



# MicroRNA Profiling of a Familial Primary Pulmonary Enteric Adenocarcinoma

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- pulmonary enteric adenocarcinoma (PEAC) is a rare lung adenocarcinoma with a predominant component (>50%) of intestinal differentiation and tumor cells positive for at least one intestinal marker (CDX2, CK20, MUC2)
- only 31 cases in literature and occurrence in two family members has never been reported before
- PEAC should be distinguished from metastatic colorectal carcinoma because of the different treatment strategy and prognosis
- due to its rarity and peculiarity, no specific guidelines for therapy exist

## **Aim of the study**

- molecular and histological characterization of a PEAC from a patient with another family member affected by the same tumor

# Methods

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**ORIGINAL PAPER**

## MicroRNA profiling for the identification of cancers with unknown primary tissue-of-origin

Manuela Ferracin,<sup>1,2</sup> Massimo Pedriali,<sup>1</sup> Angelo Veronese,<sup>1,3</sup> Barbara Zagatti,<sup>1</sup> Roberta Gafà,<sup>1</sup> Eros Magri,<sup>1</sup> Maria Lunardi,<sup>1</sup> Gardenia Munerato,<sup>1</sup> Giulia Querzoli,<sup>1</sup> Iva Maestri,<sup>1</sup> Linda Ulazzi,<sup>1</sup> Italo Nenci,<sup>1</sup> Carlo M Croce,<sup>3</sup> Giovanni Lanza,<sup>1</sup> Patrizia Querzoli<sup>1\*</sup> and Massimo Negrini<sup>1,2\*</sup>

The molecular characteristics of proband's PEAC were evaluated applying a previously validated 47-miRNA diagnostic classifier to predict tumor primary origin.

It has high overall accuracy of 100% for primary cancers and 78% for metastases.

**16<sup>TH</sup> WORLD CONFERENCE ON LUNG CANCER**

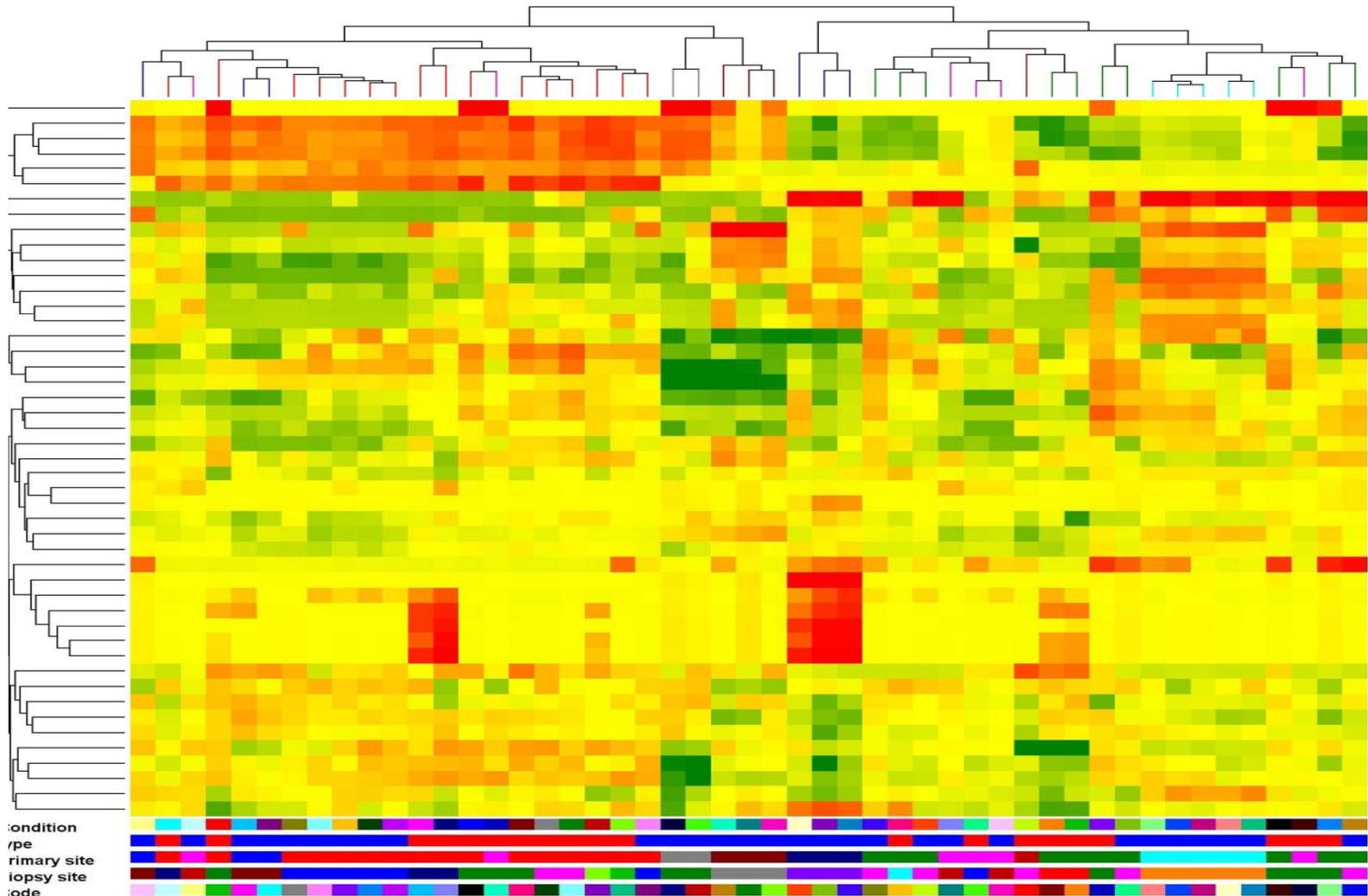
# Results

Characteristic	Proband	His sister
General		
Age at diagnosis (years)	68	71
Smoking habit	Yes	Yes
Primary localization	lung: right/lower lobe	lung: right/lower lobe
Colonoscopy	multiple benign polyps (N<10), diverticulosis	multiple benign polyps (N<10), diverticulosis
Radical surgery (R0)	Yes	Yes
Metastases/DFS (months)	bone / 1.5	lung, adrenal glands / 12
Histological findings		
Histology	PEAC	PEAC
Stage (pTNM)	pT2apN1	pT2apN0
IHC analyses		
TTF-1	NEG	NEG
Napsin A	NEG	NA
CK7	POS	NEG
CK20	NEG	POS
CDX2	POS	POS
MUC1	POS	NA
MUC2	NEG	NA
MUC5A	POS	NA
MUC6	NEG	NA
Genetic aberrations		
EGFR	wild type	wild type
K-RAS	Gly12Asp mutation	Gly12Asp mutation
ALK	wild type	wild type

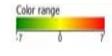
## Familial aggregation of PEAC was associated with:

- **similar clinicopathological features**
  - age at diagnosis
  - smoking-habit
  - tumor localization
  - multiple benign colon polyps
- **histologic findings**
  - TTF-1 negativity
  - CDX2 positivity
- **genetic findings**
  - K-RAS(Gly12Asp) mutation
  - no EGFR/ALK aberrations

- miRNA profiling revealed main similarities with lung cancer (75.98%), and partial overlap with pancreatic cancer (23.34%), but not with colorectal cancer (less than 0.5%)
- PEAC shares with pancreatic cancer key miRNAs associated with tumor aggressiveness:
  - miR-31
  - miR-126
  - miR-506
  - miR-508-3p
  - miR-514



hsa-miR-122  
 hsa-miR-194  
 hsa-miR-192  
 hsa-miR-215  
 hsa-miR-194\*  
 hsa-miR-552  
 hsa-miR-205  
 hsa-miR-31  
 hsa-miR-204  
 hsa-miR-30a  
 hsa-miR-30a\*  
 hsa-miR-363  
 hsa-miR-149  
 hsa-miR-181a-2\*  
 hsa-miR-340\*  
 hsa-miR-375  
 hsa-miR-135b  
 hsa-miR-141  
 hsa-miR-200c  
 hsa-miR-96  
 hsa-miR-182  
 hsa-miR-183  
 hsa-miR-126\*  
 hsa-miR-210  
 hsa-miR-342-3p  
 hsa-miR-485-5p  
 hsa-miR-873  
 hsa-miR-193a-3p  
 hsa-miR-30c  
 hsa-miR-361-5p  
 hsa-miR-31\*  
 hsa-miR-211  
 hsa-miR-510  
 hsa-miR-508-3p  
 hsa-miR-509-3p  
 hsa-miR-506  
 hsa-miR-514  
 hsa-miR-10a\*  
 hsa-miR-187\*  
 hsa-miR-650  
 hsa-miR-200a\*  
 hsa-miR-200b\*  
 hsa-miR-10a  
 hsa-miR-200a  
 hsa-miR-200b  
 hsa-miR-145  
 hsa-miR-146a



condition  
 type  
 primary site  
 biopsy site  
 mode

[t, gastric, gastric, PF085]  
 [m, colon, ovary, PF083]  
 [t, pancreas, pancreas, PF065]  
 [m, colon, liver, PF035]  
 [t, gastric, gastric, PF028]  
 [t, gastric, gastric, PF029]  
 [t, colon, colon, PF026]  
 [t, colon, colon, PF062]  
 [t, colon, colon, PF043]  
 [t, colon, colon, PF045]  
 [t, colon, colon, PF060]  
 [m, colon, ovary, PF063]  
 [m, colon, ovary, PF084]  
 [m, colon, liver, PF044]  
 [m, pancreas, liver, PF068]  
 [m, colon, liver, PF046]  
 [m, colon, lung, PF059]  
 [m, colon, lung, PF061]  
 [m, colon, peritoneum, PF102]  
 [m, colon, skin, PF101]  
 [t, colon, colon, PF064]  
 [t, liver, liver, PF030]  
 [t, liver, liver, PF031]  
 [t, kidney, kidney, PF071]  
 [t, kidney, kidney, PF027]  
 [t, kidney, kidney, PF032]  
 [t, skin(melanoma), skin(melanoma), PF076]  
 [t, skin(melanoma), skin(melanoma), PF075]  
 [t, skin(melanoma), skin(melanoma), PF077]  
 [t, lung, lung, PF052]  
 [m, lung, lymphnode, PF053]  
 [t, lung, lung, PF051]  
 [t, pancreas, pancreas, PF066]  
 [m, pancreas, colon, PF070]  
 [t, pancreas, pancreas, PF067]  
 [m, unknown, lung, B001]  
 [m, lung, brain, PF010]  
 [m, lung, brain, PF033]  
 [m, lung, liver, PF048]  
 [t, lung, lung, PF047]  
 [t, prostate, prostate, PF100]  
 [t, prostate, prostate, PF093]  
 [t, prostate, prostate, PF094]  
 [t, prostate, prostate, PF105]  
 [t, prostate, prostate, PF106]  
 [m, lung, liver, PF050]  
 [m, pancreas, liver, PF069]  
 [m, lung, liver, PF096]  
 [t, lung, lung, PF003]

type  
 m  
 t

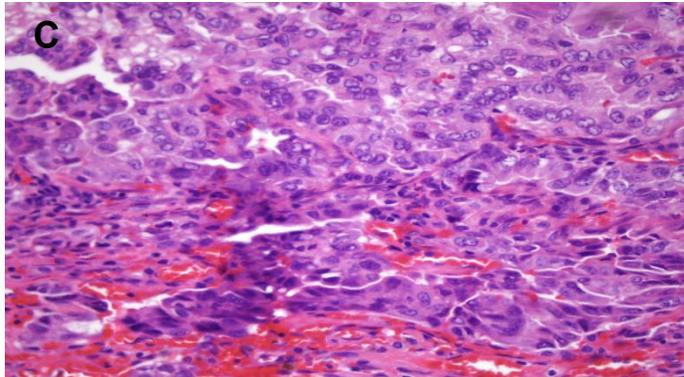
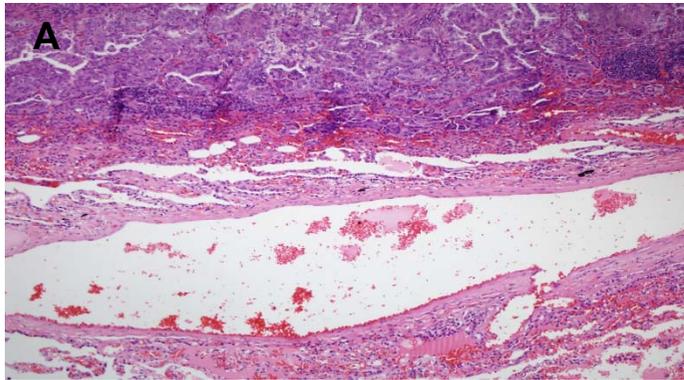
primary site  
 colon  
 gastric  
 kidney  
 liver  
 lung  
 pancreas  
 prostate  
 skin(melanoma)  
 unknown

biopsy site  
 brain  
 colon  
 gastric  
 kidney  
 liver  
 lung  
 lymphnode  
 ovary  
 pancreas  
 peritoneum  
 prostate  
 skin  
 skin(melanoma)

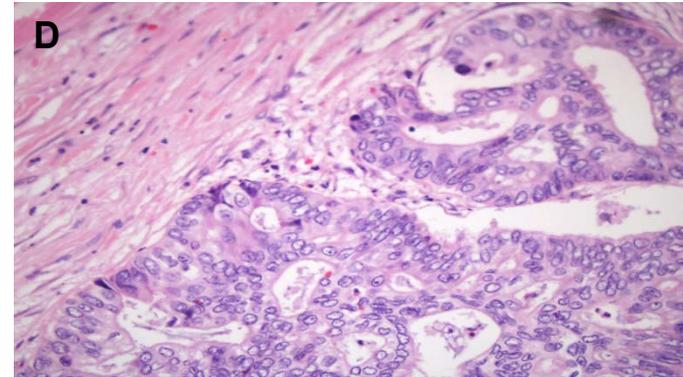
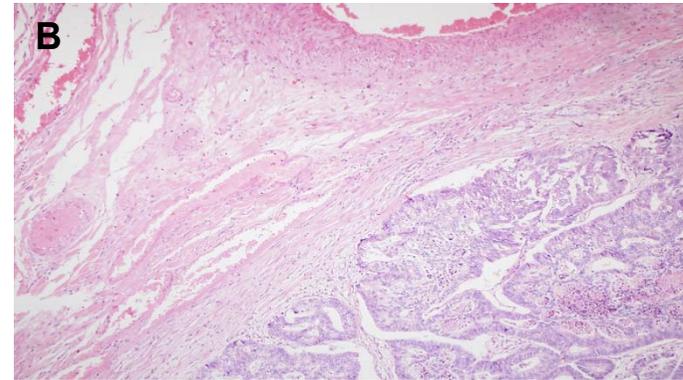


**Histological analyses revealed the presence of two different phenotypes in the lung lesion of the proband: NSCLC (30%) and PEAC (70%).**

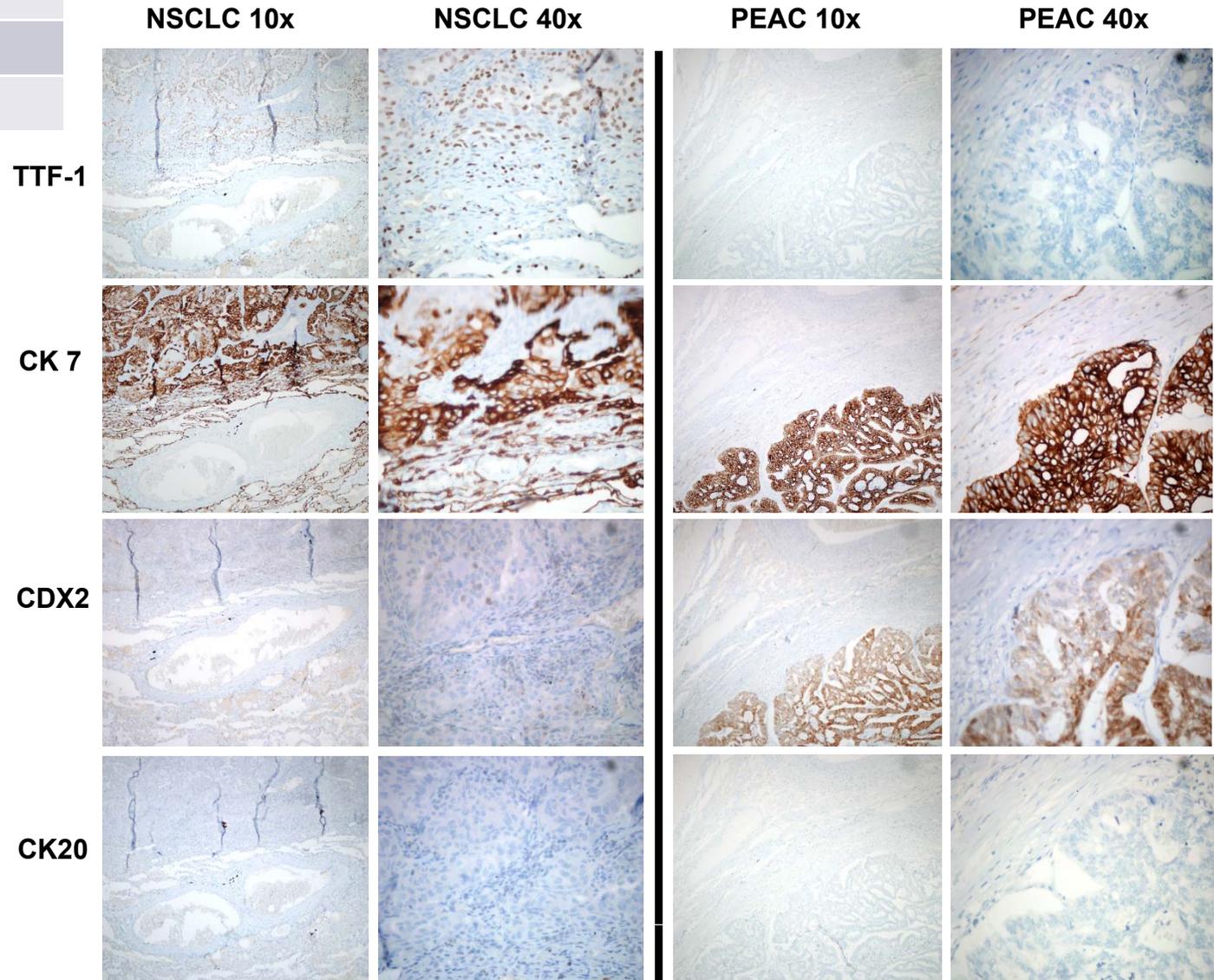
**NSCLC**



**PEAC**



IHC Marker	NSCLC Phenotype	PEAC Phenotype
TTF1	+	-
CK7	+	+
CDX2	-	+
CK20	-	-



## Conclusions

- we described, for the first time, a case of PEAC in two members of the same family, associated with similar clinicopathological feature and histological/genetic findings
- miRNA profiling of PEAC resembled mostly NSCLC, with a partial overlap with PDAC's pattern that could explain:
  - aggressive behavior compared to most NSCLC
  - guide future tailored-therapy approaches

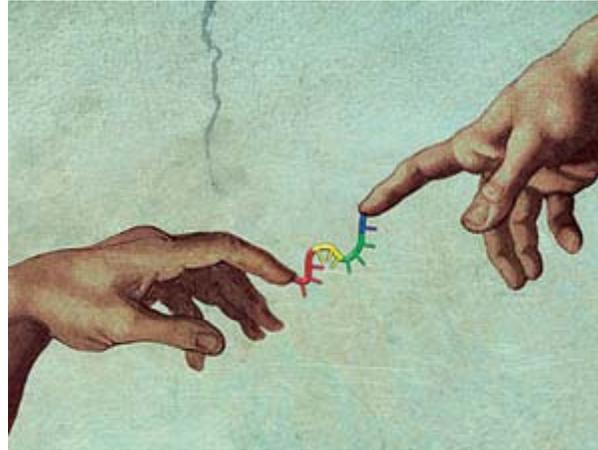
# Thank you for your attention!

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