"1000" Genomes Project

find variants
that occur in >
1% of the
population:
sequence
≈2500 genomes
at 4X coverage,
align them to
reference.

1000 Genomes Samples						6-Sept-11		
Population	Status	Available to research community (dates approx)	DNA sequenced from blood	Offspring samples from trios	First set	Second set	Third set	Total
Utah residents (CEPH) with Northern and Western European ancestry (CEU)	Available	Available	no	yes	100			100
Toscani in Italia (TSI)	Available	Available	no	no	100			100
British from England and Scotland (GBR)	Available	Available	no	no	96	4		100
Finnish from Finland (FIN)	Available	Available	no	no	100			100
Iberian populations in Spain (IBS)	Available to project	Available	no	yes	30	70		100
Total European ancestry					426	74		500
Han Chinese in Beijing, China (CHB)	Available	Available	no	no	100			100
Japanese in Toyko, Japan (JPT)	Available	Available	no	no	100			100
Han Chinese South (CHS)	Available	Available	most	yes	100			100
Chinese Dai in Xishuangbanna (CDX)	Available to project	Oct-Dec 2011	some	no		100		100
Kinh in Ho Chi Minh City, Vietnam (KHV)	Available to project	Oct-Dec 2011	yes	some		100		100
Chinese in Denver, Colorado (CHD) (pilot 3 only)	Available	Available	no	no				0
TOTAL East Asian ancestry					300	200		500
Yoruba in Ibadan, Nigeria (YRI)	Available	Available	no	yes	100			100
Luhya in Webuye, Kenya (LWK)	Available	Available	no	no	100			100
Gambian in Western Division, The Gambia	Collecting samples	Mar-May 2012	no	yes			100	100

http://www.1000genomes.org/about#ProjectSamples