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Orientation about actual trends and results in our understanding of the organization, evolution, and function of the plant genome at the level of the DNA (gene), the level of chromatin, and the level of the karyotype. 3. Presentation of hypotheses and models which may be stimulating for further research. Moreover, younger students should have the possibility to present their results and to

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This supports a recent assertion that the chromatin organization establishes functional and spatial biases on

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specific regions of the genome [3,6], and opens up a possibility that variation in mutation rate is adaptive [3], although this has been debated [4].

Chromatin structure and evolution in the human genome ...

Conclusion: We conclude that, overall, non-CpG mutation rates are lowest in open regions of the genome and that regions of the genome with a closed chromatin structure have the highest background mutation rate.

Chromatin structure and evolution in the human genome

The availability of a map of long-range chromatin structure across the human genome allows us to assess this idea and, through the comparison of various measures of neutral variation, we have identified those forms of chromatin structure associated with the highest rates of background mutation.