





Optimizing extraction technologies for efficiency in high throughput genotyping data generation

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Dr. Frank Schubert Business Unit Manager Nucleic Acid Preparation (NAP)

Amy Pierce US Lab Director





# LGC Genomics – part of the LGC Group

- Established in 1842 as a scientific testing laboratory for the UK Government
  - Privatised in 1996
- Retains role as UK National Measurement Institute
  - Standard bearer for chemical
     & bio-analytical measurement
- Global presence
  - Over 30 locations
  - 1800+ staff
  - 3-fold growth since 2004

#### **Five Operating Divisions**



Home of the UK National Measurement Institute





## LGC Genomics – solutions for genomic research

Extraction

Proprietary and standard extraction kits and services, developed and optimized over many years in our development laboratories

- sbeadex<sup>®</sup>
- mag
- · Kleargene spin
- Kleargene silica

#### Genotyping

Unique PCR-based genotyping chemistry and fit-for-purpose instruments developed and optimized in our service laboratories

- KASP genotyping chemistry
- SNPline instrumentation & validated procedures
- Kraken workflow man agement software
- Laboratory services

#### Sequencing

Sequencing services using traditional Sanger and Next Generation (NGS) sequencing methods. Backed by 20+ years of direct experience

- · Sanger sequencing
- NGS Illumina
- NGS Roche 454
- Covaris shearing instrumentation

#### Accurate, flexible, highly cost-effective solutions





# LGC Genomics in summary

- Deliver accurate, reproducible analytical science solutions
- We use what we sell and sell what we use
- We use the instruments and reagents that we sell to also drive a profitable service business







Extraction

## **Extraction Products and Services**



- Patented sbeadex<sup>®</sup> technology
- Mag<sup>™</sup>





# **Optimised solutions for any application - Just a few examples**



#### Food, saliva, plasmids, prokaryotes etc.





# **Analytical pathway in DNA analysis**







## How to cope with m(illions of) any samples

- Parallel processing to increase throughput (MTPs)
- Decrease scale (Miniaturization)
- Automation



Increase data output / time and money

- Throughput (samples per run, day etc.)
- Investment
- Running cost
- Reliability of chemistry and instrumentation
- Flexibility of protocols regarding

sample material, scale





# **Portfolio – Nucleic Acid Preparation (NAP)**

## **Kits for automated DNA extractions**

- sbeadex<sup>®</sup> kits
- mag<sup>™</sup> kits

## Instruments

Magnetic particle manipulators









## **Magnetic bead based SPE of DNA**







# General benefits of magnetic beads technology:

- Reduced labor, *cost effective* and usually highly scalable
- Facilitate high-throughput extraction, minituriazation & easy to automate
- Reduced set up costs
- Ongoing running costs are low







## sbeadex<sup>®</sup> principle

- a change of binding mechanism during extraction -



now incorporating



## sbeadex<sup>®</sup> - advantages / benefits

- No compromise in lysis conditions
- No organic solvents in wash buffers Price, convenience
- No drying Speed
- In principle, no chaotrophs for binding required
- Fast extractions







# sbeadex<sup>®</sup> - plant kit

### **DNA extraction from different plant (leaf material)**

Material	ID	ng/ μL	260/280	260/230	Total [µg]	RNase								
Pepper	1/1	68.3	2.1	1.5	5.5	-	(er 1	er 2	er	ato		u		
plant <sup>1)</sup>	1/2	66.0	2.1	1.7	5.3	-	Mark	<b>l</b> ark	epp	omá	tape	Cotto	lice	
Tomato <sup>2)</sup>	2/1	29.1	2.2	1.7	1.5	-		2		1961 1992	LE.	0	Ľ.	11-10-
	2/2	26.7	2.0	1.6	1.3	-	isti i	-	-		-	-		
Rape <sup>1)</sup>	3/1	39.9	1.9	0.8	3.6	+				周藤				
	3/2	21.0	1.9	1.2	1.9	+	-							
Cotton plant <sup>1)</sup>	4/1	84.3	1.8	1.2	7.6	+	-							
	4/2	91.7	1.8	1.1	8.3	+	No.							
Rice <sup>1)</sup>	5/1	9.5	1.8	0.8	0.9	+	-							
	5/2	9.1	1.5	0.7	0.8	+					B. S			





# sbeadex<sup>®</sup> - plant kit

## **Examples of validated protocols**

- Cotton
- Chicory
- Grapewine
- Hops
- Lettuce
- Maize
- Oak
- Onion

- Pepper
- Pine
- Rapeseed
- Rice
- Sunflower
- Tomato
- Wheat









# No contractual / technical obligation!

## Automated liquid handling

oktopure<sup>™</sup>





## **Co-operations with manufacturers of magnetic bead** handling systems

- TECAN
   King Fisher; Thermo Scientific
- Hamilton
   Beckman
- PerkinElmer
  - C Genomics



# **Applications & analysis**

- Molecular biology: e.g. PCR, SNPs
- Plant & animal breeding: e.g. Genotyping & Sequencing
- Diagnostics:

e.g. pathogens, viruses etc.









# The sbeadex<sup>®</sup> toolbox

## **Standard ingredients**

- 2. Binding buffer 5. Wash buffer 2
- 3. sbeadex beads 6. Elution buffer
- 1. Lysis buffer 4. Wash buffer 1



## Individual adds on / Tailor made solutions

- Customized bulk kit for almost all robotic systems (home made included)
- Challenging samples / protocol development
- **DNA** concentration
- Total amount of DNA





# oKtopure™

- 96 channel pipetting head
- Optimized for magnetic bead based extractions
- Max. 8 plates x 96-well platess per run
  - = 768 samples (extraction, only)

### <u>Run time</u>

mini scale 1h 10min

maxi scale ca. 2h











Genotyping

## **Genotyping Products and Services**



- KASP genotyping chemistry
- SNPline workflow





# **KASP™ Genotyping System**

- Novel Kompetitive Allele-Specific PCR
- Designed to overcome high cost of dual-labeled probes
- Homogeneous fluorescent assay no separation steps
- Scalable single-plex PCR: 96/384/1536 plates
- Delivers accurate SNP and InDel genotyping data while providing maximum flexibility and cost savings.
  - Improved genotype clustering, especially with plant and polyploid organisms; fewer primer-dimer (PD) issues.
  - Run as few or as many assays as you like; repeat what you want
  - Price per data-point remains consistently low





# **KASP Chemistry**



Allelic discrimination achieved through competitive annealing of two allele-specific forward primers, each containing a unique tail sequence that corresponds with a distinctly labeled FRET cassette in the master mix.

- Excellent data quality
- Maximum flexibility
- Cost-effective
- Minimal DNA material required
- Leading SNP to assay conversion rate (>90%)







# **KASP workflow**







## **SNPline workflow – process overview**







## **SNPline workflow:** DNA stamping - plate replication & reformatting



#### **SNPline Lite: Kpette – manual**

- Manually stamp DNA into 96/384/1536 plate formats
- Tip guides make for quick processing
- Optional tip regeneration station with optimized protocol can virtually eliminate tip costs



#### **SNPline XL: repliKator – fully automated**

- 24 x 1536 well plates in 15 minutes
- Onboard tip washing eliminate tip costs
- 96/384/1536 integrated into Kraken Software
- Integration friendly





## **SNPline workflow** Assay assembly – reagent dispensing

#### **SNPline Lite & XL: Meridian Dispenser**

- Single or 8 Channel configurations
- Dispense volume: 300nl to 100ul
- Unique aspirate function greatly reduces waste of expensive reagents
- Standalone PC control interface, and integrated control through Kraken software
- Angled tips to avoid cross contamination
- Peltier cooled plate locator
- Fill 24 x 1536 plates per hour









# **SNPline workflow: Plate Sealing**





#### **SNPline Lite: Kube Plate Sealer**

- Automated heat sealer
- Hermetically seals 96 & 384 plates
  - Standard height and deep well
- Advanced onboard control interface
- Integration friendly

### **SNPline XL: Fusion Plate Sealer**

- Automated laser sealer patented
- Hermetically seals 96, 384, & 1536 plates
- Ability to re-seal up to 50 times!
- Intuitive software control interface
- Integration friendly





## **SNPline workflow:** High throughput PCR – water bath thermal cycling





#### **SNPline Lite: Hydrocycler-4**

- Process 4 x 96, 384, or 1536 plates at a time
- Up to 30% faster than Peltier based systems
  - Typical KASP PCR run = 1.15 hours
  - 6 full runs in an 8 hour day
- Automated, robust and consistent

#### SNPline XL – Hydrocycler-16, 32, or 64

- Process 16 to 64 x 384 PCR plates at a time; or 14 to 56 x 1536 PCR plates.
  - Up to 500,000 data points/day!
- Energy efficient; throws off less heat
- Eliminates need for special AC environment





# **SNPline workflow: Fluorescent Plate Reading**



- Kraken interface for both models

   automatic file generation, export, and scoring
- Optional 36-plate stacker available for both models.

#### SNPline Lite: BMG FLUOstar Omega

- Single excitation/emission
- Three passes, three reads
- Reads KASP assays in 96 & 384 plates
- Speed: 1.5 minutes/384 well plate

### **SNPline XL: BMG PHERAstar**

- Simultaneous dual-emission technology
- Two passes, four reads
  - Faster with greater sensitivity
- Reads KASP assays in 96, 384, & 1536 plates
- Speed: 1.2 minutes/1536 well plate





# **Scalability and Cost Savings**



Reagent consumption is the primary cost driver
Two-fold savings from 96 to 384, Ten-fold from 96 to 1536
Next is equipment: cost, throughput, flexibility
Hydrocycler 64 will process 64 x 384, or 52 x 1536 plates in 1.2 hours!

•Plate and seal costs are a wash when analyzed per well





# **Driving scalability and cost savings**







Novel primer approach with KASP Minitaturisation to enable 1µL reaction volumes The flexibility and scalability of SNPline

All this means we can drive cost per data point to as low as 1 or 2 cents per data point, which is back where we started and why we do all this, which is to enable our customers to do more with their research dollars





# **Thank You For Your Time**



