Molecular Haplotyping

global

haplotypes

cost/labor

Chromosome sorting (FACS, microfluidics, ...)
StrandSeq

3rd-gen. sequencing (PacBio, ONT, ...)

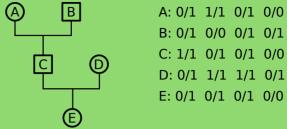
synthetic long reads

(TruSeq, 10xGenomics, ...)

2nd-gen. sequencing (Illumina, ...)

Read-base

Genetic Haplotyping

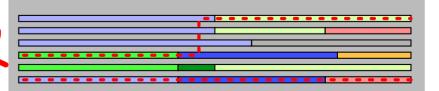


Use genotypes across a pedigree

Supported by most packages for statistical phasing (SHAPEIT, beagle, etc.)

SHAPEIT extension
(Delaneau et al., 2013)

Statistical Phasing



Use genotypes/haplotypes from a reference population