

After hybridization, the BeadXpress® Reader is used for microbead code identification and fluorescent signal detection. During scanning, a laser beam penetrates the digitally inscribed VeraCode microbead to generate a unique code image, which allows for rapid and highly specific detection. Data generated using the BeadXpress Reader can be analyzed with Illumina's GenomeStudio™ data analysis software, which performs automated genotype clustering and calling.

Typical Results

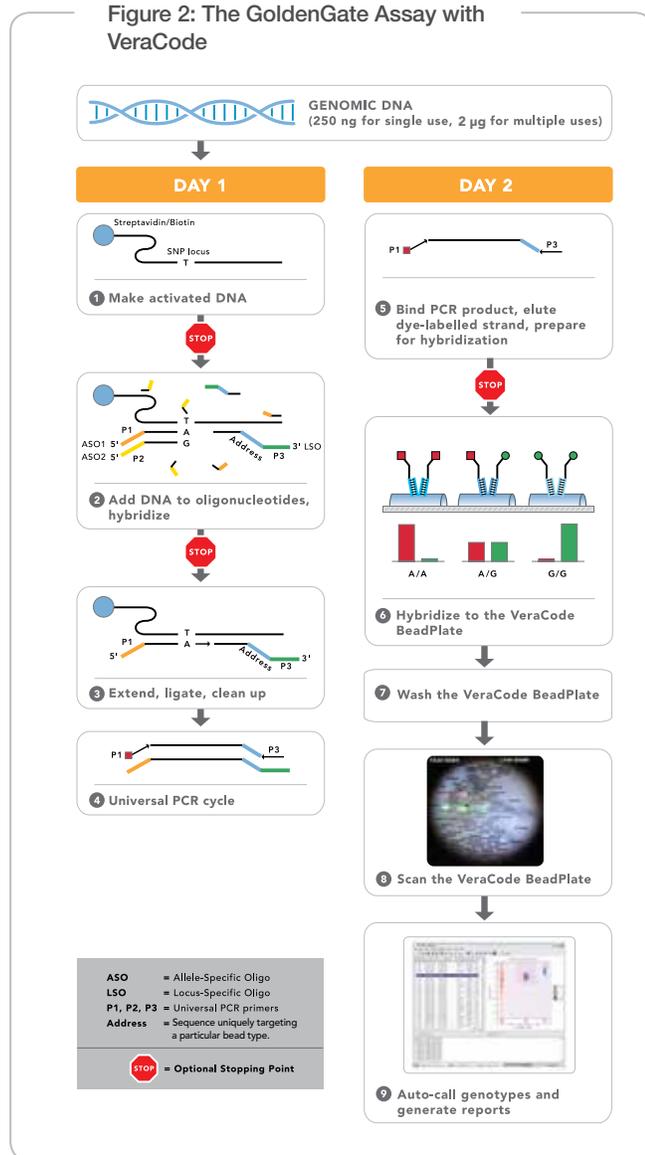
The GoldenGate Assay and the highly specific VeraCode technology exhibit superior consistency, reproducibility, and success rate. Results of laboratory testing of multiple plates at 96-plex and 384-plex scale are shown in Table 1.

Table 1: Typical GoldenGate Assay Performance

96-plex	
Call Rate	> 99.9%
Heritability	> 99.9%
DNA Success Rate	> 99%
Locus Success Rate	> 99%
384-plex	
Call Rate	> 99.9%
Reproducibility	> 99.9%
Heritability	> 99.9%
DNA Success Rate	> 99%
Locus Success Rate	> 98%

GoldenGate Assays at 48-, 144-, 192-, and 384-plex meet the same performance specifications.

Figure 2: The GoldenGate Assay with VeraCode



GoldenGate Assay Controls

The GoldenGate Genotyping Assay includes 48 assay controls, lending a high level of confidence and the ability to troubleshoot errors such as PCR and hybridization failures (Figure 3). VeraCode microbead digital coding serves a dual role of enabling built-in assay controls as well as internal tracking controls. Illumina's GenomeStudio data analysis software provides a dashboard for simple viewing of controls performance.

References

1. Fan JB, Gunderson KL, Bibikova M, Yeakley JM, Chen J, et al. (2006) Illumina universal bead arrays. *Methods Enzymol*, 410: 57–73.
2. The International HapMap Consortium. (2003) The International HapMap Project. *Nature* 426: 789-796.
3. Lin CH, Yeakley JM, McDaniel TK, Shen R. (2009) Medium- to High-Throughput SNP Genotyping Using VeraCode Microbeads. *Methods Mol Biol*. 496:129–42.
4. Kulathinal RJ, Bennett SM, Fitzpatrick CL, Noor MAF. (2008) Fine-scale mapping of recombination rate in *Drosophila* refines its correlation to diversity and divergence. *PNAS* 105(29): 10051–10056.
5. Cooper GM, Johnson JA, Langae TY, Feng H, Stanaway IB, et al. (2008) A genome-wide scan for common genetic variants with a large influence on warfarin maintenance dose. *Blood* 112(4): 1022-7.
6. Is Genetic Testing the Next Revolution in Agriculture? http://www.illumina.com/emailers/icomunitySep08/iC_Sep08_GeneticTesting.pdf.
7. Using Genetics to Uncover Cat Secrets. http://www.illumina.com/emailers/icomunitySep08/iC_Sep08_CatSecrets.pdf.

