

### European User Group Meeting OncoDEEP® Kit

21<sup>st</sup> - 22<sup>nd</sup> October 2024

Documenting the value of CGP a Belgium Society of Medical Oncologists (BSMO) study in collaboration with Belgian NGS Laboratories: The BALLETT Study



#### Brigitte Maes, MD, PhD





On behalf of the BALLETT investigators, the NGS lab consortium and the nMTB

## **Disclosure information**

**Illumina** is supporting the BALLETT study by providing TSO500 kits and logistical assistance; B.Maes has been an invited speaker for Illumina

OncoDNA is providing the OncoKDM sequencing data analysis software

Velsera is providing the CGW sequencing data analysis software

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Funded by the European Union

Funded by the European Union. Views and opinions expressed are however those of the author(s) only and do not necessarily reflect those of the European Union or <u>HaDEA</u>. Neither the European Union nor the granting authority can be held responsible for them. Two parallel precision oncology projects in Belgium







**Belgian Society of Medical Oncologists** 







## Belgian Approach for Local Laboratory Extensive Tumor Testing

## BALLETT: the concept



Collaboration of 9 Belgian NGS labs
 ✓ Fully standardized CGP testing
 ✓ Pooling of samples to reduce TAT
 ✓ Exchange of Expertise

Broad access to CGP in Belgium

Matched treatment recommendation

national Molecular Tumor Board (nMTB)

- ✓ Multidisciplinary nationwide virtual
- ✓ Personalized treatment selection
- ✓ Standardized reporting

Awareness and knowledge of clinicians





# Main objectives

To organise broad *access* to 'Comprehensive Genomic Profiling' for patients with metastatic cancer in Belgium

To evaluate the *clinical value* of CGP in "real-world" practice for offering more therapeutic options to patients

To support decision for *reimbursement* of CGP





# Study design

## Identification of study participants

 Investigator and local APO lab check availability of tissue biopsy (< 2 years old)</li>

#### Comprehensive Genomic Profiling

Performed by the BALLETT consortium of 9 NGS labs
Fully standardized

CGP guided therapy recommendation by national

**Molecular Tumor Board** 

#### Collection of subsequent treatment info

- nMTB recommendation followed or not?
- PFS-ratio

#### Pre-MTB laboratory meeting

- Classification of variants
- Preparation of nMTB





## **Comprehensive Genomic Profiling**









<ul> <li>BALLETT</li> <li>Dashboard</li> <li>Per patient data</li> <li>ZNA05</li> </ul>	Gender: Fema Tumortype: E Diagnosis dat 1 metastatic : Liver metasta Bone metasta Brain metasta Other metasta	ale , Age: 23 Grain Glioma e: 31-07-2019 Sites at inclusion asis: No asis: No asis: No asis: No asis: No		Number of previous treatme Including: Radio-chemother Most recent: Chemotherapy Period: 06-04-2022 to 16-02 Best response: Stable Disea	ent lines: 1 rapy 7, Cisplatin, vincristine, lomustine -2021 se		
Overview	PATIENT INFO CGP RESULTS DR	UGS MTB					
	INCLUSION INFO Physician: Dr. Signed ICF version n°: 4 ICF signature date: 27-03-2022	DEMOGRAPHICS Gender: Female Age: 23 ECOG at inclusion: 0 Smoking status: No	CGP SAMPLE Lab reference Sampling dat Metastatic sit Tumor cell %	E INFO e of FFPE block: 22B264761 te: 17-02-2022 te, Lymph node : 80	PATHOLOGY Tumortype: Brain Glioma Subtype: Other Diagnosis date: 31-07-2019 1 metastatic sites at inclusion Liver metastasis: No Lung metastasis: No Bone metastasis: No Brain metastasis: No Other metastasis: lymph nodes	IHC/ISH MARKERS Markers HER2 ER PR ALK	MARKERS Presence Not available Not available Not available Not available
DESIGNED BY	TREATMENT HISTORY Medulloblastoma 05/11/2019 untill 17/12/2019: adjuvante o craniospinale as, gevolgd door boostbest gecombineerd met wekelijkse perifere too 6/4/2022: start Cisplatin, vincristine, lom	hemoradiotherapie (36 Gy in 20 fract raling van 18 Gy in 10 fracties op het r ediening van vincristin 2mg) ustine	ies op de resectiegebied;			ROS1 pan-TRK HPV MMR PDL1	Not available Not available Not available Not available >? No PDL1 IHC lot applicable

LETT ent data	Gende Tumor Diagno 3 meta Liver r Lung r Bone r Brain Other	r: Female , Age: 41 type: Cholangiocarcino osis date: 06-08-2021 Istatic sites at inclusion netastasis: Yes netastasis: Yes netastasis: No metastasis: /	oma	0	Number of prev Including: Cher Most recent: Ch Period: 23-08-2 Best response:	rious treatment lines notherapy emotherapy , cispla 021 to ? ?	:: 1 tinum-gemcitabi	ine		
CGP RES	SULTS PATIENT INFO	МТВ								
	FUSION GENES			в	MSI	CANCER	PREDISPOSING GEN	E	Sampling dat	e: 06-08-2021
sions	Fusion/Splice variant	🕴 Clinical class* 🌲		ns riants/Mb	Negative	Germli	ne variant like	ly? Yes	Primary tumo Tumor cell %:	90
2	FGFR2-KIAA1598	TIER IA					VARIANTS			
Q	AMPLIFICATION		Gene 🔶	HGVSc	HGVSp ♦	Shortcode	<b>VAF</b> \$	Biological class	Validation \$	Clinical class*
	MDM4 2	TIER III	PTPRD	c.4187G>A	p.(Arg1396Gln)	R1396Q	94%	VUS	Consensus	TIER III
			ATM	c.6067G>A	p.(Gly2023Arg)	G2023R	50%	VUS	Consensus	TIER III
N	NUMBER OF VARIANTS PER BIC	LOGICAL CLASS	KMT2A	c.5665G>T	p.(Asp1889Tyr)	D1889Y	43%	VUS	Consensus	TIER III
		Pathogonic	SPTA1	c.6617T>C	p.(Ile2206Thr)	I2206T	31%	VUS	Consensus	TIER III
		Pathogenic	ARID1B	c.980G>C	p.(Gly327Ala)	G327A	7%	VUS	Consensus	TIER III
1		Likely pathogenic	Showing 1 to 6 o	of 6 entries					P	revious 1 Ne
5		VUS	*AMP/CAP/ASCC	/ACMG clinical class	sification system of sequenc	e variants in cancer. L	i et al, J Mol Diagr	n, 19(1):4-23 (2017) .		
0	10 20 Number of variants	30								



. BALLETT	Report preview	
🤣 Dashboard	A Download report as PDF	
Per patient data		î
ZNA05 👻		
💵 Overview		
Report	BALLETT	r -
MTB discussions		
💫 Patients by MTB date	15-04-2022	
	To Dr.	
	BALLETT patient ZNA05	
A Reload data	Dear colleague,	
	Your patient was discussed by the BALLETT laboratory working group on 15-04-2022 and during the Molecular Tumor Board on 15-04-2022.	
	Please find below the summary of the patient data, the results of the 'Comprehensive Genomic Profiling' (performed by the TSO500 kit of Illumina) as well as the therapy recommendation(s) based on the genomic	
	results. Please consider these recommendation(s) in view of all detailed patient factors, treatment history, contraindications, other therapy options and patient's preferences. The final treatment decision (including clinical trial eligibility assessment) is the full and sole responsibility of the treating physician.	
	Patient summary	
DESIGNED BY	Gender: Female Metastasis: Yes	
20553	Age: 23     1 metastatic sites at inclusion       Tumor type: Brain Glioma     Liver metastasis: No	

BALLETT	Dashboa	ard												
Dashboard	PATIENTS	TESTING	TUMORTYPES	ALL ABERRATIONS	SNP/MNP	AMPLIFICATIONS	FUSIONS/SPLICE	VARIANTS TM	B MSI	CLINICAL	SIGNIFICANCE	THERAPY R	ECOMMENDAT	ΓΙΟΝ
	GERMLINE													
Per patient data	NUMBER OF UI	NIQUE VARIANTS thogenic variants	IN GENES			VARIANTS REPORTED I	N BALLETT STUDY					PATIENTS W	ITH VARIANT	
Enter a patientID 🔹	Minimal numb	per:		Reset selection							BRCA1	c.2311_2317	del   p.(Pro77	3Leufs*)
Overview			183	now 10 🗸 entries										
Report	5 24 42 81 Click on a bar	79 98 118 135 to see the varia	153 172183 ants of	Gene 🗧 Ho	GVSc 🗘	HGVSp	Shortcode	Biological class	; n	\$	2000			Clinical
MTP discussions	the correspon	ding gene.		BRCA1 🛞 All		All	All	All	All		Patient_ID	† Tumorty	be ÷	class*
		3 TP - AF	253 — PC	BRCA1 c.1	164A>C	p.(Arg388Ser)	R388S	VUS	1		All	All		All
Patients by MTB date	33	- RE	31 FN	BRCA1 c.1	231G>A	p.(Asp411Asn)	D411N	VUS	1		AZD071	Cholangic	carcinoma	TIER IIC
	33	AF	RID1A	BRCA1 c.127	6_1279del	p.(Ser426Argfs*3)	S426Rfs*3	Pathogenic	1		UZA46	Kidney ca	ncer	TIER IIC
	26 25	- PII - NF	<3CA	BRCA1 c.1	1390del	p.(Thr464Profs*11)	T464Pfs*	Pathogenic	1		Showing 1 to 2 c	f 2 entries	Previous	1 Next
🖪 Reload data	<u>222</u> 119	- BA	VP1	BRCA1 c.1	660G>A	p.(Glu554Lys)	E554K	VUS	1					
	ig	AT	M	BRCA1 c.1	865C>T	p.(Ala622Val)	A622V	VUS	1					
	iE iE	AT	RX	BRCA1 c.190	7_1913del	p.(Cys636Tyrfs*13)	C636Yfs*13	Likely path	1					
	110 116	- BF - TE	RCA2	BRCA1 c.:	199G>A	p.(Asp67Asn)	D67N	VUS	1					
	æ	LF	P1B	BRCA1 c.2	086A>G	p.Thr696Ala	T696A	VUS	1					
	Ē	KF	RAS	BRCA1 c.231	1_2317del	p.(Pro773Leufs*)	P773Lfs*	Pathogenic	2					
	E	- EG - FB	BFR SXW7 Sh	nowing 1 to 10 of 23 entries	(filtered from 6,	,492 total entries)		Previous 1	2 3	Next				
DESIGNED BY	F	SN	ARCA4 *A	MP/CAP/ASCO/ACMG clinica	al classification	system of sequence var	iants in cancer. Li et a	al, J Mol Diagn, 19(1):	4-23 (2017) .					
	0 50 10	00 150	DH1											

BALLETT	Gender: Female , Age: 44		Number of previous	s treatment lines: 1			
hboard	Tumortype: Ovarian carcinoma Diagnosis date: 31-08-2017 1 metastatic sites at inclusion Liver metastasis: No		Including: Chemoth Most recent: Chemo Period: 04-08-2023 Best response: 7	ierapy otherapy , carboplatinum + p to ?	aclitaxel		
r patient data	Lung metastasis: No Bone metastasis: No						
H0 <b>-</b>	Brain metastasis: No Other metastasis: trachea						
ew	PATIENT INFO CGP RESULTS DRUGS MTB						
discussions	HRD Negative	4 variants/Mb	MSI Negative	CANCER PREDISPOSII	NG GENE	Sampling date Metastatic site Tumor cell %: S	: 19-07-2023 , Other 50
s by MTB date	FUSION GENES None	Show 7 🗸 entries		VARIAN	rs		
Reload data	AMPLIFICATION None	BRAF C.1405G>A	HGVSp	G469R 14%	Pathogenic	Consensus	Clinical class*
PATIENT INFO	CGP RESULTS DRUGS MTB	AKUTA CAUYEN	D (HIST 37 ASD)	HI3/I) 56%	VIS	Consensus	TIER III
		SUG	GESTED DRUGS				
Gene	Drug					link OncoKB	
BRAF	Belvarafenib, Encorafenib, Lifirafenib, Naporafenib, Sor	rafenib, Vemurafenib				BRAF	
Showing 1 to 1 of 1 e	entries					Prev	vious 1 Next
Suggested drugs are	e based on NCT-POT Drugs. NCT POT Drugs is a data package cov	rering drug-related information in the contxt of	precision oncology workflo	ows. The data is intended fo	r application in a scientific conte	ext. It must not be used for clin	ical decision-making
without decision ver	anneation by canicians with appropriate experience in precision c	sicology, here of blugs of oldub.					
ESIGNED BY	Number of variants						
18553							

### Tertiary NGS data analysis and Decision Support Systems



### **Results** Tumor types and patient characteristics (n = 813)





# Study flow





L1 L2 L3 L4 L5 L6 L7 L8 L9 Sequencing lab

### All aberrations in all cases

1465 SNVs and small indels, 579 amplification, 83 fusions



Genes with 5 or more aberrations





### Results Actionability





**83 %** Actionability

### **Results** Clinical classification of biomarkers



Li et al, J Mol Diagn 2017

### **Results** Clinical classification of biomarkers per tumor type



## **83 %** Actionability

clinical class

### Results Actionability: per tumor type



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all patients n=756

n=616

patients with actionable findings in ComPerMed

n=175

patients with actionable findings using CGP

### Results Actionability: CGP versus ComPerMed



Immunotherapy and PARPi biomarkers: TMB, MSI and HRD





### Results Treatment recommendations



### Treatment recommendations per tumor type



treatment options

Cooccurence of variants – multiple treatment recommendations per patient



Incidental findings of likely germline variants

Likely germline variant of cancer susceptibility gene (CSG) in 12 % of cases (n=90):

Advice to refer for germline genetic testing and genetic counseling



Reanalysis according to the recently updated ESMO recommendations for germline-focused analysis of tumor-only sequencing: 121 CSG variants in 111 patients (15 %) (considering all the 40 ESMO CSG genes and all tumor types) Kuzbari, Z. et al. Annals of oncology **34**, 215-227 (2023)

### Uptake of treatment recommendations per hospital







### Uptake of treatment recommendations per tumor type



23 %

### Results Turn around times



- median TAT from inclusion to the nMTB report = 29 days
- 95% of the reports available within 66 days
- median TAT differed significantly between the hospitals (range: 18 days - 45 days, p < 0.0001, Anova).</p>

# Conclusions

- Access of patients to CGP
- Consortium of NGS labs working closely together and exchanging expertise
- Standardization of CGP in Belgium
- **nMTB** is a valuable framework for close collaboration of lab people and clinicians contributing to optimal patient management
- BALLETT app
- Study data analysis shows that:
  - 83 % of cases has at least one (potentially) actionable variant and/or an immunotherapy biomarker
  - 69 % of patients received a treatment recommendation based on CGP and nMTB discussion
  - 12 % of patients received advice for germline genetic testing and counselling
  - 23 % of patients were treated according to the CGP based treatment recommendation





# Limitations and challenges

- Standardization of clinical variant classification
  - Consistency
  - Evolving evidence
- Standardization and prioritization of therapy recommendation
- Consistency in the treatment uptake across hospitals
  - recommendations for selecting patients for CGP and dealing with treatment advises
- Access to treatments Belgian DRUP-like trial desirable
- Clinical benefit yet to be determined based on the follow-up of the patients
- Limitations of the TSO500 CGP assay (HRD add-on, DRAGEN, cost, deletions, limited fusion panel)





### Study of the comparison of OncoDeep versus TSO500

Cant 2024			-	TSO500 (Illumina)	OncoDEEP (OncoDNA)
Sept. 2024	ISO500 (Illumina)	OncoDEEP (OncoDNA)			Pre-analytics
Total size	1.9 Mb	1.8 Mb	Bacommondod input	DNA: 40 ng	DNA: 40 ng
			Recommended input	RNA: 40 ng	RNA: 80 ng dried
Detection at DNA lev	vel				Library prep
	#	genes	DNA Fragmentation	Shearing	Enzymatic
SNVs and indels	523	638	Use of UMIs	Yes	No
	525	630	Normalisation	With beads	Quantification and dilution
CNV	514 ° *	614			Hybridization capture
LOH	514 <sup>°,</sup>	41	Pooling before hyb	No	Yes (8 samples)
	pan-tum	or biomarkers	# Hybridization times	Overnight + 2.5h	Overnight
MSI	Yes	Yes			Automation
TMB	Yes	Yes	Instrument	MOA STAR	STARlet
HRD	Yes <sup>&amp;,</sup> *	Yes	max # samples	96	24
<sup>&amp;</sup> using DRAGEN and	lucic		Hands-on-time	1.5 h	2 h
* as an add-on to the					Sequencing on a NextSeq550
us un uuu-on to the	ussuy		Read length	2 x 101 bp	2 x 74 bp
			#Samples per run	8; DNA + RNA	24; DNA + RNA
Detection at RNA level			Flowcell NextSeq550Dx	HO v2.5 -300 cycles	HO v2.5 -150 cycles
	# dri	ver genes			Data analysis
Fusions	55	13	Sec and tert analysis	ICI (add-on)	OncoKDM
Splice variants	3	9			
				Comir	a coon

# diagnostic DNA samples	234
DNA extraction method	
# reference DNA samples	11
Based on TSO500 results	
# SNVs and indels	674
# amplifications (FC ≥6)	31
# MSI-High (>20%)	9
# TMB-High (>16)	22
# HRD pos (GIS ≥ 42)	0
# diagnostic samples with rearrangements	67
RNA extraction method	
# reference RNA samples	8
Based on TSO500 results	
# gene fusions	172
# splice variants (AR, EGFR, MET)	20

Coming soon













### Study of the comparison of OncoDeep versus TSO500

#### Analysis of Comprehensive Genomic Profiling of Solid Tumors with a Novel Assay for Broad Analysis in Clinical Diagnostics

Guy Froyen<sup>1,2,3,\*</sup>, Pieter-Jan Volders<sup>1,3,4</sup>, Ellen Geerdens<sup>1</sup>, Severine Berden<sup>1</sup>, Joni Van der Meulen<sup>5,6,7</sup>, Aaron De Cock<sup>5</sup>, Stefanie Vermeire<sup>8</sup>, Jacques Van Huysse<sup>8</sup>, Marie de Barsy<sup>9</sup>, Gabriela Beniuga<sup>9</sup>, Wendy W. J. de Leng<sup>10</sup>, Anne M. L. Jansen<sup>10</sup>, Imke Demers<sup>11</sup>, Zeliha Ozgur<sup>12</sup>, Hendrikus Jan Dubbink<sup>13</sup>, Ernst-Jan M. Speel<sup>11,13,\$</sup>, Wilfred F.J. van IJcken<sup>12,\$</sup> and Brigitte Maes<sup>1,2,3</sup>

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Submitted

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A Study to Examine the Clinical Value of Comprehensive Genomic Profiling Performed by Belgian NGS Laboratories: a Belgian Precision Study of the BSMO in Collaboration With the Cancer Centre - Full Text View - ClinicalTrials.gov: NCT05058937

UZ Gent