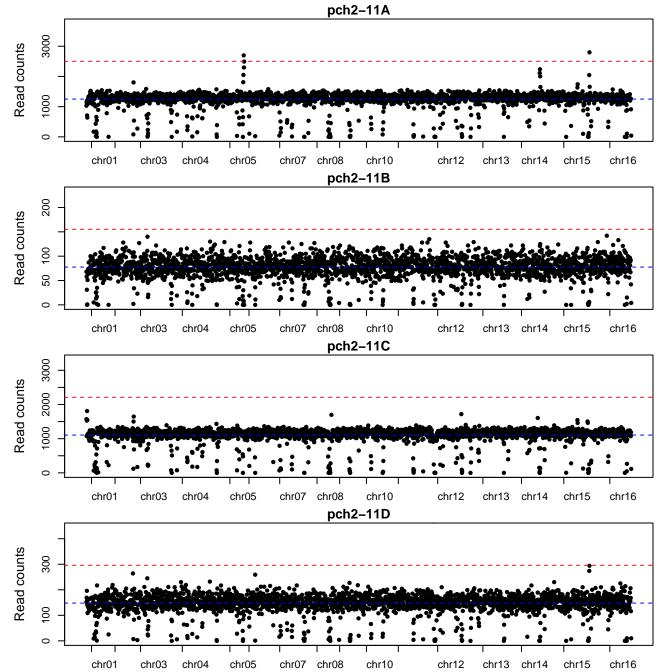


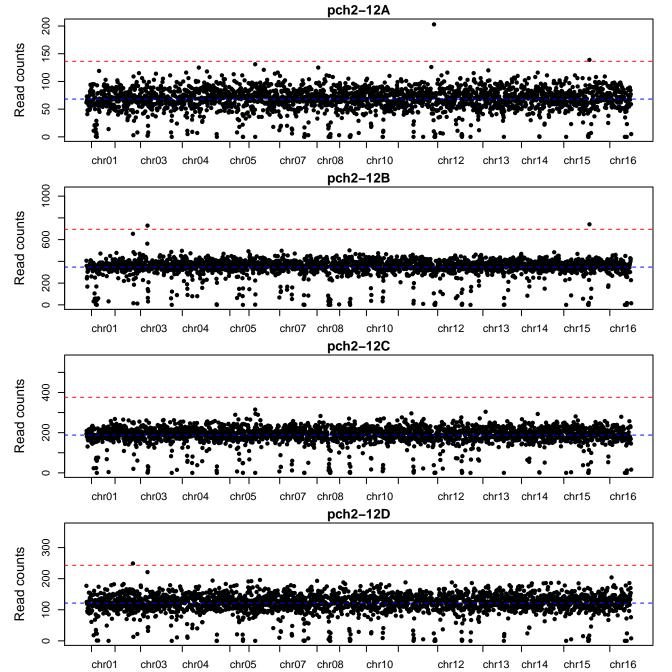


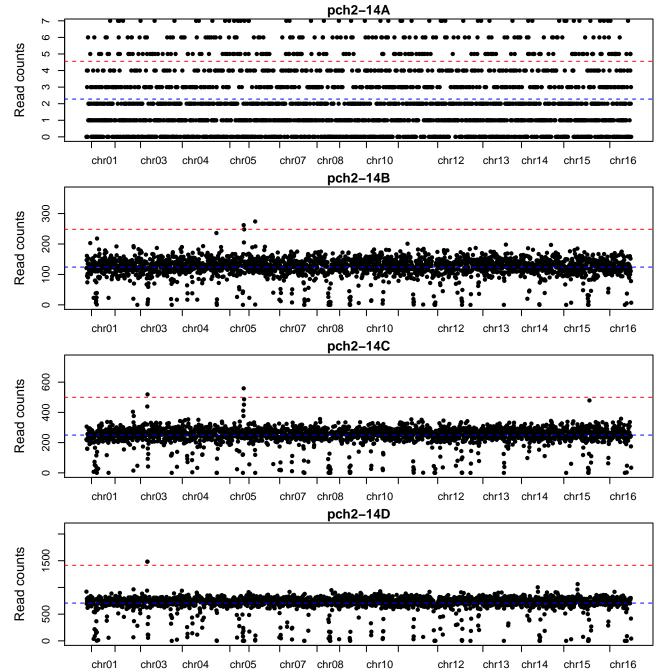


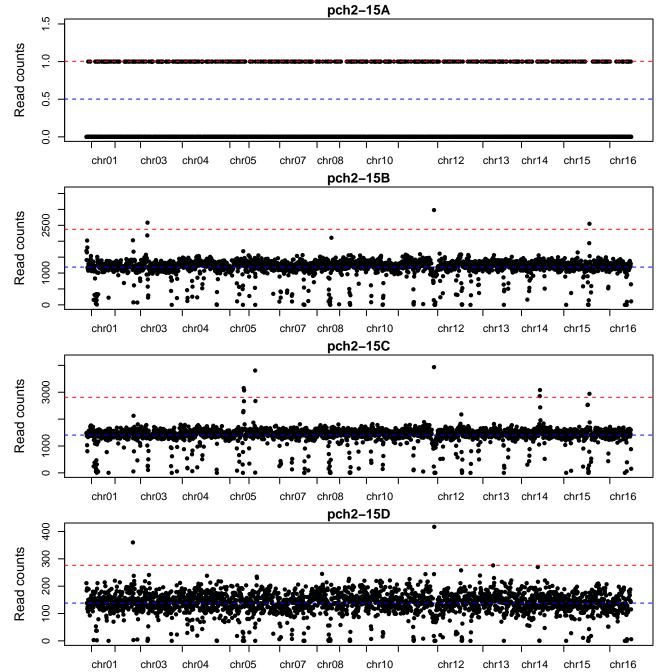


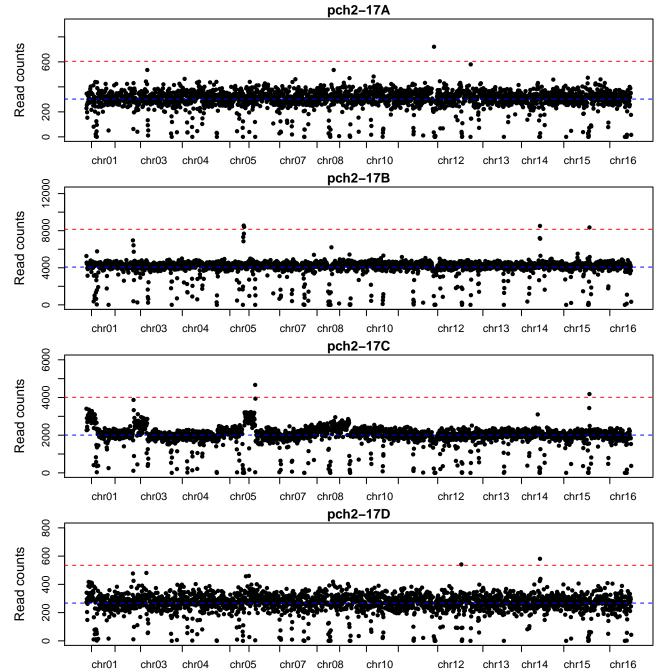
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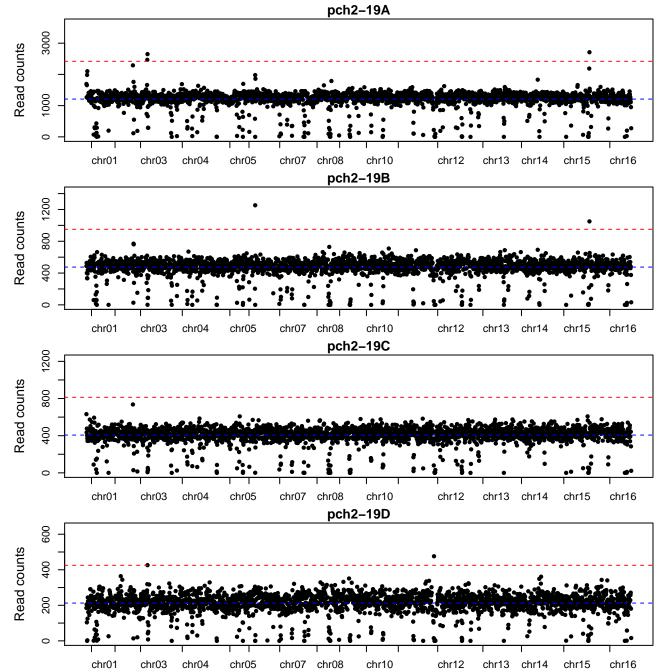


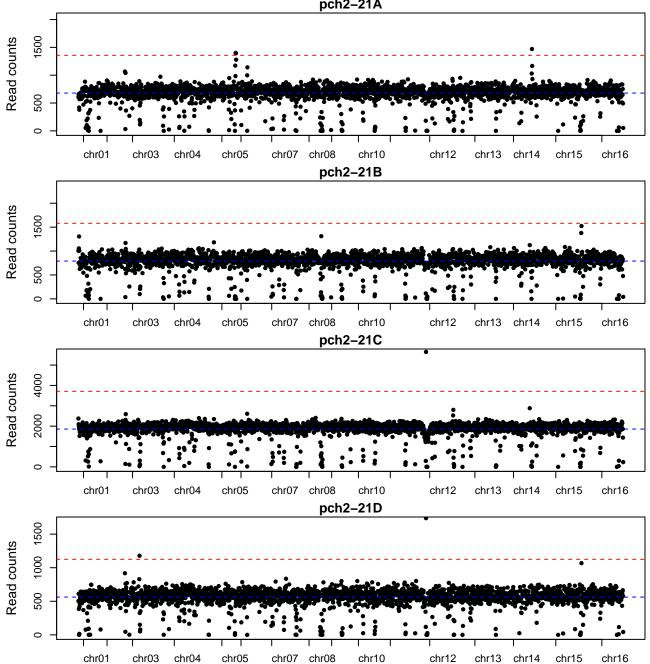




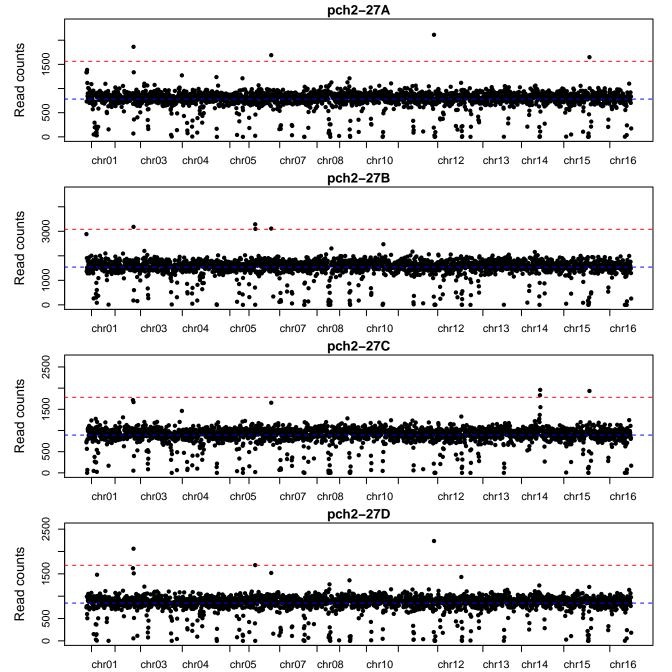


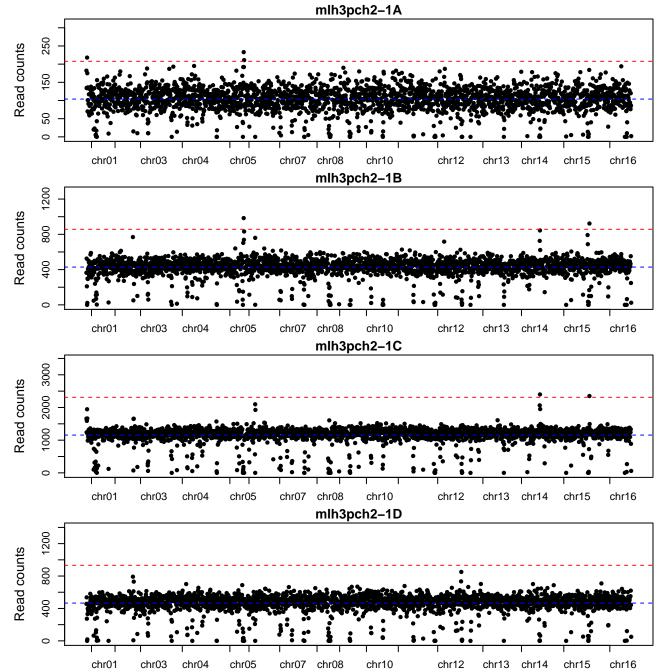


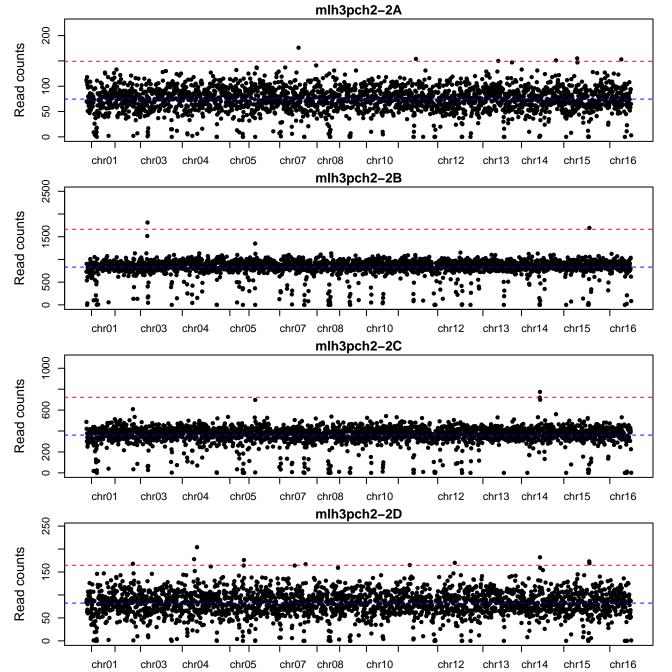


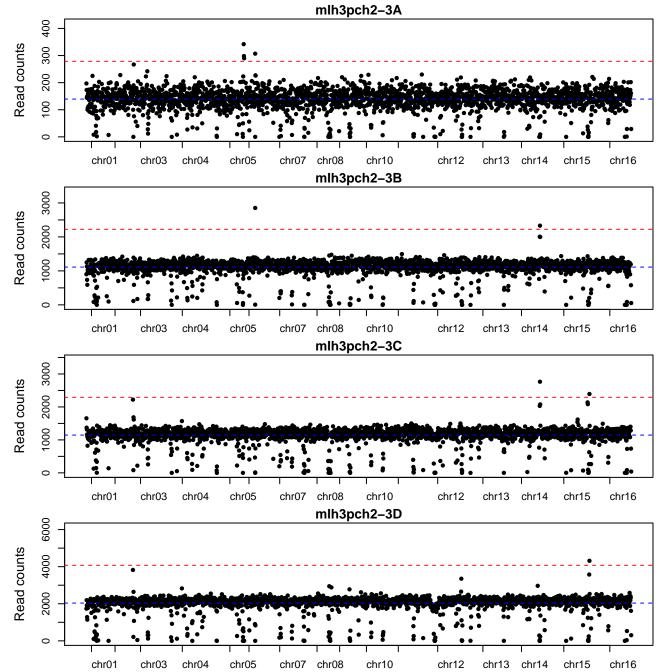


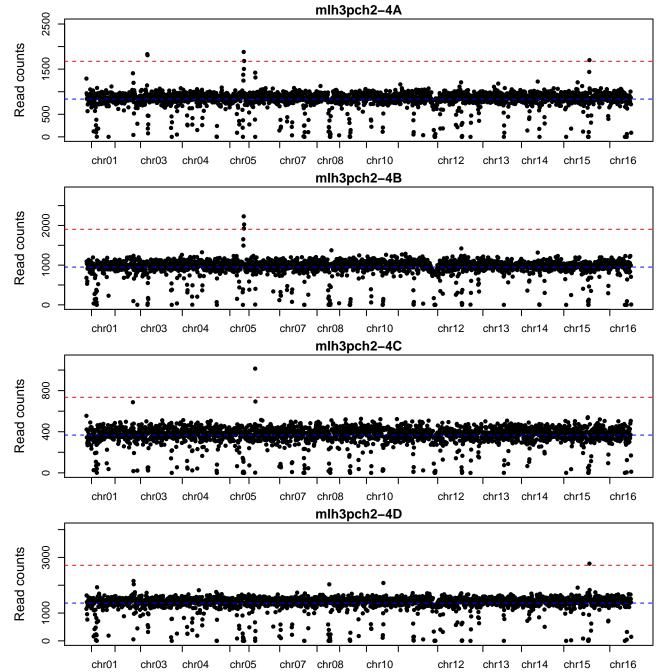
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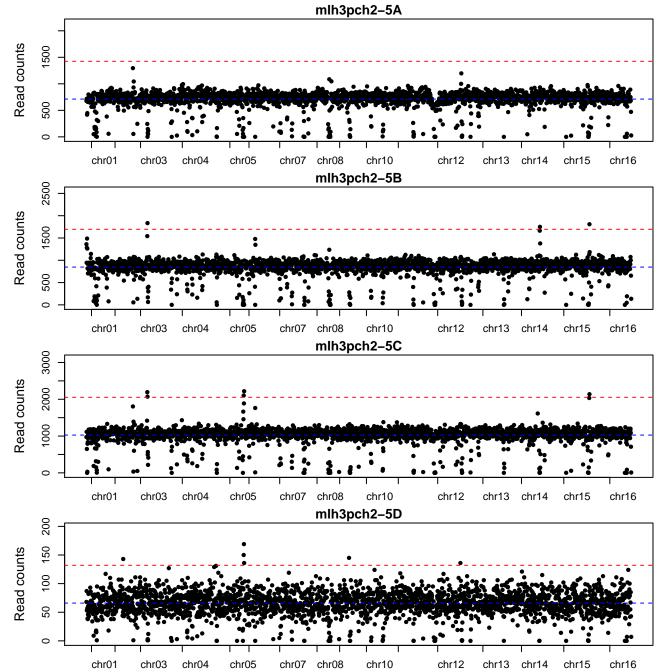


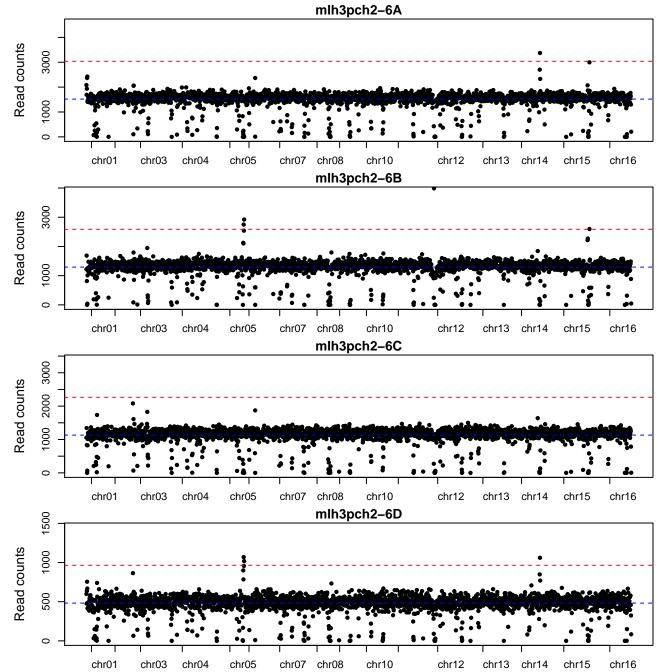


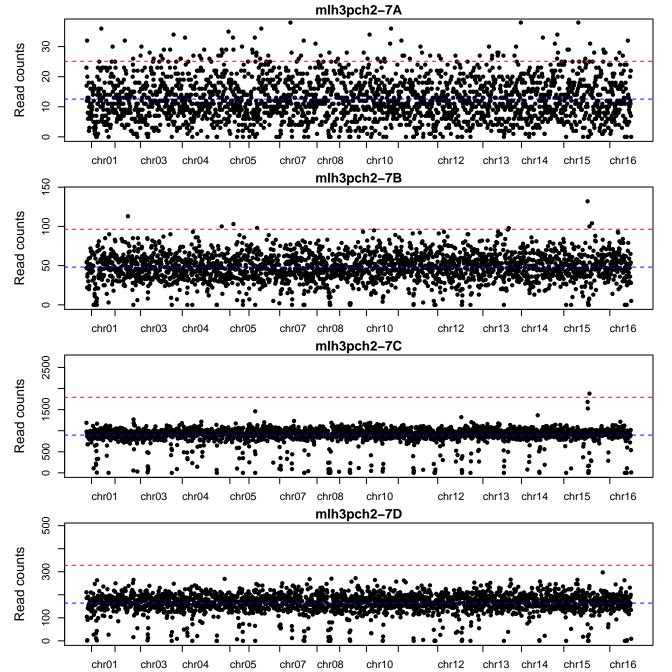


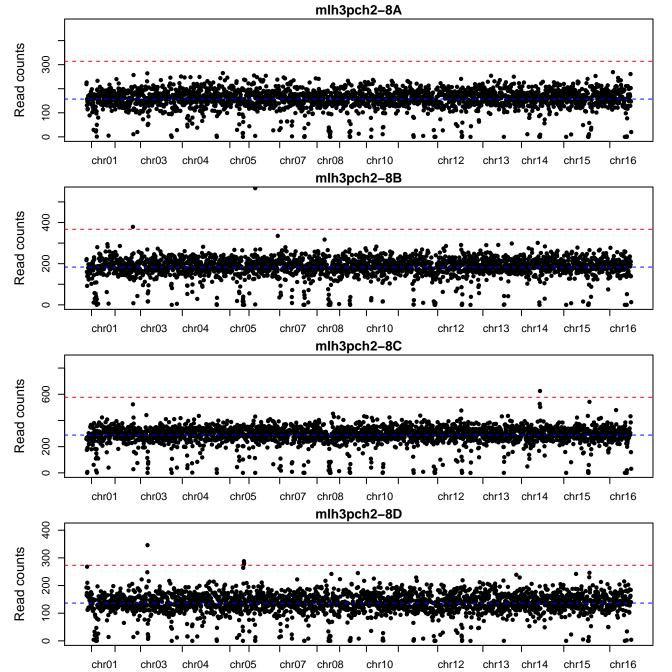


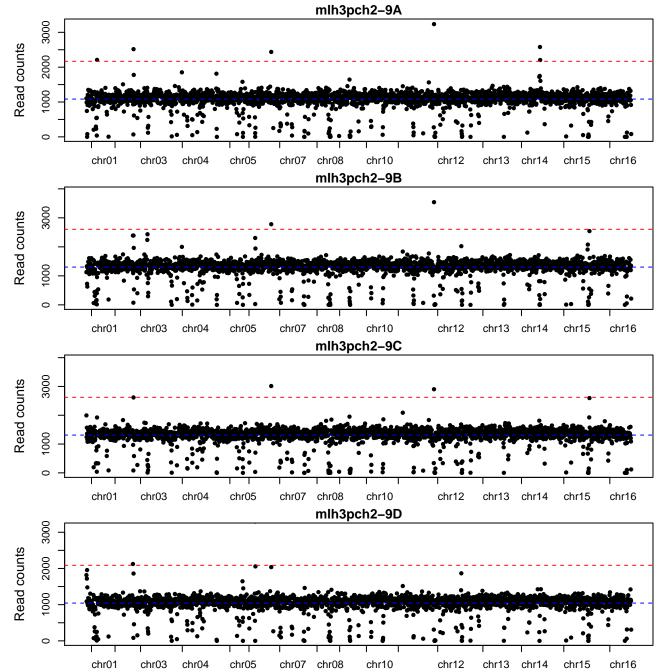


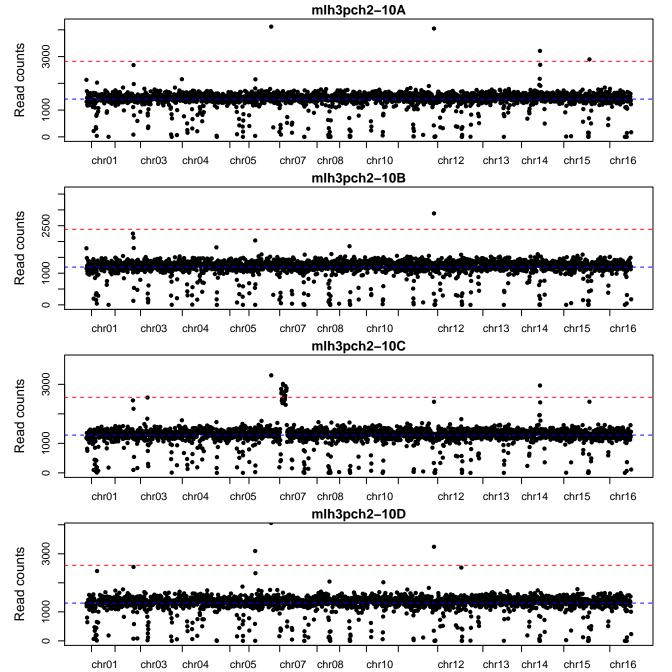


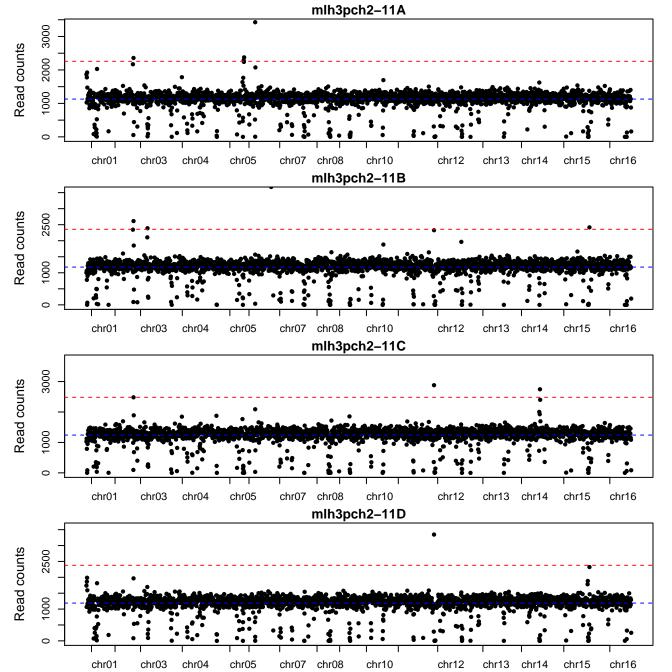


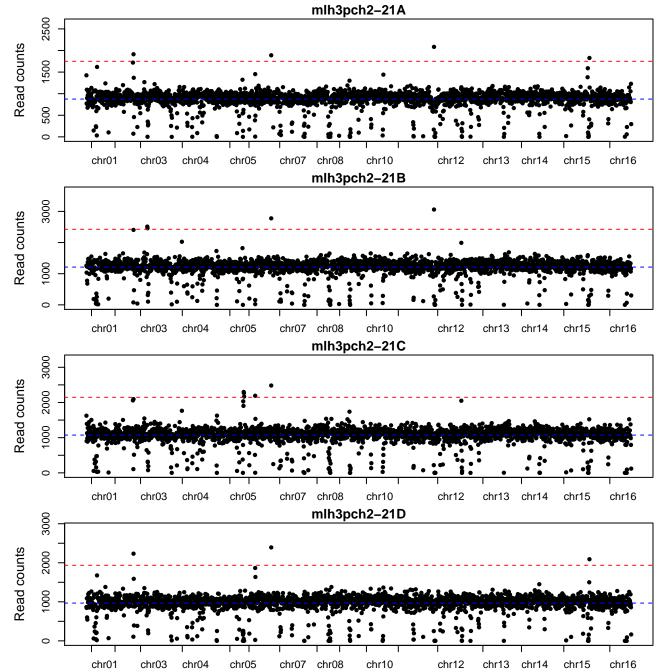


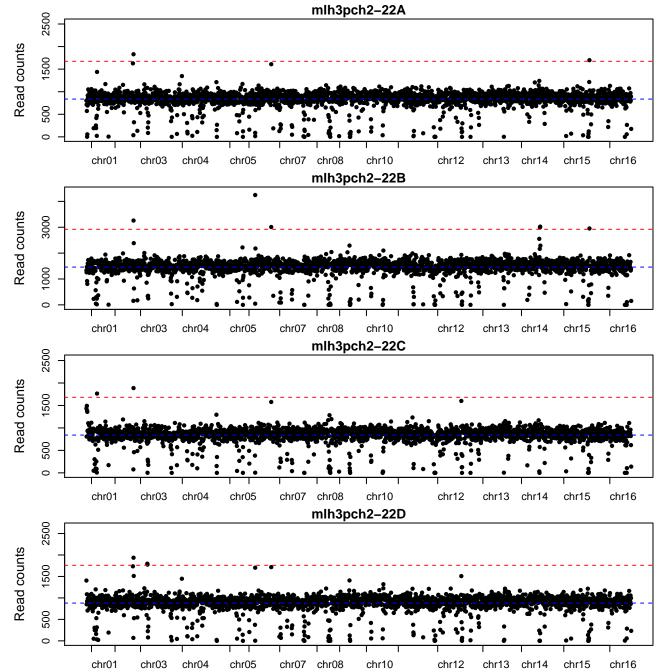


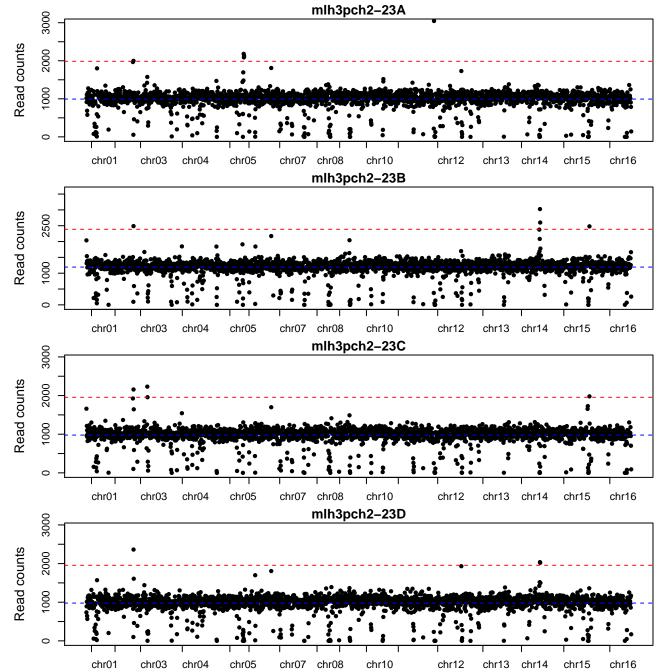


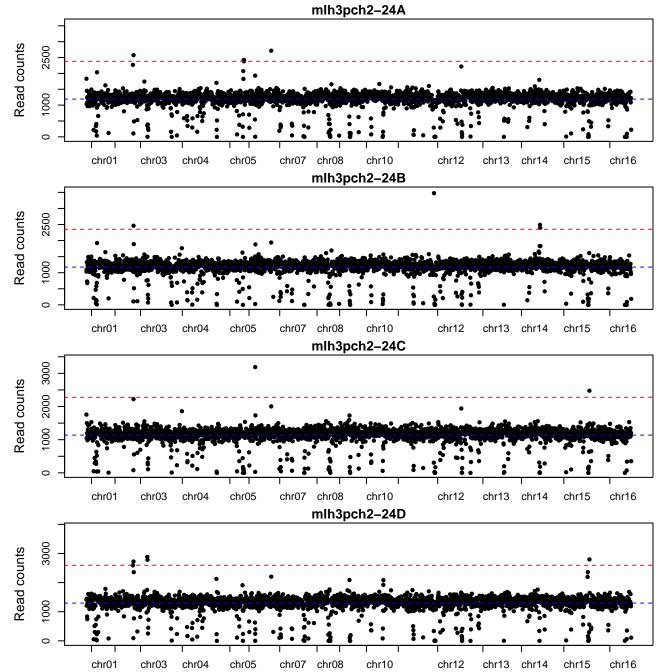


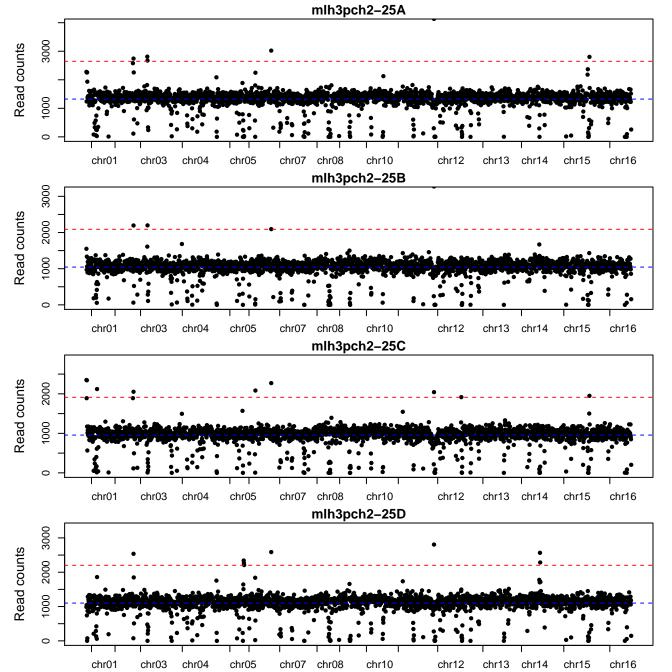


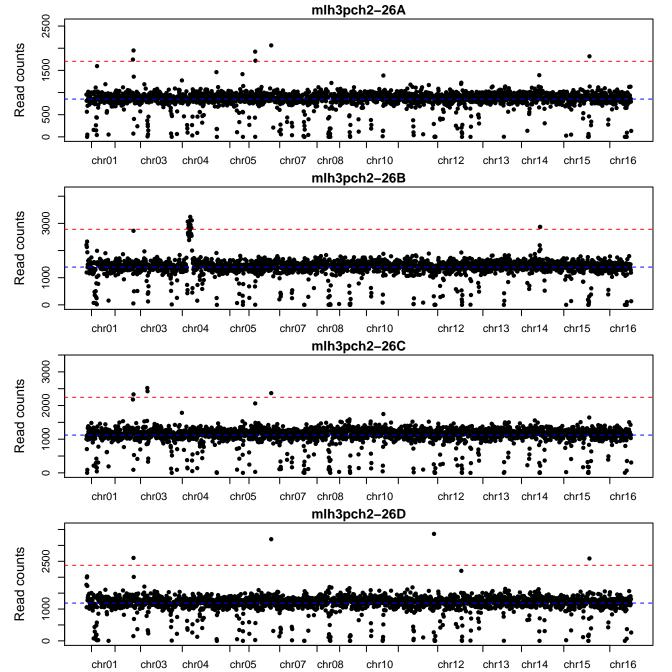


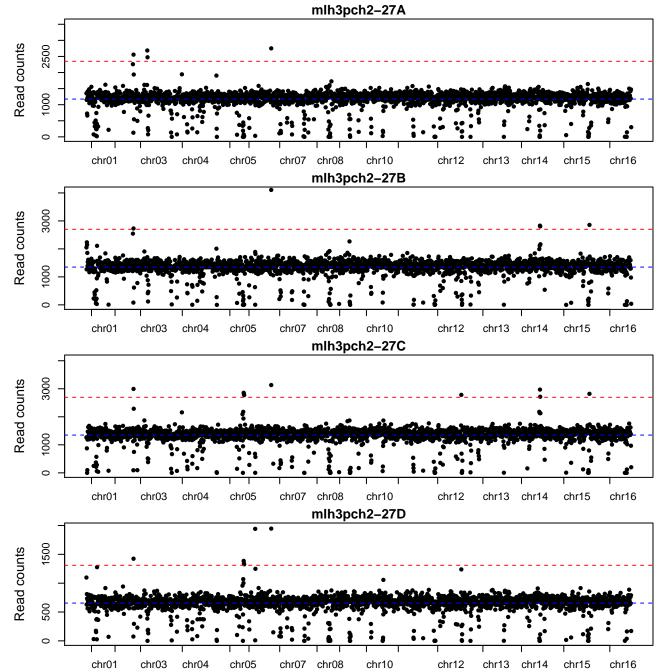


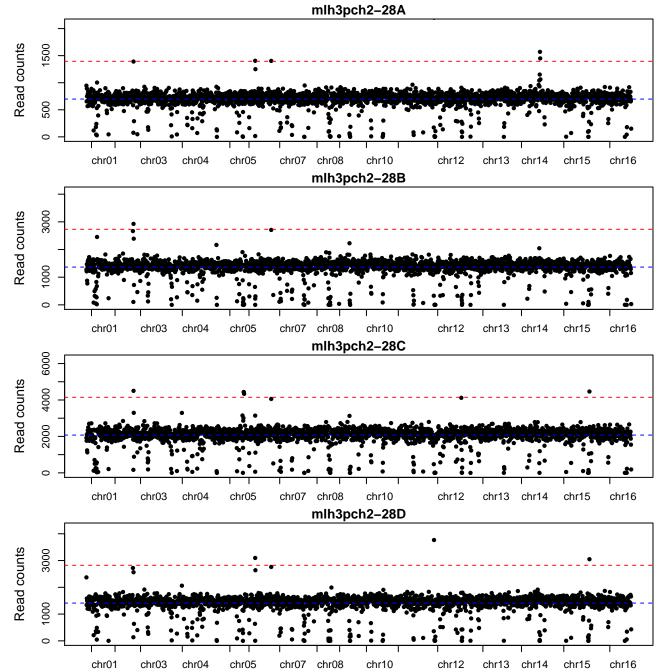


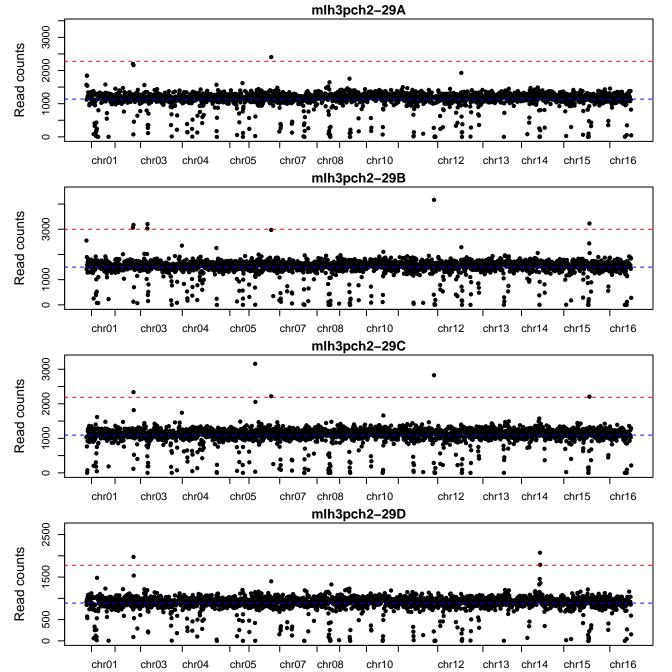












**Figure S9** Genome wide coverage plot to detect an euploidy in sequenced spores from  $mlh3\Delta$ ,  $pch2\Delta$  and  $mlh3\Delta$   $pch2\Delta$  tetrads. Each dot indicates the total read count in 5 kb bins. Average read count is shown as dotted blue line. Red line indicates 2X average read count.