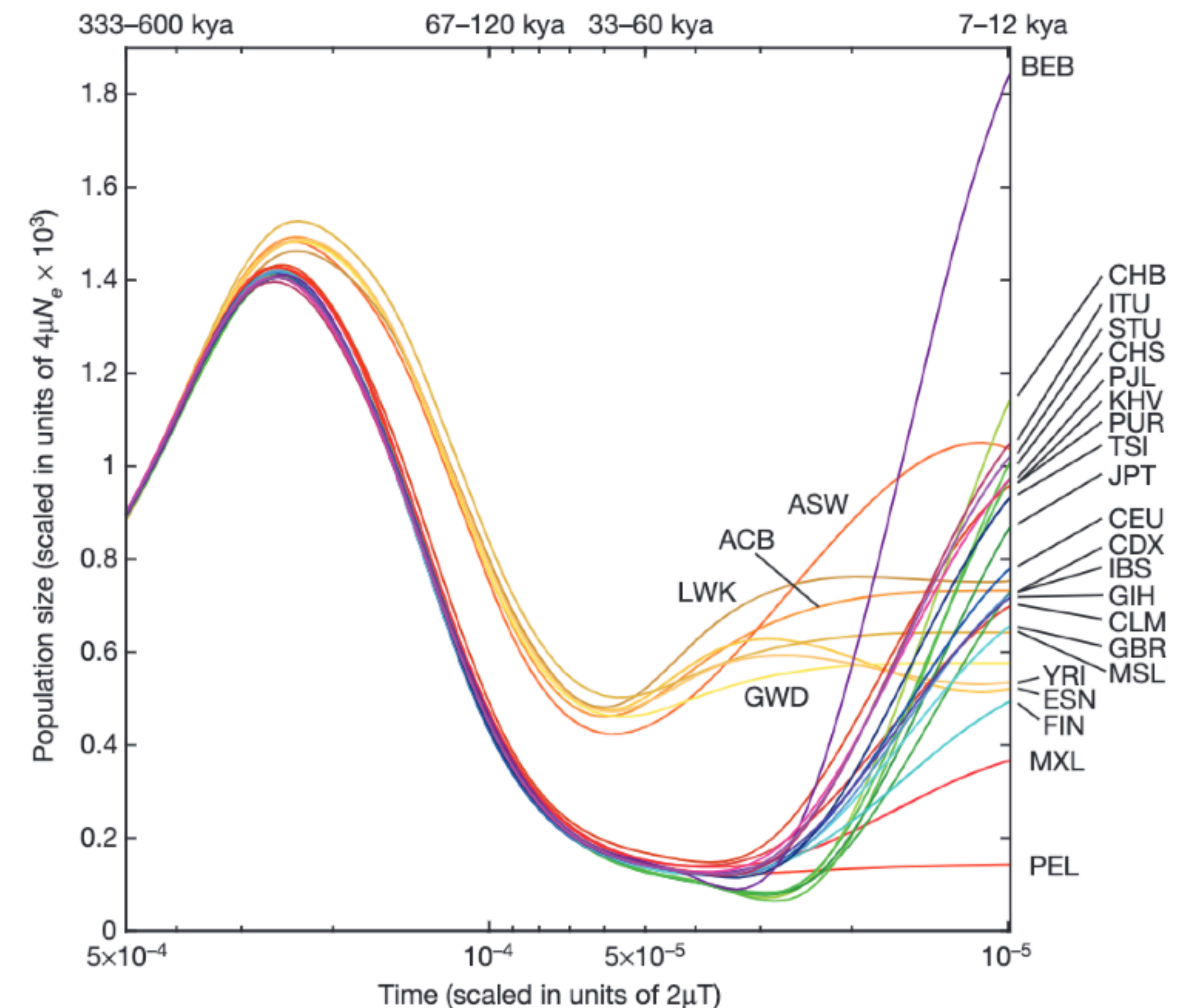
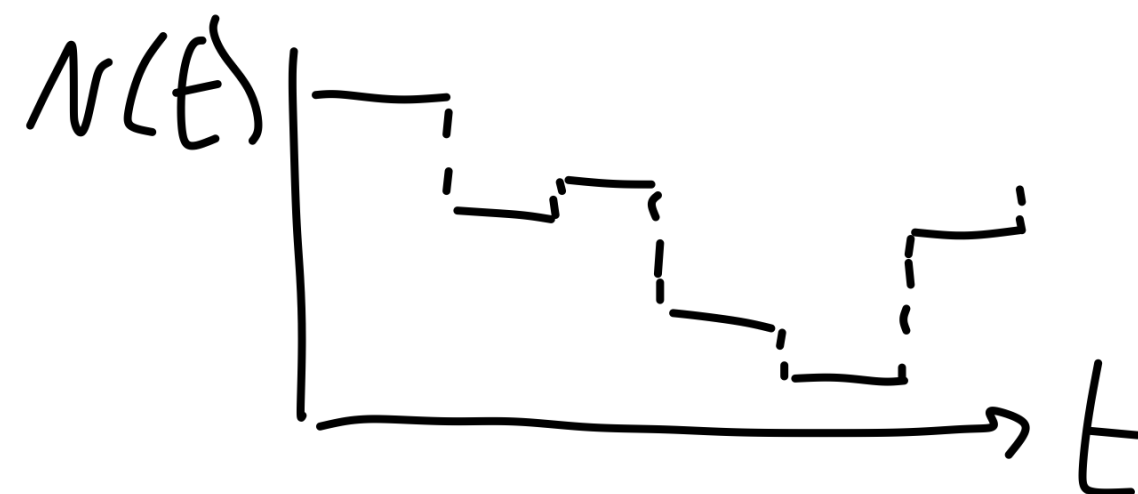


A global reference for human genetic variation

The 1000 Genomes Project Consortium*

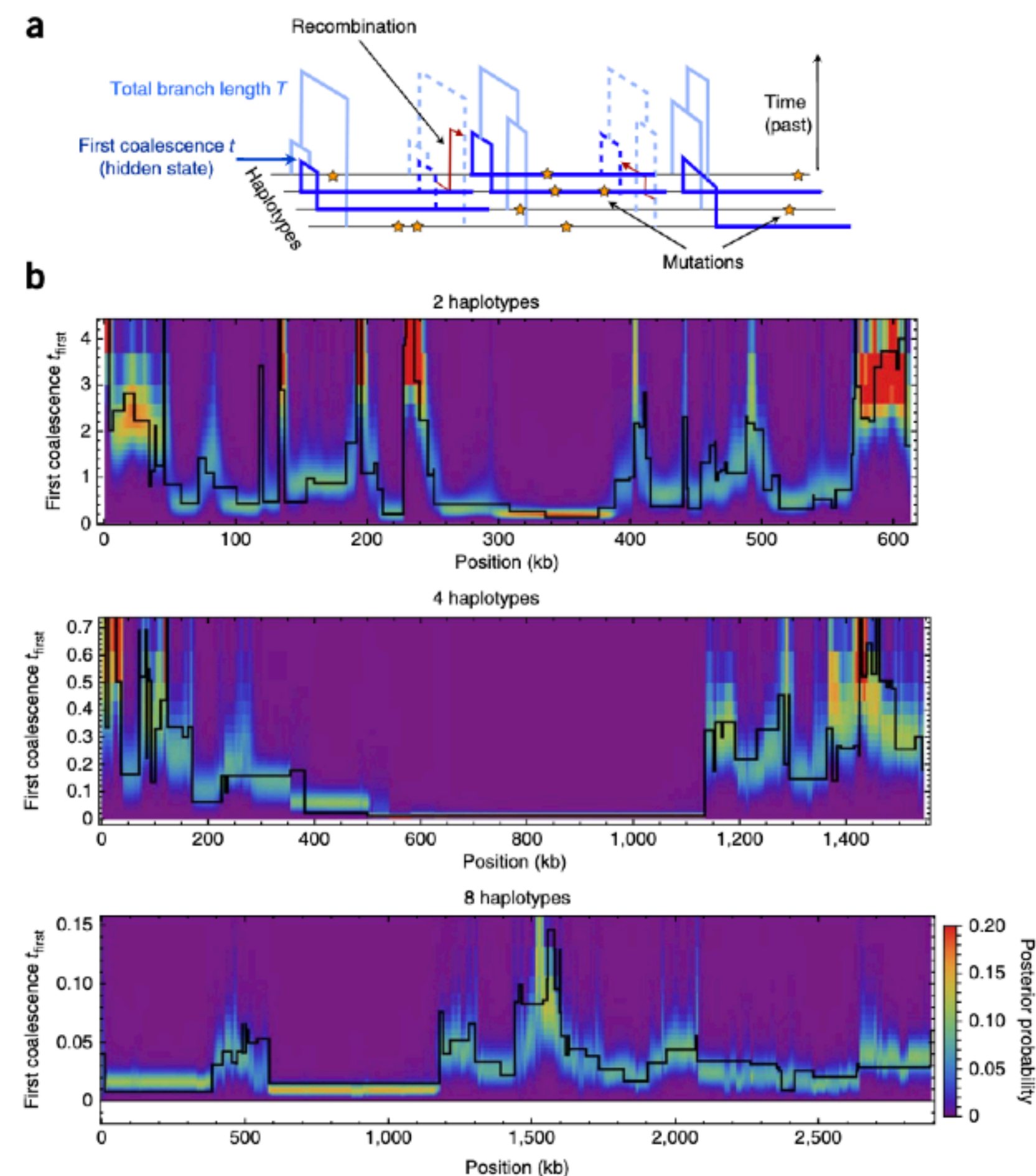
Averaging indiv. estimates
to get pop estimate



Generalizing beyond diploids

Inferring human population size and separation history from multiple genome sequences

Stephan Schiffels & Richard Durbin



Robust and scalable inference of population history from hundreds of unphased whole genomes

Jonathan Terhorst¹, John A Kamm^{1,2} & Yun S Song¹⁻⁴

