

	DS1	DS2	DS2p	DS3	DS3p	DS4	DS5	DS6	DS7
Analyses	Worldwide Context	Phasing	Local Ancestry	Phasing	Local Ancestry	European Ancestry	African Ancestry	Native American Ancestry	<i>F-stats</i>
Present Study	✓	✓	✓			✓(a)	✓(a)	✓(a)	✓(a)
(Homburger et al. 2015)	✓			✓	✓	✓(a)	✓(a)	✓(a)	✓(a)
(de la Fuente et al. 2018)	✓	✓	✓					✓(a)	✓(a)
(Reich et al. 2012)	✓							✓	✓(b)
(The 1000 Genomes Project Consortium 2015)	✓(c)	✓(d)	✓(d)	✓(d)	✓(d)	✓(e)	✓(f)		✓(g)
(Nelson et al. 2008)(h)						✓			
(Pagani et al. 2012)							✓(i)		
(Schlebusch et al. 2012)							✓(i)		
(Patin et al. 2017) and references therein							✓(i)		
(Raghavan et al. 2015)								✓	✓
(Moreno-mayar et al. 2017)								✓	✓
(Moreno-mayar et al. 2018)								✓	✓
(Lindo et al. 2018)								✓	✓
(Posth et al. 2018)								✓	✓
# SNPs without LD-pruning	64,531	608,501	608,501	694,626	694,626	29,347	137,136	47,003	88,564
# SNPs with LD-pruning	59,237	–	–	–	–	27,634	128,086	39,423	–
# Individuals	2,076	1,645	1,645	1,772	1,772	1,612	1,742	505	632

- a. Masked data for the specific ancestry analyzed
- b. Including individuals from Mixe population in Mexico
- c. Individuals from South America, Africa, Europe, and Afro-Americans
- d. Individuals from America, Africa and Europe
- e. Individuals from Africa
- f. Individuals from South America
- g. Only individuals from Yoruba population
- h. Only individuals identified in Novembre et al. 2008
- i. Removing individuals with less than 99% of African ancestry estimated through Admixture analyses ($K=7$)