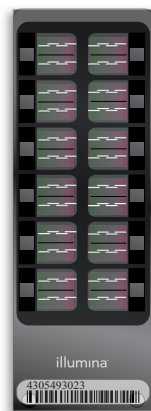


Infinium® HumanCore BeadChips

Customizable array for cost-effective large-scale genotyping and screening studies.

Figure 1: HumanCore BeadChip



The HumanCore BeadChip enables genotyping of markers informative across diverse world populations, delivering high-quality data that can be used for a variety of downstream applications.

Overview

Customizable HumanCore BeadChips offer the most economical way to perform and support large genetic studies, especially large-scale genotyping studies. Developed in collaboration with several leading research institutions, HumanCore BeadChips contain more than 240,000 highly-informative genome-wide tag SNPs and over 20,000 high-value markers, including indels and updated exome-focused content. Furthermore, the HumanCore+ BeadChip has the added capacity to include up to 200,000 semi-custom markers. In addition to performing cost-effective large-scale genotyping studies, HumanCore BeadChips can be used to quickly and easily obtain baseline sample datasets for a variety of downstream applications, including common variant, mtDNA, ancestry, sex confirmation, loss-of-variant, indel, and CNV detection studies.

HumanCore and HumanCore+ BeadChips are based upon the trusted Infinium assay. Using the proven iScan or HiScan™SQ System, these twelve-sample BeadChips combine affordability with high-throughput sample processing to deliver high-quality, genome-wide information.

HumanCore BeadChip Product Information

Feature	Description
Number of tagSNP markers	250,421
Number of additional high-value markers (indels, exomes)	> 20,000
Headroom for additional custom markers	200,000
Number of samples per HumanCore BeadChip	12 samples
Number of samples per HumanCore+ BeadChip	12 samples
DNA Requirement	200 ng
Assay	Infinium HD
Instrument Support	HiScanSQ, HiScan, or iScan
Sample Throughput**	> 1,400 / week
Scan Time / Sample	5 minutes

% Variation Captured† ($r^2 > 0.8$)	1kGP† MAF > 5%	1kGP† MAF > 1%
CEU	0.57	0.44
CHB + JPT	0.61	0.49
YRI	0.26	0.15

Data Performance	Value‡ / Product Specification
Call Frequency	99.9% / > 99.9% avg.
Reproducibility	99.9% / > 99.9%
Log R Deviation	0.17 / < 0.30§

Spacing	Mean
Spacing (kb)	1 SNP / 10 kb

** Estimate assumes one iScan system, one AutoLoader2, one Tecan Robot, and a five-day work week.

† Compared against the June 2011 1kGP data release.

‡ These are representative values for Infinium performance.

§ Value expected for typical projects, excluding tumor samples or any samples prepared not following standard Illumina protocols.

