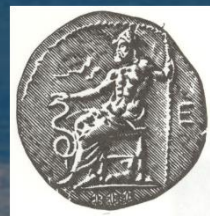


George Pentheroudakis
Associate Professor of Oncology
Medical School, University of Ioannina



Cancer of Unknown Primary: Diagnosis and Treatment AND BIOLOGY

CYE Session
ESMO Conference, Madrid 2014

What we will talk about

- Definition of CUP
- CUP Epidemiology and Statistics
- CUP Pathology and Molecular Assays
- CUP Staging
- CUP Clinicopathologic Subsets
- CUP Prognostication
- CUP Therapy
- CUP Biology
- CUP Clinical Research

Definition of CUP

- **Cytologic/pathologic diagnosis of malignancy (systemic metastases) in the absence of an identifiable primary tumour after a standardised diagnostic work up**



CUP: What is it?

- Metastases from a primary we simply cannot locate?
- Tumours with not only a primary tissue-specific biology but also with a distinct biological signature, common for most CUPs?



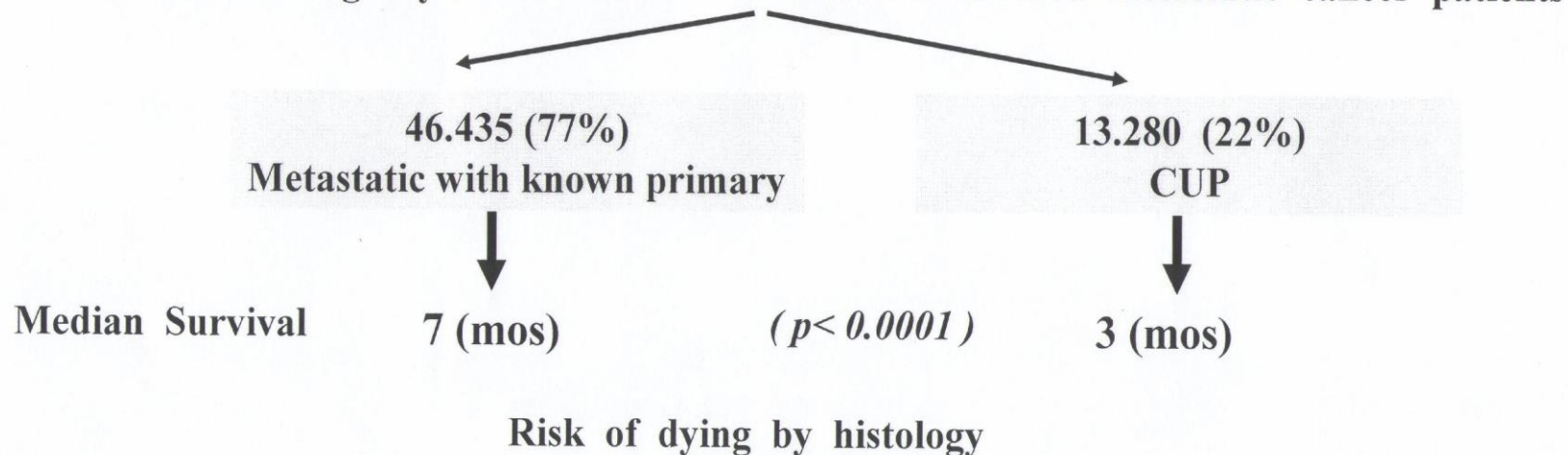
**IS THERE EVIDENCE TO
SUGGEST THAT CUP HAS
PECULIAR AND DISTINCT
BIOLOGY COMPARED TO METS
FROM KNOWN PRIMARY
TUMOURS (KPM)?**

PROGNOSIS OF SUB-TYPES OF CUP COMPARED TO METASTATIC CANCER

Bishop JF et al [New South Wales, Australia]

ASCO 2007 (abs 21010)

❖ Cancer Registry : 1980 – 2004 ❖ 65.239 confirmed metastatic cancer patients

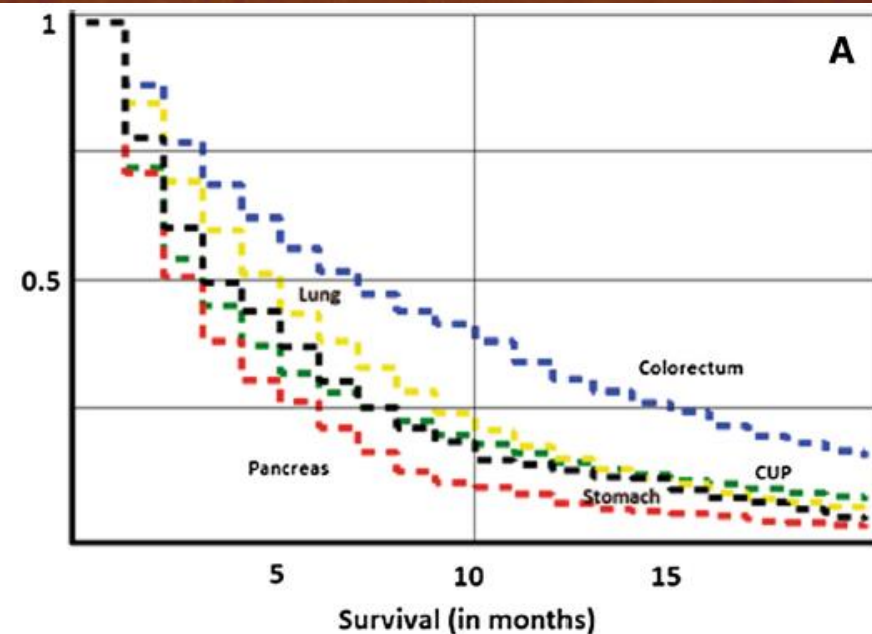


Risk of dying by histology

| | |
|------------------|----------------------|
| Adenocarcinoma | 80% greater (HR 1.8) |
| Squamous | 50 % less (HR 0.49) |
| Small cell | 20% less (HR 0.77) |
| Undifferentiated | 15% less (HR 0.84) |

Survival: CUP vs KPM

- Swedish Cancer Registry: 2881 CUP vs 6745 KPM



| Primary site | N | HR |
|----------------------------|------|-------------|
| <i>CUP (reference)</i> | 2881 | 1 |
| <i>Colorectal cancer</i> | 1438 | 0.61 |
| <i>Pancreatic cancer</i> | 460 | 1.71 |
| <i>Stomach cancer</i> | 322 | 1.16 |
| <i>Liver cancer</i> | 188 | 1.58 |
| <i>Lung cancer</i> | 2453 | 0.98 |
| <i>Kidney cancer</i> | 284 | 0.71 |
| <i>Bladder cancer</i> | 139 | 0.93 |
| <i>Prostate cancer</i> | 1259 | 0.24 |
| <i>Breast cancer</i> | 202 | 0.53 |
| <i>All known primaries</i> | 6745 | 0.69 |

CUP characteristics

- Regression/dormancy of the primary
Autopsies fail to identify the primary in 30-40% of cases or identify small nodules in lung, pancreas. Why does it not grow?

- Early uncommon systemic metastases

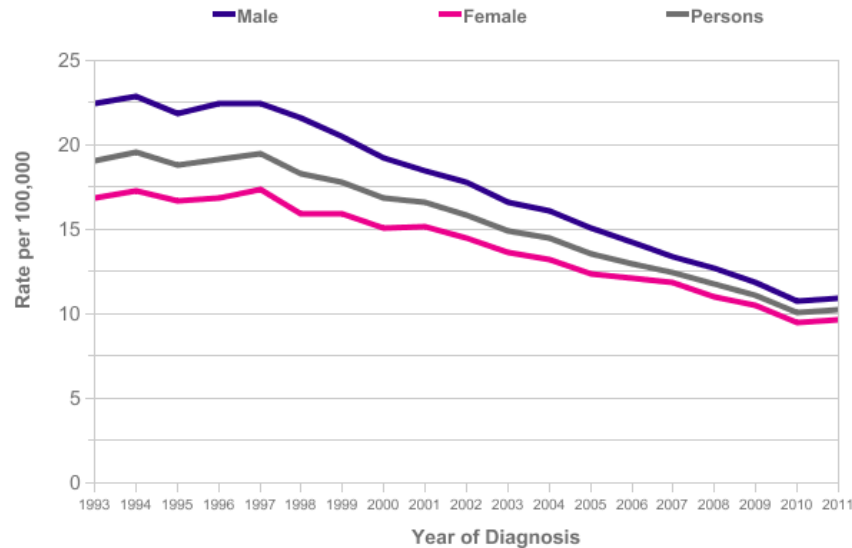
High volume mets in multiple sites.

Viscera, skin, heart, scalp, kidneys, distant nodes

EPIDEMIOLOGY OF CANCER OF UNKNOWN PRIMARY

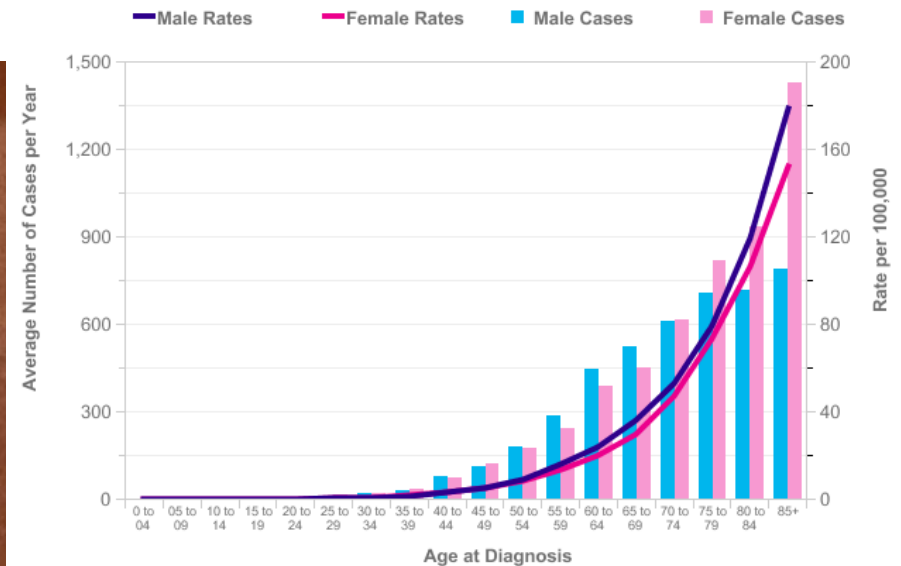
| Geographical area | Source | Frequency (%) | Period |
|-------------------|------------------------------|---------------|-----------|
| USA | SEER | 2.3 | 1973-1987 |
| Australia | New South Wales Registry | 4.2 | 1970-1990 |
| Netherlands | Eindhoven Cancer Registry | 4.0 | 1984-1992 |
| Finland | IARC | 2.5 | - |
| Germany | - | 7.8 | 1968-1984 |
| Russia | - | 3.6 | - |
| Switzerland | Local registries | 2.3 | 1984-1993 |
| Japan | IARC | 3.0 | - |

Incidence



1993-2011: European Age-Standardised Incidence Rates per 100,000

Average Number of New Cases Per Year and Age-Specific Incidence Rates per 100,000, UK 2009-2011



CUP Pathology

- Biopsy needed
- Generous material wellcome (excisional biopsy or tru-cut rather than FNA)
- Classical histo work up with H&E stain

Primary markers

Additional markers

CK 7-/CK 20+



Colorectal
and Merkel
cell carcinoma



CEA and CDX-2

CK 7+/CK 20-



Lung, breast, thyroid,
endometrial, cervical, and
pancreatic carcinoma
and cholangiocarcinoma



TTF-1, ER, PR,
GCDFP-15,
and CK 19

CK 7+/CK 20+



Urothelial, ovarian, and
pancreatic cancer and
cholangiocarcinoma



Urothelin and WT-1

CK 7-/CK 20-

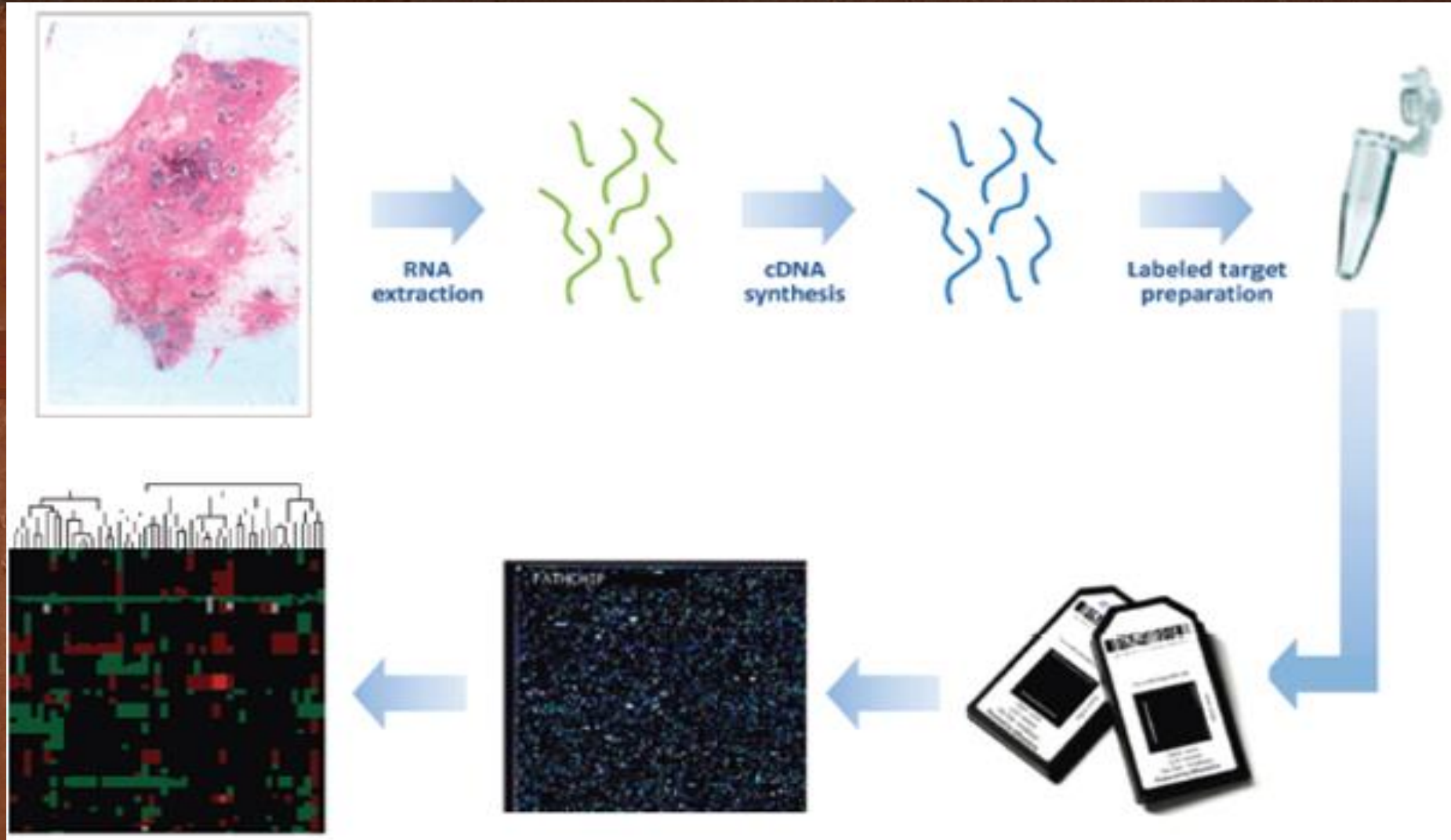


Hepatocellular,
renal cell, prostate,
squamous cell



Hep Par-1 and PSA

Molecular Assays



Molecular assignment of primary: Gene expression arrays

Assays

| Assay | Platform | Tissue | No. of Tumor types | Number of genes | Accuracy (%) |
|---|------------------|-----------------|-------------------------|-----------------|--------------|
| Veridex | RT-PCR mRNA | FFPE | 6 and “other” | 10 | 76 |
| Pathwork Diagnostics Tissue of Origin test | cDNA microarray | Frozen/ FFPE | 15 | 1500 | 89 |
| Rosetta Genomics MiRview mets-2 | Microarray miRNA | FFPE | 42 | 64 miRNAs | 92 |
| bioTheranostics CancerType ID | RT-PCR mRNA | FFPE | 39 (including subtypes) | 92 | 86 |

IDENTIFICATION OF PRIMARY SITE BY GENETIC PROFILING (MICROARRAYS) FROM ALL PUBLISHED CUP SERIES

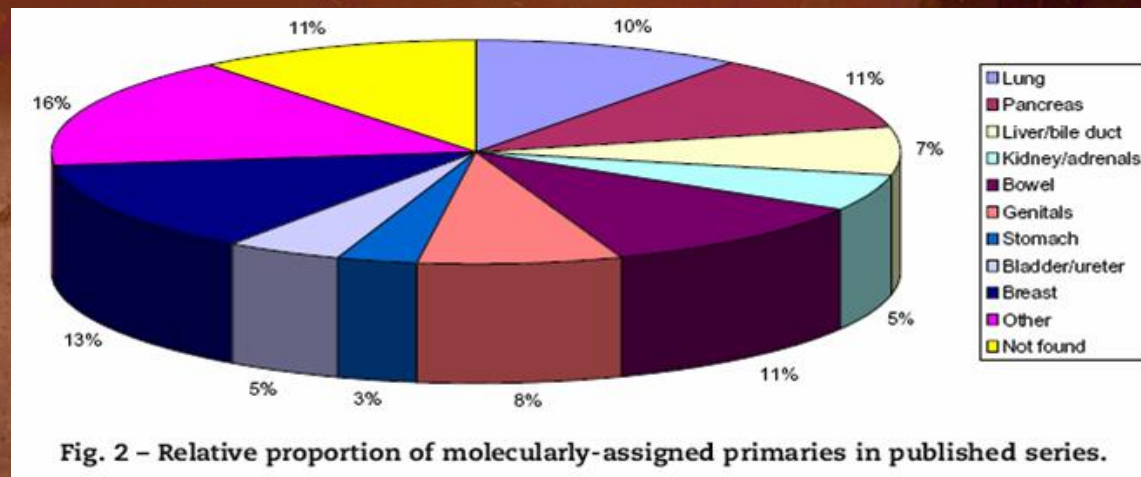
Years of Publications : 2005- 2007

No of Samples : > 500 (cDNA)

Biological Assignment of Primaries (Accuracy) : 50 – 87 %

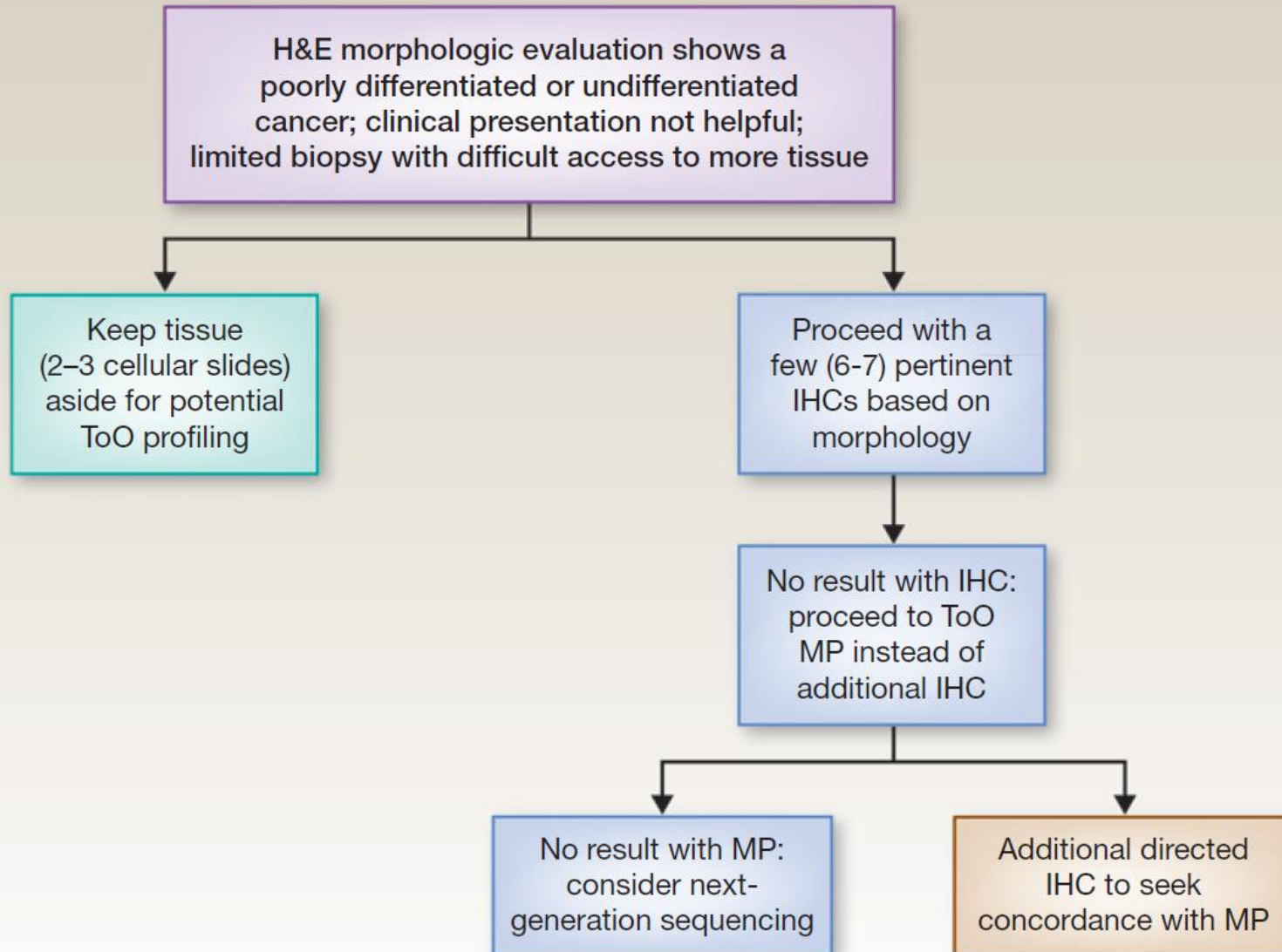
Primary Sites Identified :

| | |
|-------------------|--------|
| Breast | 15 % |
| Pancreas | 12.5 % |
| Bowel | 12 % |
| Lung | 11.5 % |
| Genital system | 9 % |
| Liver/bile duct | 8 % |
| Kidney / adrenals | 6 % |
| Bladder / ureter | 5 % |
| Stomach | 3 % |
| Other | 18 % |



Suggested use of Molecular Assays

Clin Cancer Res 2013;19:4027-4033



CLINICAL AND THERAPEUTIC UTILITY OF GENE AND PROTEIN MICROARRAY TECHNOLOGIES

QUESTION 1

DOES MOLECULAR ASSAYS, INCREASE THE ACCURACY OF IDENTIFYING THE PRIMARY SITE?

ANSWER 1

YES : UP TO 90% ACCURACY

QUESTION 2

DOES THIS DIAGNOSTIC AID RESULTS IN IMPROVEMENT OF PATIENT OUTCOME ?

ANSWER 2

?

CUP Staging

| Assessment suggested | Target patient population |
|--|--|
| Thorough medical history and physical examination | All patients |
| Basic blood and biochemistry survey | All patients |
| CT scans of thorax, abdomen and pelvis | All patients |
| Mammography | Female patients |
| Work-up for CUP subsets | |
| Breast MRI | Female with axillary adenocarcinoma |
| Serum α -fetoprotein and human chorionic gonadotropin | Patients with midline metastatic disease |
| Serum prostate-specific antigen | Male with adenocarcinomatous bony metastases |
| Head and neck CT/PET scan (optional) | Cervical squamous carcinoma |
| Endoscopies | Sign/symptom/lab-oriented |
| Octreoscan and plasma chromogranin A | Patients with neuroendocrine tumor CUP |

SERUM TUMOR MARKERS

- ❖ Routine evaluation of current commonly used markers have not been proven of any prognostic or diagnostic assistance
- ❖ A non – specific multiple overexpression of the adenocarcinoma markers (CEA, CA 125, CA 15-3, CA 19-9) has been observed in the majority of CUP patients.
- ❖ Worthwhile to request :

| | |
|-------------|--|
| PSA | in men with bone metastatic adenocarcinoma |
| B-HCG & AFP | in men with an undifferentiated tumor |
| AFP | in patients with hepatic tumors |
| CA 125 | women with papillary adenocarcinoma of peritoneal cavity. |
| CA 15-3 | women with adenocarcinoma involving only axillary lymph nodes. |

CUP

```
graph LR; CUP --> Favourable[FAVOURABLE OR GOOD PROGNOSIS SUBSETS 20%]; CUP --> Unfavourable[UNFAVOURABLE OR POOR PROGNOSIS SUBSETS 80%];
```

**FAVOURABLE OR
GOOD PROGNOSIS SUBSETS**

20%

**UNFAVOURABLE OR
POOR PROGNOSIS SUBSETS**

80%

Favourable Subsets

1. Poorly differentiated carcinoma with **midline distribution** (extragonadal germ cell syndrome).
2. Women with **papillary** adenocarcinoma of peritoneal cavity.
3. Women with adenocarcinoma involving only **axillary** lymph nodes.
4. **Squamous** cell carcinoma involving cervical lymph nodes
5. Poorly differentiated **neuroendocrine** carcinomas.
6. Men with **blastic bone** metastases PSA+ (adenocarcinoma).
7. Adenocarcinoma with a **colon-profile** (CK 20⁺, CK 7⁻, CDX 2⁺)
8. Isolated **inguinal** adenopathy (squamous carcinoma).
9. Patients with a **single**, small, potentially resectable tumor.

CHARACTERISTICS OF PATIENTS WITH POORLY DIFFERENTIATED CUP

| | |
|---------------------------|--|
| GENDER / AGE | : Men / < 50 yrs |
| TUMOR INVOLVEMENT | : Mediastinum Retroperitoneum Lungs Lymph nodes |
| TUMOR MARKERS | : Elevated serum levels of β -HGC or AFP |
| CLINICAL EVOLUTION | : Rapid tumor growth |
| RESPONSE TO Rx | : Favourable response to Cisplatin - based chemotherapy. RR 50% (CRs: 15-25%) |
| SURVIVAL | : Median : 13 months 15% long – term survivors |

PERITONEAL CARCINOMATOSIS IN FEMALES

THE NATURAL HISTORY

| | |
|-------------------------|--|
| <i>Incidence</i> | 10% of CUP patients |
| <i>Mean Age (yrs)</i> | 60 (25 – 80) |
| <i>Clinical Picture</i> | Abdominal distension, pelvic masses, ascites |
| <i>Surgical Picture</i> | Abdominal masses, peritoneal disease, ascites, with normal ovaries |
| <i>Histology</i> | Papillary serous carcinoma (\pm psammoma bodies) |
| <i>Serum CA-125</i> | Often abnormal or markedly elevated. |

WOMEN WITH PAPILLARY ADENOCARCINOMA OF PERITONEAL CAVITY

(*Peritoneal Adenocarcinomatosis*)

Treatment :

- As FIGO III ovarian cancer.
- Surgical cytoreduction.
- Platinum – based chemotherapy.

Response Rate :

40 – 60 % (CR : 30 %)

Survival :

Median : 16 months

Long – term survival :

5-yr: 10 %

Axillary nodal metastases from carcinoma of unknown primary (CUPAx): a systematic review of published evidence

George Pentheroudakis · George Lazaridis ·
Nicholas Pavlidis

| | |
|-------------------------|---|
| Years | : 1975 – 2006 (24 studies) |
| N | : 689 patients |
| Mean Age | : 52 yr |
| Menopause status | : Postmenopausal 66% Premenopausal 34% |
| Histology | : Ductal adenocarcinoma 83%, ER/PR 40 - 50/%, HER2 31% |
| Nodal status | : N1 : 48% > N1 : 52 |

Treatment and Outcome

- ✓ **Mastectomy / axillary dissection** : **59 %**
- ✓ **Primary breast irradiation** : **26 %**
- ✓ **Observation** : **15 %**
- ✓ **Logoregional recurrence rate** : **25 %** (mostly in observation cases)
- ✓ **5-yr Survival** : **72 %** (similar to stage II-III breast cancer)
- ✓ **No survival difference** between conservative management (breast preservation + RT) and mastectomy

SQUAMOUS CELL CANCER INVOLVING CERVICAL LYMPH NODES

- Treatment :**
- As locally advanced head-neck cancer.
 - Surgery alone is inferior except **pN1** neck disease with no extracapsular extension.
 - **Radiation** : both sides of neck and mucosa (entire pharyngeal axis and larynx).
 - **Chemotherapy** remains undefined (despite encouraging results with Platinum-based).

- Survival :**
- 5-year survival **35–50%**.
 - Documented **long term** disease – free survivors.

POORLY DIFFERENTIATED NEUROENDOCRINE CARCINOMAS

Treatment:

Platinum – based or
paclitaxel / carboplatin – based
chemotherapy

Response :

50 – 70% (CR : 25%)

Survival :

Median : 14.5 months

3-yr : 24%

UNFAVOURABLE SUBSETS

1. Adenocarcinoma metastatic to the **liver or other** organs
2. **Non-papillary** malignant ascites (adenocarcinoma)
3. Multiple **cerebral** metastases (adeno or squamous Ca)
4. Multiple **lung/pleural** metastases (adenocarcinoma)
5. Multiple **metastatic bone** disease (adenocarcinoma)
6. **Squamous** – cell carcinoma of the **abdominal cavity**

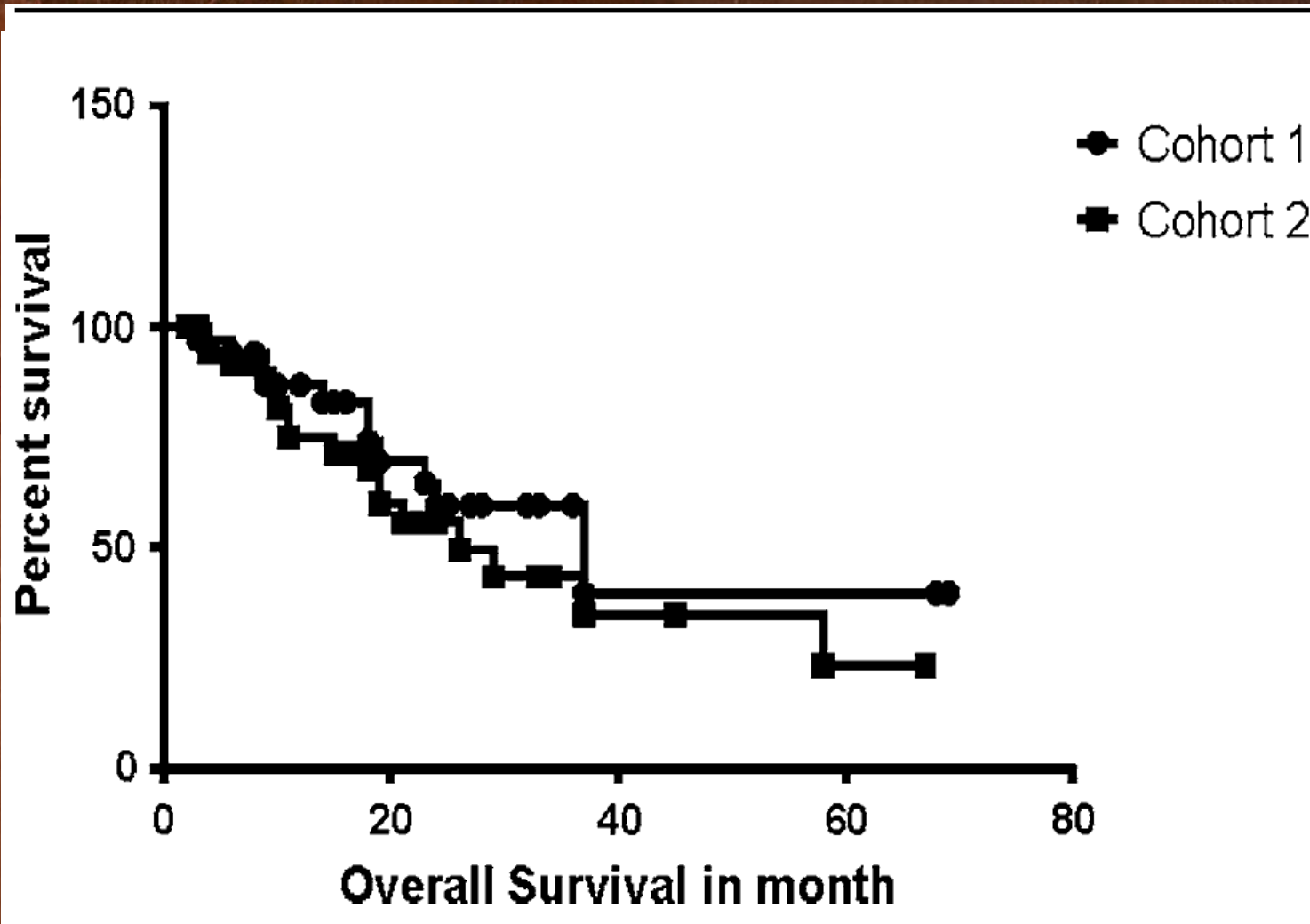
Table 4. Long-Term Survival in Patients With Unknown Primary Carcinoma and Unfavorable Prognostic Factors

| Author and Year of Publication | No. of Patients | Regimen | Median Survival (mo) | 1-Year Survival (%) | 2-Year Survival (%) | 3-Year Survival (%) |
|---|-----------------|---------------------------------|----------------------|---------------------|---------------------|---------------------|
| Briasoulis et al, 2000 ³⁴ | 33 | PCb | 10 | 25 | 5 | NR |
| Dowell et al, 2001 ³⁵ | 34 | P5FUL (17) CbE (17) | 8.3 6.4 | 26 | NR | NR |
| Balaña et al, 2003 ³⁸ | 30 | GCE | 7.2 | 36 | 14 | NR |
| Park et al, 2004 ⁴⁰ | 37 | PC | 11 | 38 | 11 | NR |
| Piga et al, 2004 ³⁹ | 102 | CbDoxE | 9 | 35.3 | 18 | 11 |
| Pouessel et al, 2004 ⁴¹ | 35 | GD | 10 | 43 | 7 | NR |
| El-Rayes et al, 2005 ⁴³ | 22 | PCb | 6.5 | 27 | NR | NR |
| Pittman et al, 2006 ³⁶ | 51 | GCb | 7.8 | 26 | 12 | NR |
| Palmeri et al, 2006 ⁴⁴ | 66 | GPC (33) GVC (33) | 9.6 13.6 | 30 52 | NR NR | NR NR |
| Berry et al, 2007 ⁴⁶ | 42 | PCb | 8.5 | 33 | 17 | NR |
| Briasoulis et al, 2007 ⁴² | 47 | Oxlr | 9.5 | 40 | NR | NR |
| Schneider et al, 2007 ⁴⁵ | 33 | GCaCb | 7.6 | 35.6 | 14.2 | NR |
| MPCRN (5 trials) 1997-2008 ^{1,21-24} | 396 | Multiple regimens (see text) | 9.1 | 38 | 19 | 12 |
| Total | 928 | | 8.9* | 34.6* | 13* | 12* |

OVERALL RESULTS OF CHEMOTHERAPY IN CUP PATIENTS WITH LIVER METASTASES

N° of trials : 5 (1991, 1998, 2002, 2005, 2008)
N° of patients : 711
Response rate : < 20%
Median survival : 5.5 months

A poor-risk CUP subgroup that may not be so

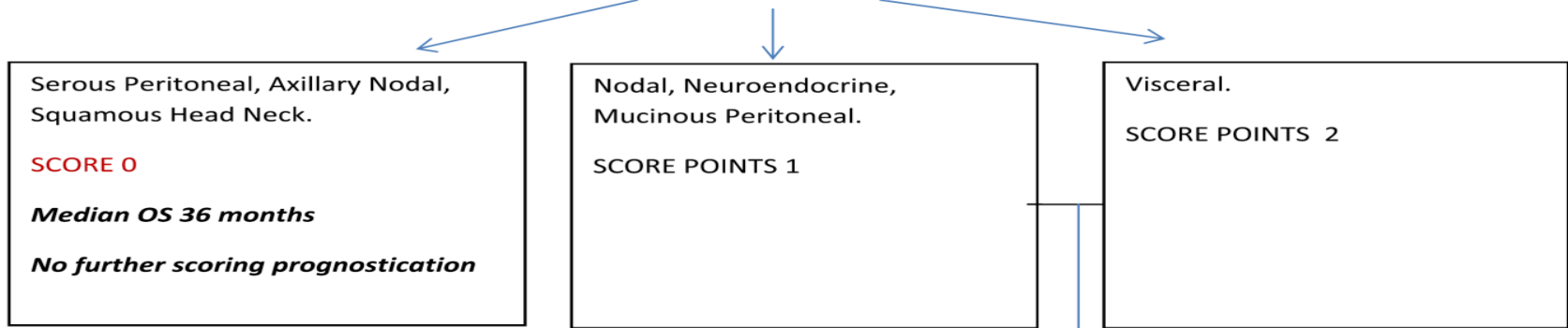


MedOS
37 m
21 m

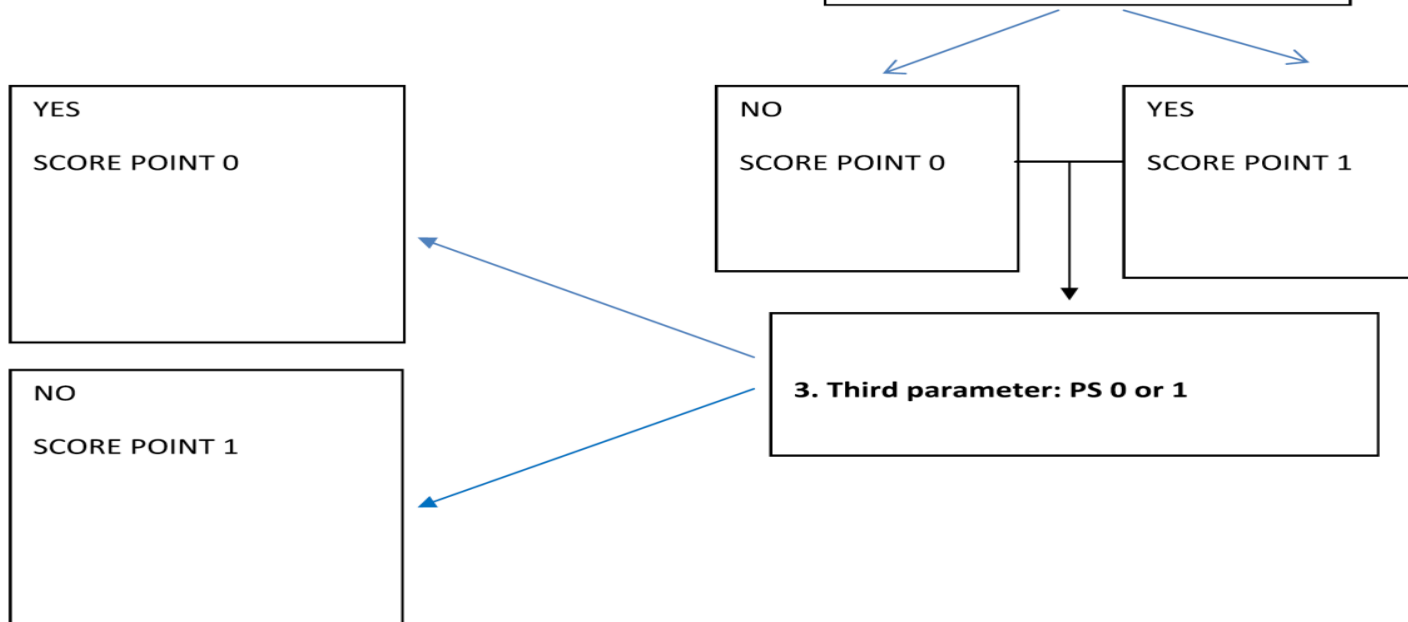


PROGNOSTICATION AND TREATMENT

1. First Parameter: CUP Clinicopathologic Subgroup



2. Second parameter: Leykocytosis (WBC>10,000/mm³)



3. Third parameter: PS 0 or 1

SUMMARY OF RISK GROUPS AND MEDIAN SURVIVAL

- | | | |
|-----------|---------------------|--------------|
| • SCORE 0 | MEDIAN OS 36 months | LOW RISK |
| • SCORE 1 | MEDIAN OS 14 months | INTERMEDIATE |
| • SCORE 2 | MEDIAN OS 11 months | RISK |
| • SCORE 3 | MEDIAN OS 8 months | HIGH |
| • SCORE 4 | MEDIAN OS 5 months | RISK |

| Chemotherapy (mg/m ²) | Time | Interval | Comments |
|-----------------------------------|-----------|-----------|--|
| Cisplatin 60–75 | Day 1 | Q 3 weeks | Fit patients, adequate hydration |
| Gemcitabine 1000 | Day 1 + 8 | | |
| Cisplatin 75 | Day 1 | Q 3 weeks | Fit patients with neuroendocrine feature-CUP, adequate hydration |
| Etoposide 100 | Days 1–3 | | |
| Paclitaxel 175 | Day 1 | Q 3 weeks | Convenient outpatient regimen, monitor neurotoxicity |
| Carboplatin AUC 5 | | | |
| Docetaxel 75 | Day 1 | Q 3 weeks | Convenient outpatient regimen, monitor neurotoxicity |
| Carboplatin AUC 5 | | | |
| Irinotecan 160 | Day 1 | Q 3 weeks | Outpatient regimen, monitor for neurotoxicity and diarrhoea |
| Oxaliplatin 80 | | | |
| Oral Capecitabine 2000 ± | Days 1–14 | Q 3 weeks | Outpatient regimen, risk for diarrhea and neurotoxicity |
| Oxaliplatin 85–130 | Day 1 | | |
| Gemcitabine 1000/Irinotecan 100 | Day 1+8 | Q 3 weeks | Convenient outpatient regimen, monitor diarrhoea |



Meta-analysis

Golfinopoulos et al, Cancer Treat Rev 2009

Multiple-treatments meta-analysis results for death.

| | HR ^a | 95% CrI |
|-----------------------------------|-----------------|-----------|
| nPnTc vs. nPnTm | 1.01 | 0.59–1.72 |
| Platinum vs. nPnTm | 0.69 | 0.39–1.28 |
| Taxane vs. nPnTm | 0.66 | 0.22–2.08 |
| Platinum plus taxane vs. nPnTm | 0.81 | 0.34–1.89 |
| Platinum vs. nPnTc | 0.69 | 0.43–1.15 |
| Taxane vs. nPnTc | 0.66 | 0.23–2.00 |
| Platinum plus taxane vs. nPnTc | 0.80 | 0.39–1.67 |
| Taxane vs. platinum | 0.95 | 0.37–2.5 |
| Platinum plus taxane vs. platinum | 1.16 | 0.56–2.38 |
| Platinum plus taxane vs. taxane | 1.22 | 0.36–4.00 |

^a A hazard ratio (HR) above one means that the risk of death is higher with the first rather than second listed regimen; nPnTm, non-platinum, non-taxane monotherapy; nPnTc, non-platinum, non-taxane combination.



- **SHOULD WE TARGET CUP
GENETIC ABERRATIONS?**

**SHOULD WE ADMINISTER
TARGETED THERAPY
ACCORDING TO THE
PRIMARY SITE?**

Actionable genetic alterations in CUP-1

- **N=1350 patients with CUP**
- **Caris Life Sciences, Phoenix, USA**
- Mutational analyses, in-situ hybridisation, immunohistochemistry, (RT)-qPCR.
- Actionable biomarkers were identified in **77%**:
 - Steroid receptor IHC expression
 - MET IHC expression
 - PTEN loss of IHC expression,
 - Activating mutations in EGFR, BRAF, PIK3CA
 - Gene copy number variations in HER2, TOP2A and MET.

Targeted next-generation sequencing (NGS) of carcinoma of unknown primary site (CUP): Actionable genomic alterations (GA) and new routes to targeted therapies. Jeffrey S. Ross, Kai Wang, Geoff Otto, Gary A. Palmer, et al
J Clin Oncol 32:5s, 2014 (suppl; abstr 11048)

- Hybridization capture of exons/introns from 236 cancer-related genes was applied to DNA extracted from 200 CUP FFPE specimens.
- **169 (85%) CUP had at least 1 actionable GA (2.00 actionable GA/CUP).**
- The most common actionable GA were **KRAS (25%), CDKN2A (19%), MCL1 (8-10%), PTEN (8%), PIK3CA (8%), BRAF (6% in ACUP and 2% in non-ACUP) and NF1 (5%).**
- Mutations, amplifications and rearrangements of ERBB2 (10%), EGFR (8%) and BRAF (6%) were common in ACUP but not present in non-ACUP.

Molecular gene expression profiling to predict the tissue of origin and direct site-specific therapy in patients with carcinoma of unknown primary site (CUP): Results of a prospective Sarah Cannon Research Institute Trial

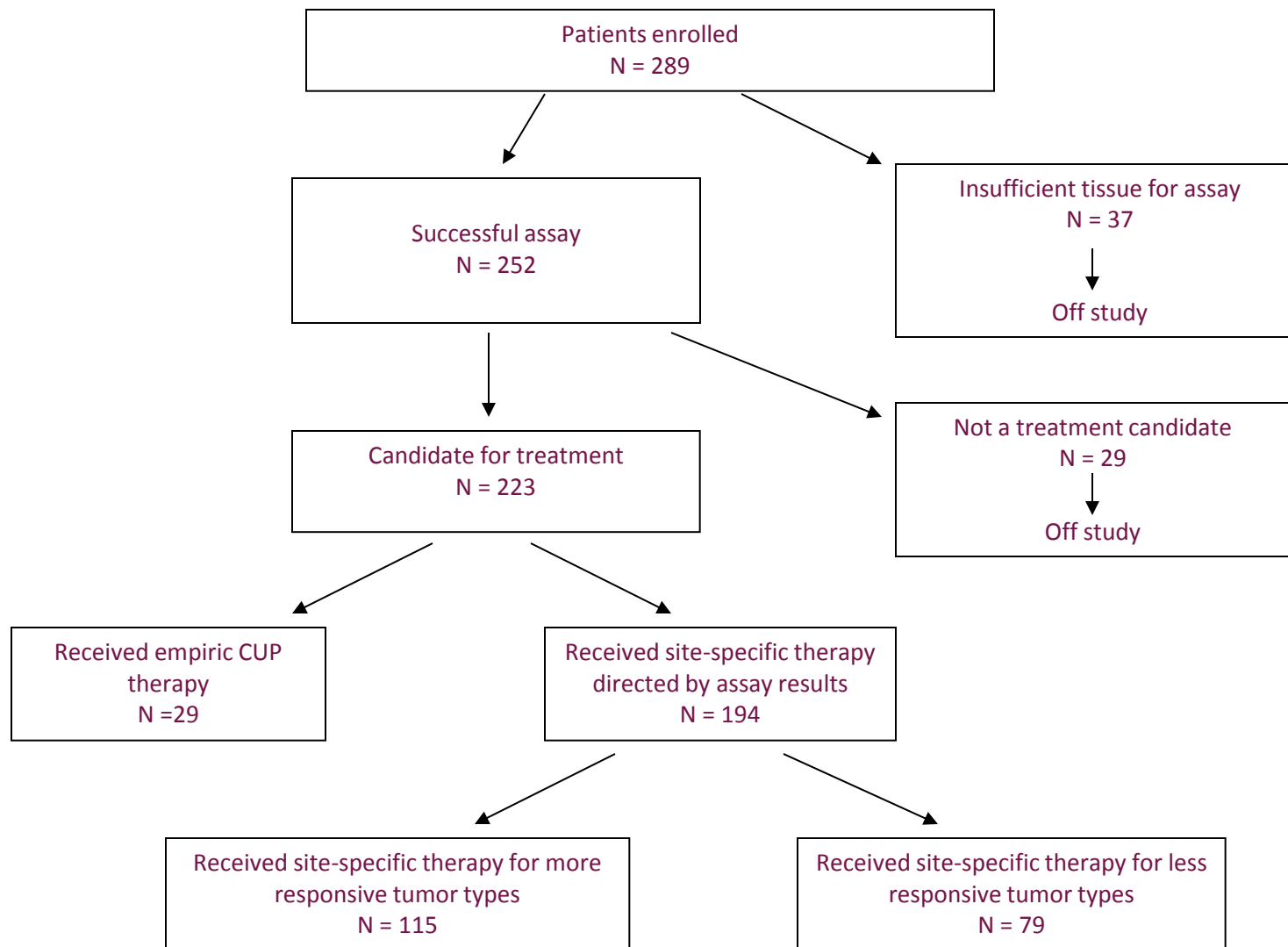
F. Anthony Greco, MD^{1,2}; Mark S. Rubin, MD^{1,3}; David R. Spigel, MD^{1,2}; Samuel Raby¹; Thabiso Chirwa¹; Raven Quinn, MS¹; Catherine A. Schnabel, Ph.D.⁴; Mark G. Erlander, Ph.D.⁴; John D. Hainsworth, MD^{1,2}

¹Sarah Cannon Research Institute (SCRI), Nashville, TN ; ²Tennessee Oncology, PLLC, Nashville, TN; ³Florida Cancer Specialists/SCRI, Ft Myers, FL; ⁴bioTheranostics, Inc., San Diego, CA

bioTheranostics CancerTypeID 92-gene test



Patient Flow Diagram



Survival in 223 Treated Patients and in Subsets

| | <u>Patient Group</u> | <u>Number</u> | <u>Median survival (mo.)</u> |
|--------------------------|---------------------------|---------------|------------------------------|
| All treated | | 223 | 10.8 |
| Assay-directed treatment | | 194 | 12.5, p=0.02 |
| Empiric treatment | | 29 | 4.7 |
| Tumor type* | | | |
| | Treatment responsive | 115 | 13.4, p=0.04 |
| | Less treatment responsive | 79 | 7.6 |
| Individual tumor types | | | |
| | Biliary tract | 45 | 6.8 |
| | Pancreas | 12 | 8.2 |
| | Colorectal | 26 | 12.5 |
| | NSCLC | 23 | 15.9 |
| | Ovary | 10 | 29.6 |
| | Breast | 10 | NYR (>24) |

NYR = not yet reached; *Includes 194 patients who received assay-directed treatment

Real life questions

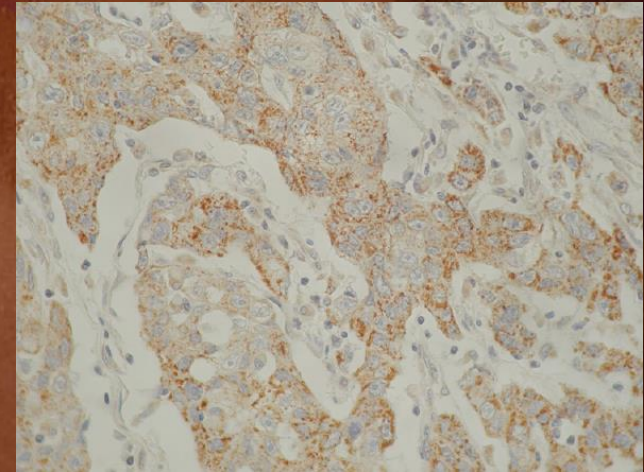
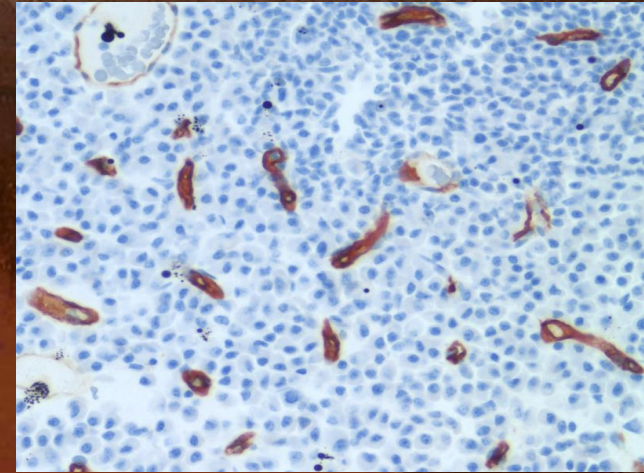
- Will Molecular Profiling Assays improve patient outcome by administration of Primary Tissue-specific therapy?
- Does a CUP biologically assigned as X Cancer behave similarly to Typical Metastatic X cancer?
- What about comparing CUP metastases to matched Known Primary Metastases (KPM)?

More

- Should we screen for actionable Genetic Aberrations in CUP?
- If we find such, how do we gain access to the targeted agents?
- Even if we have access to the targeted agents, how do we know they will work?
- How can we ignore the microenvironment and systems biology?
- Example: Targeting mutant BRAF in melanoma vs Colorectal Cancer

Angiogenesis in CUP

- Rashid et al, ASCO 2005:
IHC VEGF expression in 49% of 76 CUP.
- Karavasilis et al, BMC Cancer 2005:
N=81.
IHC VEGF expression in 100%.
- **Angiogenesis is active in CUP, though this is a feature common for metastatic solid tumours in general.**



Mutational profiling in CUP

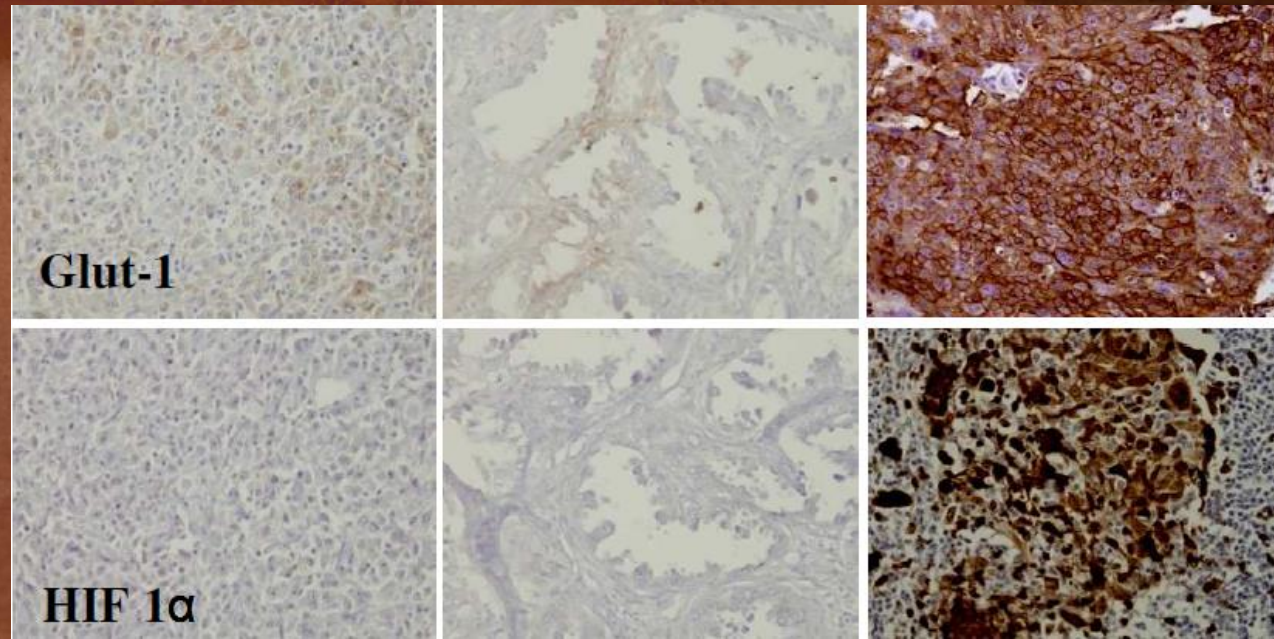
| N=87 | | Parameter | Category (Reference: Activating mutation) | HR | p |
|--------------------|----------------------------|---|--|------|-------|
| N (%) | | | | | |
| KRAS exon 2 | Activating Inactivating | | | | |
| BRAF exon 15 | Activating Inactivating | | | | |
| PIK3CA exons 9, 20 | Activating Inactivating | MET_exon_18 | inactivating mutation | 0.43 | 0.34 |
| CTNNB1 exons 1,3,5 | Activating Inactivating | | Wild type | 0.20 | 0.001 |
| MET exon 18 | Activating Inactivating | CTNNB1_exons_1_3_5 | inactivating mutation | | |
| | | | Wild type | 0.46 | 0.46 |
| | | | | 0.70 | 0.27 |
| | | Activating MET or CTNNB1 mutation | no activating mutation | | |
| | | | | 0.47 | 0.02 |

Kotteas et al, Clin Exp Metas 2014, submitted

Kotteas et al, Clin Exp
Metas 2014, submitted

Hypoxia

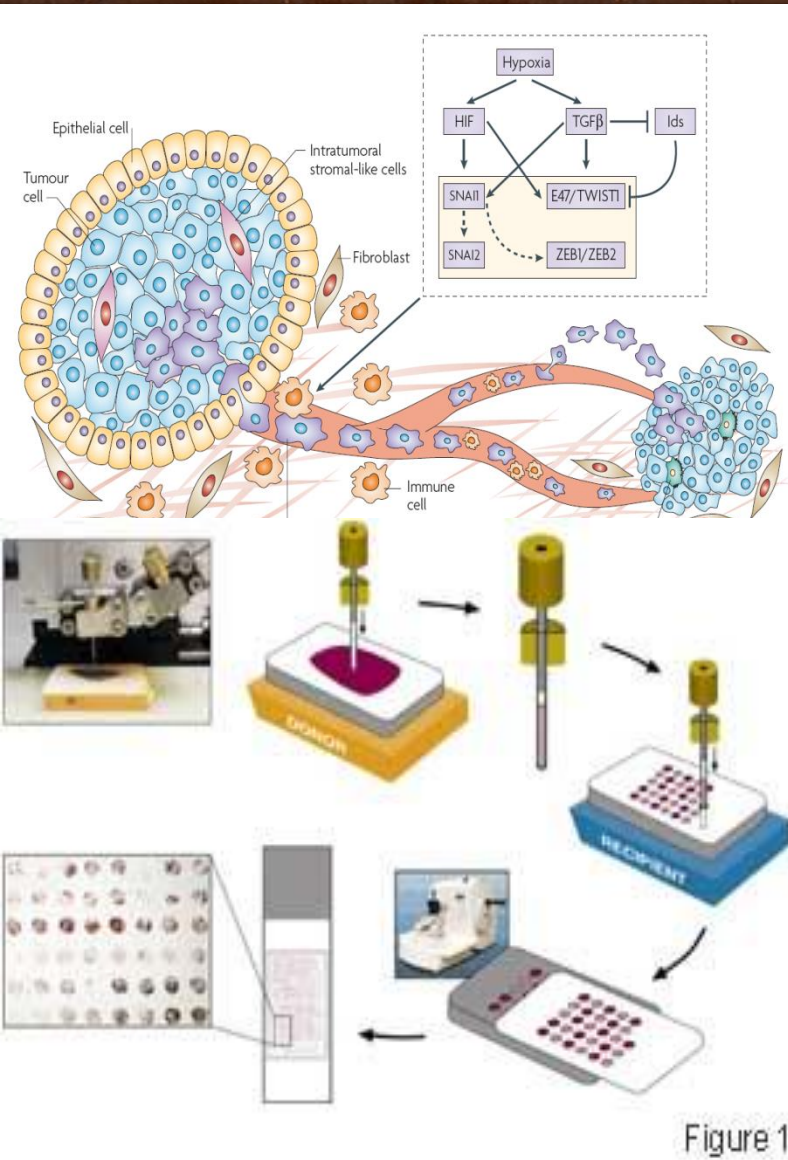
- Koo J et al, Cancer Cell Biol 2010
- TMA IHC study of hypoxia related proteins in 69 CUP
- Hypoxia phenotype (GLUT1, HIF1 α , COX2) present in 25% of CUP, associated with poor prognosis



Metabolic phenotypes of CUP

- N=77 CUP
- TMA IHC expression of metabolic enzymes
 - GLYCOLYSIS Glut-1, CA IX, MCT4;
 - GLUTAMINOLYSIS GLS1, GDH, ASCT2;
 - MITOCHONDRIAL ATP synthase, SDHA, SDHB
- **More visceral metastases with glutaminolytic activity in tumour stroma ($p=0.003$)**
- Kim HM, *Journal of Translational Medicine* 2014, 12:2

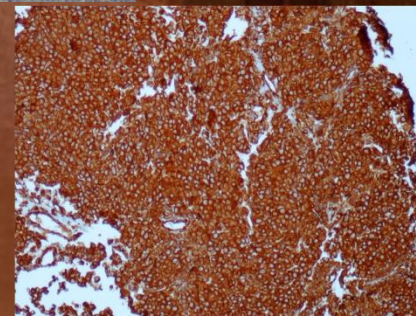
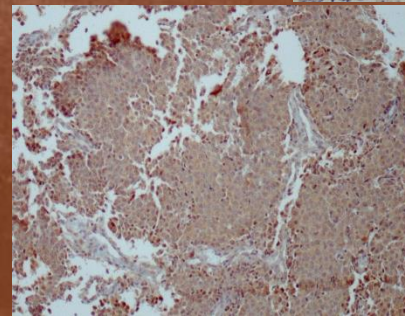
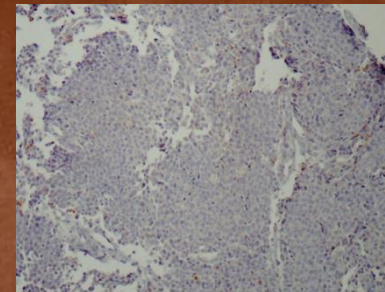
3. EPITHELIAL MESENCHYMAL TRANSITION AND STEMNESS in 100 CUP



• IHC:

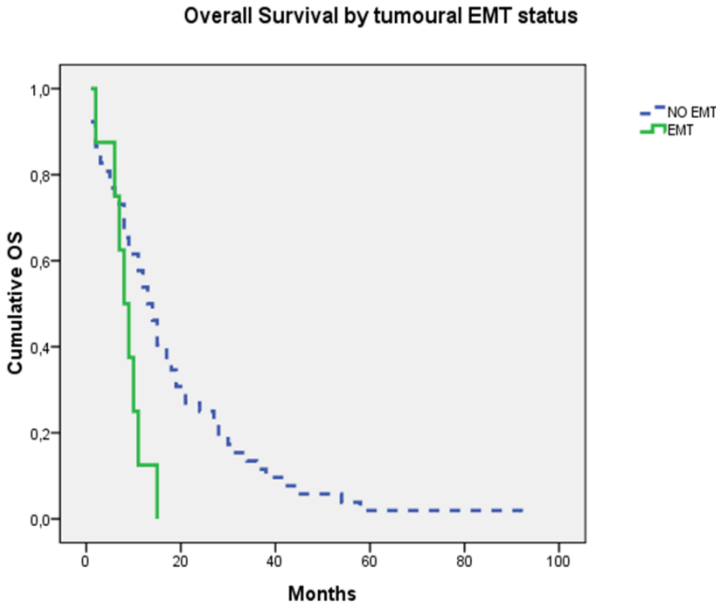
EMT phenotype:

E-Cadherin,
Snail, N-Cadherin, Vimentin



| | EMT present | EMT absent | p |
|-------------------------|-------------|------------|-------|
| Characteristics | | | |
| Male gender | 8(17.4) | 38(82.6) | 0.002 |
| Female gender | 0 (0) | 52 (100) | |
| High grade | 6(13.6) | 38(86.4) | 0.05 |
| Low grade | 2 (4.5) | 42 (95.5) | |
| Visceral metastases | 2(13.3) | 13(86.7) | 0.05 |
| Non-visceral mets | 3 (5.1) | 55 (94.9) | |
| | | | |
| Response | | | |
| CR or PR | 3(7.5) | 37(92.5) | 0.893 |
| NO CR or PR | 3(8.3) | 33(91.7) | |
| | | | |
| PFS(months) | | | |
| Median | 5 | 8 | 0.112 |
| 95% CI | 3.0-7.0 | 5.3-10.7 | |
| | | | |
| Survival(months) | | | |
| Median | 8 | 13 | 0.023 |
| 95% CI | 5.2-10.7 | 9.8-16.1 | |
| 1-year (%) | 0 | 75 | |
| | | | |
| EMT H-SCORE | | | |
| | | | |
| PFS | | | |
| Median (months) | 5 | 8 | 0.541 |
| 95% CI | 5.6-6.3 | 5.4-10.5 | |
| | | | |
| Survival | | | |
| Median | 9 | 14 | 0.074 |
| 95% CI | 7.2-10.7 | 10.6-17.3 | |
| 1-year (%) | 7.7 | 72.3 | |

Stoyianni A et al,
Anticancer Res.
2012;32(4):1273-81



SHALL WE FIND ANYTHING WE DO NOT SEE IN METASTATIC TUMOURS?

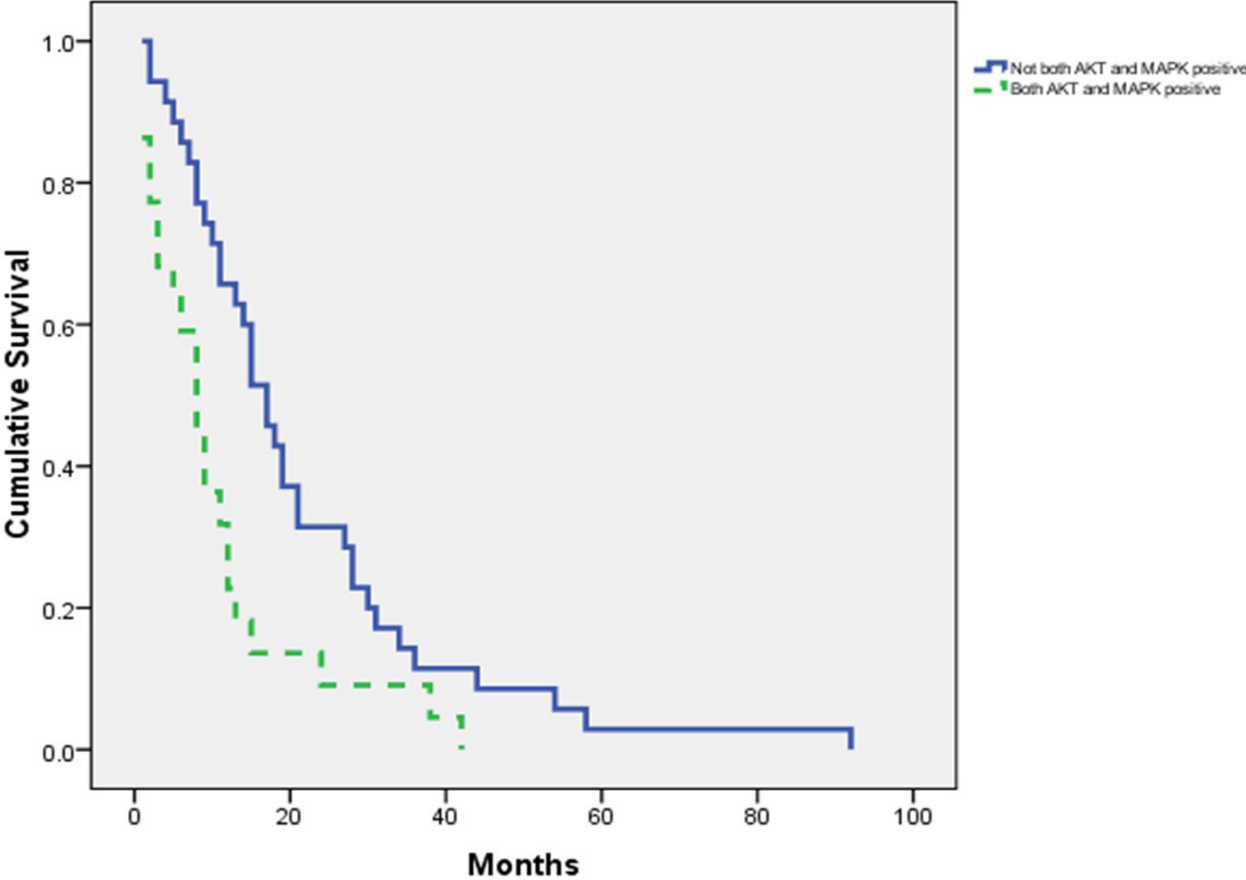
TMA PROFILING

- Profiling 100 CUP cases for protein and phosphoprotein expression of **12 key membrane receptors, signal transducers, transcription factors hypoxia and cell cycle regulators**



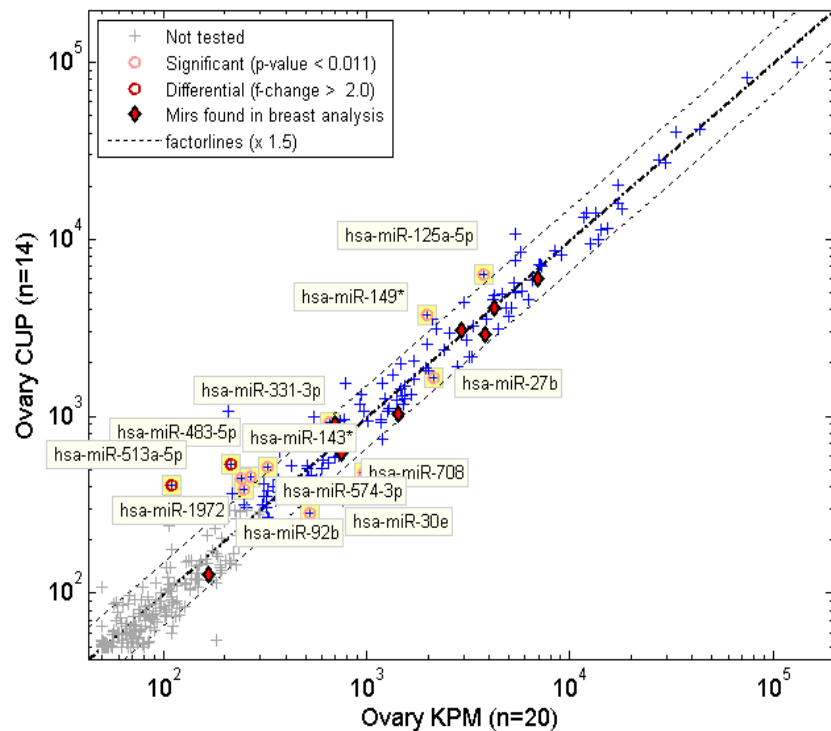
Golfinopoulos et al, Ann Oncol. 2012;23(10):2725-30
Krikelis et al, Clin Exp Metastasis. 2012;29(6):603-14

Overall survival in patients with CUP with both high pAKT and pMAPK IHC expression



| Parameter | Hazard Ratio for risk of death | 95% CI | 2-sided p |
|------------------|--------------------------------|-------------|-----------|
| P21 | 0.34 | 0.16 – 0.73 | 0.005 |
| cMET | 0.48 | 0.26-0.91 | 0.025 |
| pAKT Thr308 | 2.39 | 1.23 – 4.66 | 0.01 |
| pS6RP Ser235/236 | 2.76 | 1.31 – 5.84 | 0.008 |

Ovary CUP vs. Ovary KPM



- Metastases from primary tumours (K)

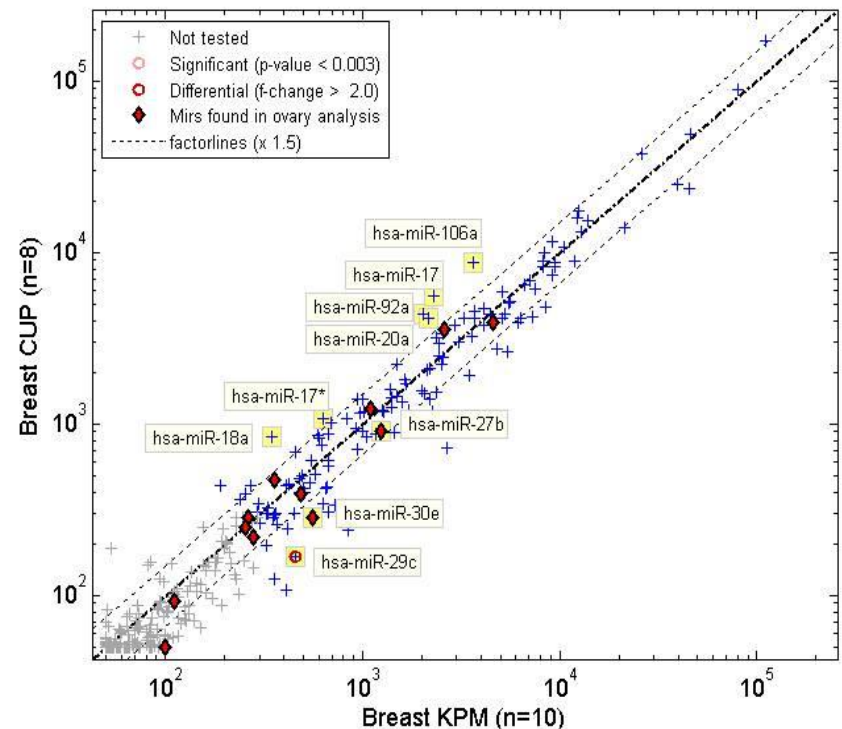
• IS THERE A miRNA C

Pentheroudakis et al, Clin Exp Metastasis 2013;30(4):431-9

GNATURE

ession of a library of
ween:

Breast CUP vs. Breast KPM



Conclusions

- **No trace of a «CUP biologic signature», distinct from KPM**
- **Activated pathways in CUP:**
 - Angiogenesis with concurrent hypoxia
 - Stromal glutaminolytic activity
 - AKT/S6RP axis with deficient apoptosis
 - b-Catenin/Wnt axis
 - MET axis
 - EMT activity

Target them!



- ***Is it simply metastases from a missed primary?***
Find the primary, administer primary tumour-specific therapy.
- ***Or is it truly a disease with a unique molecular signature that defines its metastatic propensity?***
- Target the genetic aberrations.



Current Clinical Research: CUP Trials

| Trial | Sponsor | Design | Arms |
|------------------|--|--------------------------------|--|
| CUP ONE | NCRI/CRUK | <i>RCT Phase II</i> | ECX vs ECX+Vandetanib |
| MIRVIEW | TEVA/ROSETTA | <i>Parallel Cohort</i> | ClinPath work up vs ClinPath work up+MirView |
| UNKPRI20 | Sarah Cannon RC, USA | <i>Single arm Phase II</i> | Therapy based on molecular profiling |
| UNKPRI21 | Sarah Cannon RC, USA | <i>RCT Phase II</i> | Carboplatin Paclitaxel vs CP+Belinostat |
| GEFCAPI04 | GEFCAPI, France | <i>RCT Phase III</i> | CDDP+Gemcitabine vs Pathwork test-based therapy |
| PACET-CUP | Ruprecht-Karls- University of Heidelberg | <i>RCT Phase II</i> | Carboplatin Paclitaxel vs CP+Cetuximab |
| CUP02 | CRUK | <i>RCT Phase III</i> | ECX vs Molecularly Assigned Therapy |

Evaluation and Management of Possible CUP Patient

