

***BRCA Tumour Testing
Masterclass – Hadassah
Medical Center
January 29th, 2018***

***Institute for Pathology
Center for Integrated Oncology
CIO – Cologne
Sabine Merkelbach, Carina
Heydt, Reinhard Buettner***



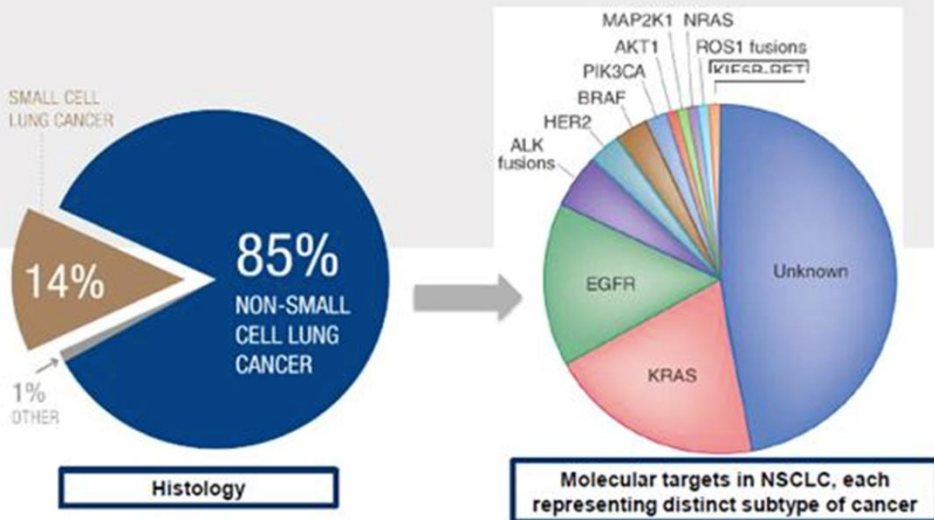
welcome!

Agenda

- 08:30-09:00 **Gathering & Breakfast**
- 09:00-09:30 **Introduction for sBRCA testing** - Prof. Reinhard Büttner
- 09:30-10:15 **Review of methodologies for tumour testing; highlighting experience from University Hospital, Cologne** - Dr. Carina Heydt
- 10:15-10:45 **gBRCA testing - classification of genomic variations and clinical implications**
Prof. Vardilella Meiner - Director of Center for Clinical Genetics
- 10:45-11:15 **Coffee break**
- 11:15-12:45 **Determining Pathogenicity & Interpretation of Results with the aid of the challenging samples from Cologne**
Group A - Dr. Sabine Merkelbach – Bruse
Group B - Dr. Carina Heydt
- 12:45-13:15 **Challenging case session from ISRAEL**
Dr. Yaniv Zohar - (Rambam medical center), Dr. Ariel Erental - (Hadassah medical center) and panel members
- 13:15- 14:15 **Lunch**
- 14:15- 14:45 **The Future: Panel testing** - Prof. Reinhard Büttner
- 14:45- 15:15 **Summary of key processes for tumour BRCA testing Q&A session**
Dr. Sabine Merkelbach – Bruse, Prof. Reinhard Büttner,
Dr. Carina Heydt & Prof. Eli Pikarsky (Hadassah medical center)
- 15:15- 16:00 **Meeting close & on demand private discussion on sBRCA testing**

Molecular Classification of Cancer

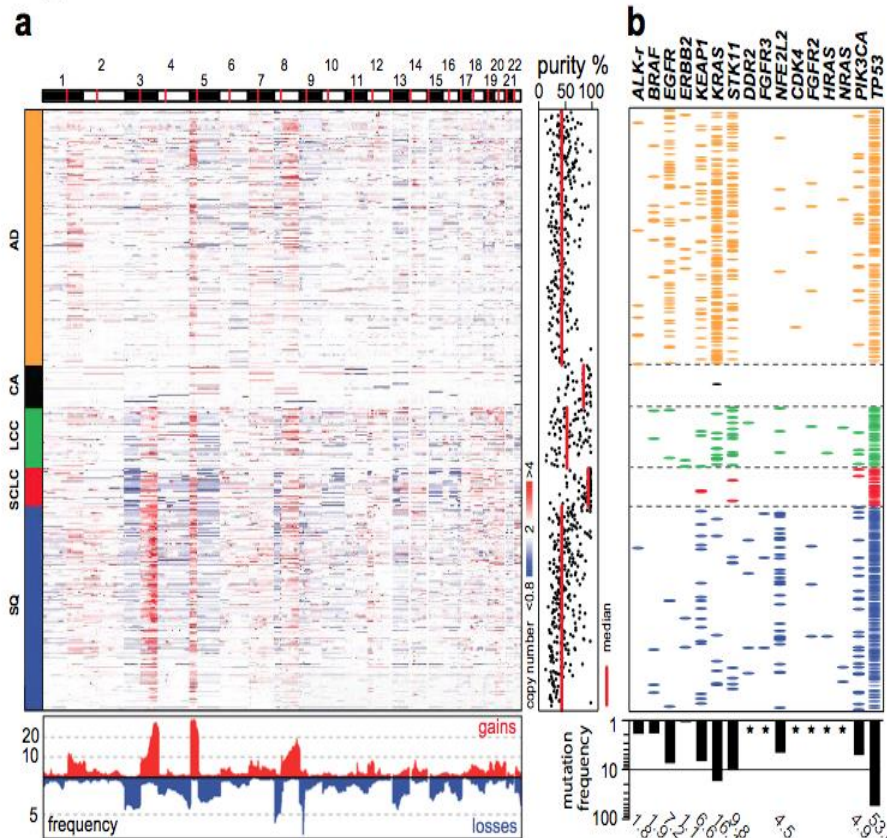
Lung cancer: from histological to molecular classification



Pao & Hutchinson Nature Medicine 2012

Pao W, Nature Medicine 2012

Figure 1



Seidel D, Wolf J, Buettner R, Thomas RK:
Science Transl Med Oct30th, 2013

~5,000 Lung Cancer Genomes connected to clinical data (NGM-L)

The Network Genomic Medicine NGM A Platform for Molecular Diagnostics and Precision Medicine www.ngm-cancer.com

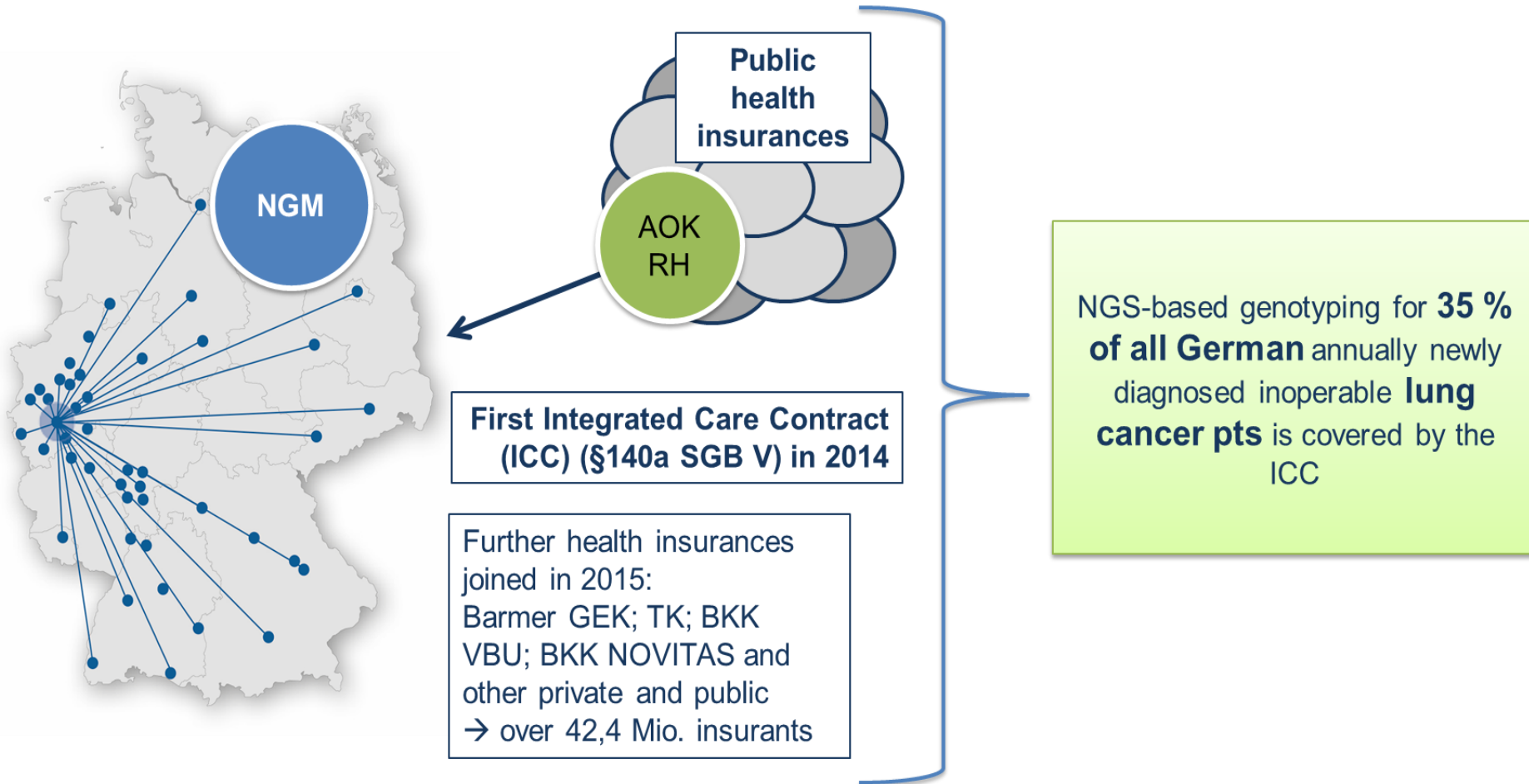
Onkologisches Spitzenzentrum
seit 2008



The Center for Integrated Oncology – CIO Köln Bonn

Reimbursement: Integrated Care Contract (ICC)

The Network Genomic Medicine



NGS-based genotyping for **35 % of all German** annually newly diagnosed inoperable **lung cancer pts** is covered by the ICC

Gene panels: Multiplex PCR versus Hybrid Capture

Multiplex PCR panel

NGS_LUN4-QIAG_#39

Gen	Exon
ARAF	7, 10, 15
BRAF	11, 15
CTNNB1	3
DDR2	4 - 19
EGFR	18 - 21

NGS_GCGC3-QIAG_#33

Gen	Exon	Codon
BRAF	15	582 - 612
DDR2	4 - 19	1 - 856
ERBB2/HER2	8, 19 - 21	302 - 338, 737 - 883
HRAS	2 - 4	1 - 93, 98 - 150
KEAP1	2 - 6	1 - 625
KRAS	2 - 4	1 - 73, 98 - 150
NFE2L2	2	16 - 104
NRAS	2 - 4	1 - 37, 41 - 92, 98 - 150
PIK3CA	1, 4, 7, 9, 20	58 - 117, 304 - 353, 418 - 468, 514 - 554, 980 - 1069
PTEN	1 - 9	1 - 404
RHOA	2, 3	1 - 92
TP53	5 - 9	126 - 331

NGS_MEL2QIAG_#

Gen	Exon
BRAF	11, 15
CDK4	2
CDKN2A	1, 2
GNA11	5
GNAQ	5
HRAS	2, 3, 4
IDH1	4
KIT	9, 11, 13, 17, 18
KNSTRN	1
KRAS	2 - 4
NRAS	2 - 4
OXA1L	1
PDGFRA	12, 14, 18
PIK3CA	9, 20
PTEN	1 - 7
RAC1	2
TP53	5 - 9

NGS_PROS-QIAG_#49

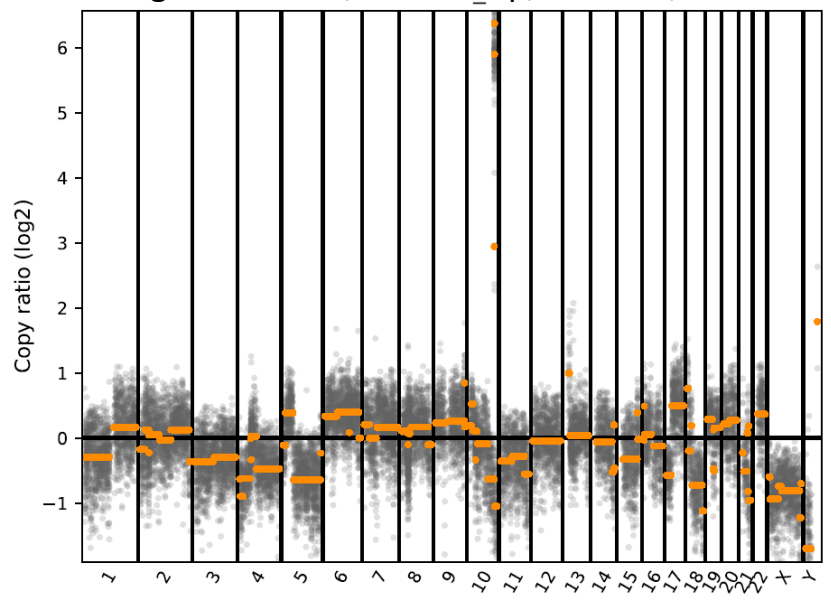
Gen	Exon
AR	1 - 8
ATM	2 - 63
CTNNB1	3
PIK3CA	1, 4, 7, 9, 20
PTEN	1 - 9
RAD51C	1 - 9
TP53	5 - 8

Hybrid Capture CAIO Panel 92 Gene und 14 MSI Marker

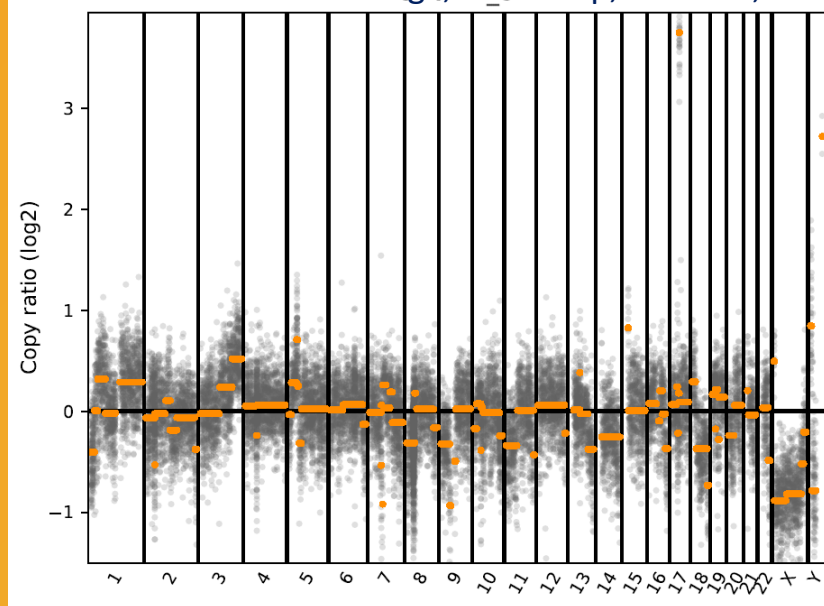
Gene	Target	Gene	Target	Gene	Target
ABL1	Exons	IDH1	Exons	RAC1	Exon 2
ALK	Breakpoints and Exons	IDH2	Exons	RAD50	Exons
APC	Exons	IGF2R	Exons	RAD51C	Exons
AR	Exons	JAK2	Exons	RB1	Exons
ARAF	Exons	KDR	Exons	RET	Breakpoints and Exons
ATM	Exons	KEAP1	Exons	RHOA	Exon 2,3
ATR	Exons	KIF5B	Breakpoints	RICTOR	Exons
BCL6	Exons	KIT	Exons	ROS1	Breakpoints and Exons
BRAF	Breakpoints and Exons	KNSTRN	Exon 1	RPTOR	Exons
BRCA1	Exons	KRAS	Exons	SMO	Exons
BRCA2	Exons	MAP2K1	Exon 2	STK11	Exons
CCND1	Exons	MET	Breakpoints and Exons	TERT	Exons
CCNE1	Exons	MSH3	Exons	TGFB2	Exons
CD74	Breakpoints	MTOR	Exons	TMPRSS2	Breakpoints
CDK4	Exons	MYC	Exons	TP53	Exons
CDK6	Exons	MYCL1	Exons	TSC1	Exons
CDKN2A	Exons	MYCN	Exons	TSC2	Exons
CDKN2B	Exons	NF1	Exons	VHL	Exons
CTNNB1	Exons	NF2	Exons		
EGFR	Exons	NFE2L2	Exons		
EML4	Breakpoints	NOTCH1	Exons		
ERBB2	Exons	NOTCH2	Exons		
ERG	Breakpoints	NOTCH3	Exons		
FGFR1	Whole Gene	NR3C1	Exons		
FGFR2	Breakpoints and Exons	NRAS	Exons		
FGFR3	Whole Gene	NRG1	Whole Gene		
FLT1	Exons	NTRK1	Breakpoints and Exons		
FLT4	Exons	NTRK2	Breakpoints and Exons		
GNA11	Exons	NTRK3	Breakpoints and Exons		
GNA13	Exons	OXA1L	Exon 1		
GNAI2	Exons	PDGFRA	Breakpoints and Exons		
GNAQ	Exons	PDGFRb	Breakpoints and Exons		
GNAS	Exons	PIK3CA	Exons		
GNAT2	Exons	POLD1	Exons		
GNG2	Exons	POLE3	Exons		
HDAC2	Exons	PTCH1	Exons		
HRAS	Exons	PTEN	Exons		

Amplificon Detection via Hybrid Capture Enrichment

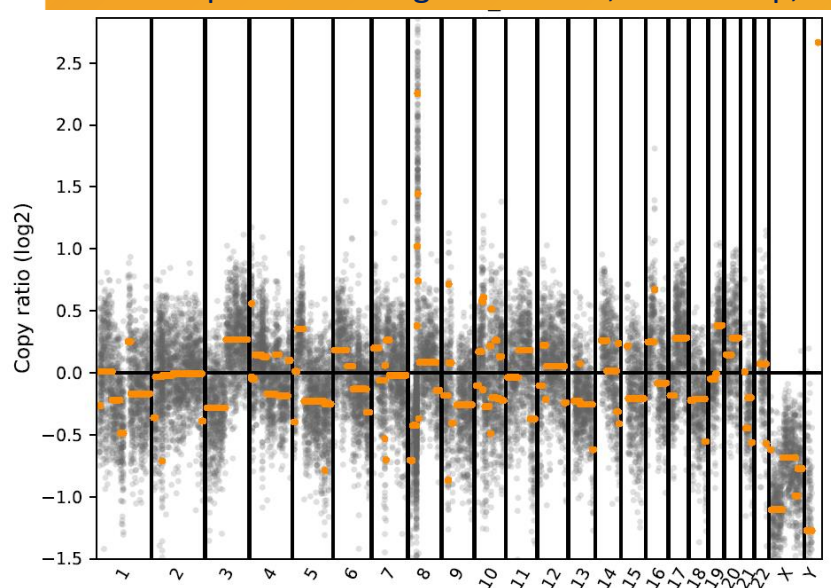
Magenkarzinom, FGFR2amp, Ratio 45,5



Adenokarzinom Lunge, HER2amp, Ratio 11,55



adenosquamöses Lungenkarzinom, FGFR1amp, Ratio 3,67

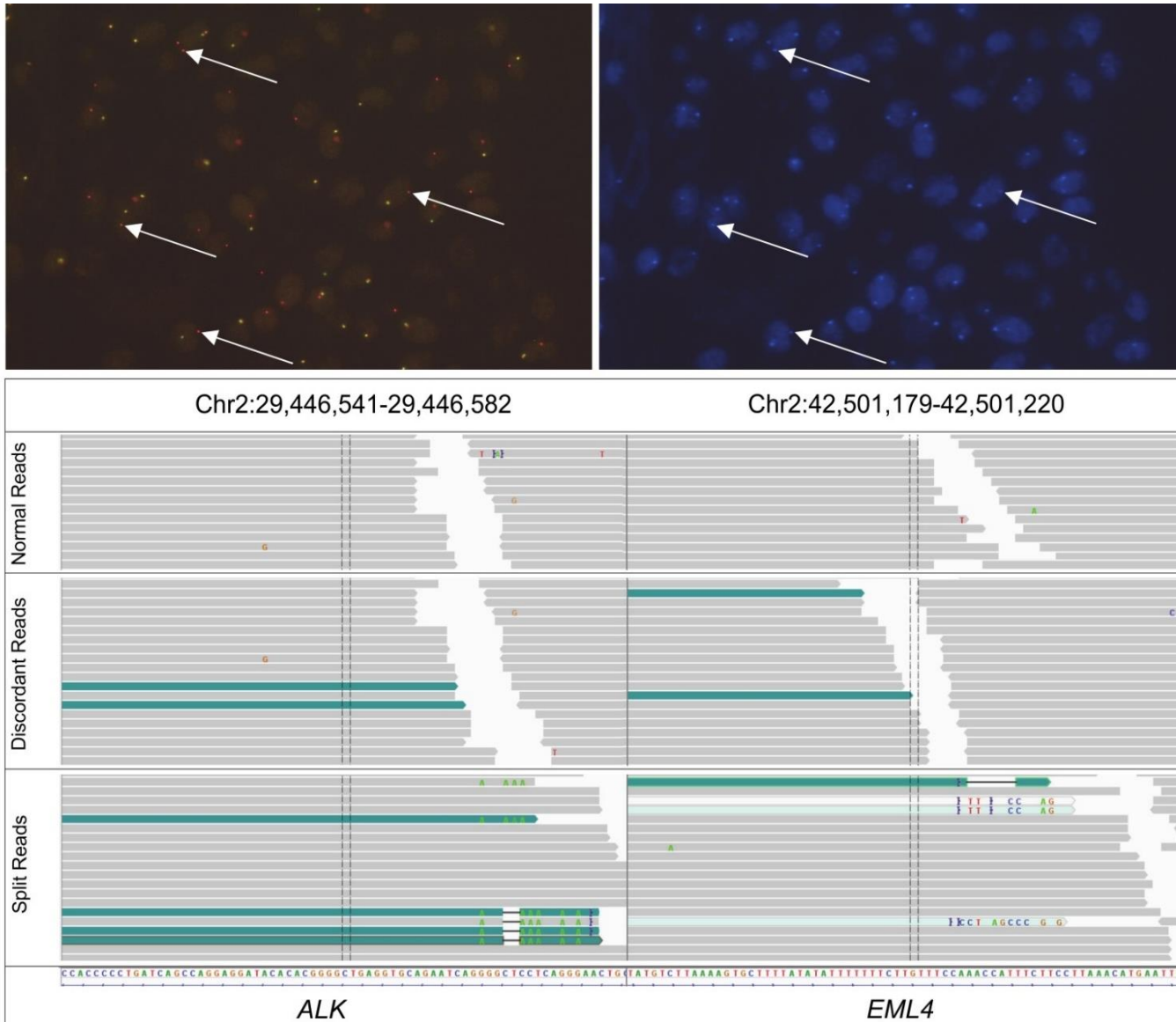


- analysed by CNVkit:

<https://cnvkit.readthedocs.io/en/stable/>

- Reference: WT DNA from FFPE

Detection of Gene Fusions by Hybrid-Capture



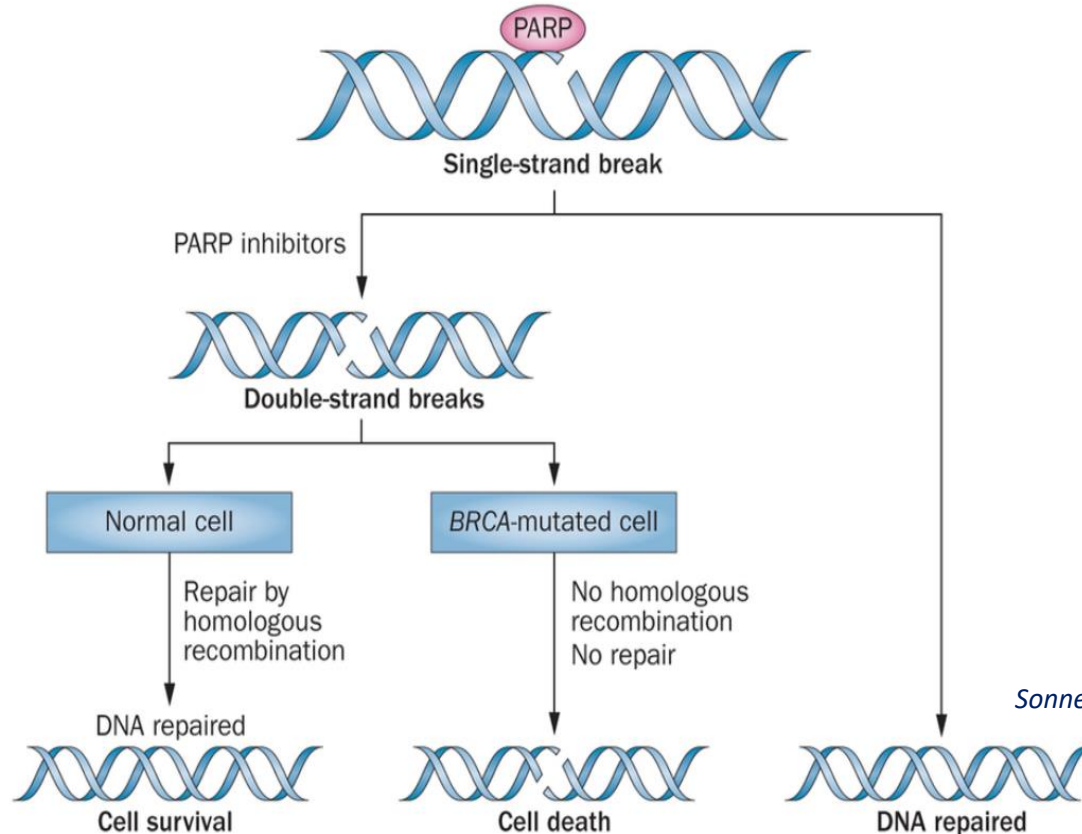


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CIO-KölnBonn

BRCA-Diagnostics in Ovarian Carcinomas

- 2015: Approval of the first PARP-inhibitor (Olaparib) for *BRCA*-mutant, relapsed, Platinum-sensitive, serous HG carcinomas of ovary, Fallopian tube or peritoneum.
- in ~25-40% of *BRCA*-mutant ovarian carcinomas there is a somatic mutation (*Cancer Genome Atlas Research, Nature 2011; Hennessy BT et al., J Clin Oncol 2010*)

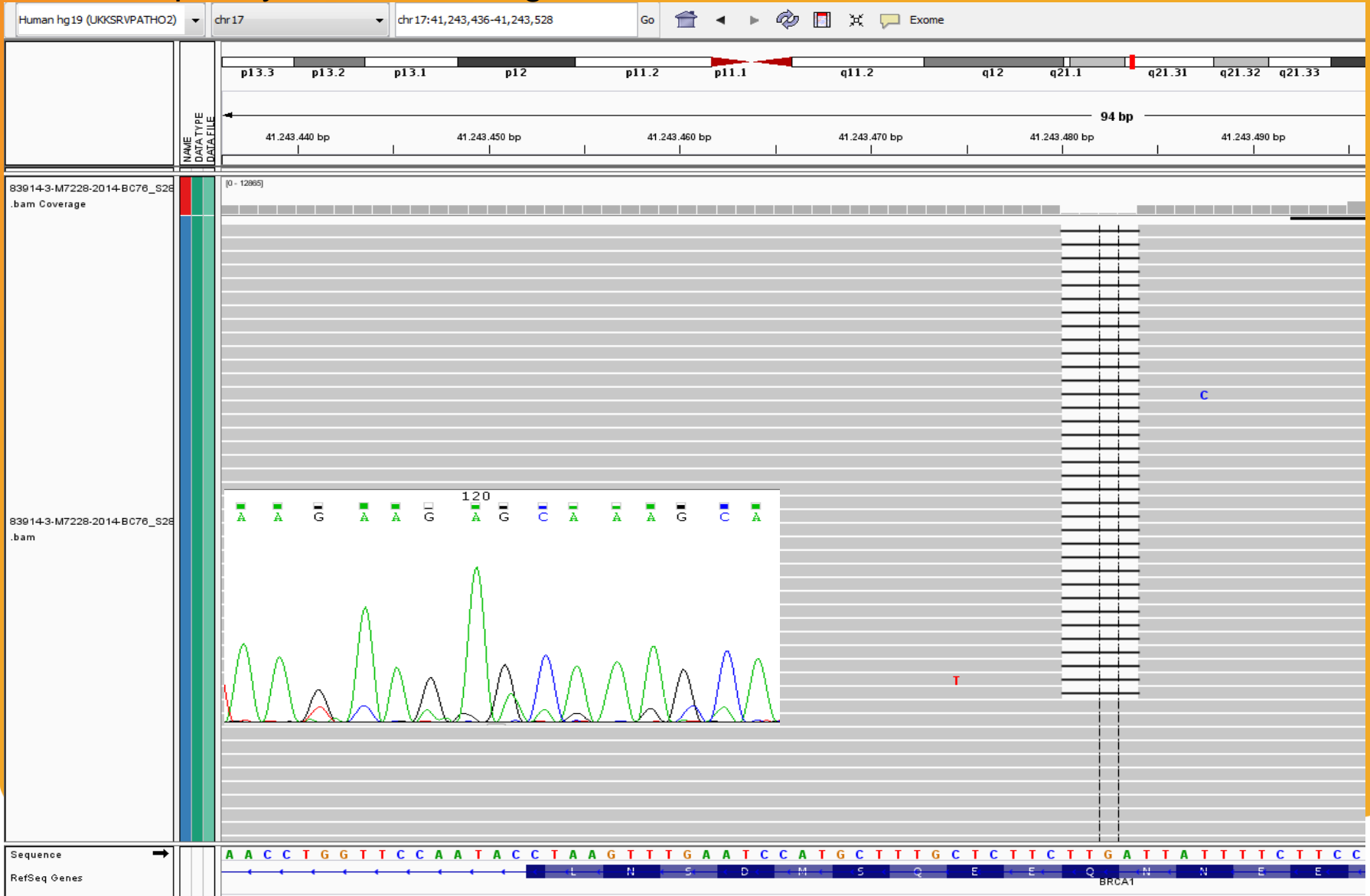
Mechanism of PARP-inhibition in *BRCA*-mutant tumors :



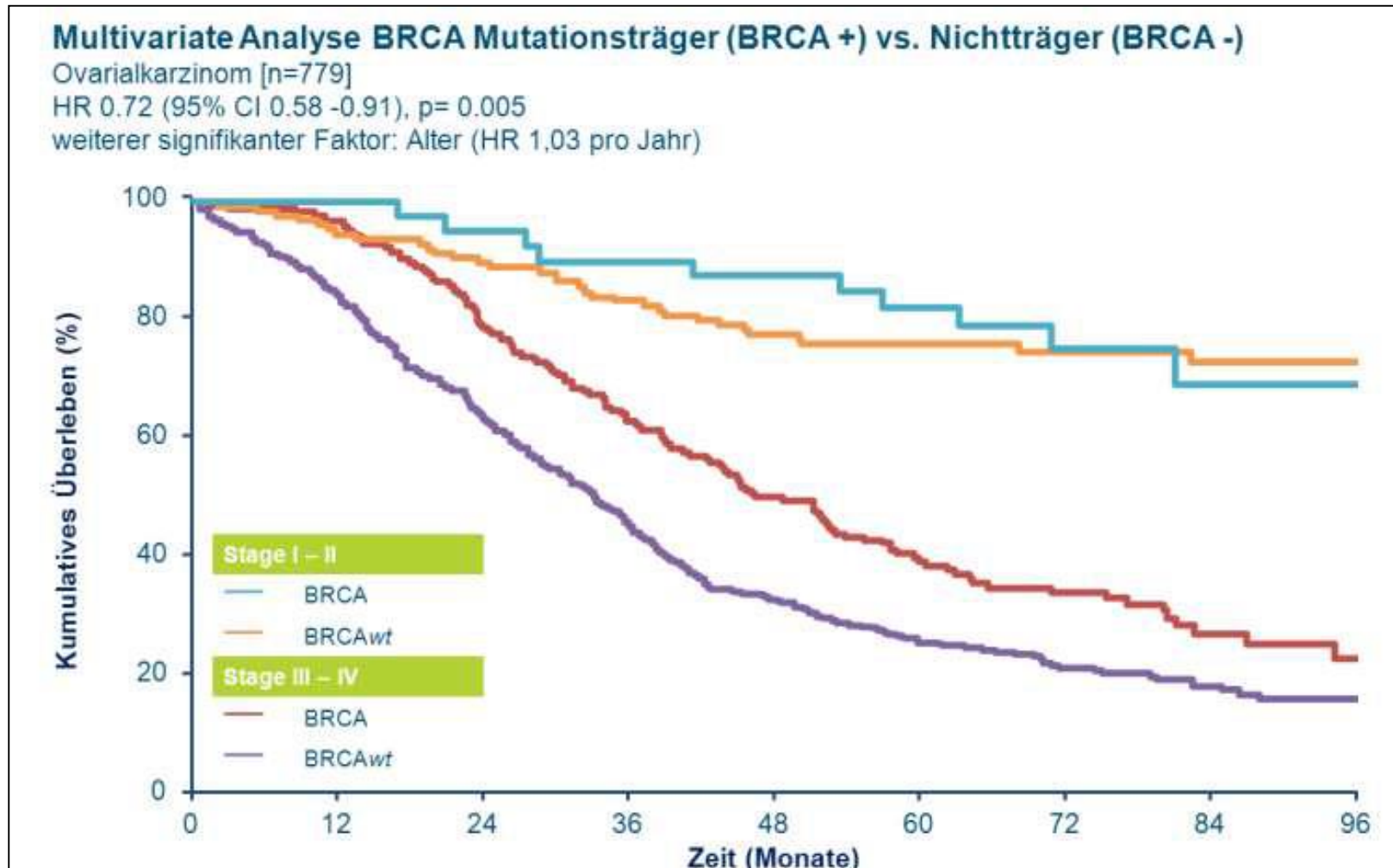
BRCA1 Mutation in Ovarian Cancer (Run 5): Sample 7228_14

Exon 11: **c.4065_4068del** leads to frameshift mutation p.N1355Kfs*10

Allele frequency: 82.32%, Sanger validated

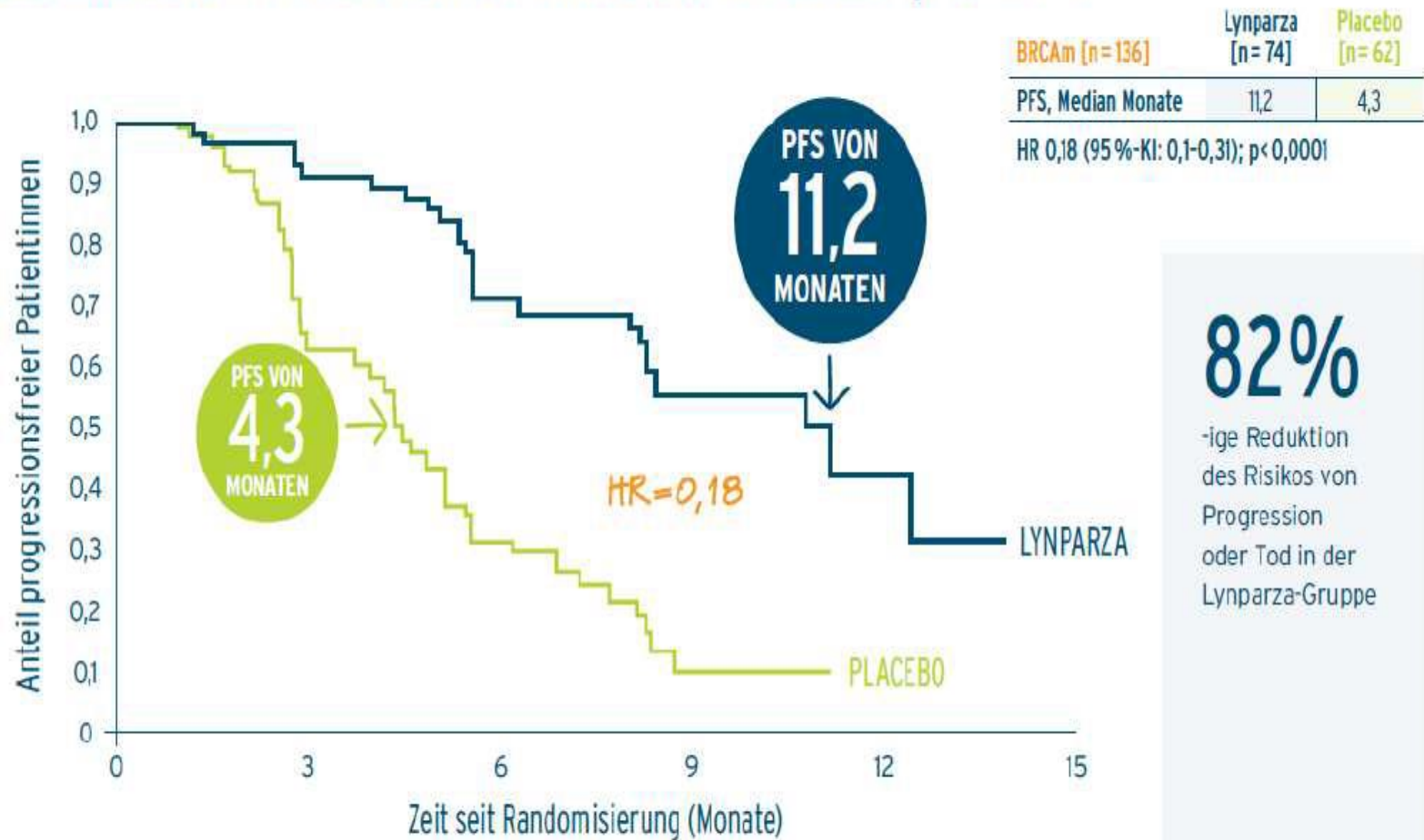


BRCA mutations influence OS of patients with ovarian cancer



Olaparib perlongs PFS in stage IV ovarian cancer patients with BRCA mutations

Ovarialkarzinom-Patientinnen mit BRCA Mutation profitieren



Number at risk

Lynparza BRCAm	74	59	34	15	5	0
Placebo BRCAm	62	35	13	2	0	0

Results of BRCA testing in Cologne, first 21 months:

Cases	N =	%
Total	668	100,0
n.a.	26	3,9
Wild-type	474	72,4
Mutant	168	25,6
all Categories 4/5 (based on databases)	114	17,4
Categories 4/5	80	12,2
Categories 5 (truncating muts))	34	5,2
Category 3	51	7,8

Stand 19.10.2017

Datenbanken (DB):

Kategorie 1: neutral

UMD-DB

Kategorie 2: wahrscheinlich neutral

ARUP-DB

Kategorie 3: unklare Signifikanz

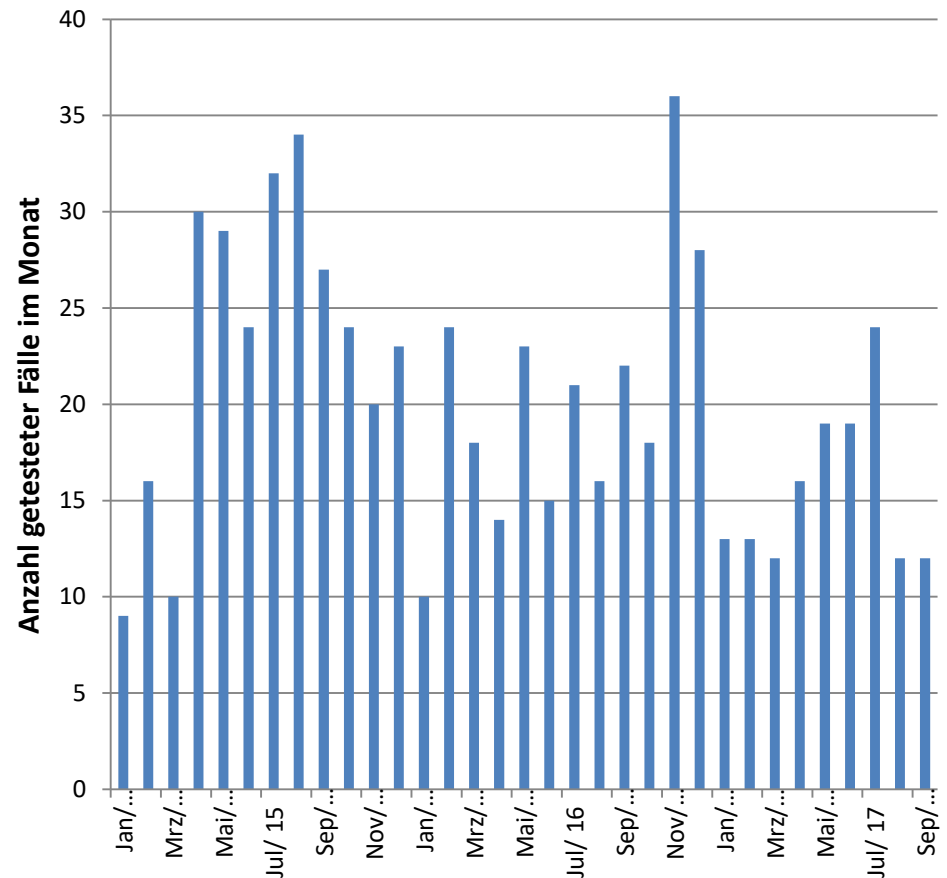
LOVD-DB

Kategorie 4: wahrscheinlich pathogen

Hogrefe-ClinVar-DB

Kategorie 5: pathogen

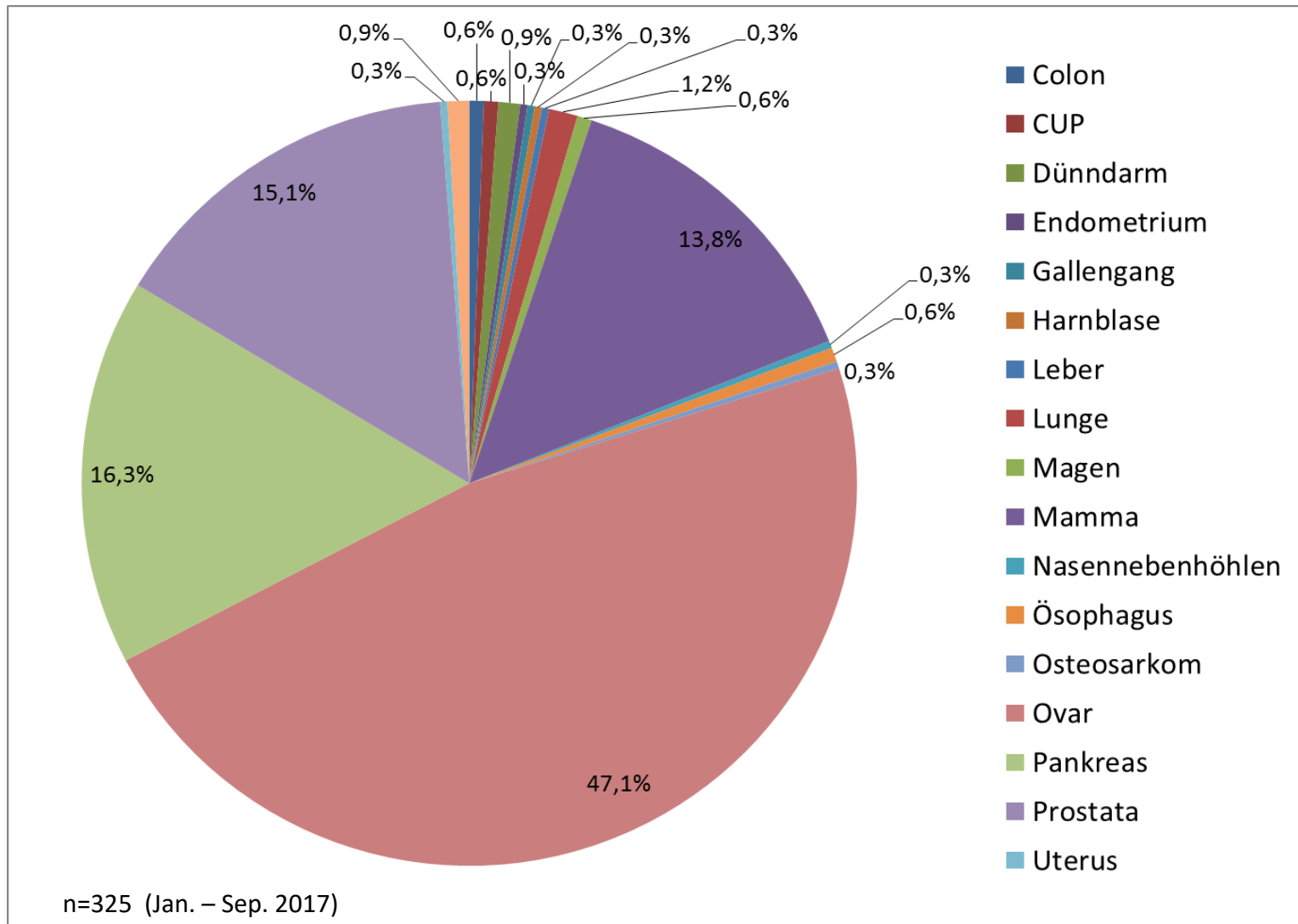
BRCA Exchange



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BRCA-diagnostics at Cologne Inst. Pathology (2017)

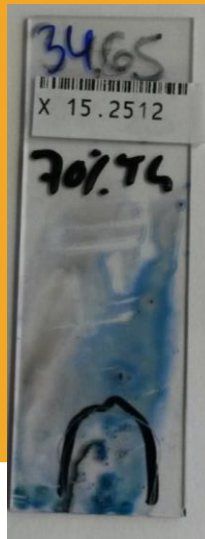
Entities:



What materials are being tested ?



- › FFPE blocks
- › Cytology specimens
- › Plasma / Blood

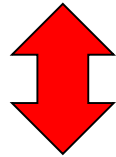


Information – Not to be shared outside of AZ.

Intention Therapy

Gyn Onkology

Platin-sensitive OvCA, stage III-IV



Institute for Pathology
Tumor Board
CIO-Oncology



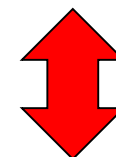
Tumorbezogene Biomarkertesting
Molekulare Pathologie
Klinisches Tumorboard

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Intention Genetic Consulting

Primary tumor, Fam+

Risc-Surveillance-Prevention



National Center for Familial Tumors
NCFT Board

Surveillance Plan
Preventive Surgery
Family screening



Keimbahntestung
Genetische Beratung

id.BRCA