The future: tree sequence inference

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Inferring whole-genome histories in large population datasets

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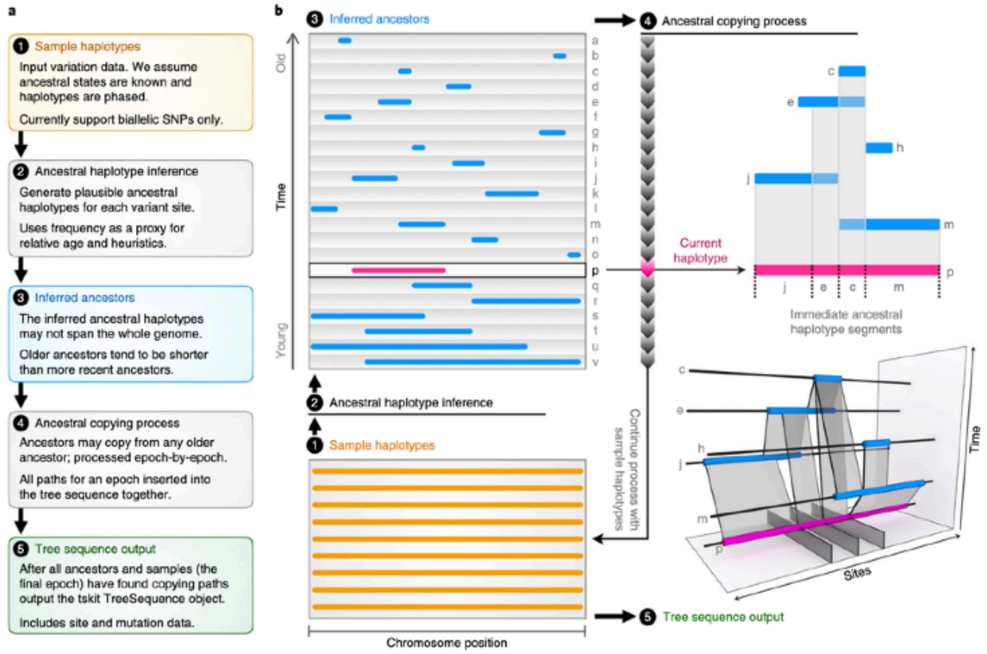

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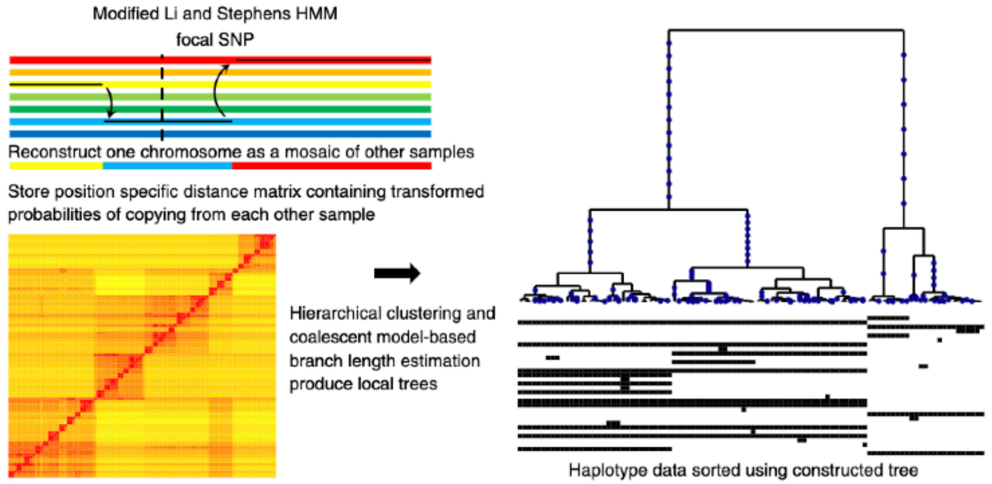
A method for genome-wide genealogy estimation for thousands of samples

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