XXV Congreso de la Sociedad Española de Anatomía Patológica y División Española de la International Academy of Pathology

Estado actual y perspectivas del estudio de Biomarcadores Desde el ámbito europeo

Fátima Carneiro President-Elect of the European Society of Pathology IPATIMUP & Medical Faculty/Hospital SJoão Porto, Portugal







Several initiatives, both diagnostic and research driven and both private and public have been established under the auspices or with the collaboration of the European Society of Pathology



The European Molecular Genetics Quality Network



HARMONIZING GENETIC TESTING ACROSS EUROPE





KRAS-ESP External Quality Assurance (EQA) program



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REVIEW AND PERSPECTIVE

KRAS mutation testing for predicting response to anti-EGFR therapy for colorectal carcinoma: proposal for an European quality assurance program

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KRAS-ESP European Quality Assurance (EQA) program

The European Society of Pathology (ESP) established a European QA program for testing KRAS mutations in colorectal cancer. This program aims to ensure optimal accuracy and proficiency in KRAS mutation testing across all countries or institutions in the European Union.

A first European pilot KRAS EQA scheme was running May - June 2009. Based on these experiences Regional and European EQA schemes were organized in different countries in 2009 and 2010.







Regional KRAS EQA scheme 2010

(IPATIMUP was one of the 6 Regional Laboratories responsible for the preparation and validation of the samples)







LOGO Institute/Laboratory

EQA ID Institution - Department - Address Contact person laboratory Tel - Fax

Leuven, 10-10-2010

Referring clinician: Doctor X Street City, Country

Molecular genetic analysis for KRAS

Name:	John Doe (*)
Date of birth:	01/01/1980 (*)
Reason for analysis:	KRAS mutation present?
Sample received:	01-10-2010
Sample type:	Paraffin embedded section of adenocarcinoma
Your reference:	CF07-2
Our reference:	KRAS 10.199

RESULT:

- MICROSCOPY: Microscopic analysis of the sample showed 70% tumor cells. Sample is suitable for KRAS mutation analysis
- MUTATION ANALYSIS: Mutation present in codon 12 of the KRAS gene: p.Gly12Ser (c.34G>A) -

INTERPRETATION:

In a DNA extract of the sample tissue, we detected an activating mutation in the KRAS gene: p.Gly12Ser (c.34G>A). This indicates likely resistance to anti-EGFR-therapy (cetuximab, panatimumab)

Analysis performed by

Microcopy performed by



Molecular biologist X

The method used: DNA-extraction with the Qiagen QIAamp® DNA FFPE-kit, K-RAS mutation analysis by the TheraScreen® K-RAS Mutation Kit (CE-IVD) (DxS Ltd, Manchester, UK), which combines two technologies (Amplification Refractory Mutation System, Astrazeneca, and Scorpions, DxS) to detect the most commonly reported KRAS mutations by realtime quantitative PCR. Mutations screened for: p.Gly12Asp (c.35G>A), p.Gly12Ala (c.35G>C), p.Gly12Val (c.35C>T), p.Gly12Ser (c.34G>A), p.Gly12Arg (c.34G>C), p.Gly12Cys (c.34G>T), p.Gly13Asp (c.38G>A)

These mutations represent about 98% of the KRAS mutations in colorectal cancer. The Therascreen method is highly selective. If sufficient DNA is present, presence of 1% mutant DNA can be detected in a wild type background. A negative test does not exclude the presence of a mutation.

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(*) Data might be not provided in EQA schemes: enter / or not applicable (NA)



Scores of important criteria in written reports sent by the participants. Percentages indicate in how many reports the item was present (n=75). In total **76 laboratories** participated in the EQA scheme 2010. The number of laboratories that participated (and submitted results) in 2010 is listed per country in Table 1.

	Number of laboratories that
Country	participated in the scheme and
	submitted results
Austria	7
Belgium	13
Denmark	8
France	12
Germany	3
Greece	2
Italy	6
The Netherlands	16
Norway	2
Portugal	1
Spain	2
Sweden	2
Switzerland	1
Turkey	1
TOTAL	76

Table 1: Number of laboratories that participated in the Regional KRAS EQA scheme, per country, in 2010.

Table with the individual genotype results:

Subscheme	EQA IDriab	KRASRc10.301	KRASRe10.302	KRASPc10.303	KRASRc10.304	KRASPCI 0.305	KRASPert 0.306	KRASRc10.307	KRASRc10.308	KRASRc10.309	KRASRc10.310	Total score genotyping	
		c.35G>A,	c.34G>T,		c.38G>A,		c.34G>A,		c.35G>C,	c.35G>T,			í
SCHEME A	Result	p.Gly12Asp	p.Gly12Cys	No mut	p.Gly13Asp	No mut	p.Gly12Ser	No mut	p.Gly12Ala	p.Gly12Val	No mut		1
	,	a	a	a	a	а	a	a	a	a	a	10,0	1
		a	a	a	a	a	a	a	a	a	a	10,0	1
		a	a	a	a	a	a	a	a	a	a	10,0	
		a	a	a	a	a	w (no mut)	a	a	a	n	8,0	
		a	a	a	a	a	a	a	n	a	a	9,0	1
		a	a	a	a	а	a	a	a	a	a	10,0	1
		a	a	a	n	а	a	a	n	a	a	9,0	1
		a	a	a	a	а	a	a	w (no mut)	a	a	9,0	1
		a	а	a	a	а	a	a	a	a	a	10,0	1
		a	а	a	a	а	a	a	a	a	n	9,5	1
		a	а	a	a	а	a	a	a	a	a	10,0	1
		a	а	a	a	а	a	a	a	a	a	10,0	1
		a	a 	a 	a	а	a	a 	a	a 	a /a 25Cb T	10,0	1
		w (no mut)	w (c.36G>A, p.Gly12Asp)	w (c.34G>1, p.Gly12Cys)	w (no mut)	а	w (no mut)	w (c.34G>A, p.Gly12Ser)	w (no mut)	p.Gly12Ala)	w (c.30G>1, p.Gly12Val)	1,0	
		a	а	a	n	а	a	а	n	a	a	9,0	
		a	а	a	a	а	a	a	a	a	a	10,0	1
		c.35G>A,	c.34G>T,		c.38G>A,		c.34G>A,		c.35G>C,	c.35G>T,			1
SCHEME B		p.Gly12Asp	p.Gly12Cys	No mut	p.Gly13Asp	No mut	p.Gly12Ser	No mut	p.Gly12Ala	p.Gly12Val	No mut		
		a	а	a	a	а	a	a	a	a	a	10,0	
		a	а	a	a	n	a	a	a	a	a	9,5	
		a	а	a	a	n	a	a	a	a	a	9,5	
		a	а	a	a	n	a	a	a	a	a	9,5	
								w (c.38G>A,					1
		a	a	a	a	а	a	p.Gly13Asp)	a	a	a	9,0	1
		a	a	a	a	а	а	а	a	a	a	10,0	
		a	a	a	a	а	а	a	a	a	a	10,0	
		a	а	a	а	а	a	а	a	a	a	10,0	
		a	а	a	a	n	a	a	a	a	a	9,5	
									w (c.35G>C, p.Gly12Ala + c.35G>A,				
		a	а	a	a	а	a	а	p.Gly12Asp)	a	а	9,0	1
		а	а	а	а	а	а	а	a	а	a	10,0	
		а	а	а	a	n	а	n	a	w (no mut)	а	8,0	
		а	а	а	a	n	a	а	а	а	а	9,5	1

			, <i>(</i>		/		, ,					
		c.35G>A,	c.34G>T,		c.38G>A,		c.34G>A,		c.35G>C,	c.35G>T,		
SCHEME E	Result	p.Gly12Asp	p.Gly12Cys	No mut	p.Gly13Asp	No mut	p.Gly12Ser	No mut	p.Gly12Ala	p.Gly12Val	No mut	
	PT0001	а	а	а	а	а	а	а	а	а	а	10,0
	ES0001	a	а	а	а	а	а	а	а	а	а	10,0
	ES0003	a	а	а	а	а	а	а	а	а	а	10,0
	IT0001	a	а	а	а	а	а	а	а	а	а	10,0
	IT0002	a	а	а	а	а	а	а	а	а	а	10,0
	1000										w (c.35G>A,	
	IT0003	a	а	а	а	а	а	а	а	а	p.Gly12Asp)	9,0
	IT0004	a	а	а	а	а	а	а	а	а	а	10,0
	IT0005	a	а	а	а	а	а	а	а	а	а	10,0
	IT0006	a	а	а	а	а	а	а	а	а	а	10,0
	1.1.1	c.182A>G,		c.34G>T,		c.38G>A,	c.35G>C,	c.183A>T,	c.35G>A,	c.34G>A,	c.35G>T,	
SCHEME F	Result	p.GIn61Arg	No mut	p.Gly12Cys	No mut	p.Gly13Asp	p.Gly12Ala	p.GIn61His	p.Gly12Asp	p.Gly12Ser	p.Gly12Val	
	FR0001	b	а	а	а	а	а	b	а	а	а	10,0
	FR0002	a	а	а	а	а	а	а	а	а	а	10,0
	FR0003	b	а	а	а	а	а	b	а	а	а	10,0
	FR0004	b	а	а	а	а	а	b	а	а	а	10,0
	FR0005	b	а	а	а	а	а	b	а	а	а	10,0
	FR0006	b	а	а	а	а	а	b	а	а	а	10,0
	FR0007	b	а	а	а	а	а	b	а	а	а	10,0
	FR0008	b	а	а	а	а	а	b	а	а	а	10,0
	FR0009	b	а	а	а	а	а	b	а	а	а	10,0
	FR0010	b	а	а	а	а	а	b	а	а	а	10,0
	FR0011	b	а	а	а	а	а	b	а	а	а	10,0
	FR0012	а	а	а	а	а	а	а	а	а	а	10.0



European external quality assessment for the improvement of KRAS testing

- Bellon E.¹, Ligtenberg M.J.L.², de Hertogh G.³, de Stricker K.³, Edsjö A.³Laurent-Puig P.³, Machado JC.³, Rouleau E.³, vandenberghe P.³, vander Borght S.³, van Krieken J.² and Dequeker E.¹
- ¹ University of Leuven, Centre for Human Genetics, Biomedical Quality Assurance Research Unit Leuven, Belgium . ² Department of Pathology, Radboud University Nijmegen Medical Centre, The Netherlands. ³ Regional scheme organizers of the 2010 ESP KRAS Regional EQA scheme (alphabetically).
- In total 76 laboratories from 14 different European countries participated.
- From the 76 participating laboratories, 50 reported all 10 genotypes correctly (66%).



http://kras.eqascheme.org

Agenda invitational conference on quality assurance in molecular pathology June 9 2011, Office of the ESP, Rue Bara 6, Brussels

12.00 informal, sandwiches 13.00 Han van Krieken 13.10 Els Degueker 13.20 Marjolein Ligtenberg 13.30 Frik Thunnissen 13.40 Simon Patton 13.50 Frank Opdam 14.00 Hans Kreipe 14.10 Andreas Jung 14.20 Etienne Rouleau 14.30 Sandi Deans 14.40 Nicola Normanno 14.50 Ed Schuuring 15.00 15.30 Peter Collins 15.40 Sabine Tejpar 15.40 Nicola Normanno 15.50 Scott Patterson 16.00 Ivonne Marondel 16.10 discussion on statements

Welcome, what is the ESP, goal of the conference The ESP KRAS EQA program: organization and results The ESP KRAS EQA program: practical aspects EQA for EGFR testing in lung cancer The vision of the European Thoracic Oncology Platform Development and potential of artificial samples The German approach to EQA for molecular pathology Experiences from the German EQA The French approach to EQA for molecular pathology The NEQAS approach to EQA for molecular pathology The Italian approach to EQA for molecular pathology The Dutch approach to EQA for molecular pathology Tea break Test regulatory approval and registration The view of the clinician The approach in Italy and the view of ESMO The view of industry: Amgen The view of industry: Pfizer

IPATIMUP

(FOUR EXAMPLES OF BIOMARKERS ROUTINELY TESTED)

- EGFR mutations and EML4/ALK translocations in lung carcinoma
- KRAS mutations in colorectal carcinoma
- KIT/PDGFRA mutations in GIST tumours
- HER-2 amplification in breast carcinoma

DNA extraction from paraffin-embedded material (Biopsy and surgical specimens) Crucial role of the pathologist

KRAS mutations in colorectal carcinoma

KRAS mutations	Resi	ults - IPA	TIMUP	Results from literature		
	N	%	Rank	%	Rank	
12Asp	169	37%	1	31-47	1	
12Val	125	28%	2	22-29	2	
13 <i>As</i> p	73	16%	3	14-21	3	
12Cys	43	9%	4	6-9	4	
12Ser	20	4%	5	4-6	5	
12Ala	17	4%	6	3-6	6	
12Arg	4	1%	7	1-2	7	
13Cys	3	1%	8	0,5-1	8	
1 mutation	454					
>1 mutation	9					
Total (mutations)	463	38%			30-50%	
Wild-type	763	62%				
Total (cases)	1226					



EGFR mutations	Results - IPATIMUP		Results from literature
	Ν	%	%
Exon 18	3	7%	5-10
Exon 19	17	41%	45-50
Exon 20	7	17%	5-14
Exon 21	14	34%	20-45
otal (mutations)	41	12%	10-20
Wild-type	295	88%	
Total (cases)	336		

KIT/PDGFRA mutations in GIST tumours

GIST mutations	Results	- IPATIMUP	Results from literature			
	Ν	%	%			
KIT						
Exon 9	4	14%	10%			
Exon 11	12	43%	70%			
Exon 13	3	11%	1%			
Exon 17	1	4%	1%			
PDGFRA						
Exon 12	1	4%	1%			
Exon 14	1	4%	1%			
Exon 18	6	21%	6%			
Total (mutations)	28	57%	90%			
Wild-type	21	43%	10%			
Total (cases)	49					

TAKE HOME LESSONS

- Quality control is a must-have! The crucial words are competence and harmonization.
- Oncologists, Pathologists and Geneticists have to be perfectly sychronyzed for patients to take full advantage from available technology!
- The involvement of Pathologists in these ventures will increase the **quality and the public awareness** (clinicians and patients) **of our profession**.

Thanks for your attention

25th European Congress of Pathology



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sociedade Portuguesa de Anatomia Patológica

First week of September 2013 Lisbon, Portugal