1.1 Some Very Useful Numbers for human population genetics.

When you're thinking about genomes and genomic data it's often useful to have a sense of rates and scales¹.

Genome properties.

Genome size: 3.1 Gb (haploid size). Number of chromosomes: 23 pairs Number of coding genes: ~20,000 Exons per gene: 8 (median) Number of genes per megabase: 6.5 (mean) Total in protein-coding exons: 1% of genome Total in genes (introns+exons): 40% of genome Active chromatin (per cell type): 1% of genome Active chromatin (all cell types): 13% of genome

Length scales. (Orders of magnitude.) Transcription factor binding site: 10 bp Enhancer: 100 bp – 1 Kb Exon (coding): 150 bp Coding length per gene: 1200 bp (median) Intron: 1 – 50 Kb Gene (pre-mRNA): 10 – 100 Kb Extent of LD: 10 Kb – 1 Mb (varies by locus & population) Enhancer–promoter interactions: 1 Kb – 1 Mb Chromatin topological domains (TADs): ~1 Mb

Genetic variation.

Heterozygosity: $0.5-1.0 \times 10^{-3}$ (varies by population) Human-Chimpanzee divergence: 1.4×10^{-2} Number of common SNPs: ~8 million (at > 5% MAF in global sample) Number of SNPs for genome-wide SNP tagging: ~1 million Fst between populations: ~0.10-0.15 between continents

Population genetic parameters.

Mutation rate per generation: 1.3×10^{-8} per basepair (at parental age 30) Mutation rate per year: 4.0×10^{-10} per basepair Number of mutations per child: \sim 70 Recombination rate: 1.2 centiMorgans per Mb (mean, sex-averaged) Cross-overs per egg: \sim 42 Cross-overs per sperm: \sim 26 Effective population size: 10,000–20,000 $(H/4\mu)$

Timescales of population divergence. (Take with grain of salt) Human to chimpanzee: 6.5 MY Human to Neanderthal/Denisovan: 600 KY Deepest human population splits: ~200 KY Out-of-Africa migration: 65–100 KY Deepest non-African splits: 65 KY

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Notes and References.

¹Here's an excellent book-length treatment of this topic, with a focus on cell biology, free online [Link]: Milo R, Phillips R. Cell biology by the numbers. Garland Science; 2015