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Overview of Circulating Biomarkers





4th Milan NET Conferences 2018

Eva Tiensuu Janson Professor of Medicine Endocrine Oncology Uppsala University







Network Adult Cancers (ERN EURACAN)

Uppsala University ospital — Swede



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ENETS Guidelines Biochemical Markers



Table 1. General and specific biomarkers currently used for the management of patients with neuroendocrine tumors

General tumor markers	Related indications
Chromogranin A	Almost all NETs (follow-up, limited in diagnosis)
Neuron-specific enolase	Atypical carcinoids, lung NEC, microcytoma
Pancreatic polypetide	Pancreatic NET
α-Subunit, β-hCG	Pancreatic, lung NET
Specific tumor markers	Related indications
Serotonin, 5-HIAA	Well differentiated NET
Gastrin	Zollinger-Ellison syndrome
Insulinoma	Insulin-secreting pancreatic NET
Glucagon, VIP, somatostatin	Well differentiated pancreatic NET
Catecholamines	Pheocromocytoma/paraganglioma
Calcitonin	Medullary thyroid cancer and pancreatic NET
PTHrp, ACTH, CRH, GHRH	Syndromes from (ectopic) mainly lung or pancreatic NET
NTpro-BPN (marker of ventricular dysfunction)	Carcinoid syndrome (carcinoid heart disease)

NET, neuroendocrine tumor; NEC, neuroendocrine carcinoma; 5-HIAA, 5-hydroxyindolacetic acid; VIP, vasoactive intestinal peptide; PTHrp, parathormone-related peptide; ACTH, adrenocorticotropin hormone; CRH, corticotropin-releasing hormone; GHRH, growth hormone-releasing hormone; NTpro-BPN, N-terminal pro-brain natriuretic peptide.



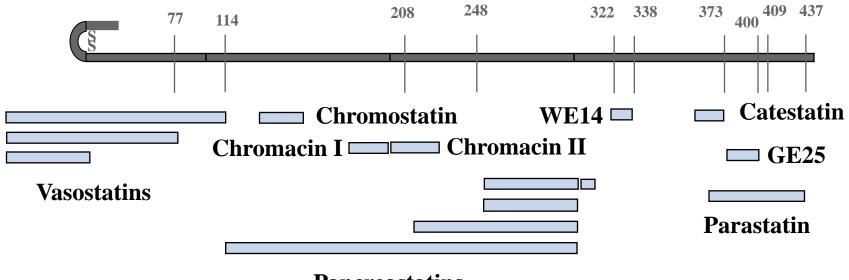


 A general marker can be found in many different (neuroendocrine) cells— indicates that the cell is of neuroendocrine type

 A specific marker can be found in one certain type of cell – indicates which kind of cell it is. (EC-cell, beta-cell, alpha-cell...)



Belongs to a family of acidic proteins which are produced in neuroendocrine cells and stored and secreted together with the specific hormones

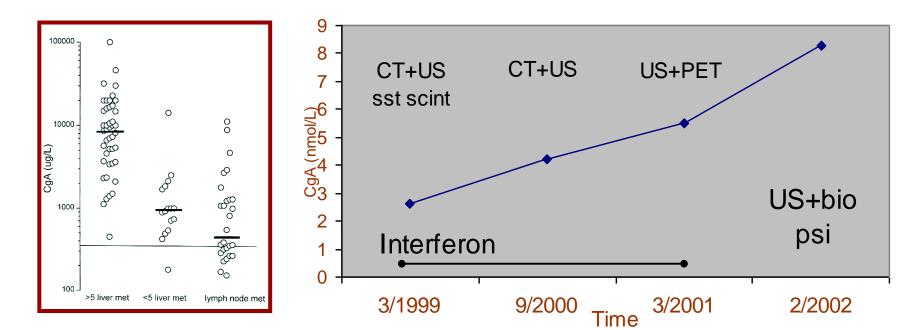


Pancreastatins









Chromogranin A levels are related to metastatic spread in untreated patients The chromogranin A level in plasma is a sensitive marker for recurrent disease in radically operated patients (8 mo *vs.* 32 mo)

Janson et al Ann Oncol 1997

Welin et al, Neuroendocrinology 2009



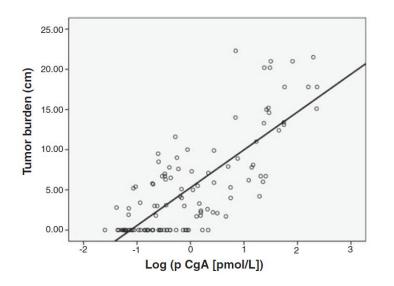
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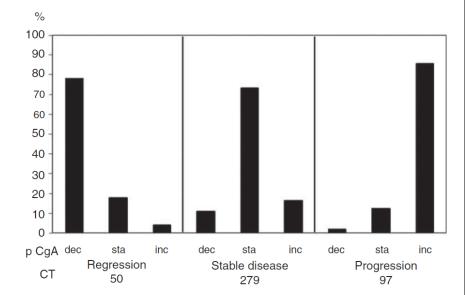


Chromogranin A is a sensitive marker of progression or regression in ileo-cecal neuroendocrine tumors

KENNETH HØJSGAARD JENSEN^{1,4}, LINDA HILSTED², CLAUS JENSEN³, TOMMIE MYNSTER^{1,5}, JENS F. REHFELD² & ULRICH KNIGGE¹

106 SI NET patients





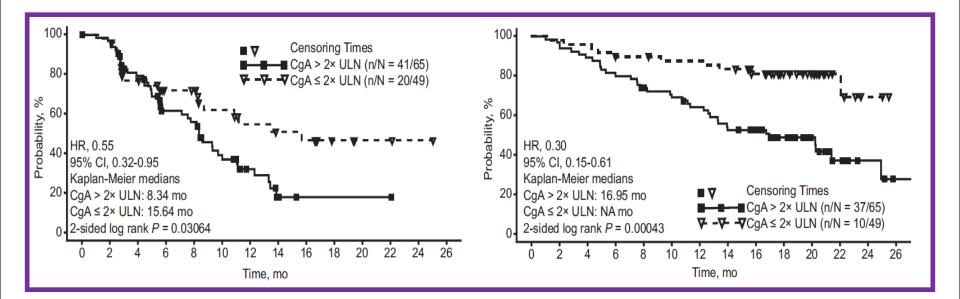
Scandinavian Journal of Gastroenterology. 2013; 48: 70-77



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Plasma Chormogranin A as prognostic marker in pNET - Radiant-1 study





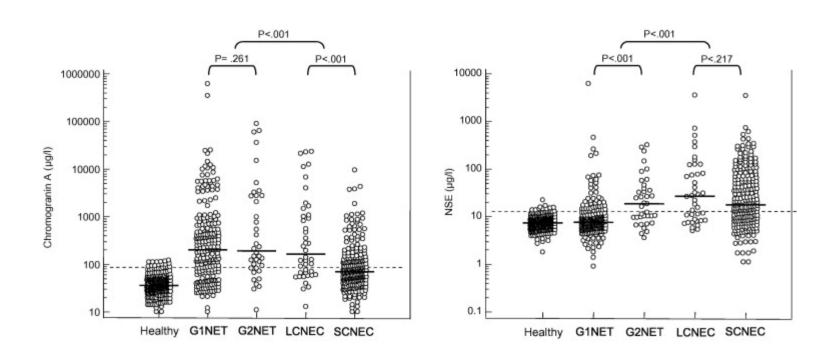
Progression-free survival

Overall survival

Yao et al. J Clin Endocrinol Metab, December 2011, 96(12):3741-







Chromogranin A is a good biomarker for G1-G2 NETs while NSE is better for high-grade tumors (G2 and NEC)

Korse CM et al. Eur J Cancer2012 Mar;48(5):662-71.

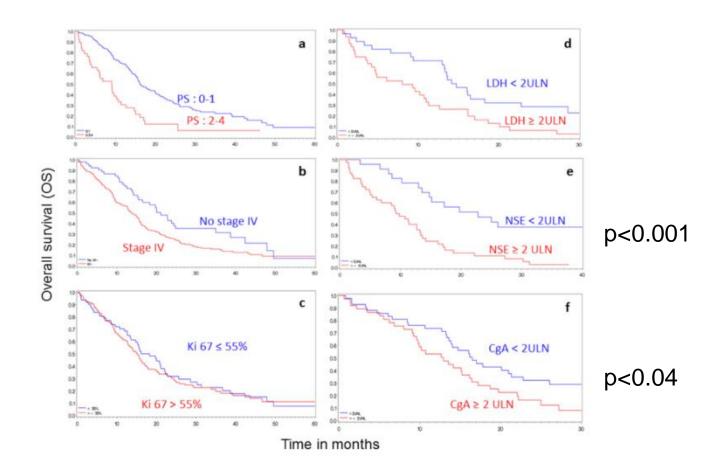


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Comparison of NSE and chromogranin A as predictors of survival in GEP-NEC





Walter T el al. Eur J Cancer 2017 Jul;79:158-165.



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- Impaired renal function
- Chronic Atrophic Gastritis
- Proton Pump Inhibitor treatment
- Impaired liver function
- Stress (increase adrenal medulla activity)
- Inflammatory bowl disease





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 A specific marker can be found in one certain type of cell – indicates which kind of cell it is. (EC-cell, beta-cell, alpha-cell...)



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Neuroendocrine cells in the GI-tract Specific markers



Cell type	Peptide hormone	Amine
Alpha	Glucagon	
Beta	Insulin	
Delta	Somatostatin	
PP	Pancreatic polypeptide	
Enterochromaffin (EC)	Tachykinins	Serotonin
Enterochromaffin-like (ECL)		Histamin
Gr	Ghrelin/obestatin	
G	Gastrin	
1	Cholecystokinin	
К	GIP	
Ν	Neurotensin	
S	Secretin	
VIP	Vasoactive intestinal peptide	
Х	Amylin	



Analytical and preanalytical validation of a new mass spectrometric serum 5-hydroxyindoleacetic acid assay as neuroendocrine tumor marker



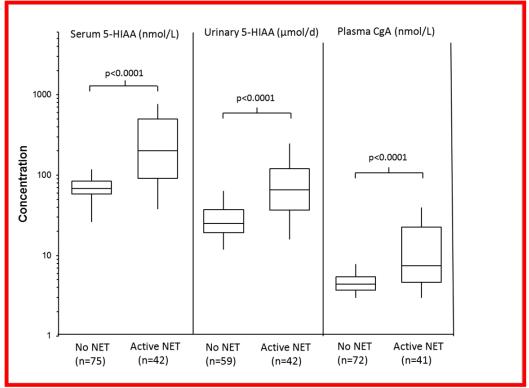
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Niina Tohmola ^{a,*}, Outi Itkonen ^b, Timo Sane ^c, Helene Markkanen ^b, Sakari Joenväärä ^a, Risto Renkonen ^{a,b}, Esa Hämäläinen ^b

Character	Mean (range)
Sex, M/F	44/44
Age, years	64 (20-85)
Origin of NET, n	
Foregut	4
Midgut	82
Hindgut	2
Clinical status of NET, n	
Clinical remission	46
Stable or progressive disease	42
Time from diagnosis of NET, years	
Patients in remission	4.6 (1-12)
Patients with stable or progressive disease	5.2 (0.5-22)

The assay for serum 5-HIAA discriminates between healthy individuals and patients with NET and is well suited for the diagnosis and follow-up of NETs

Clinica Chimica Acta 428 (2014) 38-43



Comparison betweenNET patients and healthy subjects



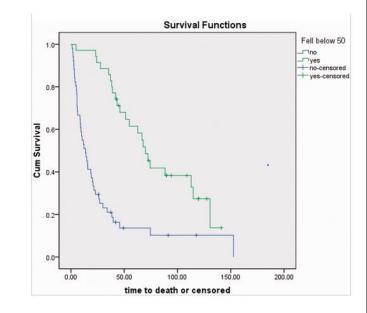




- Tachykinins have had a renaissance during the last few years produced in EC-cells (SI-NET specific biomarker)
- Tachykinins (NKA) can discriminate patients with severe carcinoid heart disease and is a specific marker for SI-NETs (Ardill et al, QJM 2016)

SI-NET patients who had a reduction in NKA from increased levels down to 50 ng/L or less during treatment had a significantly longer overall survival

	Number of patients	First NKA≥50 ng/L median (range)	Median (range) survival in months from first NKA≥50 ng/L	Significance between early and late
Total group NKA≥50 ng/L	86	91 (50-1250)	28.8 (2.0–152.6)	
Earlier period	35	116 (50-1250)	20.2 (2.1-152.6)	P=0.019
Later period	51	75 (52-1176)	39.1 (2.0-143.8)	
Earlier period (clinic patients)	32	101 (50-1250)	23.2 (2.0-152.6)	P = 0.016
Later period (clinic patients)	43	75 (52-1176)	52.1 (21.0-143.8)	



Published in: Joy ES Ardill; David R McCance; Wendy V Stronge; Brian T Johnston; *Ann Clin Biochem* 53, 259-264. DOI: 10.1177/0004563215592021 Copyright © 2015 Association for Clinical Biochemistry





- Biomarkers can be used to
 - Set a correct diagnosis
 - Estimate tumor burden
 - Identify tumor recurrence
 - Foresee prognosis
 - Predict therapy response

 So – we further biomarkers – but are those that we have any good?





- New proteins measurable in blood and in tumor tissue
- Circulating tumor cells
- miRNAs
- Genetic markers
- The NETest



UPPSALA UNIVERSITET Multiplex proximity ligation/extension assay



Novel Serum Biomarkers in Small Intestinal Neuroendocrine Tumors

Edfeldt et al Neuroendocrinology 2017

96 patients with metastasized SI-NETs + 23 controls were included
69 biomarkers were screened in serum using multiplex PLA
76 further biomarkers were analyzed by multiplex PEA

Results were confirmed in an extended cohort using IHC and ELISA

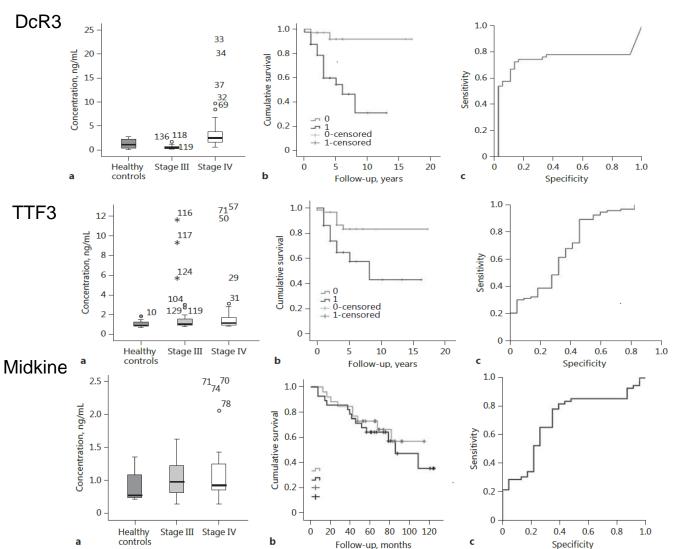


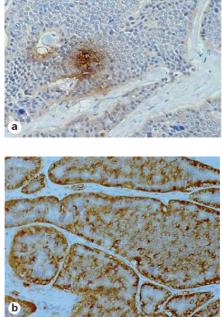
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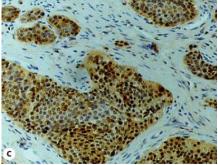
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Three proteins were of interest DcR3, TTF3 and Midkine







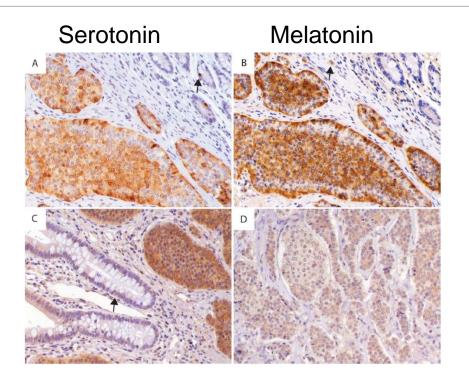




Melatonin in SI-NETs



- Melatonin IR was found in all 26 SI-NET patients examined
- Melatonin receptor MT1 was low or absent in all tumors
- Melatonin receptor MT2 was highly expressed in primary tumors but low in metastases
- Plasma levels ranged between 4.5 – 220.0 pg/L
- High plasma levels were associated with nausea (p= 0.027) and flush (p=0.020)



Melatonin Receptor 2 Primary Tumor Liver Metastases



So, where are we as far as proteins takes us?



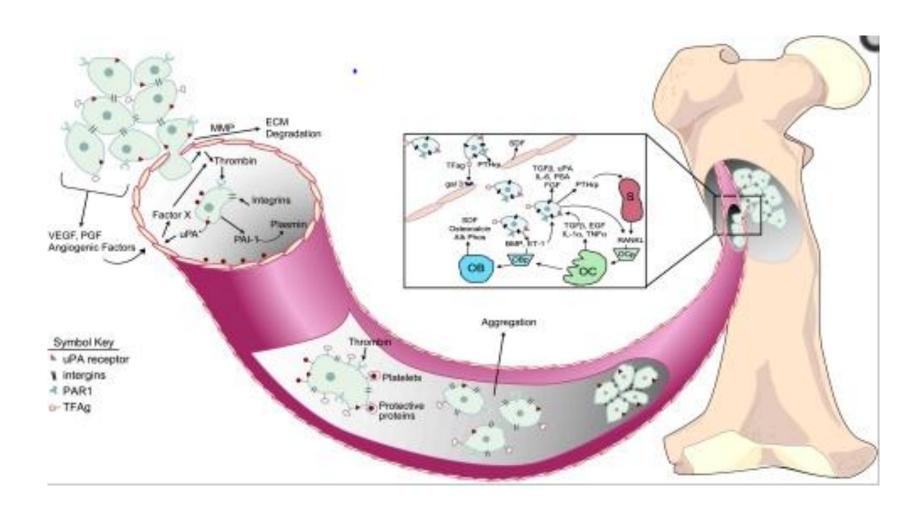
- DcR3
- TFF3
- Midkine
- Melatonin
- ...and many moore

- Several new potential protein biomarkers have been suggested
- Pros proteins are relatively easy to measure
- Cons none of the candidates have shown true clinical benefit yet





Circulating Tumor Cells







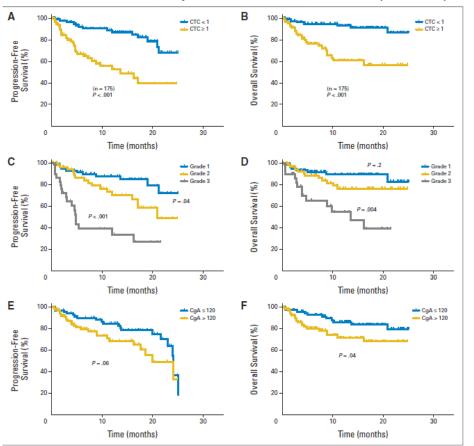
- CTCs can be correlated to PFS and OS in breast cancer (and other cancers)
- The CellSearch platform can detect CTCs in a 10
 ml blood sample
- NETs express EpCAM which is required for detection of CTCs by this platform



The number of CTCs which can be found in blood is correlated to survival



PFS and OS in patients with NET (n 175)



	No. of		PFS		OS		
Risk Factor	Patients	HR	95% CI	Р	HR	95% CI	Р
Baseline CTC count							
<1	50	1.0		_	1.0		_
≥1	33	5.0	1.3 to 18.5	(017)	7.2	1.3 to 39.4	.023
Baseline CgA, pmol/L				\smile			\sim
≤ 120	29	1.0			1.0		
> 120	54	2.4	0.6 to 9.4	.200	1.3	0.3 to 5.6	.72
Burden, %							
< 25	44	1.0			1.0		
≥ 25	39	2.8	0.8 to 9.8	.098	2.6	0.6 to 10.8	.19
ECOG PS							
0-1	78	1.0			1.0		
≥ 2	5	2.2	0.3 to 18.0	.449	3.0	0.4 to 25.0	.311
Age for every 10 years		1.2	0.7 to 2.0	.562	1.2	0.7 to 2.2	.559

Abbreviations: CgA, chromogranin A; CTC, circulating tumor cell; ECOG PS, Eastern Cooperative Oncology Group performance status; HR, hazard ratio; NET, neuroendocrine tumor; OS, overall survival; PFS, progressionfree survival.

Survival curves according to (A, B) presence of circulating tumor cells (CTCs), (C, D) grade, and (E, F) chromogranin A (CgA) demonstrating differences in (A, C, E) progression-free and (B, D, F) overall survival.



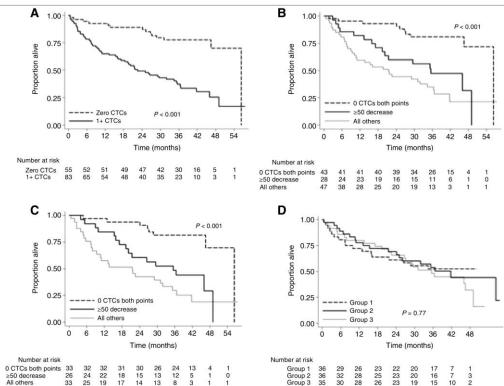
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CTCs as predictive markers for response



	Disease control*	Disease progression	
First posttreatment CTC			
0	47/49	2/49	<i>P</i> < 0.001
1-8	15/25	10/25	
>8	9/26	17/28	
Changes in CTCs			
Group A 0-0 CTCs	35/36	1/36	P < 0.001
Group B ≥50% reduction	20/24	4/24	
Group C All others	16/40	24/40	
First posttreatment CgA		•	
CgA ≤ 120	24/35	11/35	P = 0.53
CgA > 120	44/59	15/59	
Changes in CgA			
Group 1 >27% reduction	19/29	10/29	P = 0.61
Group 2 ≤27% reduction or <12% increase	24/32	8/32	
Group 3 ≥12% increase	24/32	8/32	



□•Stable disease or partial response.

The Kaplan–Meier survival curves demonstrating

- (A) effect of the presence of baseline CTCs on OS
- (B) OS dependent on changes in CTCs at first posttreatment time point (3–5 weeks) compared with baseline CTC
- (C), OS dependent on changes in CTCs at second posttreatment time point (10–15 weeks) compared with baseline CTC in groups
- (D), OS dependent on changes in CgA at first posttreatment time point

Mohid S. Khan et al. Clin Cancer Res 2016;22:79-85

Clinical AAGR Machine

©2016 by American Association for Cancer Research





Pros

- Useful in other kinds of malignancies
- Seems to be interesting for NETs (of any origin?)
- Quite easy to take a sample

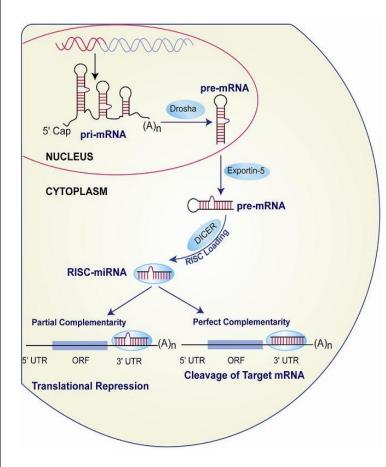
Cons

- Not a routine assay the CellSearch instrument is not available everywhere
- Further studies are needed to verify the results



MicroRNAs





miRNAs are expressed by non-coding parts of the genome and are post-transcriptional regulators, which control cell proliferation, differentiation and apoptosis in a variety of cells by degradation or translation inhibition of specific mRNAs

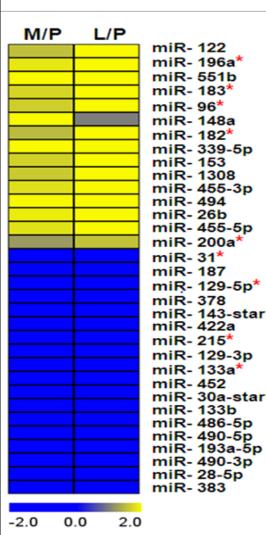
miRNAs have been identified as potential oncogenes or tumor suppressors

From, Bioinformation, 2010; 5(6): 271-276





uppsala miR



- The expression of miR-96, -182, -183, -196a and -200a is significantly upregulated in **SI-NET** cells compared to normal EC cells
- The expression of miR-31, -129-5p, -133a and -215 revealed significant down-regulation in SI-NET cells compared to normal EC cells
- Comparison of miRNA expression between **pNET**, normal pancreatic islets and serum samples
 - miR-624 correlated with Ki67 index
 - miR-201 correlated with metastatic disease
 - miR-193b was higher in neoplastic than in normal pancreatic islets, and also increased in serum

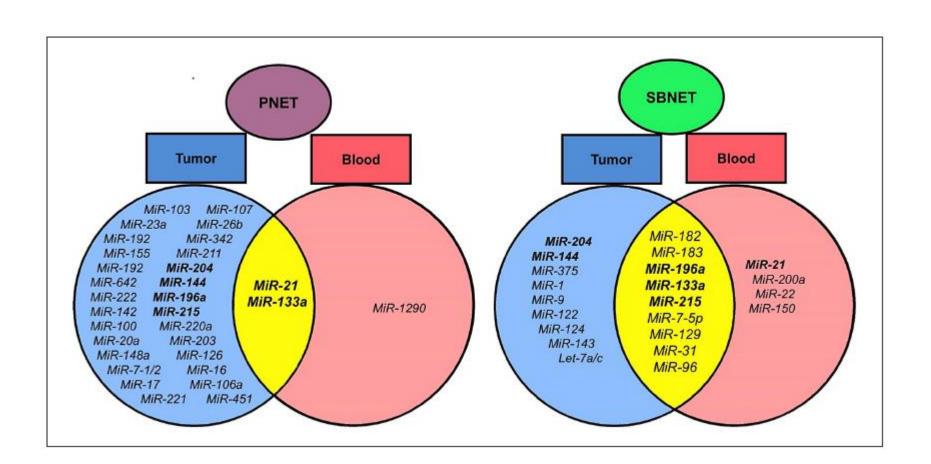
Difficult to find any predictive or prognostic miRNAs



.

Systematic overview of miRNA expression in NET









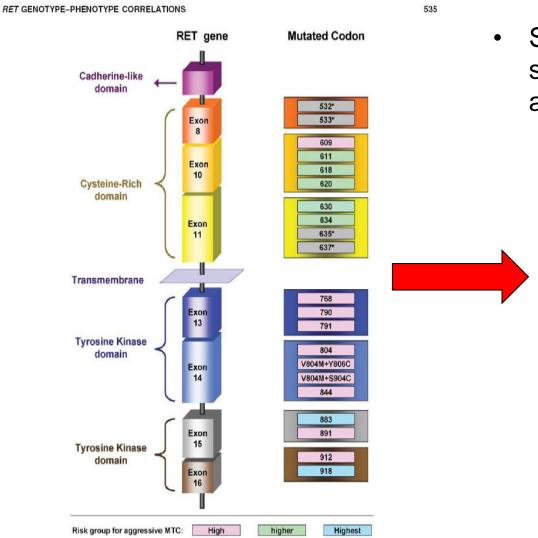


- An enormous amount of effort has been put into miRNAs to define their role in tumor development
- Their place in the diagnostic work-up of NETs is still quite uncertain



Genetic testing





- Some inherited NETs have specific genetic changes and should be tested
 - MEN 1; *MENIN*, no phenotype/genotype correlation
 - MEN 2; *RET*, with phenotype/genotype correlation
 - Von Hipple-Lindau; no phenotype/genotype correlation

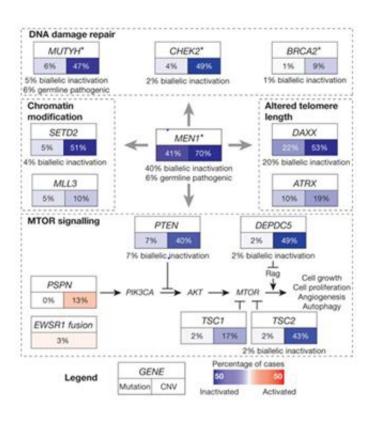
* Risk category not clear, probably low to intermediate





UPPSALA UNIVERSITET The genetic landscape of Pan-NETs

Core pathways in PanNETs



- MENIN often mutated in both hereditary and sporadic cases
- DAXX and ATRX have been identified as predictors of prognosis
- mTOR pathway mutations may be of importance for treatment
- Mutations in DNA-repair pathways genes a new finding



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Loss of DAXX and ATRX Are Associated With Chromosome Instability and Reduced Survival of Patients With Pancreatic Neuroendocrine Tumors



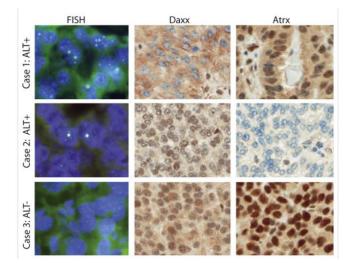
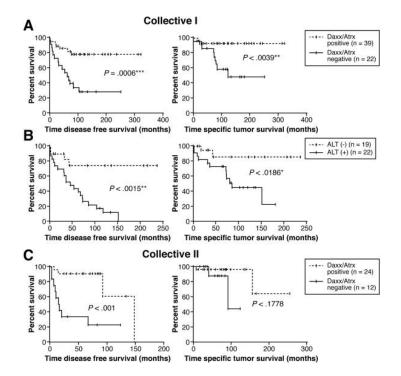


Table 2.

Correlation Between ALT Activation, CIN, and DAXX and ATRX Nuclear Expression

	CIN-	CIN+	Pvalue	ALT +	ALT -	Pvalue
ALT						
Positive	4	16	.0095			
Negative	13	7				
DAXX/ATRX nuclear staining						
Positive	14	9	.0361	3	17	.0003
Negative	6	16		17	7	

Figure 2 (*A*) Kaplan–Meier curves of collective I depicting a significantly shorter relapse-free survival and a shorter tumor-specific survival in DAXX- or ATRX-negative pNETs compared with DAXX- and ATRX-positive pNETs.



DAXX/ATRX and Alternative Lengthening of Teolmeres (ALT) associate with Chromosomal Instability (CIN) in turn associating with worse outcome Marinoni et al

Gastroenterology 2014



SI-NETs Sporadic and inherited



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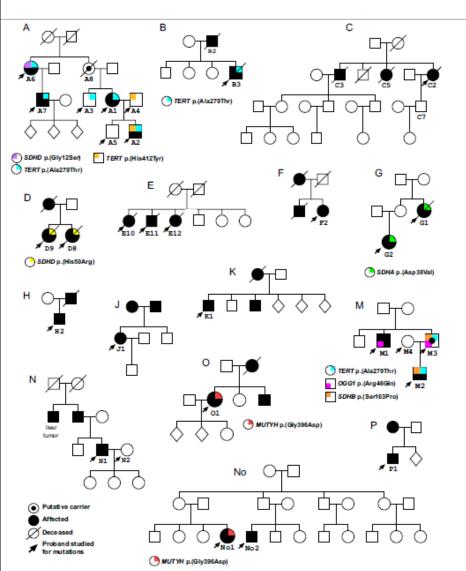
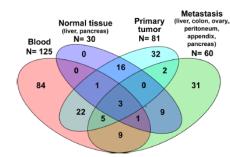


 Table 1
 Summary of clinical data comparing hereditary and sporadic patients.

Patients	Hereditary (n=26)	Sporadic (n=215)
WHO grade 1 ^a	10	115
WHO grade 2 ^a	8	45
Unknown tumor grade	8	55
TxNxM0 ^b	5	61
TxNxM1 ^b	20	147
Unknown tumor stage	1 '	7
Dead with disease	13	111
Alive with disease	11	96
No follow-up information	2	8
Median age at diagnosis	57 (34-68)	61 (23-90)
Median survival (months)	83 (40–348)	92.5 (2-348)



Germline mutations TERT SDHA SDHD MUTYH OGG1

Loss of chromosome 18 CDKN1B mutations DNA repair – *MUTYH, OGG1*

Dumanski JP et. al Endocrine-Related Cancer 2017



Genetic tests can be used in the clinic (?)



- Clinical sequencing of tumors are becoming increasingly used
- The challenge is to know how to interpret the results
- In some cases, mutations may be useful for prognosis
- For patients with inherited syndromes, genetic testing is valuable for relatives
- More research is needed to define how mutation screening (or whole genome sequencing) can be included in the clinical practice
 - The problem with data storage capacity is fundamental and needs a solution

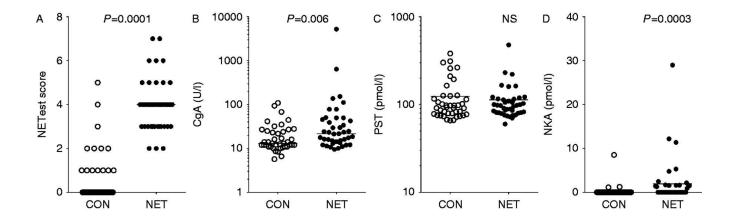


The NETest



- A PCR based test containing 51 neuroendocrine tumor gene transcripts
- Can be used for both SI-NETs and P-NETs

- Can define stable and progressive disease
- Can identify NET disease
 recurrence prior to imaging



Irvin M Modlin et al. Endocr Relat Cancer 2014;21:615-628



Can the NETest be used to predict response?



- A study of 49 NET patients undergoing ⁶⁸Ga-DOTA PET/CT
- SUV_{max} , CgA, Ki67 and NETest were analyzed
- A combination of circulating transcript levels, particularly MORF4L2, and imaging effectively differentiated progressive from stable disease.





- Still few published studies
- We don't really know what is being measured
- However, the test is already commercially available – patients may ask for it so we need to understand it's usefulness and clinical value







There are several new potential biomarkers, but data is missing on their usefulness in the clinic

When we are evaluating new potential candidates special attention should be given to:

How easy is the sampling?

How reproducible is the analysis?

How sensitive and specific is it?

The cost/benefit of the analysis.



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Thank you for your attention



