cfDNA analysis in a molecular pathology laboratory

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Ipatimup – Porto - Portugal





Ipatimup

- Founding member of i3S
- Leading cancer research institute in Portugal
 - Founding partner of Porto Comprehensive
 Cancer Center





i3S

- Biggest research in health institute in Portugal (1250 researchers);
- Cancer, Neurosciences and Host-Pathogen interactions research lines



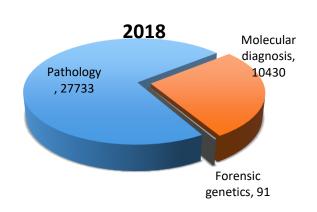
IPATIMUP diagnostics

Provide services in the areas of surgical pathology and cytopathology, genetic diagnosis and genetic identification and parentage, in order to improve Portuguese citizens life quality.

Molecular diagnosis

Laboratory accreditation

- Tumor mutation screening 4220 cases
- Genetic diagnosis 2960 cases
- Pre-natal screening 3260 cases



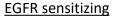






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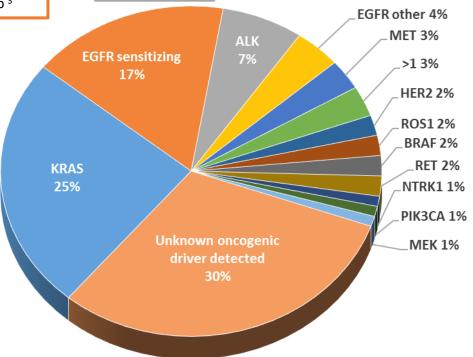
Lung cancer clinically relevant alterations



- Gefitinib ⁴
- Erlotinib ⁴
- Afatinib ⁴
- Osimertinib ⁴
- Necitumumab ⁴
- Rociletinib³

<u>ALK</u>

- Crizotinib 4
- Alectinib ⁴
- Ceritinib ⁴
- Lorlatinib ²
- Brigatinib²



ROS1

- Crizotinib ⁴
- Cabozantinib²
- Ceritinib ²
- Lorlatinib ²
- DS-6051b¹

patimup

<u>KEY</u>

- 1 Phase I 3 Phase III
- 2 Phase II 4 Approved



Lung cancer clinically relevant alterations

ALK

7%

EGFR sensitizing

- Gefitinib 4
- Erlotinib 4
- Afatinib 4
- Osimertinib 4
- Necitumumab 4
- Rociletinib³

KEY

1 - Phase I

2 - Phase II

3 - Phase III

4 - Approved

ALK

- Crizotinib 4
- Alectinib 4
- Ceritinib 4
- Lorlatinib²
- Brigatinib²

EGFR sensitizing

17%

KRAS

25%

MET

- Crizotinib²
- Cabozantini²

HER2

- Transtuzumab emtansine²
- Afatinib ²
- Dacomitinib²

ROS1

- Crizotinib 4
- Cabozantinib²
- Ceritinib²
- Lorlatinib ²
- DS-6051b 1

BRAF

- Vemurafinib²
- Dabrafenib²

RET

- Cabozantinib²
- Alectinib 2
- Apatinib²
- Vandetanib²
- Ponatinib²
- Lenvatinib 2

MEK1

Unknown oncogenic

driver detected

- Trametinib²
- Selumetinib³
- Cobimetinib 1

PIK3CA

EGFR other 4%

MET 3%

>13%

HER2 2%

ROS12% BRAF 2%

RET 2%

NTRK1 1%

PIK3CA 1%

MEK 1%

- LY3023414²
- PQR 309 ¹

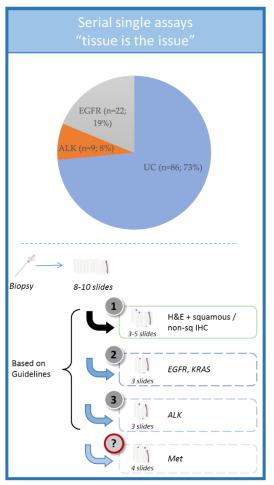
NTRK1

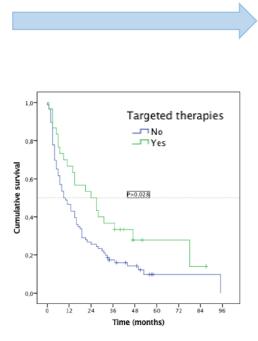
- Entrectinib ²
- LOXO-101²
- Cabozantinib²
- DS-6051b 1

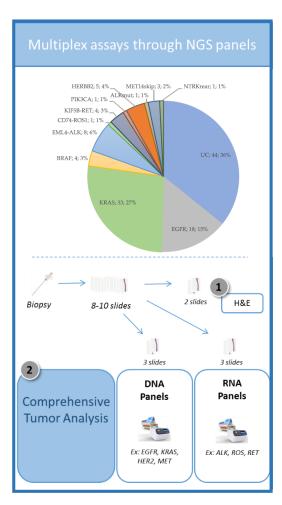




Comprehensive biomarker characterization





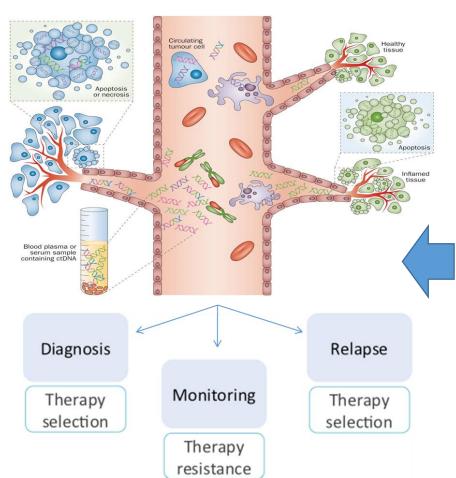


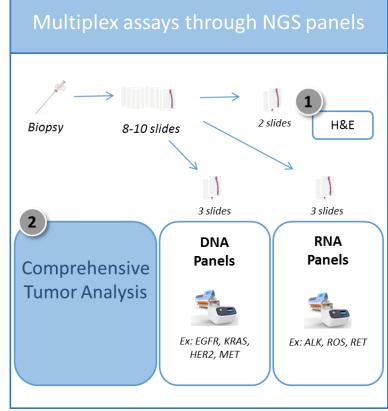
Comprehensive biomarker testing leads to better future patient care





Lung cancer characterization ctDNA analysis





Tissue may not be available for all assays or situations



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Validated strategy

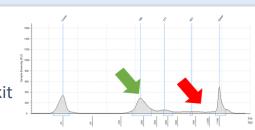
Plasma collection 1. Plasma collection

BD Vacutainer PPT (K2EDTA)



2. cfDNA isolation

MagMax cfDNA extraction kit



cfDNA isolation

NGS **Variant** Discovery

dPCR validation

3. NGS variant identification



- Ion AmpliSeq Colon & Lung Cancer Research Panel
- Ion AmpliSeq RNA Fusion **Lung** Cancer Research Panel
- Oncomine cfDNA Lung assay



Ion S5xl System

4. Digital PCR validation





TaqMan Assays





ISO 15189 accreditation

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OncoNetwork multicentric study sensitivity and specificity

Lab	Sensitivity	Specificity	NPV	PPV
ARCNET	89,6%	99,6%	99,6%	87,8%
CROM	95,8%	100,0%	99,8%	100,0%
IPATIMUP	95,8%	100,0%	99,8%	100,0%
Radboud	89,6%	99,8%	99,6%	95,6%
Queens	97,9%	99,9%	99,9%	97,9%
St James	95,8%	100,0%	99,8%	100,0%
UHCW	95,8%	99,8%	99,8%	93,9%
Charite	95,8%	99,8%	99,8%	93,9%
Viollier*	97,5%	99,8%	99,9%	95,1%
HEGP	93,7%	99,7%	99,8%	91,8%
Kindai*	95,8%	99,7%	99,8%	92,0%
Total	94,8%	99,8%	99,8%	95,9%

Multiplex I cfDNA Reference Standard Set (5%, 1%, 0,1%, 0%)

- **EGFR** p.L858R
- EGFR p.E746 A750delELREA
- **EGFR** p.T790M
- EGFR p.V769 D770insASV
- **KRAS** p.G12D
- **PIK3CA** p.E545K







OncoNetwork multicentric study

Benchmarking at 0.1% AF

		OncoNetwork study	FOUNDATION ACT	GUARDANT 363 ™	
Genes (n)		11	62	73	
SNVs	sensitivity	83.9%	67.3%	63.8%	
	PPV	99.1%	93.6%	92.1%	
InDels	sensitivity	83.9%	86.2%	67.8%	
	PPV	99.1%	100%	88.4%	

www.foundationmedicine.com www.guardanthealth.com

Confident detection of variants at 0.1% AF





Concordance analysis of liquid and tissue biopsy in NSCLC patients: a multiinstitutional molecular pathology study

Multi-institutional study, 159 NSCLC patients who underwent tissue and plasma-NGS analysis:

- 1. 94 patients had concurrent tissue and plasma NGS analysis at diagnosis
- 2. 65 patients were monitored through plasma-NGS analysis followed-up over time and compared to tissue biopsy at diagnosis

Alborelli I., Costa J.L. et al.: data from IPATIMUP (Porto) and the Institute of Pathology of Basel





Cohort Characteristics

Characteristics	No. (%)
Patients	159 (100)
Age, y	
Median (Range)	67.5 (38 - 89)
Sex	
Male	77 (48.4)
Female	82 (51.6)
Histology	
Adenocarcinoma	158 (99.4)
Squamous cell carcinoma	1 (0.6)
Matched tissue biopsy	
Primary tumor	145 (91.2)
Metastasis	14 (8.8)
Metastatic status at blood draw	
Confirmed metastatic	136 (85.5)
Confirmed non-metastatic	17 (10.7)
Unknown metastatic status	6 (3.8)
Smoking status	
Never smoker	31 (19.5)
Smoker	31 (19.5)
Ex-smoker	29 (18.2)
Passive smoker	3 (1.9)
Unknown	65 (40.9)





Benchmarking

High concordance of plasma and tissue-based molecular analysis using three different commercial assays.

Study	Total	Discordant	Tissue+ AND Plasma+ (TP)	Tissue AND Plasma (TN)	Concordance %
Aggarwal et al.	128	24	31	73	81%
Li et al.	110	23	68	19	79%
Alborelli et al.	94	11	40	43	88%

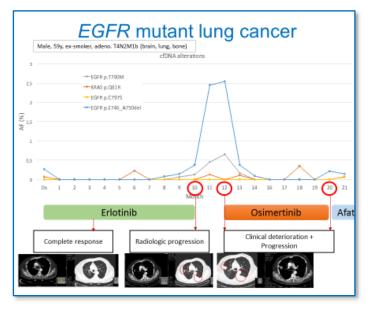
Study	TP	TN	FP	FN	Sensitivity	Specificity	PPV	NPV	Cohen Kappa
Aggarwal et al.	31	73	8	16	66%	90%	79%	82%	0.58
Li et al.	68	19	0	23	75%	100%	100%	45%	0.51
Alborelli et al.	40	43	2	9	82%	96%	95%	83%	0.77

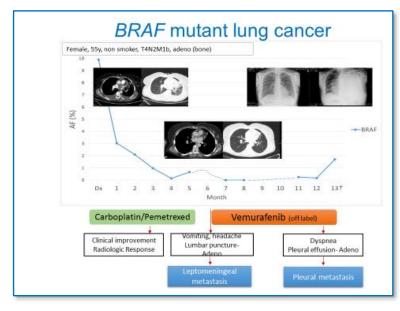
Study	Genes (n)	Depth of Sequencing	cfDNA input	In-house test
Aggarwal et al.	73	10'000X	5-30 ng	no
Li et al.	37	50'000X	Up to 100 ng	not yet
Alborelli et al.	11	25'000X	1-50 ng	yes

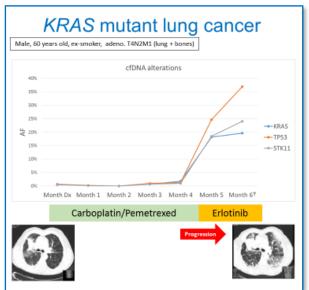
Our results highlight the feasibility of using an **in-house** plasma-NGS assay for routine molecular characterization.

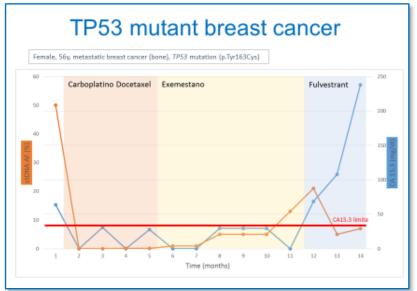




















- Informative in different tumor models
- Informative for different future therapeutic strategies
- Allows real-time study of the disease
- Allows study of future clinical relapse anticipation
- Identification of resistance mechanism







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Venceslau Hespanhol
Gabriela Fernandes
Fátima Carneiro
José Carlos Machado

Universitätsspital Basel

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First Experience from the "New World"

Thermo Fisher Scientific workshop – AMP 2019

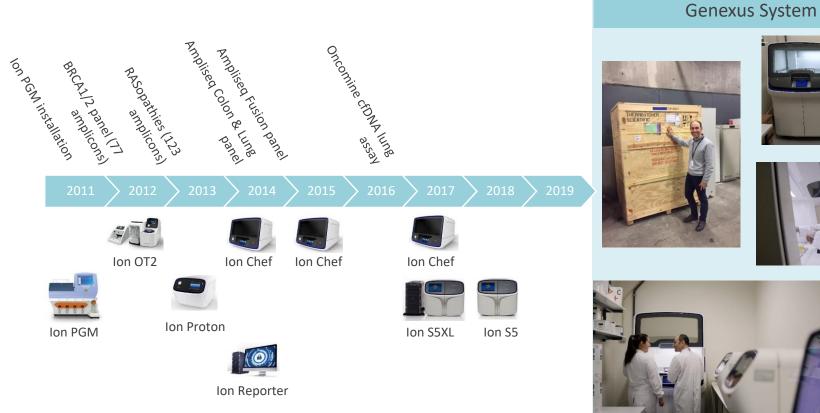
6th November 2019

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NGS clinical research timeline....



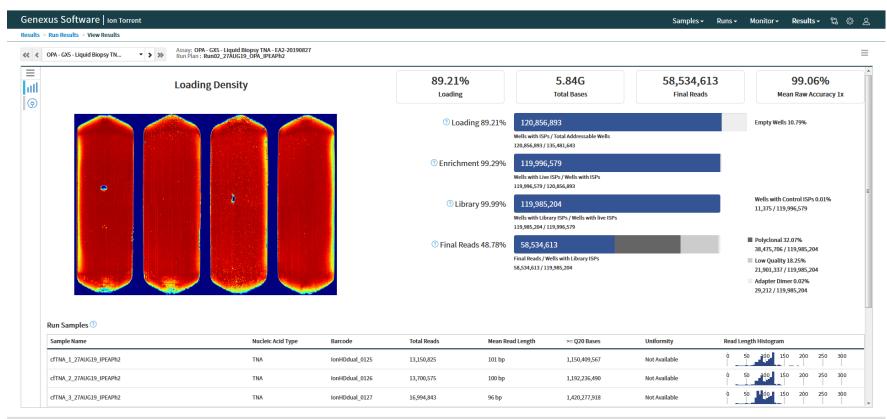


Commercial control samples

Horizon and SeraCare FFPE and cfDNA reference material

Clinical research samples

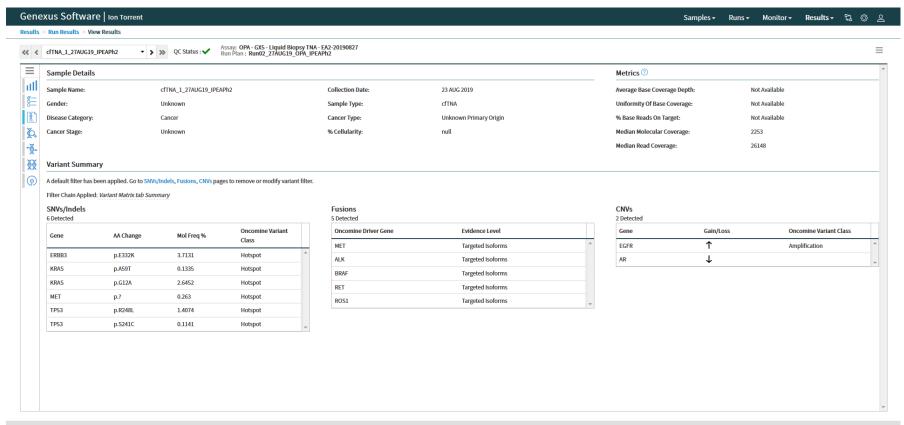
- Lung cancer tissue and liquid biopsy research samples
- Previously characterize on an Ion S5XL system



iontorrent

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Commercial control samples

Horizon and SeraCare FFPE and cfDNA reference material

Clinical research samples

- Lung cancer tissue and liquid biopsy research samples
- Previously characterize on an Ion S5XL system

Tissue biopsies – RNA and DNA

Lung cancer FFPE sample were sequenced using Colon and Lung or Lung Fusion panel on Ion S5XL system and using Oncomine Precision Assay on Genexus instruments



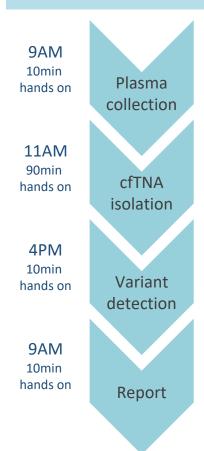
Variants covered by both panels were detected in both systems at similar allelic frequencies

Additional variants were detected using the Oncomine Precision Assay

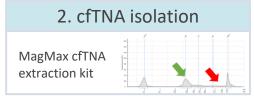
Sample 1 presented additional p.T790M mutation that had not been previously identified (intra-sample heterogeneity?)

FFPE	Gene	AA Change	Genexus	Ion S5XL
Sample 1	EGFR	p.L858R	43,8	31,3
	TP53	p.R248Q	1,8	nd
	EGFR	p.T790M	1,6	nd
	ALK	p.A1200V	1,5	nd
Sample 2	EGFR	p.L747_P753delinsS	28,6	20,1
	TP53	p.C176R	3,9	nd
	RET	p.G810S	1,6	nd
	EGFR	p.P848L	1,6	nd
	FGFR3	p.R399C	0,3	nd
Sample 3	ALK	fusion	detected	detected
Sample 4	BRAF	p.V600E	47,8	36,3
	TP53	p.R175C	13,8	nd
	FGFR2	p.A648T	6,9	nd
	EGFR	p.R836C	6,2	nd
	ERBB3	p.V104M	4,4	nd
	PDGFRA	p.T849C	4,2	nd
	PIK3CA	p.V344M	3,7	nd
	MAP2K1	p.K57N	2,8	nd
	GNAQ	p.R183Q	2,1	nd
	EGFR	p.V769M	1,8	nd
Sample 5	KRAS	p.G12D	58,3	44,7
	TP53	p.C176Y	31,4	nd
	FGFR3	p.R399C	3,6	nd
	CDKN2A	p.R58Q	3,1	nd
	RET	p.R912Q	2,5	nd
	BRAF	p.D594N	2,3	nd

Liquid biopsies – Lung cancer plasma sample at recurrence



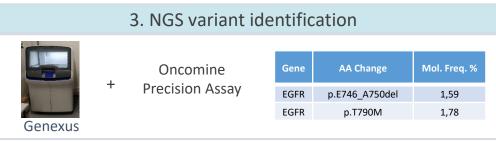


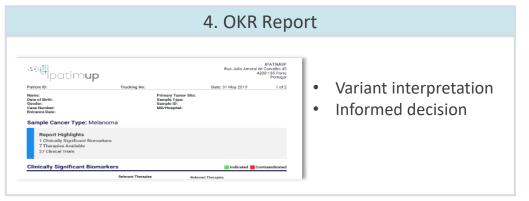


Sample

< 24h

Report





Summary

- ALL-IN-ONE solution from nucleic acids to variants;
- ACCESSIBLE: no a priori NGS expertize needed;
- SIMPLE: all I needed was a pipette to add my samples to be sequenced;
- ROBUST: vision-system checks all your steps;
- FAST: allowed a turn around time of 24h from plasma to report.
- This system allows a wider implementation of NGS for genomic profiling, potentially bringing precision medicine closer to clinical practice.

First Experience From New World





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Venceslau Hespanhol Cabriela Fernandes Carneiro Conceição Souto Moura nurses Carneiro

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Sohaib Qureshi?

Jussi Vanhatalo?

Ian Grinsell?

Rosella Petraroli?

Andy Felton?



















