Infinium[™] XT Genotyping Solution

The accuracy and robustness of proven Infinium chemistry taken to the next level of scalability

- Enable production-scale genotyping with an optimized assay and 96-sample BeadChip
- Customize targeted single- or multispecies assays of 100s to 50K single nucleotide polymorphisms (SNPs)
- Access a streamlined workflow with user-friendly enhancements and less hands-on and total assay time

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Introduction

The Infinium family of genotyping assays harnesses proven chemistry and a robust BeadChip platform to produce high data guality, call rates, and reproducibility. Infinium XT technology builds on this success by featuring a 96-sample BeadChip and optimizing the Infinium workflow to provide the highest throughput genotyping solution to date from Illumina. The product was developed for customers who need to perform large-scale screening applications (up to 50K SNPs) on any species with user-defined content. The Infinium XT solution simplifies the custom assay design process, reduces overall handson time, and enhances automation robot performance and utilization. The workflow offers an option to reduce the overall turnaround time, if the time to answer is more important than maximum weekly output. Also, the Infinium XT solution introduces an enhanced data analysis software solution for real-time data generation and on-demand quality control (QC) report functionality so production issues can be identified and corrected earlier.

Production-scale genotyping

The Infinium XT solution features a 96-sample BeadChip (Figure 1) for efficiently genotyping large numbers of samples, scaling from hundreds of thousands to more than one million samples per year. It supports as few as several hundred SNPs, providing a production-scale solution for targeted genotyping applications, including animal parentage, sample traceability, or sample QC. This product also supports up to 50,000 SNPs, which are required for key agricultural applications like genomic selection for both diploid and polyploid organisms.

Human applications, such as biobanking, population screening, and personalized medicine initiatives also benefit from this technology. The low- to mid-range number of SNPs and low cost are ideal for labs that require simple QC checks on large sample sets or need to perform a more thorough population stratification before proceeding with deeper genomic testing. The versatility of the Infinium XT BeadChip makes it a cost-effective, reliable solution for sample QC, tracking, and validation throughout an entire workflow.



Figure 1: Infinium XT 96-sample BeadChip—The Infinium XT BeadChip offers the proven power and flexibility of the Infinium assay in the highest throughput available from Illumina.

Flexible, application-specific content

The Infinium XT BeadChip fully supports custom iSelect[™] arrays, consortia, or commercial product opportunities within human or agrigenomics applications that meet the content and throughput requirements (≤ 50,000 markers, \geq 100,000 samples per year). To aid in the design of custom content for Infinium XT BeadChips, the online DesignStudio[™] Software includes a custom genotyping microarray probe designer, which replaces the older Assay Design Tool software. With DesignStudio Software, custom content can be developed and ordered through a guided, easy-to-use interface with a guarantee of at least a 95% custom content conversion rate. This is a significant increase over the 80% conversion rate guarantee for standard Infinium BeadChips and supports screening many samples for a focused number of critical SNPs. The Infinium XT assay maintains the input requirement of 200 ng of sample DNA with the same high-guality performance users expect from Infinium technology.

High-efficiency Infinium workflow

Each step of the fully automated Infinium HD workflow is optimized in the Infinium XT workflow, reducing the turnaround time (TAT) from three days to two days (Figure 2). On day one, DNA is amplified, enzymatically fragmented, precipitated, and resuspended. Samples are then hybridized to BeadChips during an overnight incubation, where DNA anneals to locus-specific 50-mer probes covalently linked to one of the Infinium bead types. On day two, the Infinium XT workflow continues with enzymatic base extension to confer allelic specificity, followed by fluorescent staining. The iScan[™] System detects the fluorescence intensities of the beads and Illumina software automatically performs analysis and genotype calling.

Infinium HD workflow

New efficiencies introduced into the Infinium XT workflow accommodate greater scalability with the Infinium XT BeadChip (Table 1).

- DNA amplification: increased batch size three-fold and reduced incubation time from overnight to three hours
- Enzymatic fragmentation: decreased time by 50%
- Alcohol precipitation: removed a 30-minute incubation step and reduced drying time by 75% (one hour to 15 minutes)
- DNA resuspension: reduced overall incubation time by 75% (one hour to 15 minutes)
- BeadChip hybridization: increased sample capacity three-fold using the same Tecan robotic system
- Sample extension/staining: increased capacity two-fold; up to 48 BeadChips (4608 samples) per run

Infinium XT supports a flexible workflow where maximum weekly outputs can be balanced with TAT by adjusting workflow configurations that support alternating priorities.



Infinium XT workflow

Figure 2: Infinium XT workflow improvements—Each step of the Infinium HD workflow is optimized in the Infinium XT workflow to reduce the turnaround time from three days to two days.

*Illumina recommends stopping at this point and resuming the workflow on the next day if needed.

Workflow step	Process step	Infinium HD workflow	Infinium XT automated workflow	
	Layering oil at first MA step	Present	Removed	
Amplify DNA	Incubation time	Overnight	Three hrs	
	Batch size	One plate (96 samples)	Three plates (288 samples)	
Fragment DNA	Incubation time	One hr	30 min	
	Batch size	Six plates (30 min, 576 samples)	Six plates (18 min, 576 samples)	
Amplify DNALayering oil at first MA stepPresentAmplify DNAIncubation timeOvernightBatch sizeOne plate (96 samples)Fragment DNAIncubation timeOne hrFragment DNABatch sizeSix plates (30 min, 576 samples)SPrecipitate DNAVortex before adding 2-propanolIncludedIncubation time30 mIncludedPrecipitate DNAIncubation time30 mPrecipitate DNADrying timeOne hrBatch sizeSix plates (75+ min, 576 samples)SiResuspend DNAIncubation timeOne hourResuspend DNABatch sizeSix plates (90+ min, 576 samples)SiHybridize to BeadChipHybridization chamberFour BeadChips per chamberSiHybridize to BeadChipMinimum batch (no reagent waste)Four BeadChipsSiWash and stain BeadChipNew assembly fixtureFour BeadChip capacityNew assembly fixtureFour BeadChip capacity	Vortex before adding 2-propanol	Included	Removed	
	Incubation time	30 m	Removed	
	Drying time	One hr	15 min	
	Six plates (65 min, 576 samples)			
	Incubation time	One hour	Removed Three hrs Three plates (288 samples) 30 min Six plates (18 min, 576 sample Removed Removed 15 min Six plates (65 min, 576 sample 15 min Six plates (65 min, 576 sample Three tip guides Six BeadChips per chamber (n design) 576 samples 24 BeadChips Plate-based Integrated spacers Six BeadChip capacity	
Resuspend DNA	Batch size	Six plates (90+ min, 576 samples)	Six plates (65 min, 576 samples)	
	Tip guides	Single tip guide	Three tip guides	
	Hybridization chamber			
	Batch size	288 samples	Removed Three hrs Three plates (288 samples) 30 min Six plates (18 min, 576 sample Removed Removed 15 min Six plates (65 min, 576 sample 15 min Six plates (65 min, 576 sample 15 min Six plates (65 min, 576 sample Six blates (65 min, 576 sample Six BeadChips per chamber (nondesign) Six BeadChips per chamber (nondesign) Plate-based Integrated spacers Six BeadChip capacity Provided at 20× concentration 48 BeadChips iScan System only XT	
	Minimum batch (no reagent waste)	Four BeadChips	24 BeadChips	
	X-stain reagents	Tubes	Plate-based	
	Glass back plates	Mylar spacers	Integrated spacers	
	New assembly fixture	Four BeadChip capacity Six BeadChip capacity		
	Reagent used to prepare BeadChips for hybridization	Provided at 1× concentration	Provided at 20× concentration	
	Batch size	24 BeadChips	48 BeadChips	
	Scanners supported	HiScan and iScan systems	iScan System only	
Scan BeadChip	SDF	HD	XT	
	ICS	v3.3.28 v3.4		

Abbreviations: MA, Multi-Sample Amplification; SDF, Sentrix Descriptor File; ICS, iScan Control Software.

Optimal laboratory layout

The Infinium XT BeadChip was optimized and designed for factory-scale laboratories to remove the complexities associated with process scaling and large sample number batching. Labs are encouraged to engage with Illumina for consultation about new efficiency gains. For example, a lab processing 1,000,000+ samples annually will need eight (8-tip) Tecan robots, three iScan Systems, two AutoLoader 2.x units, and ancillary lab equipment (Figure 3).

Updated automation and LIMS

The ability to process an increasing number of samples efficiently requires scalable turnkey solutions. These include integrated systems that simplify sample preparation and provide the confidence to continue to scale operations. The Illumina Automation Control (IAC) software for the Tecan liquid-handling robot has increased sample processing efficiency. This includes updates to Pre-PCR reagent transfer steps and doubling throughput capabilities at the X-stain task within the Infinium workflow.

Improved data analysis

With the high-throughput Infinium XT solution, there is a natural increase in data analysis. With this in mind, several improvements have been made to GenomeStudio[™] Software and Beeline Software. GenomeStudio Software is the Illumina visualization and analysis program for microarray-based genotyping data. It provides a tabular view for quickly accessing all the data in an experiment and allows data to be exported for use by various third-party applications. The GenomeStudio Genotyping Module supports analysis of Infinium array genotyping data with normalization, genotype calling, clustering, data intensity analysis, and more. In addition, GenomeStudio Software is necessary for creating and modifying clusters used for calling genotypes from scanned microarray signal intensities. GenomeStudio 2.0 Software speeds up genotype cluster generation, reducing the overall analysis turnaround time. Also, the Polyploid Genotyping Module is appropriate for agricultural and other applications involving polyploid organisms (Figure 4). When used in tandem with Illumina LIMS, GenomeStudio Software provides an integrated experience to view and analyze data from processed samples in real time.



Figure 3: Example lab layout for Infinium XT production-scale genotyping—An example lab layout is provided for processing 1,000,000+ samples per year and includes: three (8-tip) Tecan robots, three iScan Systems, two AutoLoader 2.x units, and ancillary lab equipment. This example layout requires approximately 1200 square feet. Layout is not to scale.



Figure 4: GenomeStudio 2.0 software—GenomeStudio 2.0 software includes a Polyploid Genotyping Module to support applications involving polyploid organisms.

After a cluster file is created, production-level analysis can be transitioned to Beeline Software. Beeline Software is the Illumina microarray data analysis tool for prefiltering and reporting data from high-throughput processing in an automated fashion. This makes it ideal for routine use with Infinium XT data and beneficial for any large-scale Infinium genotyping projects. Beeline Software offers flexible filtering capabilities, reduces experimental array data size, and identifies any samples or markers that do not meet user-defined performance specifications. Filtered data can also be imported directly into GenomeStudio Software for a more interactive analysis, if desired.

Beeline 2.0 Software optimizes its analysis and reporting capabilities for processing high-throughput genotyping studies. New features include QC on-demand, polyploid genotype calling, and faster genotype report generation. Transitioning from GenomeStudio Software to Beeline Software significantly reduces the time needed to observe quality performance specifications, generate genotyping reports, and analyze polyploid data without removing the flexibility for manual intervention.

Summary

The Infinium XT BeadChip and workflow provide a significant increase in sample throughput capabilities for microarray-based genotyping. This comprehensive solution enables large-scale genetic improvement programs in agrigenomics and supports large-scale screening for biobanks and personalized medicine initiatives. The Infinium XT workflow incorporates design software, a four-fold increase in BeadChip capacity, a 33% reduction in sample turnaround time, increased conversion rate of custom content, multispecies design capabilities, and enhanced data analysis for both diploid and polyploid organisms. The high-throughput workflow combined with low cost per sample makes the Infinium XT solution the ideal choice for commercial genotyping labs that want to scale to a factory level of throughput and efficiency without sacrificing performance and reliability.

Learn more

Infinium XT, illumina.com/InfiniumXT

Ordering information

Product	Catalog no.	Product	Catalog no.
Infinium XT Starter Kit (48 BeadChip batches)	20011069	AutoLoader 2.×, Single-Scanner Configuration, 110/220 V	SY-202-1001
Infinium XT Starter Kit (24 BeadChip batches)	20011100	AutoLoader 2.×, Single-Dual Configuration, 110/220 V	SY-202-1002
Infinium XT Upgrade Kit (24 BeadChip batches)	20011101	Infinium Automation Kit 8-tip Tecan LIMS ready, 100/220 V	SC-30- 403/404
Infinium XT Upgrade Kit (12 BeadChip batches)	20011102	Infinium Automation Kit 8-tip Tecan non-LIMS, 110/220 V	SC-30- 401/402
Illumina LIMS Annual Subscription	20073865	Infinium XT iSelect-96 Kit (1152 samples)	20006613
Illumina LIMS One-Time Onboarding	20073866	Infinium XT iSelect-96 Kit (4608 samples)ª	20006614
Infinium XT PB20 Reagent Kit	20007420	Infinium XT iSelect-96 Kit (23,040 samples)ª	20006615
iScan System, 110/220 V	SY-101-1001		

a. For high-throughput, automated processing only; contact your local sales representative for more information.

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