



The Genome-Wide Human SNP Nsp/Sty Assay Kit contains validated and qualified reagents for the most critical steps in the assay. This includes the PCR primer and adaptors, reagents to fragment and label the PCR products, and control reagents. Manual processing kits are available for either 50 or 100 reactions. An automated assay kit (for processing 96 reactions) is also available (see ordering information). Whole-genome-amplified material prepared by the Qiagen REPLI-g® kits may also be used as the starting material for the Genome-Wide Human SNP Assay Kit.

### Performance data

To validate the performance of the SNP Array 6.0, Affymetrix tested 270 samples from the International HapMap Project. In addition, two external sites tested a plate of 44 HapMap DNAs, which includes 30 unique samples, 10 trios, and five samples with multiple replicates.

The arrays that passed the QC call rate threshold were analyzed using the Birdseed algorithm at the default setting of 0.1. The average call rate for each set was greater than 99 percent, and the concordance with HapMap genotypes was observed to be greater than or equal to 99.7 percent. For the 10 trios, the Mendelian inheritance consistency was found to be greater than 99.9 percent. Reproducibility was measured at 99.9 percent (see results in Table 1).

### Genotype calls using Genotyping Console™ Software

When the Genome-Wide Human SNP Array 6.0 is used in conjunction with Genotyping Console™ Software, the following applications are enabled:

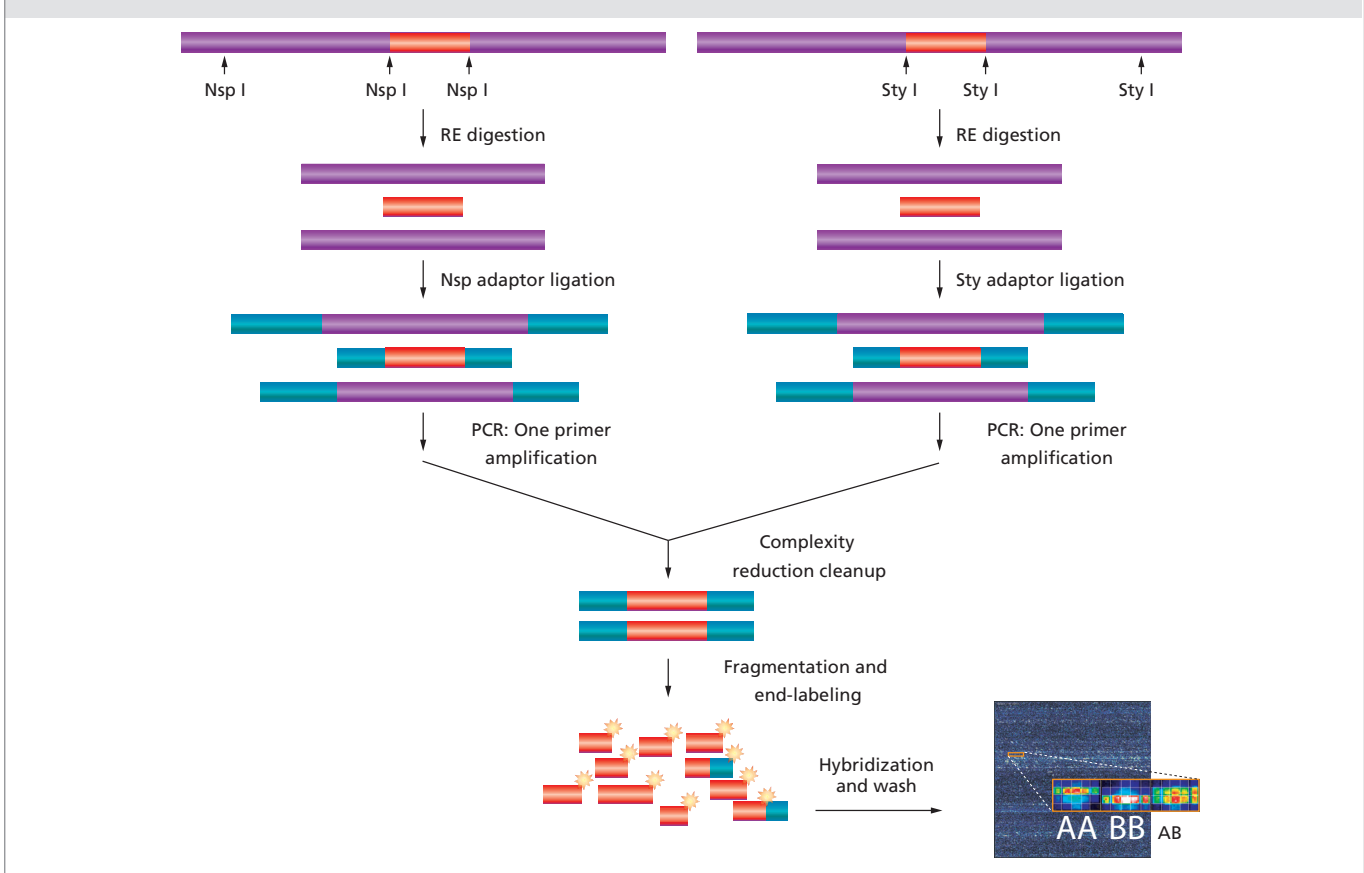
- SNP genotyping for association studies
- CNP genotyping (McCarroll, *et al.*) for association studies
- De novo CNV and LOH detection for association studies and cytogenetics research

**Table 1:** Data analyzed with Birdseed (0.1).

	270 HapMap	Site 1	Site 2
<b>Call rate</b>	99.8%	99.7%	99.7%
<b>HapMap concordance</b>	99.8%	99.7%	99.8%
<b>Mendelian consistency</b>	99.97%	99.95%	99.96%
<b>Reproducibility</b>	NA	99.9%	99.9%
<b>SNP completeness*</b>	99.9%	99.7%	99.8%

\*SNP completeness is defined as the proportion of SNPs with per-SNP call rate greater than 85 percent.

**Figure 1:** Overview of the Genome-Wide Human SNP Assay 5.0/6.0.



Genotyping Console™ Software utilizes strong quality control metrics to streamline genotyping. QC metrics include the Contrast Quality Control (CQC) metric to predict genotyping performance and the Median Absolute Pairwise Difference (MAPD) value for copy number measurements. Flexible visualization and filtering tools allow you to generate customized PDF reports for downstream statistical analysis and bridge CNP and SNP genotyping for a truly genome-wide association study.

Refer to the *Genome-Wide Human SNP Nsp/Sty Assay 5.0 or 6.0 User Guide* (P/N 702419-2, P/N 702504-3) for details on the QC call rate thresholds, as well as procedures on DNA target preparation, target hybridization, fluidics setup, array scanning, and data analysis.

**Fluidics protocol required**  
GenomeWideSNP6\_450

**Library files required**  
GenomeWideSNP\_6

Library files contain information about probe array design layout, probe use and content, scanning and analysis parameters, and other characteristics.

These files are unique for each probe array type. Library files are available from the Affymetrix website at [www.affymetrix.com/support/technical/libraryfilesmain.affx](http://www.affymetrix.com/support/technical/libraryfilesmain.affx). Affymetrix products can be purchased directly from Affymetrix in the United States and many European countries. For all other territories, refer to our list of distribution partners located at [www.affymetrix.com/site/contact/index.affx](http://www.affymetrix.com/site/contact/index.affx).

#### References

Rabbee, N., *et al.* A genotype calling algorithm for Affymetrix SNP arrays. *Bioinformatics* 22:7-12 (2006).

Affymetrix, Inc. BRLMM: An Improved Genotype Calling Method for the Mapping 500K Array Set.

Matsuzaki, H., *et al.* Genotyping over 100,000 SNPs on a Pair of Oligonucleotide Arrays. *Nature Methods* 1:109-111 (2004).

Frayling, T. M., *et al.* A Common Variant in the FTO Gene is Associated with Body Mass Index and Predisposes to Childhood and Adult Obesity. *Science* (ePub 2007).

McCarroll, S. A., *et al.* Integrated detection and population—genetic analysis of SNPs and copy number variation. *Nature Genetics* 40(10):1166-1174 (2008).

<b>Product information</b>	
Number of SNPs on the array	906,600
Number of non-polymorphic probes for copy number detection	946,000
Number of arrays	1
DNA required	500 ng
Average Birdseed call rate (0.1)	>99 percent*
Average minor allele frequency (MAF)	19.6% in HapMap Caucasians 18.2% in HapMap Asians 20.6% in HapMap Africans
Average heterozygosity	26.7% in HapMap Caucasians 24.6% in HapMap Asians 28.5% in HapMap Africans
PCR primers	1 per sample
Instrumentation	GeneChip® Scanner 3000 7G with AutoLoader
Throughput	>40 million genotypes per day, per scanner with three GeneChip® Fluidics Station 450s

\* Average scan from four independent studies

<b>Genome-Wide Human SNP Nsp/Sty Assay Kit 5.0/6.0 components</b>	
Adaptor, Nsp I or Adaptor, Sty I	Two annealed oligonucleotides specific for ligation to the Nsp I or Sty I restriction site
PCR Primer 002	PCR primer to amplify ligated genomic DNA
Reference Genomic DNA, 103	Human genomic DNA control, with consensus genotypes
GeneChip® Fragmentation Reagent	DNase I enzyme, formulated to fragment purified PCR amplicons
10X Fragmentation Buffer	Buffer for fragmentation reaction
GeneChip® DNA Labeling Reagent (30mM)	Proprietary biotin-labeled reagent for end-labeling fragmented PCR amplicons
Terminal Deoxynucleotidyl Transferase	Enzyme used to end-label fragmented PCR amplicons with the GeneChip® DNA Labeling Reagent
5X Terminal Deoxynucleotidyl Transferase Buffer	Buffer for labeling reaction
Oligo Control Reagent, 0100	Mixture of five biotin-labeled oligonucleotides, which hybridize to control regions (gridding and array controls) on the SNP Array 5.0/6.0



<b>Ordering information</b>	
Part Number	Description
<b>Genome-Wide Human SNP Array 6.0</b>	
901153	Contains 50 arrays
901150	Contains 100 arrays
<b>Genome-Wide Human SNP Nsp/Sty Assay Kit 5.0/6.0</b>	
901152	Sufficient for 50 reactions
901015	Sufficient for 100 reactions
901192	Sufficient for 96 reactions, automated processing

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