Molecular profiling-selected treatment in metastatic cancer: can we change our standard of practice

#### Dr. Ofer Purim Chief Gastrointestinal Malignancy Service Assuta Samson Hospital

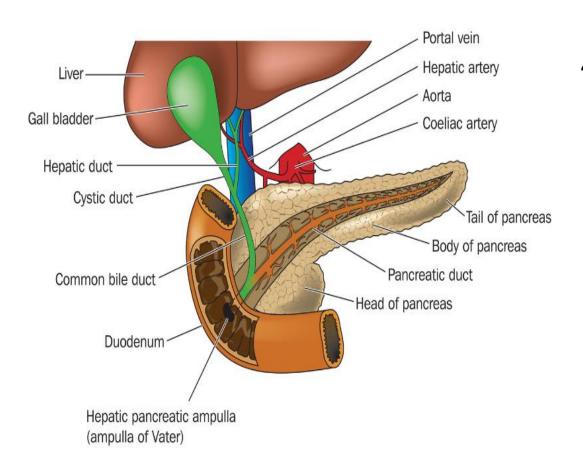
#### **Disclosure- Caris consultant**

## **Gastric & Esophageal cancer**

- Gastric cancer is the fifth most common malignancy and the third leading cause of cancer death worldwide.
- Esophageal cancer is the eighth most common malignancy and the sixth leading cause of cancer death worldwide. In 2012, there were approximately, 1.4 million new cases of gastric/esophageal cancers and 1.1 million deaths from these cancer types.

• At present, there is no internationallyaccepted consensus regarding standard-ofcare in the 1<sup>st</sup>-line metastatic setting in gastric/esophageal cancer, except for 10-25% of patients with gastric/gastroesophageal junction (GEJ) cancer whose tumors are human epidermal growth factor receptor 2 (HER2)-positive

## Pancreatic Cancer: Incidence & Mortality



- Epidemiology •
- 4th most common cause of cancer death
  - In 2015, 48,960 new cases are expected in the US, with 40,560 deaths
- Incidence much higher after 45 yrs of age

## **FOLFIRINOX vs Gemcitabine: Efficacy**

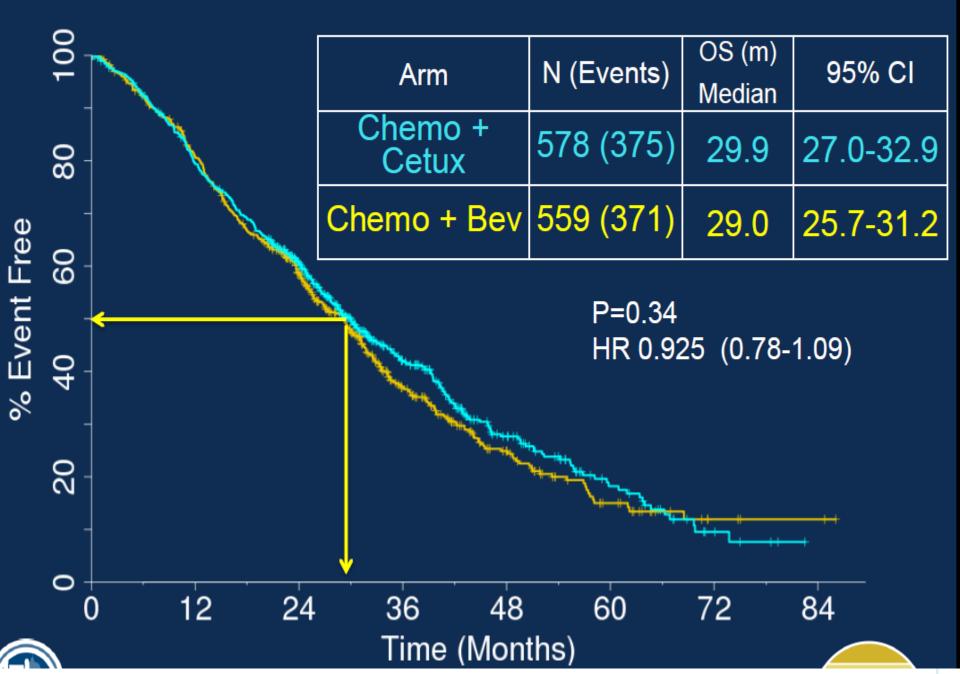
Outcome	FOLFIRINOX (n = 171)	Gemcitabine (n = 171)	
ORR, %	31.6	9.4	
Median PFS, mos	6.4	3.3	
Median survival,* mos	11.1	6.8	
1-yr survival, %	48.4	20.6	

#### **Colorectal Cancer (CRC) - Epidemiology**



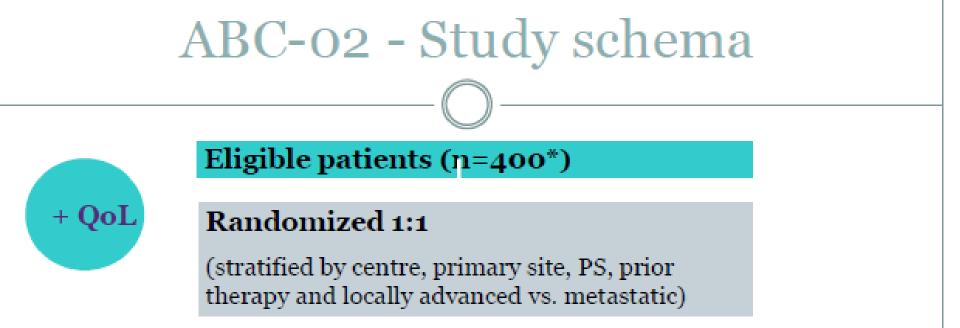
- Incidence worldwide 950,000 / yr
- Mortality: 500,000 / yr
- Cause of death in
   2 2.5% of population
- Median OS with metastasis without treatment 6 months

#### CALGB/SWOG 80405: Overall Survival



#### **CHOLANGIOCARCINOMA**

- >7,000 cases annually in 2009
- 5-year survival is <15%</li>
- Most patients present with locally advanced or metastatic disease
- Treatment commonly administered in the community, at low-volume centers.



#### Arm A

Gem 1000 mg/m² D1,8,15 q 28d, 24 weeks (6 cycles)

#### Arm B

Cisplatin 25 mg/m≧

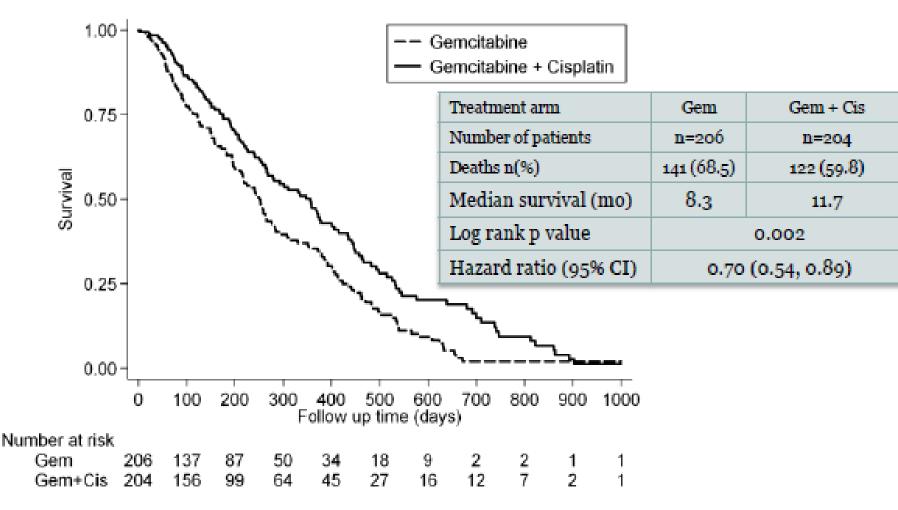
D1,8 q 21d

+ Gem 1000 mg/m

24 weeks (8 cycles)

Upon disease progression, management will be on clinician's discretion (mostly best supportive care)

## ABC-02 - Results: Overall Survival (ITT)



## **Triple Negative Breast Cancer**

- Breast cancer is the most frequent tumor worldwide; approximately 10% to 20% of these patients will be diagnosed with triple-negative breast cancers
- These group presents with a varied natural history but are collectively <u>associated with poor prognosis with high risk of</u> <u>relapse and short progression-free survival (PFS) and overall</u> <u>survival (OS).</u>
- As many as 50% of patients diagnosed with **early-stage triplenegative breast cancer** (stages I to III) experience disease recurrence, and 37% die in the first 5 years after surgery.

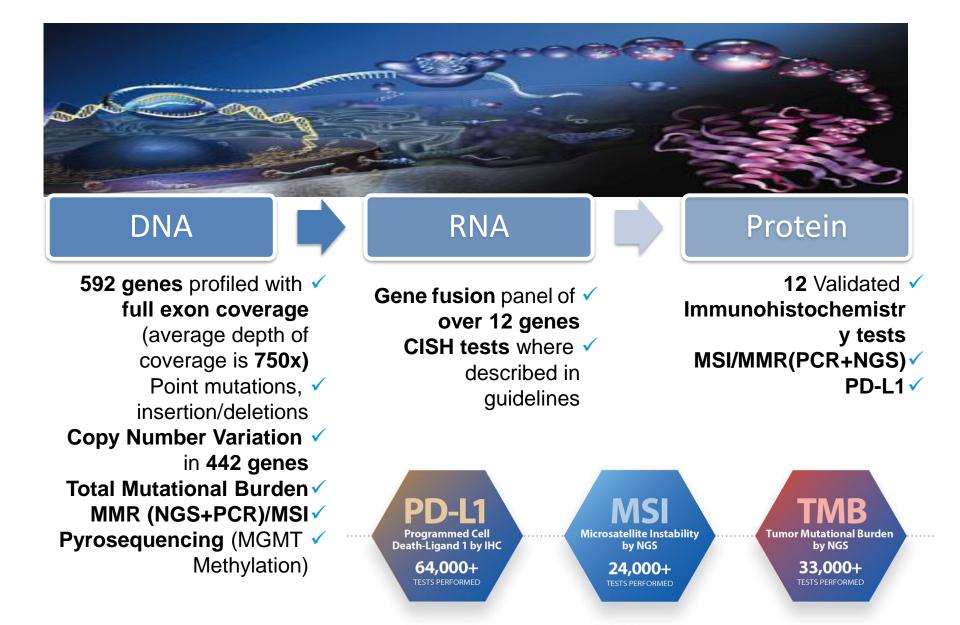
## **Triple Negative Breast Cancer**

- patients with metastatic triple-negative breast cancer have short PFS after failure of first-line chemotherapy :median PFS, 3 to 4 months.
- Current guidelines support that triple-negative breast cancer should be treated with conventional strategies primarily driven by the patient's characteristics and the toxicity profile of the treatment to be chosen.
- At present, approved targeted therapy exists and the standard remains cytotoxic chemotherapy.

## Transitional cell carcinoma(TCC)

- <u>advanced or metastatic TCC</u>
  - First-line chemotherapy regimens for advanced or metastatic TCC consists of gemcitabine and cisplatin) (GC) or a combination of methotrexate, vinblastine, adriamycin, and cisplatin (MVAC).
  - Taxanes or vinflunine have been used as second-line therapy (after progression on a platinum containing chemotherapy).
  - Immunotherapy such as pembrolizumab is often used as second-line therapy for metastatic urothelial carcinoma that has progressed despite treatment with GC or MVAC.

Change Clinical Practice with Caris Multiplatform Molecular Profiling



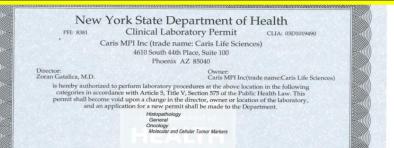
 NEW YORK STATE ,department of health recommendation: "Establish minimum criteria for depth and uniformity of coverage, i.e. number of reads, across all target areas. a minimum average of 500 reads or greater is <u>strongly</u> recommended".

 Profiling Guidelines - NCCN Guidelines<sup>®</sup> recommend "broad molecular profiling" (Full Exon ,not Hot Spot).

• Caris Type of Sequencing: 592 genes profiled with full exon coverage .average depth of coverage is 750x.

# The assay validations performed at Caris exceed CAP, CLIA, NYS and ISO 15189 regulatory standards.







Accredited

Laboratory

#### The College of American Pathologists

certifies that the laboratory named below

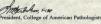
Caris MPI Inc dba Caris Life Sciences Laboratory Phoenix, Arizona Zoran Gatalica, MD,DSc

LAP Number: 7195577 AU-ID: 1460893 CLIA Number: 03D1019490

has met all applicable standards for accreditation and is hereby accredited by the College of American Pathologists' Laboratory Accreditation Program. Reinspection should occur prior to April 30, 2016 to maintain accreditation.

Accreditation does not automatically survive a change in director, ownership, or location and assumes that all interim requirements are met.

Chair, Commission on Laboratory Accreditati



American Association for Laboratory Accreditation Accredited Laboratory All Ana accredited

#### AllA has accredited CARIS MPI, INC. (D/B/A CARIS LIFE SCIENCES)

#### Phoenix, AZ

for technical competence in the field of

#### **Clinical Testing**

This laboratory is accredited in accordance with the recognized International Standard ISO 15189-2012 *Medical* laboratories - *Requirements for quality and competence*. This accreditation demonstrates technical competence for a defined scope and the operation of a laboratory quality management system (refer to joint *ISO-ILAC-LAF Communiqué dated September 2009*).



Presented this 22nd day of November 2013.

President & CEO For the Accreditation Council Certificate Number 3531.01 Valid to January 31, 2016 Revised: July 15, 2014

For the tests or types of tests to which this accreditation applies, please refer to the laboratory's Clinical Scope of Accreditation

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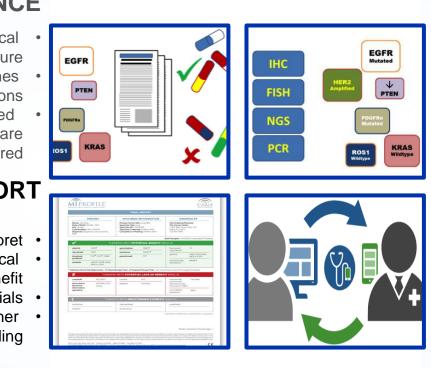
## Four critical success factors for CMI

#### ANALYSIS

- Multiplatform profiling •
- Precision IHC testing •
- High-throughput workflows: < 12 days
  - Highest Quality Standards •

#### **INTERPRETATION**

- Caris Clinical Team follows each case
  - Email with executive summary
    - Telephone call with Caris oncologist offered



#### EVIDENCE

- Continuous study of the medical literature
  - Board of experts defines associations
  - Only clinically validated predictive biomarkers are considered

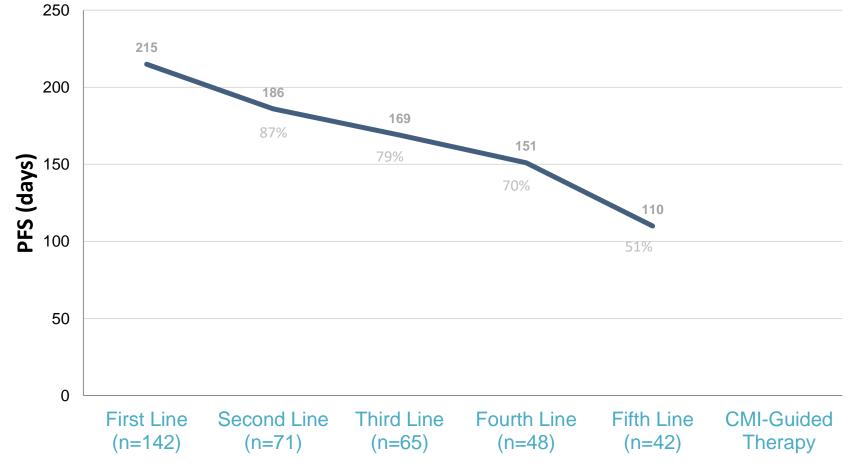
#### REPORT

- Easy to interpret .
- Drugs associated with clinical benefit and lack of benefit
  - Reference to clinical trials •
  - Detailed results and further reading

## **Bisgrove study**

- First successful publication of outcomes from molecular profiling
  - Dan von Hoff won the Karnofsky award in 2010
  - Cited over 70 times
- Molecular targets identified for 98% of patients
- Mostly targets for conventional agents were identified
- Treatment suggested by molecular profiling led to clinical benefit in 27% of treated patients

## Progression Free Survival to assess clinical benefit: PFS is getting shorter in subsequent lines of therapy



## **Bisgrove Study Primary Endpoint**

Compare progression free survival (PFS) for therapy selected by molecular profiling with PFS for the last line of therapy on which the patient progressed.

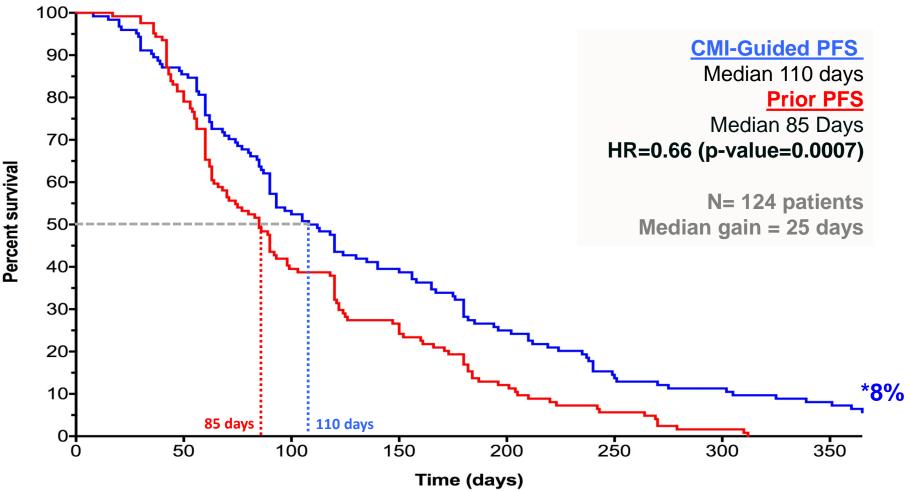


If  $PFS_b/PFS_a$  ratio was  $\geq$  1.3, MP-selected therapy was defined as having benefit for patient.

#### PFS: length of time during and after treatment in which a patient is living with a disease that does not get worse.

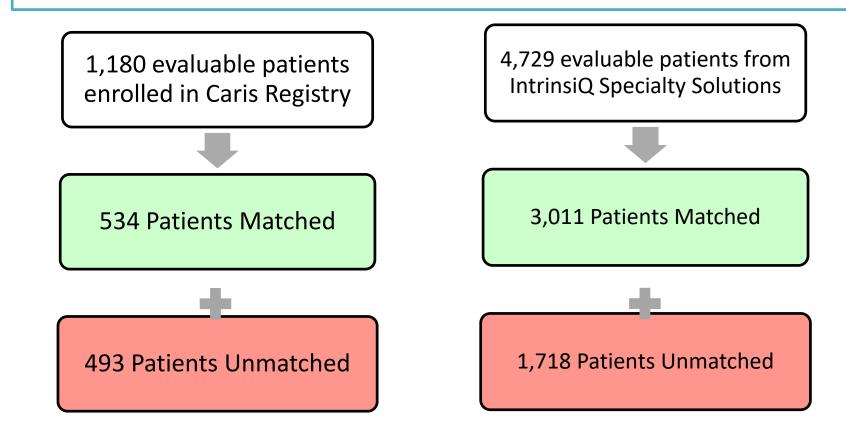
Temple, R. *Clinical Measurement in Drug Evaluation*. Ningano W. Thicker GT, eds. John Wiley and Sons Ltd: 1995; Von Hoff, D.D. 1999; Dhani et al. *Clinical Cancer Research*. 2009; 15: 118-123.

## CMI-Guided treatment leads to gain of PFS



Based on analysis of raw data from Jameson et al, Dean et al, Chahine et al and Seeber et alc

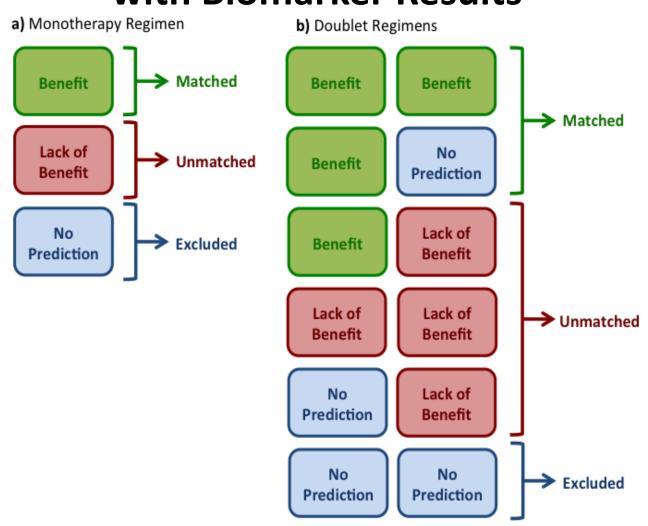
Patients Profiled using Caris Molecular Intelligence (IHC, FISH/CISH, FA, NGS) Performed under accreditation from CLIA, CAP and ISO5189:2012



Spetzler D et al. (2015) Multi-platform molecular profiling of 1,180 patients increases median overall survival and influences treatment decision in 53%

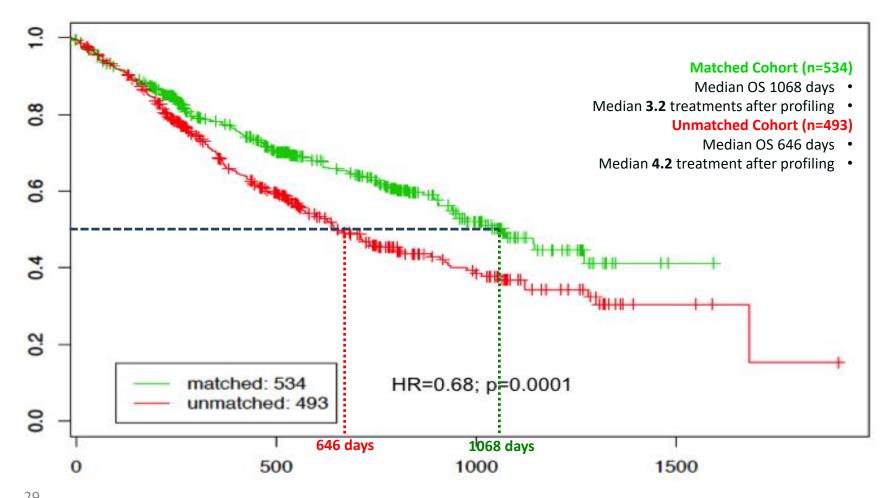
of cases. Presented at ECCO 2015 Marshall et al. (2015) Panomics validation of time to next treatment (TNT) as a surrogate outcome measure in 4729 cancer patients. J Clin Oncol 34, 2016 (suppl; abstr 11521)

#### Cohort Definitions for Monotherapies and Doublet Combinations Based on Predictive Association with Biomarker Results



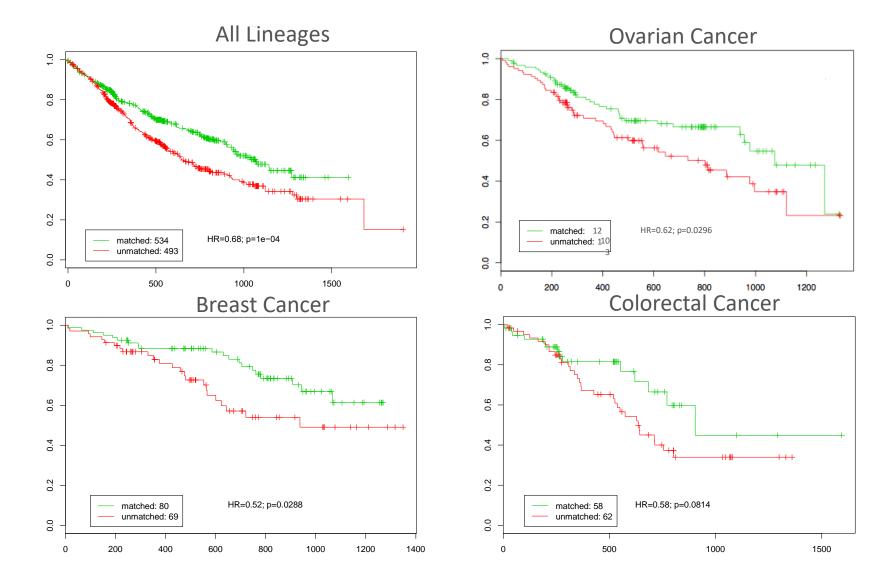
Marshall et al. (2015) Panomics validation of time to next treatment (TNT) as a surrogate outcome measure in 4729 cancer patients. J Clin Oncol 34, 2016 (suppl; abstr 11521)

## Overall Survival is higher in Matched Cohort

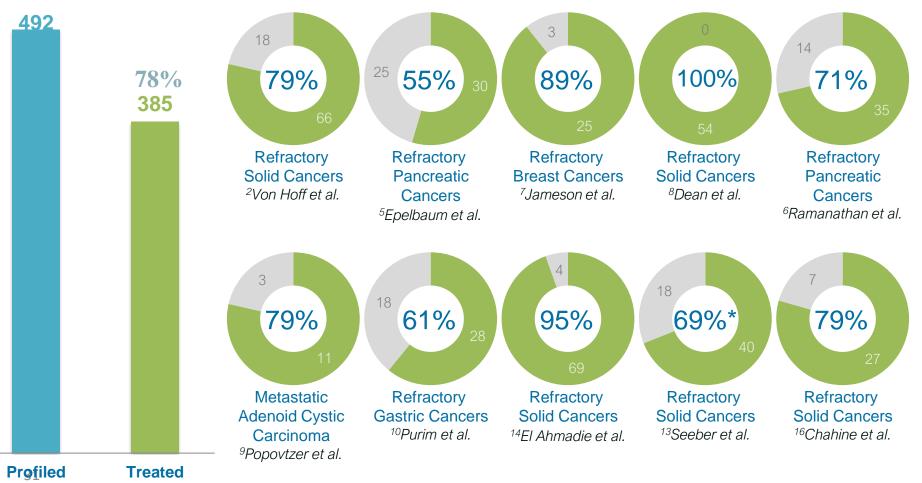


Spetzler Det al. (2015) Multi-platform molecular profiling of 1,180 patients increases median overall survival and influences treatment decision in 53% of cases. Presented at ECCO 2015

## CMI Registry: Clinical Utility Across Various Indications

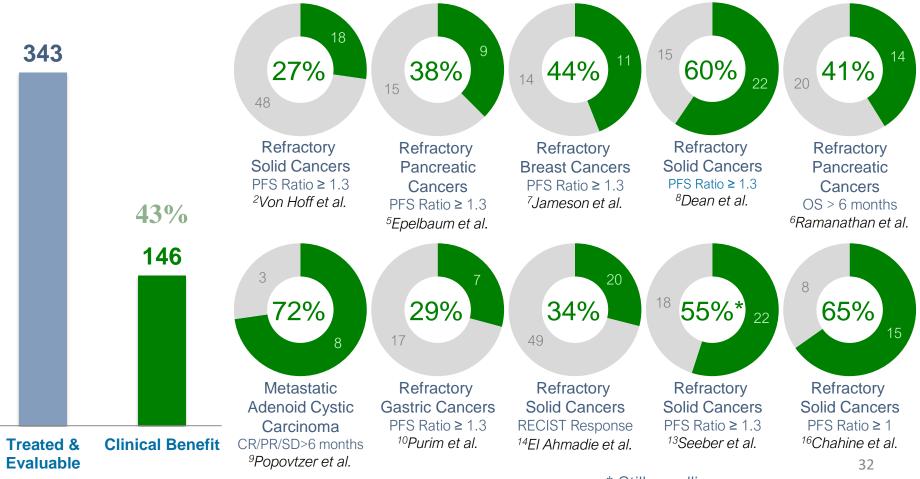


## 78% of Profiled Patients Are Treated In Line with CMI Report



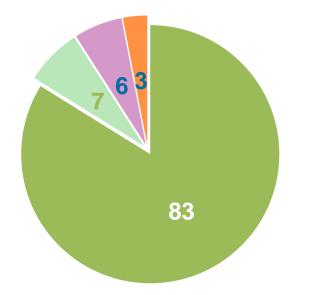
\* Still enrolling

## 43% of CMI-Guided Treatments result in Clinical Benefit



\* Still enrolling

In 10 physician-led studies, 364 patients were treated in line with the findings of the CMI report (75% of those profiled)



Conventional Chemotherapy

- Monoclonal Antobodies
- Tyrosine Kinase Inhibitors
- Hormone Therapy

The high utility is driven by accessibility of cheaper, cytotoxic chemotherapy options which are not otherwise considered

#### MULTIOMIC PROFILING OF METASTATIC LESIONS TO GUIDE TREATMENT SELECTION: THE SIDE OUT 2 TRIAL EXPERIENCE

**Study Primary Objective** 

The aim of this prospective pilot study was to explore if treatment selection based on Multi-omic Profiling (MoP) provides clinical benefits superior to empiric treatment selection in progressive metastatic breast cancers (MBC).

#### Conclusions

- ✓ This study confirmed the unique role of MoP in selecting effective treatments for MBC.
- ✓ This approach provided clinical benefits for 56% of previously treated MBC patients, which met the primary objective of the study.
- ✓ This study also suggests that irinotecan may be an under-developed drug for MBC patients.
- ✓ As such, this approach merits further investigation.

<u>Trial design</u>: The Side Out 2 trial (clinicaltrials.gov ID NCT01919749) was an open-label, multicenter pilot study which used the molecular profile of target lesions to guide treatment selection. Therapeutic regimens were selected only from FDA approved compounds.

Patient Population: Between 2014 and 2016, four US sites enrolled 32 previously treated MBC patients.

#### Key Eligibility Criteria:

- ✓ Age ≥18 years;
- ✓ ECOG of 0-1;
- Absence of symptomatic CNS metastasis;
- ✓ Adequate organ and bone marrow function;
- Documented diagnosis of metastatic breast cancer with measurable disease accessible to biopsy;
- ✓ Progression of disease on ≥ 1 prior chemotherapeutic and/or hormonal regimen(s) for advanced disease within 6 months of treatment initiation.

Response Rate Criteria: Growth Modulation Index (GMI) was used to assess patients' response to treatment based on tumor response by RECIST 1.1.

PFS prior	PFS selected	
therapy (PFS <sub>A</sub> )	by MoP (PFS <sub>0</sub> )	

 $PFS_B/PFS_A$  ratio  $\geq 1.3 =$  benefit for patient.

To meet the primary objective,  $\ge 30\%$  of patients must reach a GMI score  $\ge 1.3$  (PMID:25209003).

#### Study workflow

#### Consenting/Screening/Enrollment:

MBC patient with disease progression, clear documentation of time between treatments, and documented progression on the most recent treatment.

# Tissue Collection and Multi-Omic Analysis: Multi-omic analysis of the metastatic lesion; - RNA-Seq & Exome Sequencing; - Immunohistochemistry of 7 predictive markers\*; - LCM-RPPA based protein singling network analysis of 12 FDA approved drug targets and downstream substrates\*\*. Yes No



#### Disease Assessment Using RECIST Criteria:

Patients are assessed every 7  $\pm$  1 weeks during the GMI monitoring window until disease progression or treatment discontinuation.

If progression is not observed at the end of therapy, patients are assessed every 3 months until progression.

At disease progression an optional second biopsy may be performed.

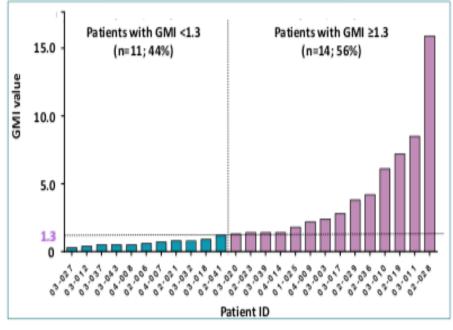
\*IHC markers: Androgen (AR), Estrogen (ER), and Progesterone (PR) Receptor; SPARC; TOP2A; TOP01, and Thymidylate Synthase (TS).

<sup>\*\*</sup>LCM-RPPA markers: ALK; pAKT \$473; pc-Abl \$735; pEGFR \$1068; pERB2 \$1248; pERB3 \$1289; pERK \$1/2 \$1202/\$204; pp7056K \$1389; pPDGFR \$151; PTEN; pRet \$905; pSrc \$527.

#### Enrollment overview

Patient Summary	Number of Patients
Enrolled	32
Treated based on MoP	29
Treated with standard of care	3
Evaluable for GMI window	25

#### Patient outcome based on GMI score



- ✓ Of the 25 patients, 14 (56%) met or exceeded a GMI of 1.3.
- The most frequently selected treatments were: Irinotecan based on TOPO1 expression (n = 12; single agent n = 5) and Capecitabine based on TS expression (n = 10; single agent n = 3).
- Seven patients received endocrine therapy, 3 of whom were treated with Everolimus and Exemestane.
- Based on HER2 amplification/pathway activation, HER2 targeted agents were given to 5 patients.

#### Molecular characteristics of metastatic lesions and treatment

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Subject ID	GMI	Receptor Status	Metastatic site	Targets	Treatment
02-03-027	0.3	ER+; PR-; HER2-	Omentum	AR; ER; TOPO1	Irinotecan; Megestrol Acetate
02-03-012	0.4	ER+;PR+;HER2-	Liver	AR; ER; TOPO1; TS	Capecitabine; Irinote can; Megestrol Acetate
02-03-037	0.5	ER+;PR+;HER2-	Liver	TOP01	Irinotecan
02-03-043	0.5	ER+; PR-; HER2-	Liver	TUBB3	Eribulin
02-04-008	0.5	ER+;PR+;HER2-	Chest wall/Skin	ER; p-p7056K	Everol imus; Exemesta ne
02-02-006	0.6	ER+; PR-; HER2-	Lymph node	p-AKT; p-ERB2; p- ERB3; p-ERK; TS	Capecitabine; Lapatinib
02-04-007**	0.7	ER+; PR-; HER2-	Chest wall/Skin	ER; p-ERB2; p-ERK; TOPO1; TUBB3	Eribulin; Irinotecan; Lapatinib; Letrozole
02-02-021	0.8	ER+; PR-; HER2-	Omentum	ER; p-p7056K	Everol imus; Exemesta ne
02-03-032	0.8	ER-;PR-;HER2-	Chest wall/Skin	TUBB3	Eribulin
02-03-018	0.9	ER+; PR-; HER2-	Liver	Thymidine Phosphorylase (TYMP)	Capecitabine
02-02-041	1.2	ER-;PR-;HER2-	Chest wall/Skin	TOP 01	Irinotecan
02-03-020	1.3	ER+; PR-; HER2-	Liver	ER; p-p7056K	Everol imus; Exemesta ne
02-02-023	1.4	ER-;PR-;HER2-	Liver & Lymph node*	EZH2*; Survivin*; TOP O1; TS; TU BB3*	Capecitabine; Irinote can; Paclitaxel
02-03-039	1.4	ER-;PR-;HER2+	Lung	TOPO1; HER2; p- ERB2; p-ERK	Irinotecan; Trastuzuma b
02-04-014	1.4	ER+;PR-;HER2-	Lung	TOP01	Irinotecan
02-01-025	1.8	ER+; PR-; HER2-	Lymph node	TS	Capecitabine
02-04-009	2.2	ER+;PR+;HER2-	Ab dominal mass	AR; ER; TS; AR; TUBB3	Capecitabine; Megestrol Acetate; Vinorelbine
02-03-003	2.4	ER+; PR-; HER2-	Liver	SPARC	Paclitaxel
02-03-017	2.8	ER+; PR-; HER2-	Liver	TS; p-EGFR; p-ERB2; p-ERB3; p-ERK	Capecitabine; Lapatinib
02-02-029	3.8	ER-; PR-; HER2- * **	Chest wall/Skin	TOP01	Irinotecan
02-02-036	4.2	ER+; PR-; HER2-	Liver	TOPO1; TS	Capecitabine; I rinote can
02-03-010	6.1	ER+;PR+;HER2-	Liver	TOP 01	Irinotecan
02-02-019	7.2	ER-;PR-;HER2+	Chest wall/Skin	p-EGFR; p-ERB2; p- ERB3/ERBB3; p-ERK; HER2; TUBB3	Docetaxel; Pertuzumab; Trastuzumab
02-03-011	8.5	ER+;PR+;HER2+	Liver	TOPO1; TS	Capecitabine; Irinotecan
02-02-028	15.9	ER+; PR-; HER2-	Chest wall/Skin	TS	Capecitabine

\* A second biopsy was collected from the same patient after recurrence; \*\* Metadatic lesion from a male breast turner; \*\*\* Data retrieved from whole exome sequencine analysis.

#### ORIGINAL RESEARCH ARTICLE



#### Biomarker-Driven Therapy in Metastatic Gastric and Esophageal Cancer: Real-Life Clinical Experience

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#### Abstract

Background Precision treatment of cancer uses biomarker-driven therapy to individualize and optimize patient care. Objective To evaluate real-life clinical experience with biomarker-driven therapy in metastatic gastric and esophageal cancer in Israel.

Patients and Methods This multicenter retrospective cohort study included patients with metastatic gastric or esophageal cancer who were treated in the participating institutions and underwent biomarker-driven therapy. Treatment was considered to have a benefit if the ratio between the longest progression-free survival (PFS) post biomarker-driven therapy and the last PFS before the biomarker-driven therapy was  $\geq 1.3$ . The null hypothesis was that  $\leq 15\%$  of patients gain such benefit.

**Results** The analysis included 46 patients (61% men; median age, 58 years; 57% with poorly-differentiated tumors). At least one actionable (i.e., predictive of response to a specific therapy) biomarker was identified for each patient. Immunohistoch emistry was performed on all samples and identified 1–8 (median: 3) biomarkers per patient (most commonly: low TS, high TOPO1, high TOP2A). Twenty-eight patients received therapy after the biomarker analysis (1–4 lines). In the 1st line after biomarker analysis, five patients (18%) achieved a partial response and five (18%) stable disease; the median (range) PFS was 129 (12–1155) days. Twenty-four patients were evaluable for PFS ratio analysis; in seven (29.2%), the ratio was  $\geq$ 1.3. In a one-sided exact binomial test vs. the null hypothesis, p = 0.019; therefore, the null hypothesis was rejected.

Conclusions Our findings demonstrated that implementing biomarker-driven analysis is feasible and could provide clinical benefit for a considerable proportion (~30%) of patients with metastatic gastric or esophageal cancer.

## Methods Study design and patient population

- Study period between January 2010 and March 2014
- The study was approved by the institutional review boards of the participating institutions:
  - Rabin Medical Center
  - Sourasky Medical Center
  - Rambam Health Care Campus
  - Hadassah Hebrew University Medical Center
  - Kaplan Medical Center
  - Wolfson Medical Center

## **Results: Patients characteristics**

#### Deomgraphics

Characteristic	N = 46	
Gender, N (%)		
Male	28 (60.9)	
Female	18 (39.1)	
Age at diagnosis, years		
Median (range)	58.4 (27.2-78.3)	
Ethnicity, N (%)		
Jewish	37 (80.4)	
Ashkenazi	16 (34.8)	
Sephardi	13 (28.3)	
Not available/mixed	8 (17.4)	
Non-Jewish	9 (19.6)	
Family history of cancer, N (%)		
Yes	25 (54.3)	
No	21 (45.7)	

# **Results: Disease characteristics**

Tumor site, N (%)	
Cardia	18 (39.1)
Gastroesophageal junction	11 (23.9)
Antrum	7 (15.2)
Esophagus	7 (15.2)
Not available	3 (6.5)
Grade, N (%)	
Well differentiated	1 (2.2)
Moderately differentiated	11 (23.9)
Poorly differentiated	26 (56.5)
Not available	8 (17.4)
HER2 status, <sup>a</sup> N (%)	
Positive	3 (7)
Negative	20 (43.5)
Equivocal	1 (2.2)
Test not performed/test failure	22 (47.8)
Metastatic at diagnosis, N (%)	
Yes	27 (58.7)
No	17 (37.0)
Not available	2 (4.3)

## **Results:**

# Treatment regimens received prior to molecular profiling

First-line treatment for metastatic disease	n=36
5-FU/capecitabine + cisplatin	11 (30.6)
5FU/capecitabine + cisplatin+ docetaxel	11 (30.6)
Other	14 (39)
	20
Second-line treatment for metastatic disease	n=20
Second-line treatment for metastatic disease FOLFIRI	n=20 7 (35.0%)
FOLFIRI	7 (35.0%)

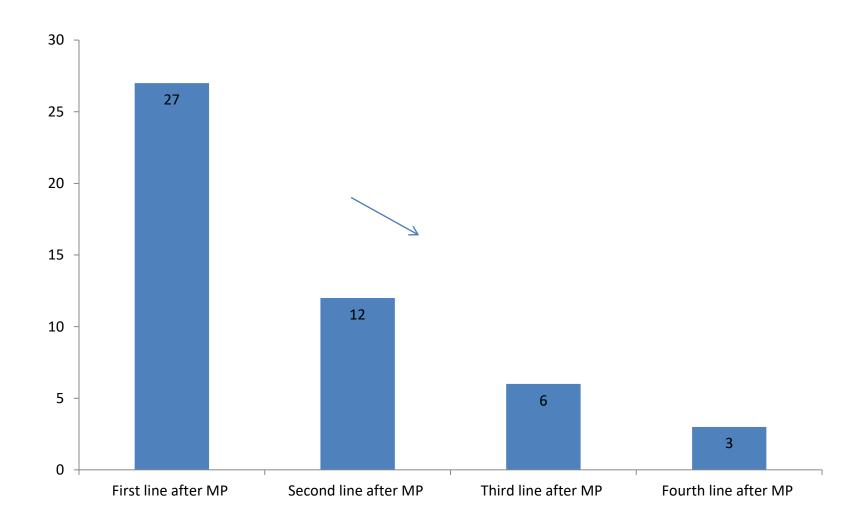
## Results: Actionable biomarkers

Actionable biomarker	Number of patients out of evaluable patients, N/N	Frequency, %	Drugs associated with clinical benefit
Negative/low TS	34/40	85.0	Fluoropyrimidines and other folate analogs
High TOPO1	27/40	67.5	Irinotecan
High TOP2A	27/41	65.9	Anthracyclines
Negative/low ERCC1	21/36	58.3	Platinum-based therapy
Negative/low RRM1	22/40	55.0	Gemcitabine
Negative/low MGMT	22/46	47.8	Temozolomide

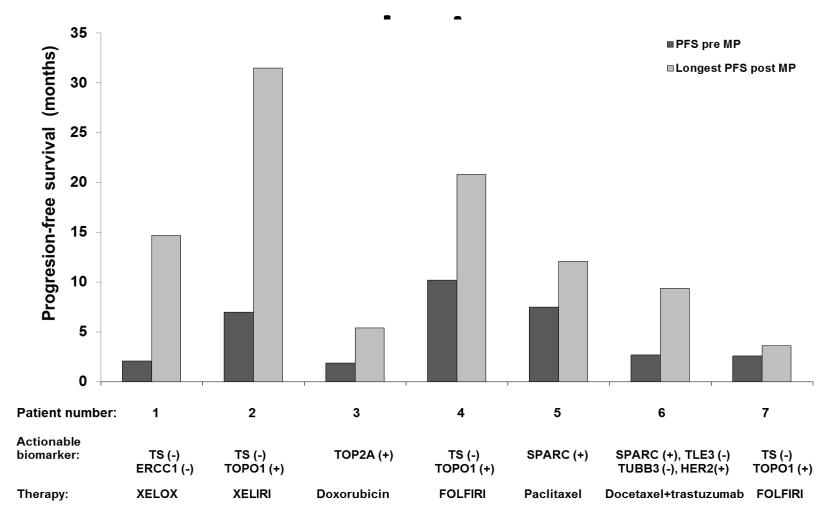
Only markers that were tested in samples of at least 35 patients are included in the table.

ERCC1, excision repair cross-complementation 1; MGMT, O-6-methylguanine-DNA methyltransferase; RRM1, ribonucleotide reductase M1 subunit; TOPO1, topoisomerase 1; TOP2A, topoisomerase IIA; TS, thymidylate synthase.

## Results: Patients treated according to MP



## :Results Patients achieving the PFS ratio



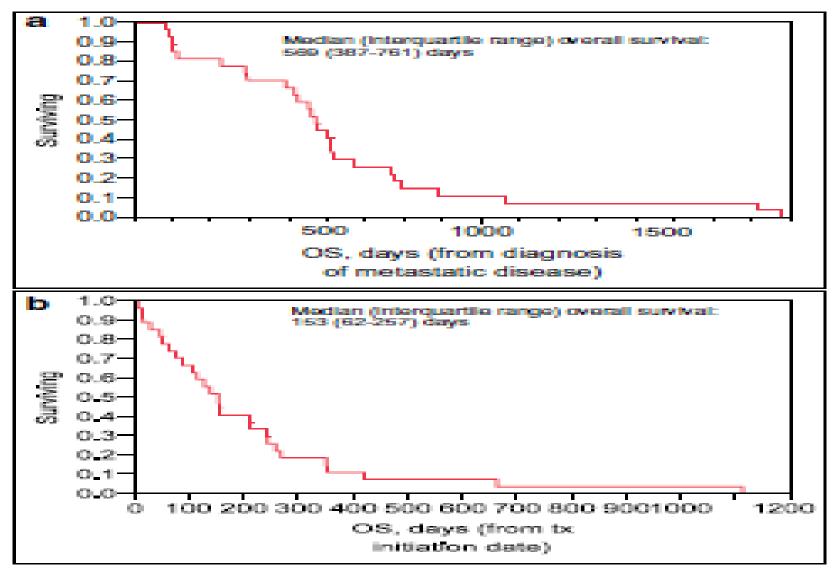


Fig. 2 Kaplan-Meier survival curve from diagnosis of metastatic disease (A) and from initiation of biomarker-driven therapy (B) for evaluable study patients who received biomarker-guided analysis (n = 2.7). Tick marks indicate censored observations

## Conclusions

In summary, this study shows in real-life clinical practice

that implementing MP is feasible and provides clinical

benefit therapy (PFS ratio of  $\geq$ 1.3) for a close to a third

of patients with metastatic gastric/esophageal cancer.

Prospective studies are warranted.

I thank the patients and .the families My co-authors And Teva and Caris

## THANK YOU