

Thermo Fisher S C I E N T I F I C

Development of AgriSeqTM targeted GBS panels for breeding and parentage applications in cats

Angela Burrell 9 July 2019

Background

- There is a need for a robust, repeatable, and unambiguous workflow needed for feline parentage and genetic trait testing.
- We developed a targeted sequencing panel, one for feline parentage/ID verification and genetic defect/trait identification.
 - The AgriSeq Feline PITD Panel
- Utilizes the AgriSeq workflow

AgriSeq™ Targeted GBS Solutions for Agriculture

AgriSeq is a powerful, customizable, flexible and cost-effective high throughput Targeted GBS workflow capable of rapidly genotyping 50 - 5000 markers across plant, aqua and animal species.

>100 custom panel designs for over 35 species and counting

Plants									
BarleyCacaoCanolaCorn (maize)CottonCucumberEucalyptusLettuce	OatsOnionPinePotatoRiceSorghumSoybeanSpinach	SpruceSunflowerTomatoWheatWatermelon							
Animals									
BovinePorcineEquineCanineDeerOvine	FelineChickenSalmonChickenShrimpTuna	TroutTurbotBlack SoldierFly							

Panel Background

 AgriSeq Feline Parentage, ID, Traits & Defects (PITD) Panel* (A43408)- 175 markers

Marker Type	SNPs	MNPs	Insertions	Deletions	Total
Parentage & ID Markers	111	0	0	0	111
Traits & Disorders Markers	43	1	4	16	64
Total	154	1	4	16	175

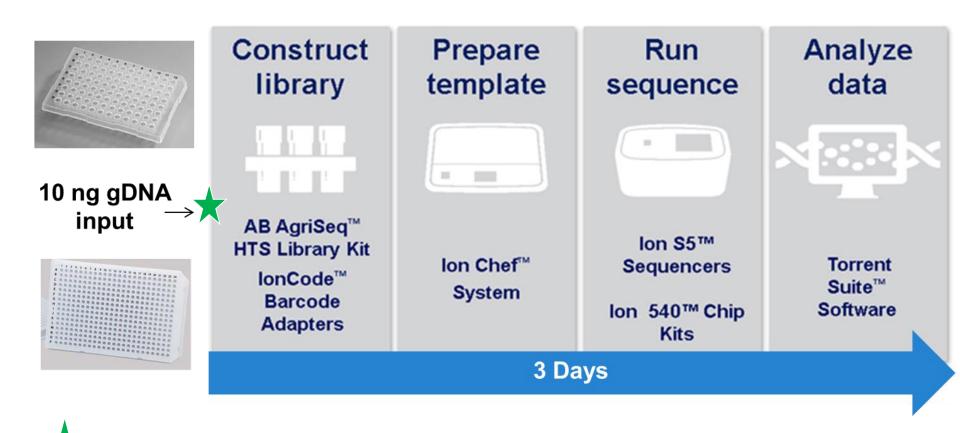


Targeted primer panel for the combined detection of feline genetic disorders/trait detection and parentage verification.



^{*}For Research Use Only. Not for use in diagnostic procedures

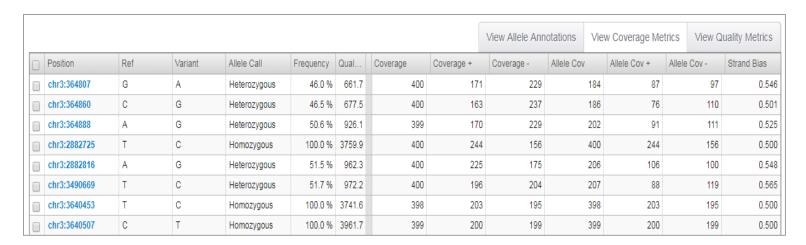
AgriSeq Sequencing Workflow



AgriSeq Feline Parentage, ID, Traits and Disorders (PITD) panel addition.

TVC Browser Output – Data for Analysis

							View Allele A	nnotations	View Coverage Met	rics View Quality Metrics		
	Position	Ref	Variant	Allele Call	Frequency	Quality	Subset Of	Variant Type	Allele Source	Allele Name	Gene ID	Region Name
	chr3:364807	G	Α	Heterozygous	46.0 %	661.7		SNP	Novel	tvc.novel.357	ASGA0100357	SP_2188.95472
	chr3:364860	С	G	Heterozygous	46.5 %	677.5		SNP	Novel	tvc.novel.358	ASGA0100357	SP_2188.95472
	chr3:364888	Α	G	Heterozygous	50.6 %	926.1		SNP	Hotspot	ASGA01003	57 ASGA0100357	SP_2188.95472
	chr3:2882725	Т	С	Homozygous	100.0 %	3759.9		SNP	Novel	tvc.novel.359	MARC0069764	SP_2190.153415
	chr3:2882816	Α	G	Heterozygous	51.5 %	962.3		SNP	Hotspot	MARC00697	64 MARC0069764	SP_2190.153415
	chr3:3490669	Т	С	Heterozygous	51.7 %	972.2		SNP	Hotspot	MARC00293	48 MARC0029348	SP_2193.195026
	chr3:3640453	Т	С	Homozygous	100.0 %	3741.6		SNP	Novel	tvc.novel.360	M1GA0025634	SP_2194.75717
	chr3:3640507	С	Т	Homozygous	100.0 %	3961.7		SNP	Hotspot	M1GA00256	34 M1GA0025634	SP_2194.75717



- Data analyzed by TVC (Torrent Variant Caller) Plugin
- All the information from the browser can be downloaded into a summary .XLS file.



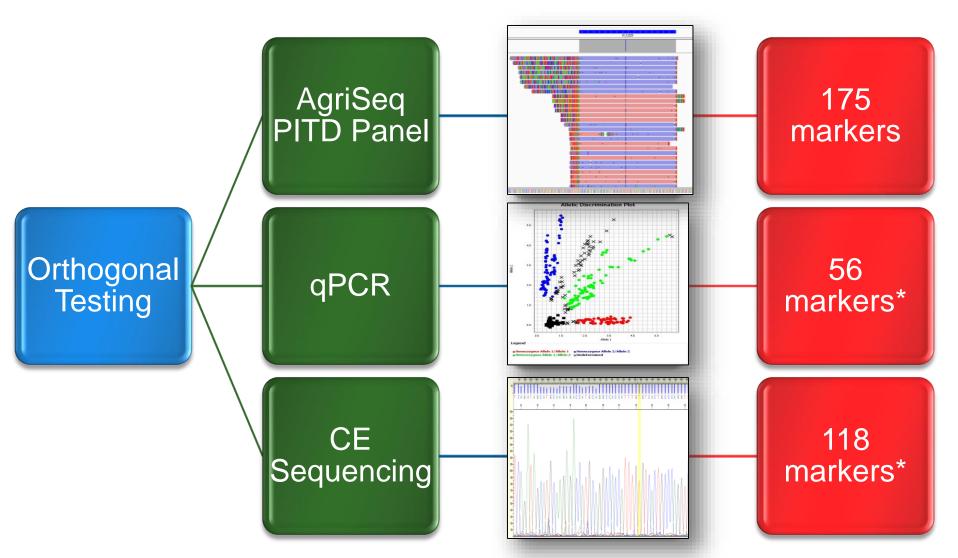
Experimental Overview

- Three general experiments were performed to validate performance of the Feline PITD panel.
 - 1. Orthogonal Testing ⇒ Evaluation of panel accuracy.
 - 2. Robustness Testing ⇒ Evaluation of panel consistency.
 - 3. Field Sample Testing ⇒ Evaluation of panel performance.

Experiment 1: Orthogonal Testing

 Purpose: To confirm that genotypes generated with the AgriSeq Feline PITD panel were accurate by testing with a separate, orthogonal technology.

Orthogonal Testing Workflow



*One marker generated a No Call with both CE and qPCR and was omitted from calculations.

Orthogonal Genotype Concordance Results

 Of the 161 markers generating a call for both technologies, no markers were discordant.

Orthogonal Method	# Concordant Markers to GBS	# Discordant Markers to GBS	# No Call Markers	Orthogonal Concordance	
CE Sequencing	105	0	13 (CE only)		
qPCR	56	0	1 (CE and GBS)	100%	

Experiment 2: Reproducibility and Repeatability

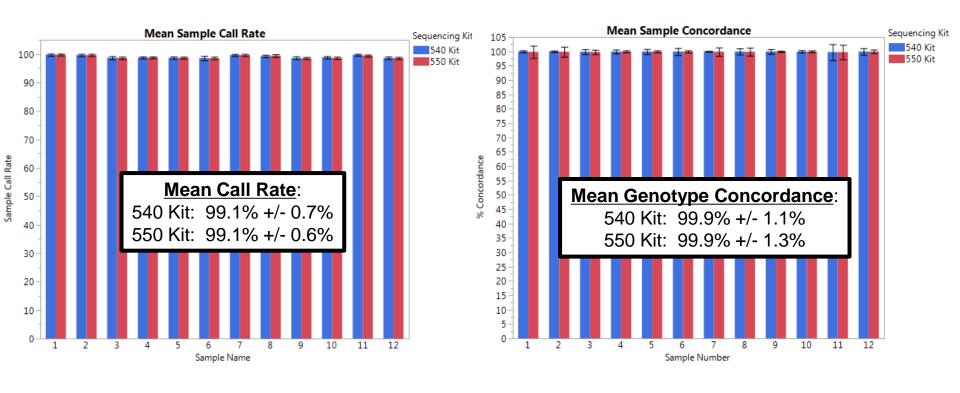
 Purpose: To test workflow robustness and genotype call consistency through multiple replicate reactions of a panel of samples.

- 12 feline DNA samples were tested in replicates (n=64) for a total of 768 barcoded libraries with the AgriSeq Feline PITD panel.
- Libraries were sequenced twice on the Ion 540 and Ion 550 chips.



Repeatability and Reproducibility

- Mean call rate was >99% with minimum variation between samples demonstrating the robustness of the genotype calls.
- The Feline PITD panel results were highly consistent with a mean genotype concordance of >99.9% for both sequencing kits between replicate samples.



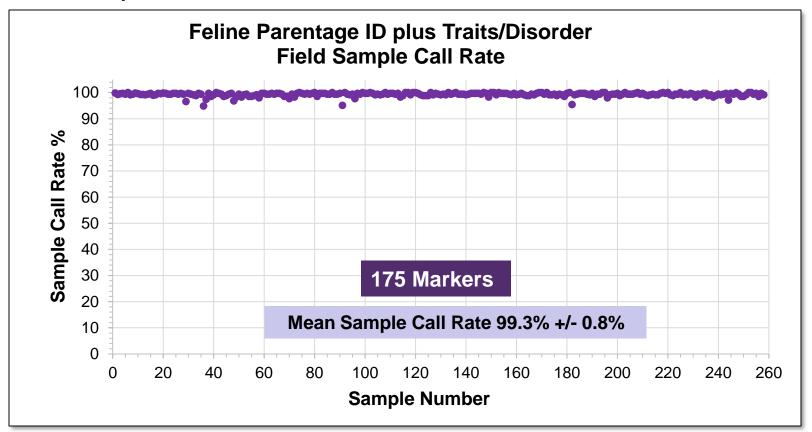
Experiment 3: Field Sample Performance

 Purpose: To determine panel performance with a diverse set of sample.

- 258 feline oral swab DNA samples from 4 different labs were tested with the AgriSeq workflow using the Feline PITD kit.
- 1ng/rxn DNA was input into library prep

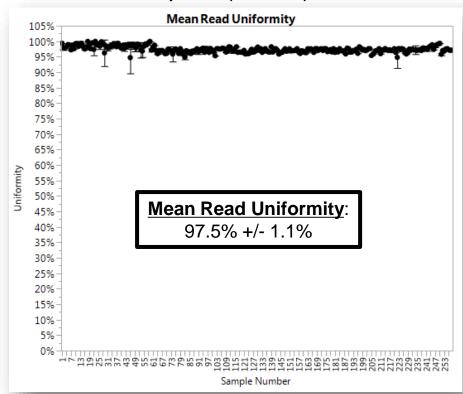
Sample Call Rates

 The mean call rate was 99.3% and all samples had call rates > 92.5% demonstrating the high performance obtained from field samples.



Read Uniformity

- Read uniformity measures how evenly you are covering target amplicons with reads.
 - Low uniformity (<90%) can lead to marker drop-off and poor call rates.
- The mean read uniformity for the panel was excellent, even when testing a set of very diverse field samples (>97%).



Conclusions

- AgriSeq library prep kit and Feline PITD panel combine into a robust and efficient workflow for feline genotyping applications.
 - Orthogonal Concordance 100%
 - Mean Field Sample Call Rate <u>>99%</u>
 - Replicate Genotype Concordance >99.9%





Experience the power of AgriSeq with 2 Enabling Options

Acknowledgements:

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Bruce Lau

Betsy Parker

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GBS EASY Jump Start bundles



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Free genotyping of your sample using AgriSeq GBS Panels

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