

4. Phäochromocytom & extraadr. Paragangliome

"The "great mimics": often sought, rarely found (Prävalenz 5/Mio)...yet, mostly discovered when it is too late... in autopsies
 Becker 01, JCEM 10; 95; NEJM 99; 340: 1872-9, Hypertension 91; 17-733-41; Lancet 05; 366: 665-75, SMF 12; 12: 66-71

Genotyp-Screening? (p22, KK-Gutsprache nötig) **RET** (MEN-II, 50% Phäo, typ A>NA), **VHL** (50% Phäo=Typ2, typ NA>A), **SDHD/B** ("maternal imprintin, dh nur Mut v Vater manifest b Kinder, Paragangliome gehäuft maligne, typ. 3-Methoxytyramin↑), **NF 1** (Neurofibromatosis Typ 1, 2% Phäo)
Aggressiv: alle Phäo (25% pos Spontan-Mutation) vs **Kons:** <40j, pos FA, bilat/extraadren. Phäo / MTC / Angiome / CVI (VHL) / Glomustu (SDHD/B)
 2x5ml EDTA Blut & informed consent ⇒ Prof. H. Neumann (neumann@med1.ukl.uni-freiburg.de) ⇒ pos ⇒ Fam.Screen / Psychol. Beratung

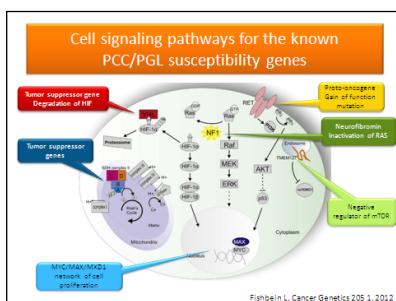
Journal of Human Hypertension (2013) 27, 141–147
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www.nature.com/jhh



REVIEW

An update on the genetics of pheochromocytoma

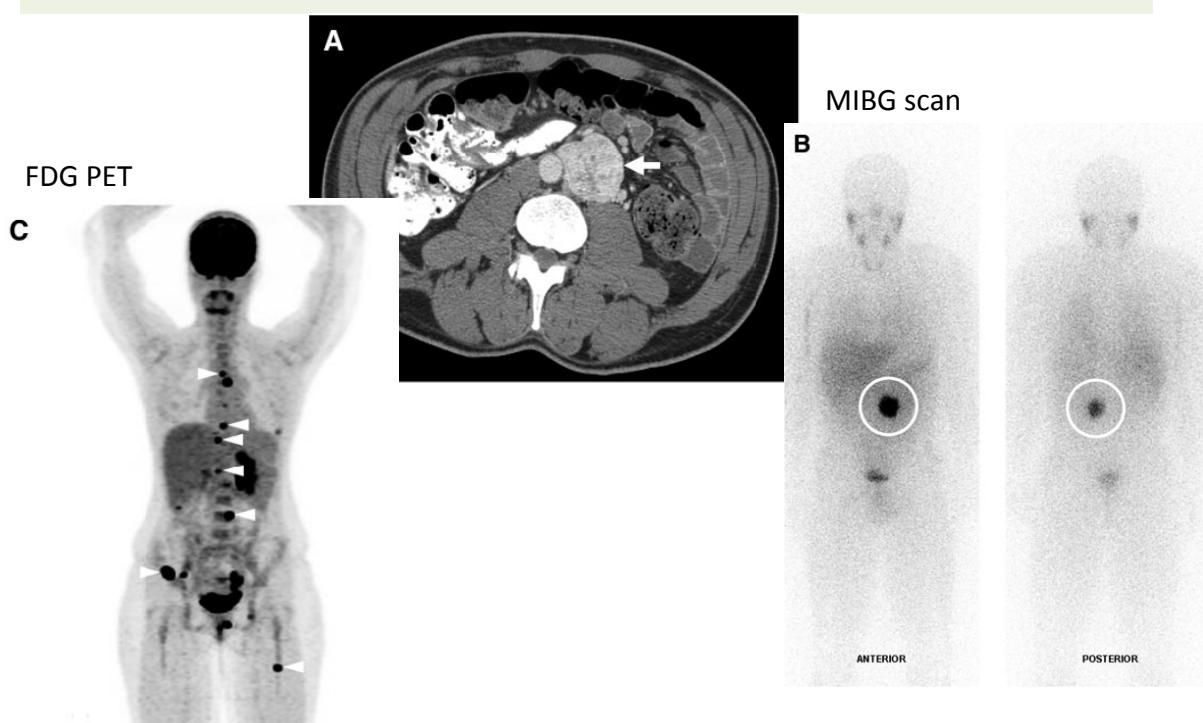
D Karasek¹, U Shah², Z Frysak¹, C Stratakis³ and K Pacak²



Phäochromozytom - Allgemeines

- aus Katecholamin-produzierenden (chromaffinen) Zellen hervorgehender Tumor
 - adrenal: Nebennieren-Mark = PHEO
 - extra-adrenal = Paragangliom (PGL)
 - Sympathikus: Thorax-Becken
 - Parasympathikus: Kopf-Hals

Paragangliom (PGL)



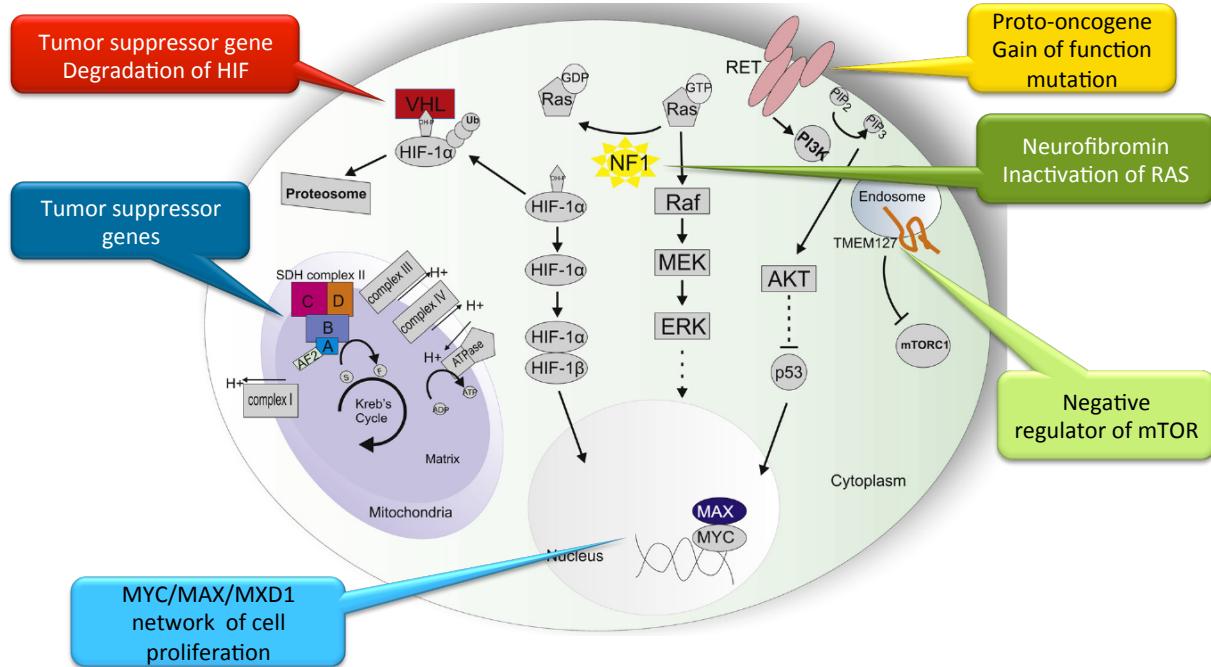
Phäochromozytom - Genetik

- ~ 1/3 of patients have a germline mutation in one of the known susceptibility genes – Vererbung autosomal dominant
- 54% of patients with head and neck paraganglioma
- Significant portion (~ 25%) of patients with apparently sporadic disease have germline mutations

Phäochromozytom - Genetik

- PHEO/PGL i.R. eines «genetischen» Syndroms
 - MEN2: RET Proto-Onkogen
 - VHL Syndrom
 - NF Typ1
- isiolerte PHEOs/PGLs
 - Familial Paraganglioma: SDH Komplex
 - neu: TMEM127, MAX

Cell signaling pathways for the known PCC/PGL susceptibility genes



Fishbein L, Cancer Genetics 205 1, 2012

MEN 2

- aktivierende Mutation im RET Proto-Onkogen
- MTC als Erstmanifestation
- PHEO in 50%, 30-40-j.
- meist adrenal, bilateral >50%, Malignität <5%
(↑bei Kindern mit MEN 2B)
- A, NA

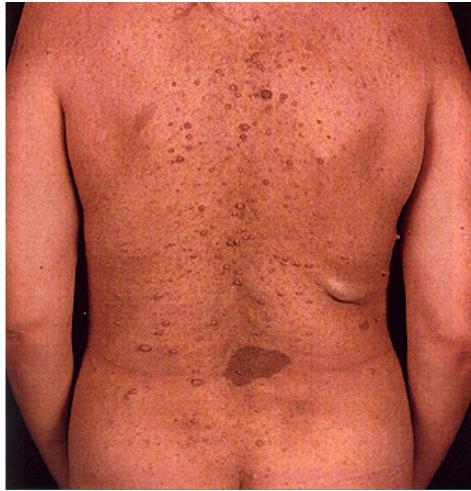
NF 1 – M. Von Recklinghausen

- aut. dom., 50% de novo Mutation
- Dx im Kindesalter
- PHEOs in 0.1-5.7%, ED 40-50-j.
- A > NA

NIH diagnostic criteria for neurofibromatosis type 1

Two or more of the following clinical features must be present:
Six or more <u>café au lait macules</u> of more than 5 mm in greatest diameter in prepubertal individuals, and more than 15 mm in greatest diameter in postpubertal individuals.
Two or more <u>neurofibromas</u> of any type or one plexiform neurofibroma
<u>Freckling</u> in the axillary or inguinal regions
Optic glioma
Two or more iris hamartoma (<u>Lisch nodules</u>)
Distinctive bony lesion such as sphenoid dysplasia, or thinning of the long bone cortex with or without pseudoarthrosis
A first-degree relative (parent, sibling, or offspring) with NF 1 based on the above criteria.





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Von Hippel Lindau (VHL)

- ↑Tumorinzidenz (benigne + maligne)
- aut. dom., 20% de novo Mutation
- VHL Typ 1: kein PHEOs/PGLs
- VHL Typ 2: häufig PHEOs/PGLs
- 30-j., mögliche Erstmanifestation
- meist adrenal, 50% bilateral; Malignität <5%
- nur NA

VHL - Manifestationen

- Hemangioblastomas of the brain (cerebellum) and spine
- Retinal angiomas
- Endolymphatic sac tumors of the middle ear
- Serous cystadenomas and neuroendocrine tumors of the pancreas
- Pheochromocytomas
- Clear cell renal cell carcinomas (RCCs)
- Papillary cystadenomas of the epididymis and broad ligament



Familial Paraganglioma Syndrome

- Succinate dehydrogenase enzyme complex (SDH)
- 4 Subunits (SDHx-Gene):
 - SDHA – SDHAF2 (Kofaktor)
 - SDHB
 - SDHC
 - SDHD
- meist PGLs, Lokalisation je nach betroffener Subunit
- oft «maternal imprinting» - Übertragung nur vom Vater
- oft «silent»; oft DA, DA + NA
- SDHB-Positivität in der Immunhistochemie: SDHx 100/84%

TMEM-127-Gen / MAX-Gen

- TMEM-127:
 - Tumor-Suppressor-Gen
 - 40-50-j.
 - adrenal (1/3 bilateral), Malignität <5%
- MAX:
 - Protein MAX (MYC-associated factor X), Transkriptionsfaktor
 - funktionell Tu-Suppressor-Gen
 - 30-j.
 - bilateral (67%), maligne (25%),

Clinical Features of hereditary PCC/PGL

	Penetrance of PCC/FPGL	Mean age at presentation	Frequency of PCC/PGL	Metabolites	Frequency of Malignancy	Frequency of bilateral PCC
RET	50	35	100/~0	MN/NMN	3	63
VHL	10-25	29	90/19	NMN	3	44
NF1	0.1-5.7	42	95/6	MN/NMN	9	14
SDHD	86	35	0/100*	NMN/MTX	4	
SDHB	77	33	25/78	NMN/MTX	31	
SDHC	U	43	0/100	--	0	
SDHA	U	40	17/83	NMN	0-14	
SDHAF2	100	32	0/100*	NMN	0	
TMEM127	U	43	95/8	NMN/MN	4	39
MAX	U	32	100/17	NMN	25	67

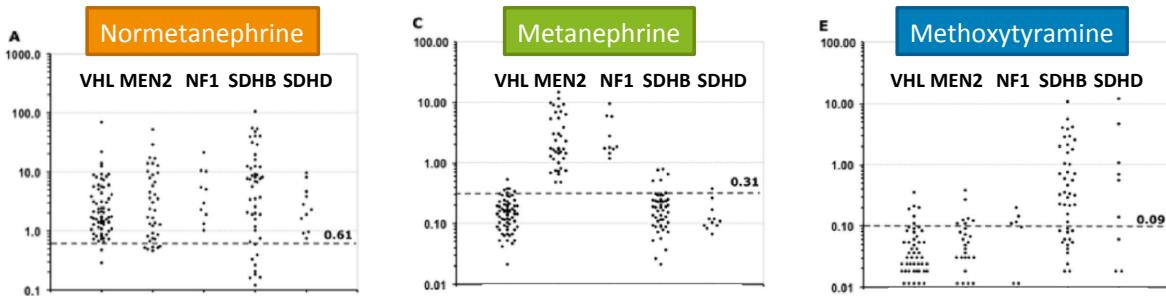
Welander J, Endocrine-Related Cancer 18 R253, 2011
 Fishbein L, Cancer Genetics 205 1, 2012
 Jafri M, EJE 166 151, 2012

Autosomal-dominant inheritance, * maternal imprinting

Malignität

- SDHB, MAX:
>25-30%
- NF1, sporadische PHEOs:
ca. 12%
- RET, VHL, SDHD, SDHC, SDHAF2, TMEM-127:
<5%

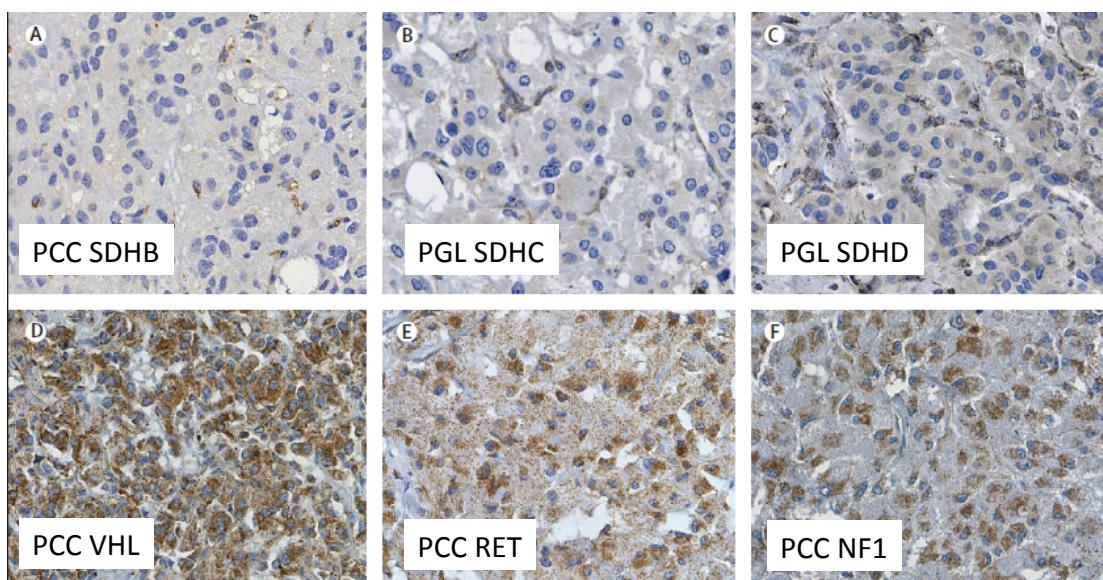
Plasma Metanephrides to guide Genetic Testing in hereditary PCC/PGL



Adrenal (bilateral):	NMN↑, MN↑, MTX↔	→ MEN2 (NF1)
Adrenal (bilateral):	NMN↑, MN ↔, MTX↔	→ VHL
Extraadrenal:	NMN↑, MN ↔, MTX↔	→ SDHD (VHL)
Extraadrenal:	NMN↑, MN ↔, MTX ↑	→ SDHB

Eisenhofer et al, Clin Chem 57: 411, 2011

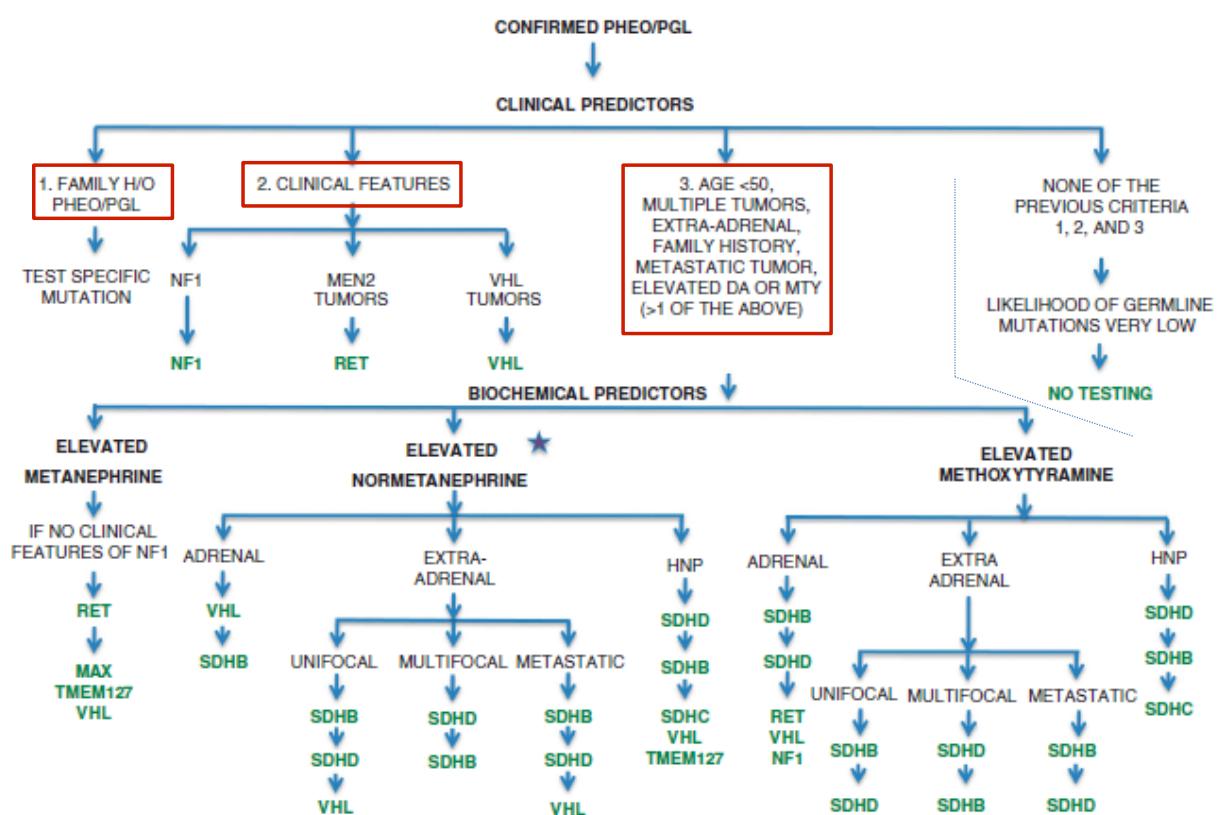
SDHB Immunohistochemistry identifies SDHB, -C and D associated tumors (Sensitivity 100%, Specificity 84%)



Van Nederveen, Lancet Oncol 10 764, 2009

Genetische Testung – warum?

1. Assoziation mit anderen (malignen) Tumoren?
2. Planung Follow-up (z.B. vermehrt Bildgebung)
3. Screening von Familienmitgliedern



Key points

- PHEOs/PGLs mit genetischem Background häufiger als bis anhin angenommen (ca. 30%)
 - mögliche Erstmanifestation eines VHL!
- genetischer Workup bestimmt durch Lokalisation, biochem. Pattern, Immunhistochemie
- In Zukunft spezifische Therapie entsprechend Genotyp(?)