

Assessing cfDNA somatic variant testing

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1st The European Liquid Biopsy Society ctDNA Technology Meeting, 23rd March 2022

IQNPath cfDNA pilot EQA Group



2016

Online survey of plasma ctDNA testing practice – 167 labs

Deans ZC, *et al.* Virchows Arch. 2017 Dec;471(6):809-813.



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2017

- Pilot EQA delivered to 32 clinical testing labs
- 5 samples for *EGFR* testing and/or 5 samples for *KRAS/NRAS* testing
- Artificial samples - 3 mL of human plasma with 20 ng/mL of fragmented ctDNA
- VAFs of 1 and 5%



Demonstrated feasibility of delivering a cfDNA EQA

Identified need for EQA due to high error rates (20%) in detecting low VAF clinically relevant variants

Keppens C, *et al.* BMC Cancer 2018 18;804



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2018

2019

- EQA provided to 264 laboratories from 45 countries
- 5 Cases with *EGFR* exon19 dels, p.(T790M), p.(L858R) & combinations - VAFs 0.5%, 1%, 5%
- Artificial plasma samples containing 250ng of fragmented ctDNA
- Genotyping accuracy high for VAFs (94%) >1%



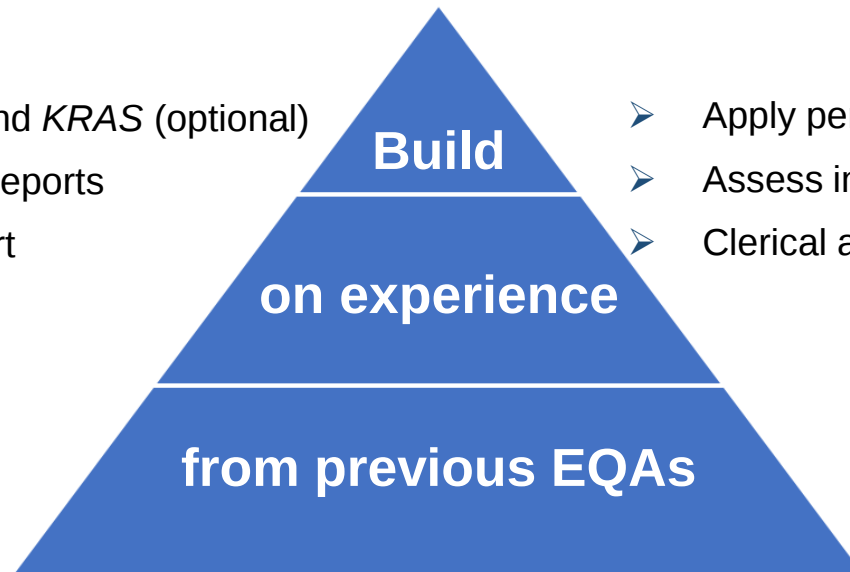
Results of a worldwide External Quality Assessment of cfDNA testing in Lung Cancer
Fairley JA, *et al.* (2022) Virchows Arch. Submitted



EQA for cfDNA testing for *EGFR* variants in Lung cancer EQA 2020 (n = 258)

- Centralised sample sourcing and validation of EQA materials
- Include common and clinically relevant variants
- Include challenging samples with low frequency allele variants
- Submission of clinical reports and data collection form

- *EGFR* (mandatory) and *KRAS* (optional)
- Individual laboratory reports
- EQA Summary Report



- Apply performance criteria for genotyping
- Assess interpretation of the results
- Clerical accuracy of the reports

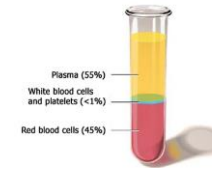
Format of EQAs



Finalise reports and
apply performance
criteria for
genotyping

Registration

Sample sourcing
& validation



Appeals
process

Individual EQA Provider
Scheme reports plus
laboratory reports



Collation of
results

Assessment

Marking criteria
defined

EQA
distribution

Results
submission

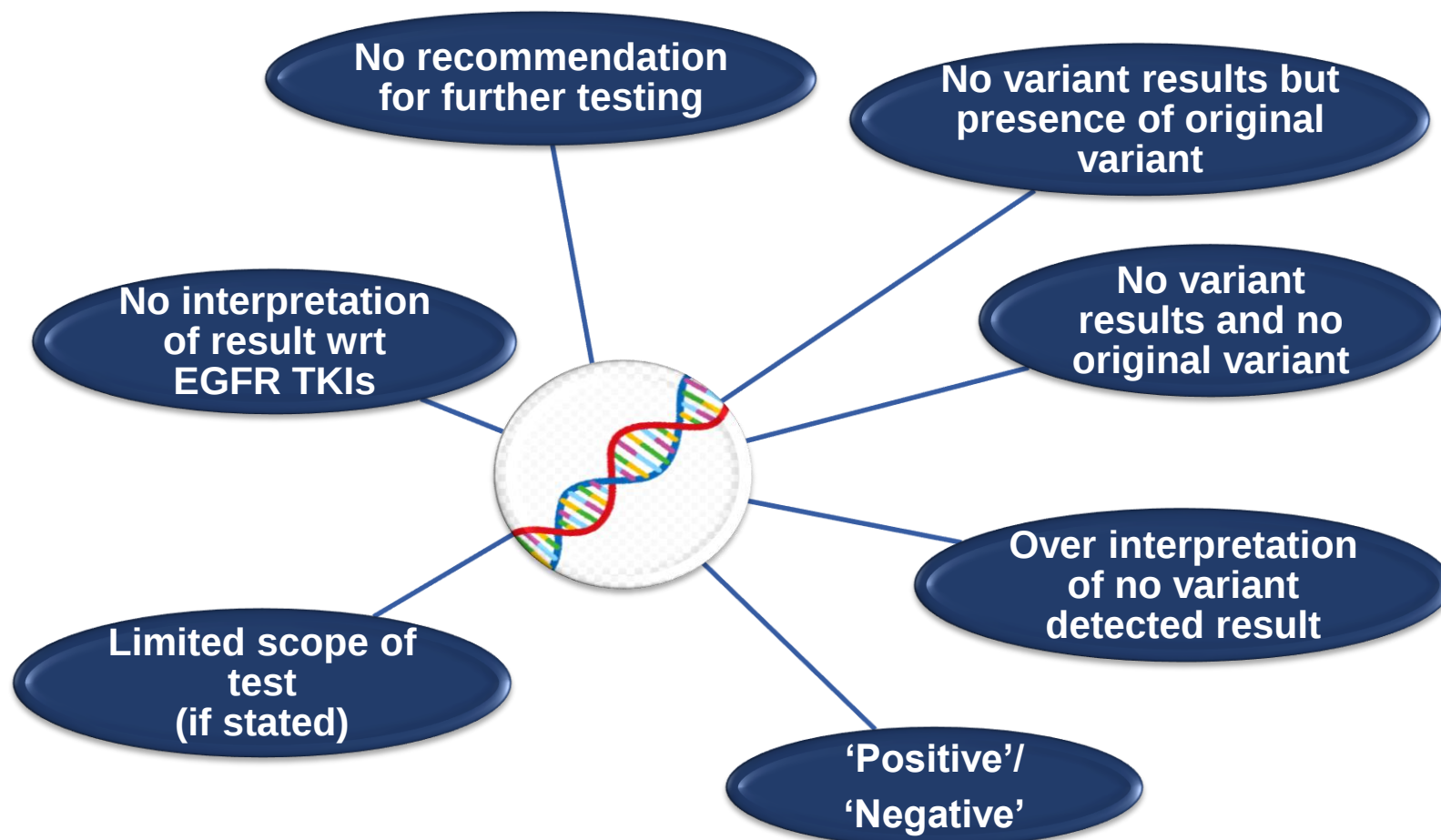


2020
2021

Genotyping

Case	Genotype	VAF	Critical genotyping errors	False positives	False negatives	Incorrect variant reported	Potential sample swap
1	No pathogenic variants	N/A	10 (4%)	9 (3.5%)	0	0	1 (0.4%)
2	<i>EGFR</i> c.2155G>A p.(Gly719Ser)	5%	27 (10%)	0	23 (9%)	3 (1.2%)	1 (0.4%)
3	<i>EGFR</i> c.2236_2250del p.(Glu746_Ala750del) and <i>EGFR</i> c.2369C>T p.(Thr790Met)	2% 1%	37 (14%)	0	17 (7%)	3 (1.2%)	2 (0.8%)
Note: 42% reported either one or no variants due to LOD of assays							

Interpretation issues





2021 EQA UPDATE

METHODS

>60 different ones!

173 labs using real-time PCR methods

- most common is COBAS EGFR mutation kit v2 (76)

97 labs used NGS with LOADS of different panels.

- most common being Oncomine lung cfDNA assay (14)
- all others <5 labs using them.

44 labs using ddPCR

There are still laboratories using kits not really suitable for cfDNA testing
– e.g. Therascreen EGFR pyro kit, Ion Ampliseq panels, Trusight Tumour 15.

Number of errors dramatically reduced from last year.

IQNPath cfDNA pilot EQA Group



AstraZeneca

NOVARTIS

MSD

Assessor	Location	Role
Rachel BUTLER	UK	Assessor
Caroline CLARK	UK	Assessor
Rachel DODDS	UK	Assessor
Hongxiang LIU	UK	Assessor
Dietmar LOHMANN	Germany	Assessor
Manisha MAURYA	UK	Assessor
Sophie MCHAFFIE	UK	Assessor
Hood MUGHLAASI	UK	Assessor
Thierry NOUSPIKEL	Switzerland	Assessor
Cathal O'BRIEN	Ireland	Assessor
Anca ONISCU	UK	Assessor
Markus RECHSTEINER	Switzerland	Assessor
Matthew SMITH	UK	Assessor
Philippe Taniere	UK	Assessor
Harri WEBB	Australia	Assessor
Melanie CHEETHAM	UK	Scheme Organiser
Jenni FAIRLEY	UK	Scheme Organiser
Simon PATTON	UK	Scheme Organiser
Nicola WOLSTENHOLME	UK	Scheme Organiser



Jenni Fairley
GenQA



Mel Cheetham
EMQN



**1st ELBS ctDNA
Technology Meeting
of 2022**
23 March 2022



Agenda

- Introduction - QuIP
- QuIP's ctDNA - EQA Program
- *EGFR* Mutation Analysis Liquid Biopsy EQA 2022



QuIP – Who are we?

The QuIP logo is positioned within a blue semi-circle at the bottom left of the slide. It consists of the word "QuIP" in a white, sans-serif font, with a stylized 'Q' that has a small dot above it.

QuIP



QuIP- Shareholders



**BUNDESVERBAND
DEUTSCHER
PATHOLOGEN e.V.**

**Federal Association of
German Pathologists e.V.
(BDP)**
Bundesverband Deutscher Pathologen e.V.

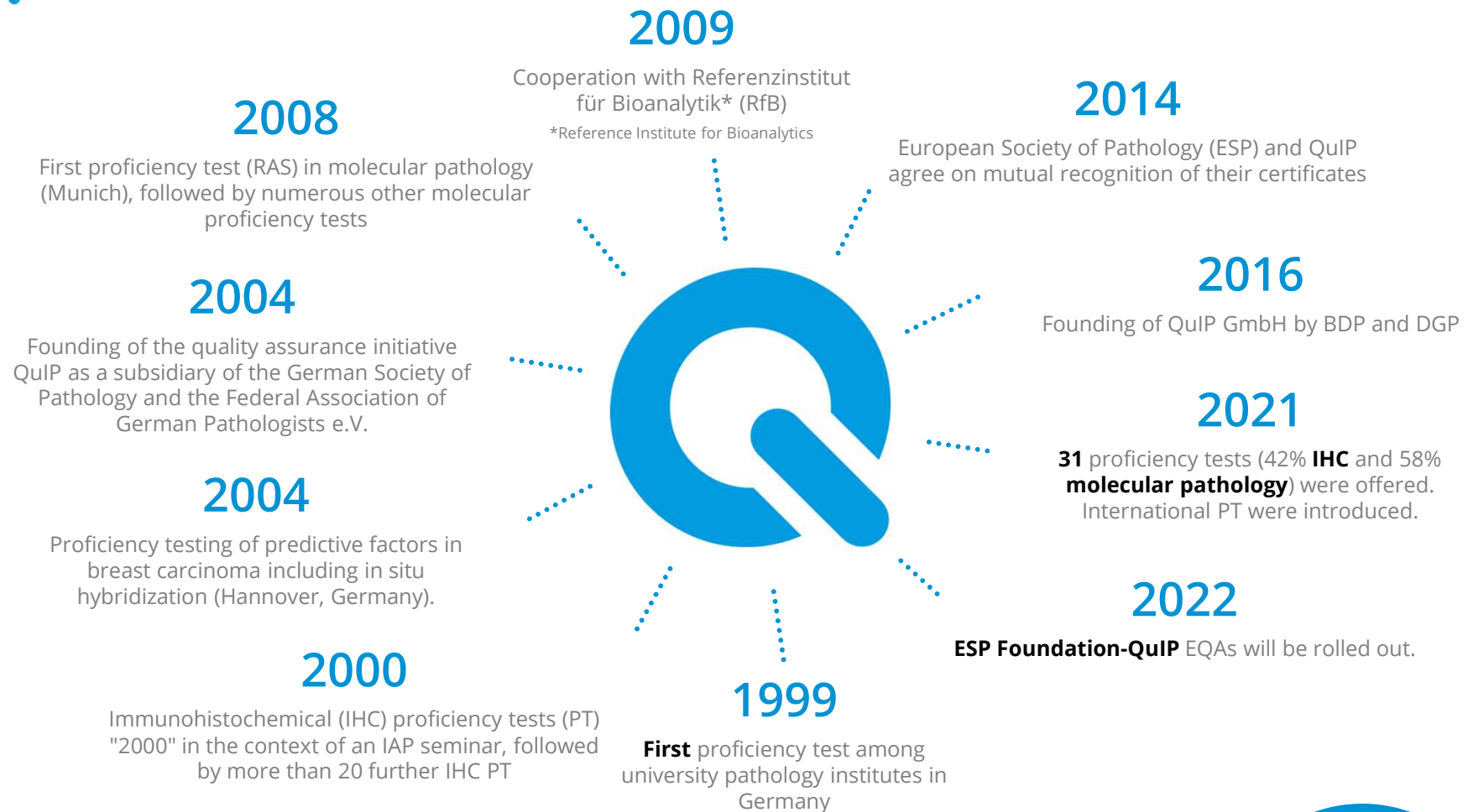


**DEUTSCHE GESELLSCHAFT
FÜR PATHOLOGIE E.V.**

Seit 1897 – dem Leben verpflichtet

**German Society for
Pathology
(DGP)**
Deutsche Gesellschaft für Pathologie e.V.

QuIP - History



QuIP's ctDNA - EQA Program

The QuIP logo is located in the bottom left corner of the slide. It consists of the word "QuIP" in a white, sans-serif font, centered within a blue semi-circular shape. The background of the slide is a solid blue color, and there are decorative white shapes: a semi-circle at the bottom left and a horizontal bar at the bottom right.

QuIP



QulP's Liquid Biopsy Program

2018

EGFR Exon 19-21

2019

EGFR Exon 19-21

2020

EGFR Exon 19-21
PIK3CA Mutation
Analysis

2021

EGFR Exon 20 Ins
PIK3CA Mutation
Analysis

2022

EGFR Exon 19-21
MET Exon 14
Skipping

23.03.2022

1st ELBS ctDNA Technology Meeting of 2022

EGFR Mutation Analysis Liquid Biopsy EQA 2022

Quip



EGFR Mutation Analysis: Design

2018 - 2020

- 9 ml blood + fragmented cell line DNA
- Min. 1 % allele frequency (exon 19-21)
- 10 samples

- + Samples resemble real-life samples
- Stability of samples
- Limited number available
- "Production" very work-intensive
- Internal testing at the same time as open proficiency test

2022

- Samples will be produced by SensID GmbH
- Min. 1 % allele frequency (exon 19-21)
- 7 samples

- + Unlimited availability
- + Stability of samples > 1 year
- + Internal testing before open proficiency test
- Artificial samples
- 1st step of the workflow (isolation of plasma) is skipped

EGFR Mutation Analysis 2018-2022:

Case-specific questions

EGFR Liquid Biopsy 1

	P1	P2	P3	P4	P5	P6	P7	P8	P9	P10
Mutationsstatus Exon 20 (T790M)										
Wildtyp	☐	☐	☐	☐	☐	☐	☐	☐	☐	☐
Mutiert	☐	☐	☐	☐	☐	☐	☐	☐	☐	☐
nicht auswertbar	☐	☐	☐	☐	☐	☐	☐	☐	☐	☐
Mutationsstatus Exon 19 (EGFR)										
Wildtyp	☐	☐	☐	☐	☐	☐	☐	☐	☐	☐
Mutiert	☐	☐	☐	☐	☐	☐	☐	☐	☐	☐
nicht auswertbar	☐	☐	☐	☐	☐	☐	☐	☐	☐	☐
Mutationsstatus Exon 21 (EGFR)										
Wildtyp	☐	☐	☐	☐	☐	☐	☐	☐	☐	☐
Mutiert	☐	☐	☐	☐	☐	☐	☐	☐	☐	☐
nicht auswertbar	☐	☐	☐	☐	☐	☐	☐	☐	☐	☐



Thank you for
your attention.



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