

Online Resource

Leveraging linkage evidence to identify low-frequency and rare variants on 16p13 associated with blood pressure using TOPMed whole genome sequencing data

Human Genetics

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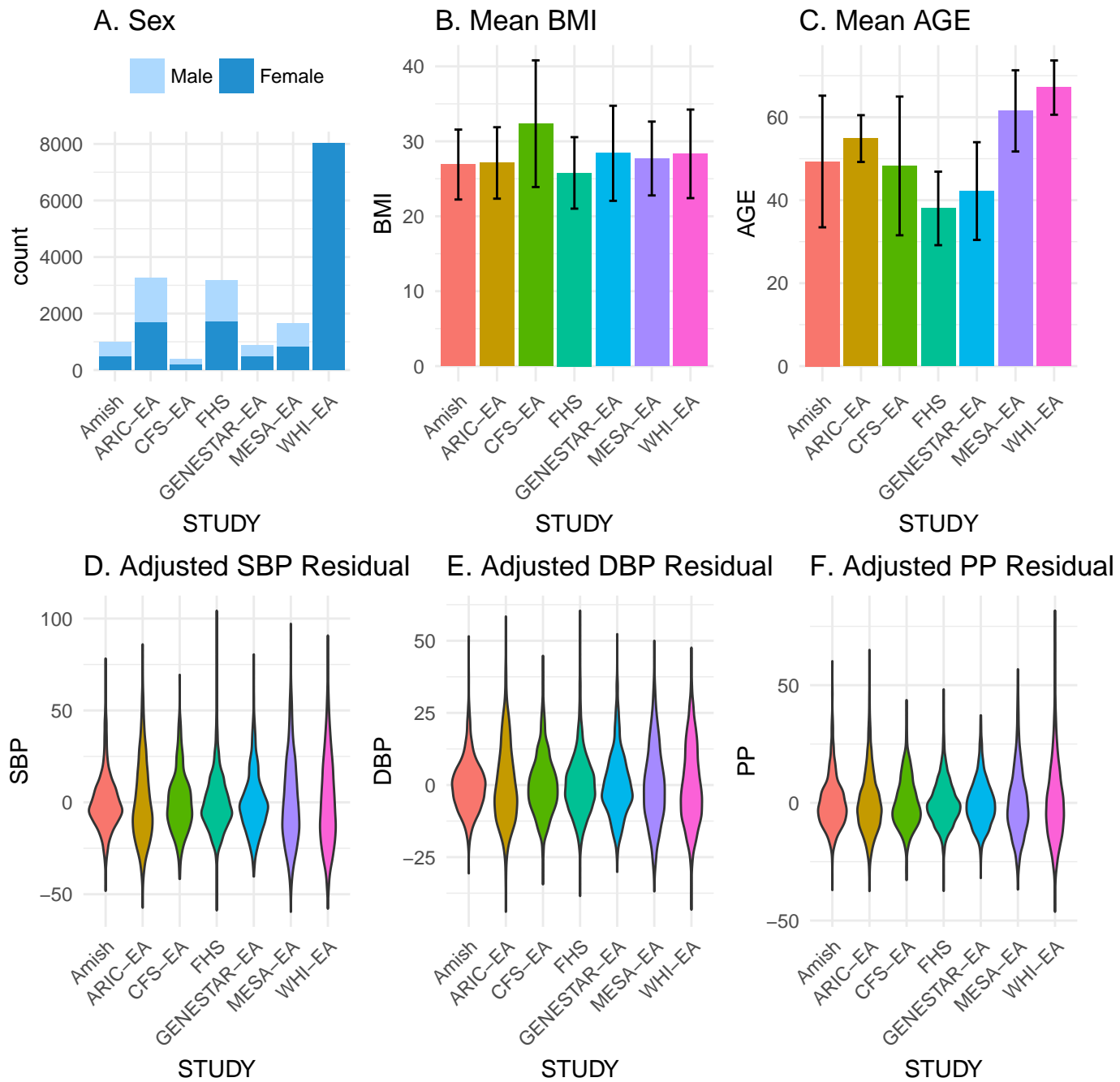
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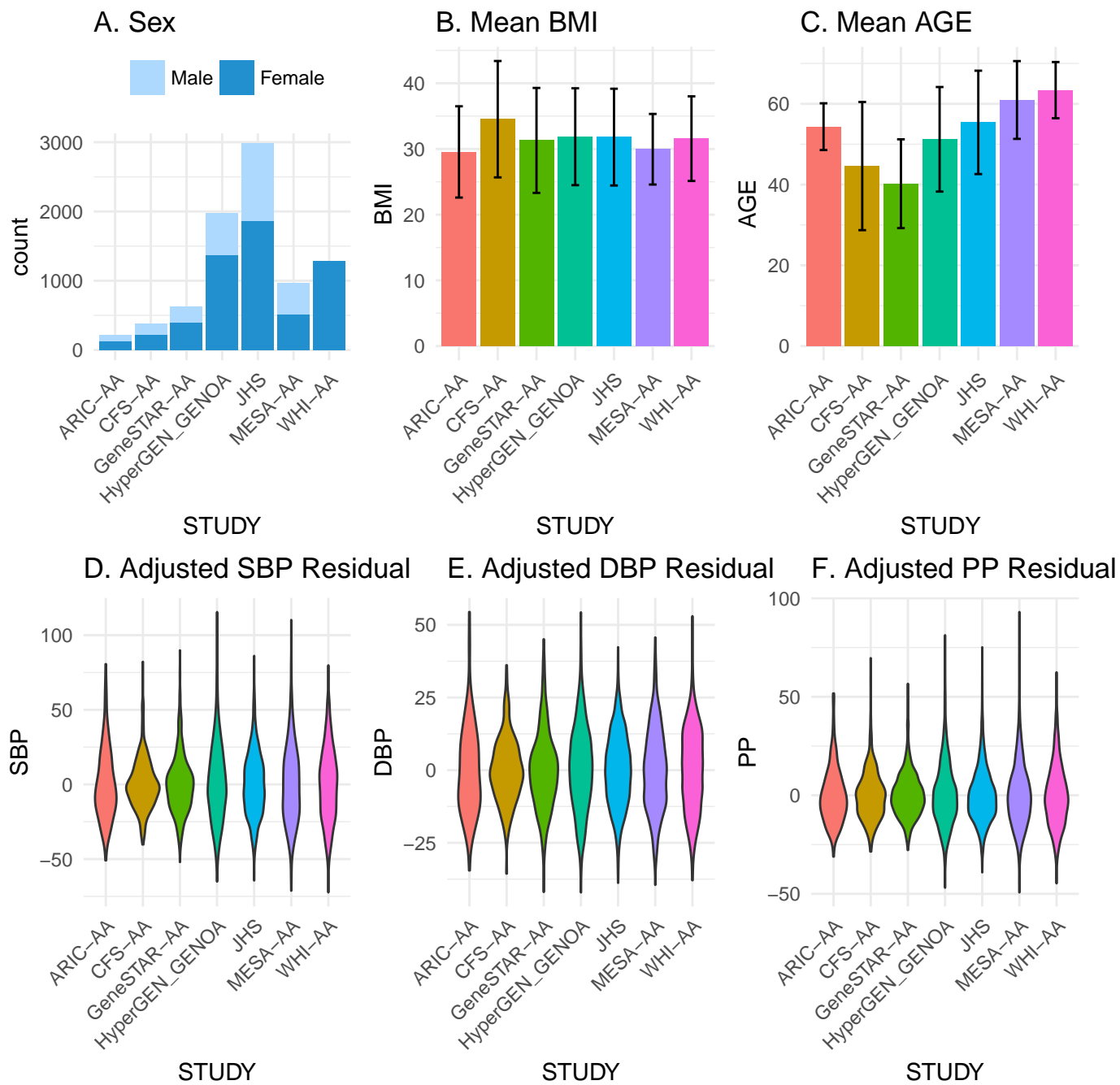
Online Resource Figures 1-4

Online Resource Note

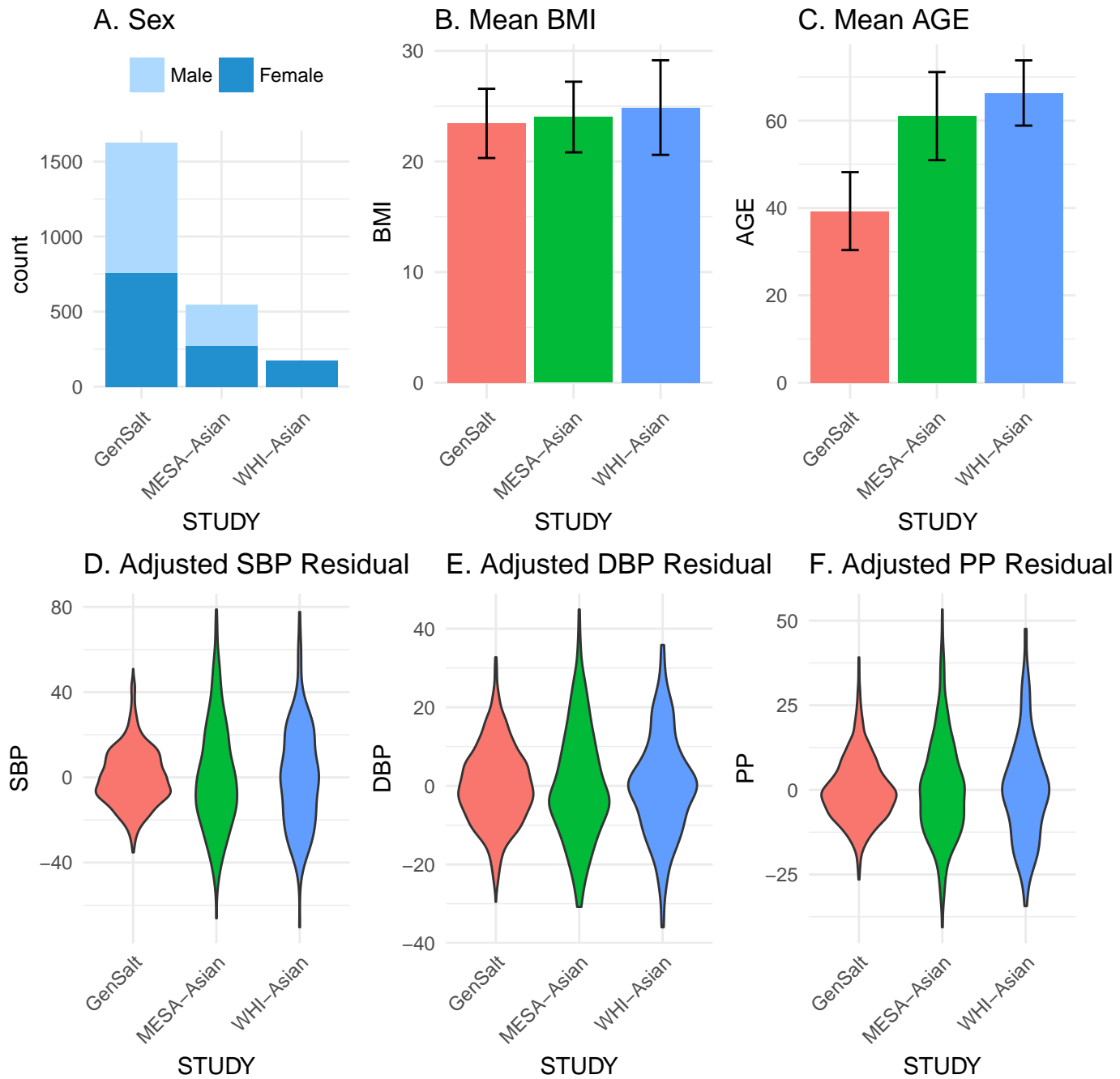
Online Resource Figure 1. Phenotypic characteristics of European-American cohorts



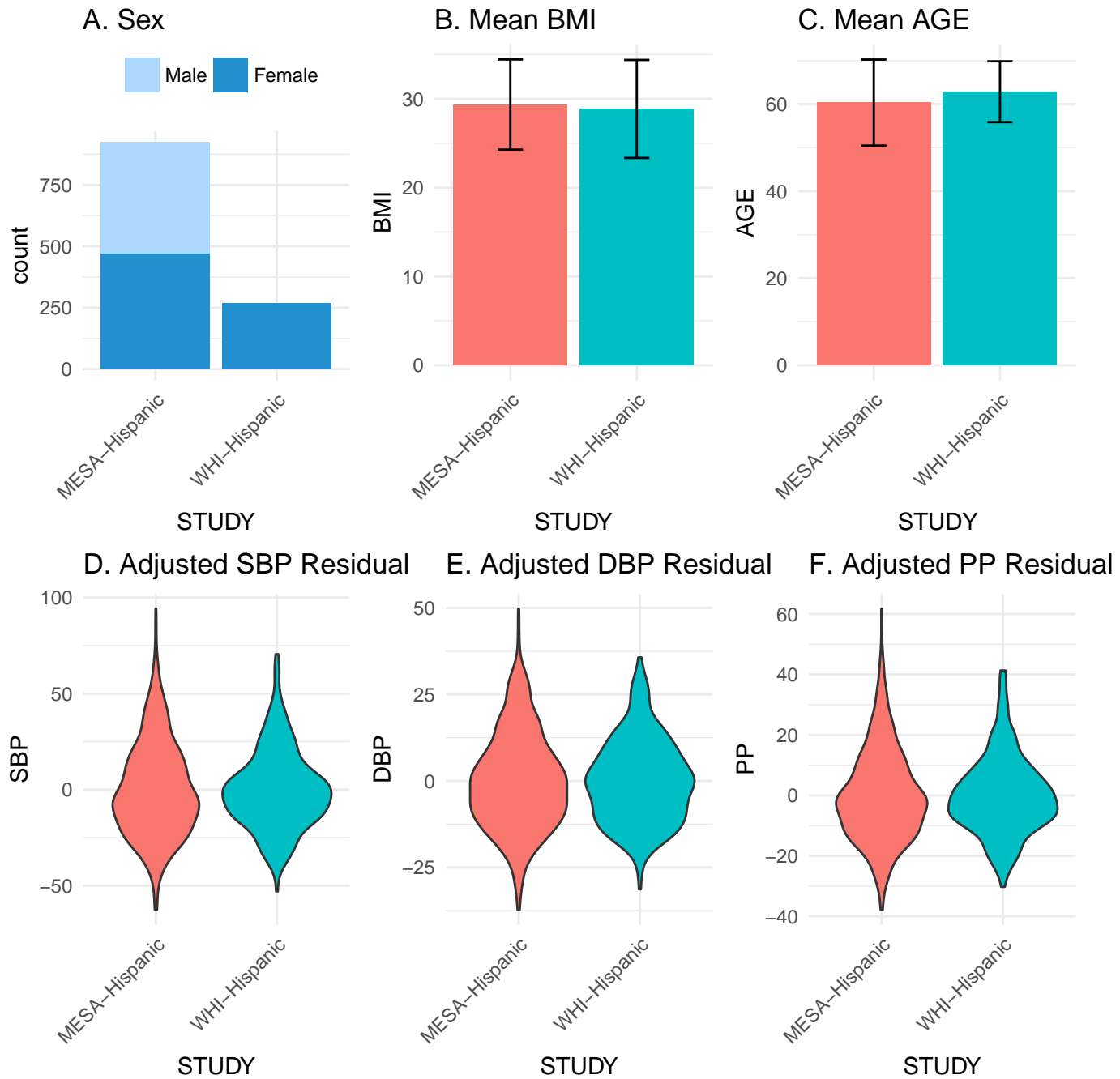
Online Resource Figure 2. Phenotypic characteristics of African-American cohorts



Online Resource Figure 3. Phenotypic characteristics of East Asian and Asian-American cohorts



Online Resource Figure 4. Phenotypic characteristics of Hispanic-American cohorts



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