

Nouvelles lames CGH array pour analyse pré- et post-natal

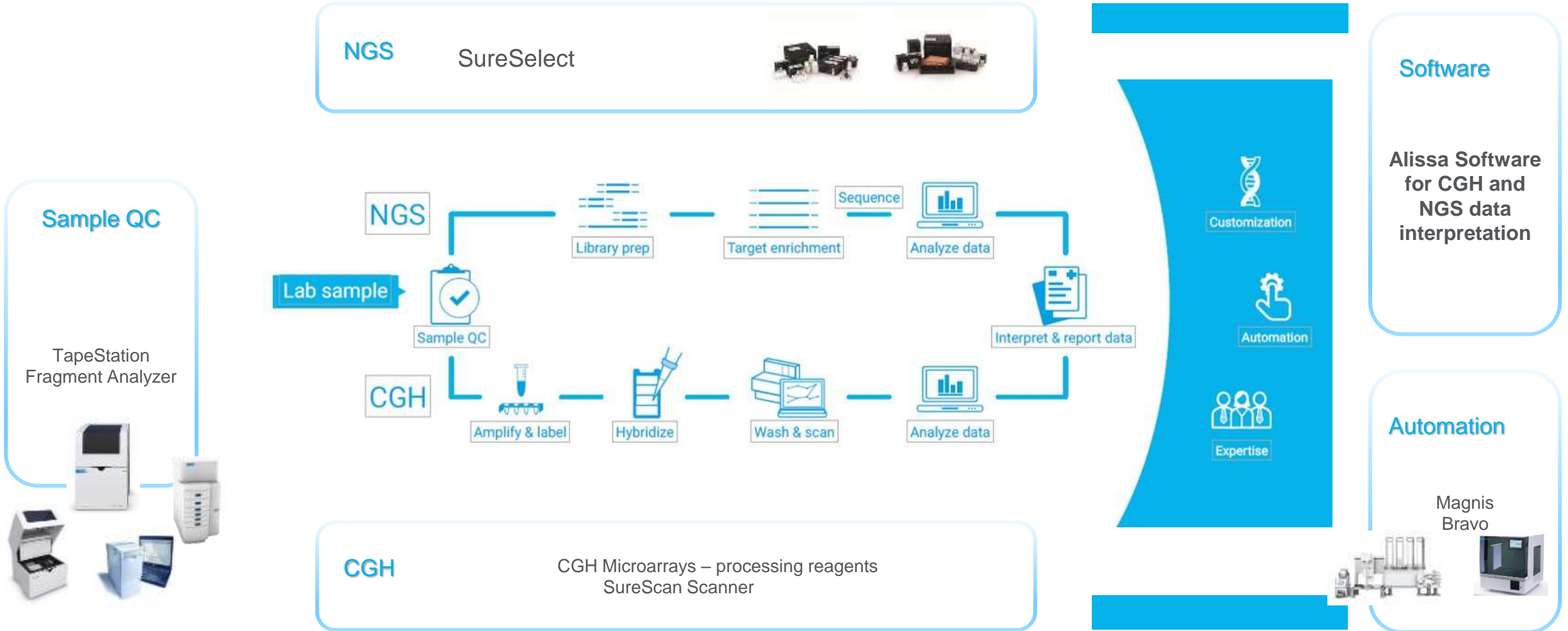
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NGS-CGH

Invest in the future without compromising quality



GenetiSure Cyto Microarrays

The perfect fit for cytogenetic and genetic labs using chromosomal microarrays to analyze constitutional samples



New design focused on clinically relevant and more recent databases:
ClinGen, ClinVar, OMIM, Development Disorder Genotype-Phenotype Database

Curated backbone with coverage on telomeres and PAR regions to improve coverage of clinically relevant regions and improve data interpretation



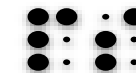
Design include only high quality probes, limiting GC bias, improving data quality and enabling more precise CN call, even for mosaic samples

Including spike-in probes for sample tracking



Free space on each array for easy customization to integrate laboratory specific gene/region of interest

Exon level coverage in genes usually tested by MLPA to improve test value and reduce need of secondary tests



Increased resolution for LOH down to 2.5Mb in autosomal chromosomes matching more updated guidelines from consortia

All

4x180K
CGH

4x180K
CGH+SNP

Focused Microarrays for Pre- and Postnatal Clinical Research

GenetiSure Cyto CGH Microarrays

	Array type	AMADID	Array format	Total genes targeted *	Median probe spacing			Free space	LOH resolution	Exon coverage	
					Target regions	Overall	Backbone				Telomere & PAR
GenetiSure Cyto 8x60K CGH	CGH	085590	8x60K	3644	7.1 Kb	50.6 Kb	67.4 Kb	31.8 Kb 13.5 Kb	~500	Na	No
GenetiSure Cyto 4x180K CGH	CGH	085589	4x180K	3619	3.5 Kb	16.5 Kb	19.8 Kb	7.8 Kb 7.5 Kb	~1500	Na	Yes on 103 selected genes
GenetiSure Cyto 4x180K CGH+SNP	CGH+SNP	085591	4x180K	3644	7.3 Kb	44.5 Kb	57.1 Kb	25.0 Kb 10.5 Kb	~1500	2.5Mb across Autosomes	No
GenetiSure Postnatal 2x400K CGH+SNP	CGH+SNP	078737	2x400K	8106	na	9.5 Kb	na	na	na	2.5Mb across Autosomes	89% of genes 3 probes/exon

* Genes are considered covered if they are targeted by 5 or more probes.
 note: 4x180K CGH is targeting "fewer" than 8x60/4x180 SNP due to the large gene cutoff which is more aggressive due to the denser backbone.

What's coming for SureSelect Human All Exon V8?

Efficient design and best-in-class performance



Same Comprehensive Content

Targets protein coding regions in major gene definition databases: CCDS, RefSeq, GENCODE



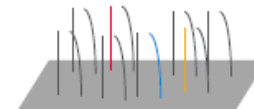
Powered by Machine Learning

Machine learning-based probe selection & placement

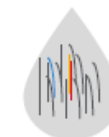


Made Using Improved Manufacturing Process

Probe printing targeting both forward/reverse strands



New probe recovery process



Design Footprint Reduction

Significantly more efficient design that is more economical to sequence

Current progress:
Design footprint <1.2



Uniformity Improvement

Deeper and more uniform coverage across all target regions

Current progress:
Best-in-class 30x coverage and fold-80



Single-Exon CNV Calling

High resolution analysis of copy number variation (CNV) to increase diagnostic yield





Agilent

Trusted Answers