



12° AME Italian Meeting 6° Joint Meeting with AACE



Bari,
7-10 novembre 2013

Carcinoma midollare tiroideo familiare

Profilo genetico e stratificazione del rischio

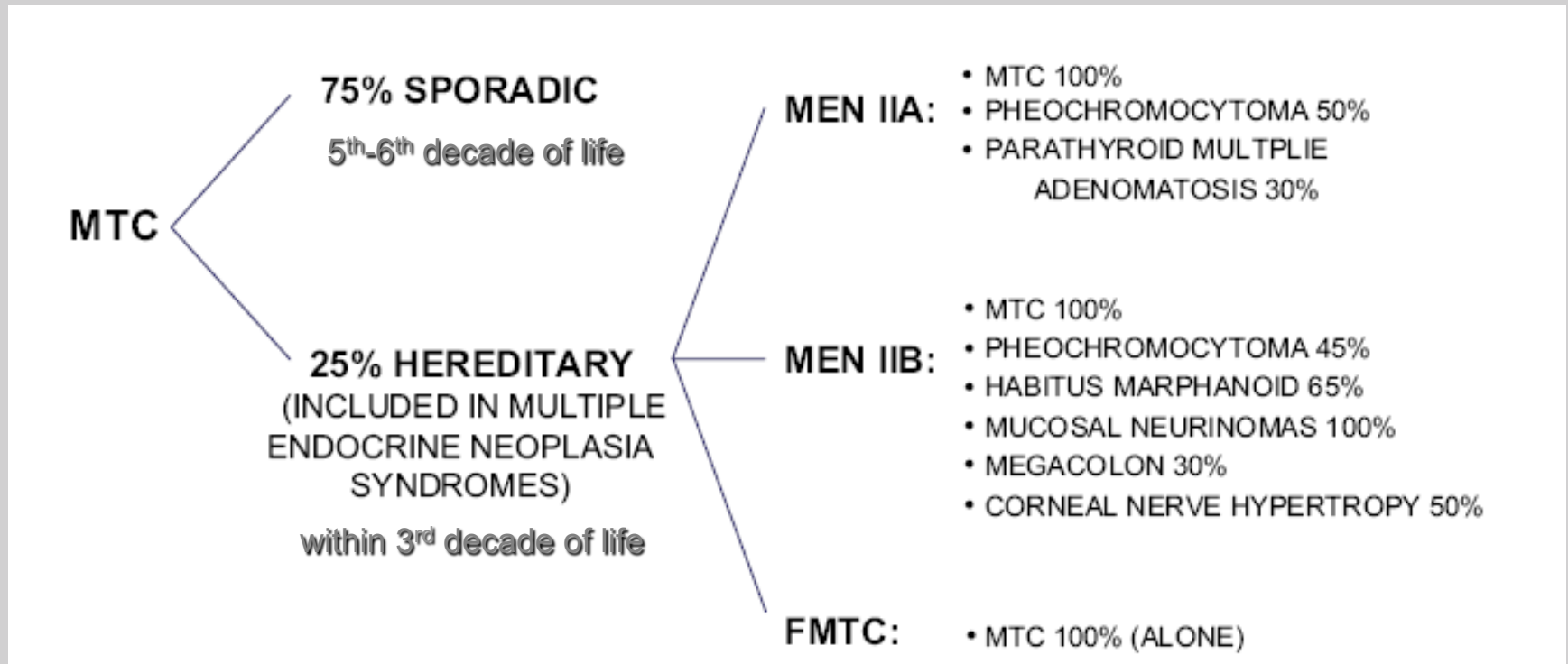
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Familial MTC: genetics and risk stratification

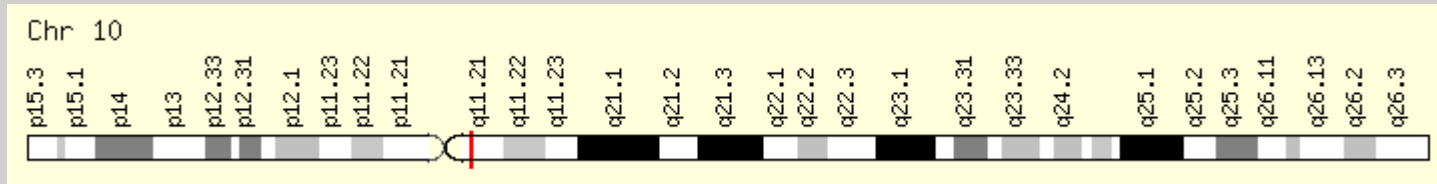
Different forms of medullary thyroid cancer



Elisei R 2008 Best Pract Res Clin Endocrinol Metab 22: 941–953

Familial MTC: genetics and risk stratification

RET (REarranged during Transfection) protooncogene



Long arm of chromosome 10
(10q11.2)

susceptibility gene for

- familial medullary thyroid cancer (FMTC)
- pheochromocytoma
- parathyroid hyperplasia/adenomas

multiple endocrine neoplasia type 2 (MEN 2)

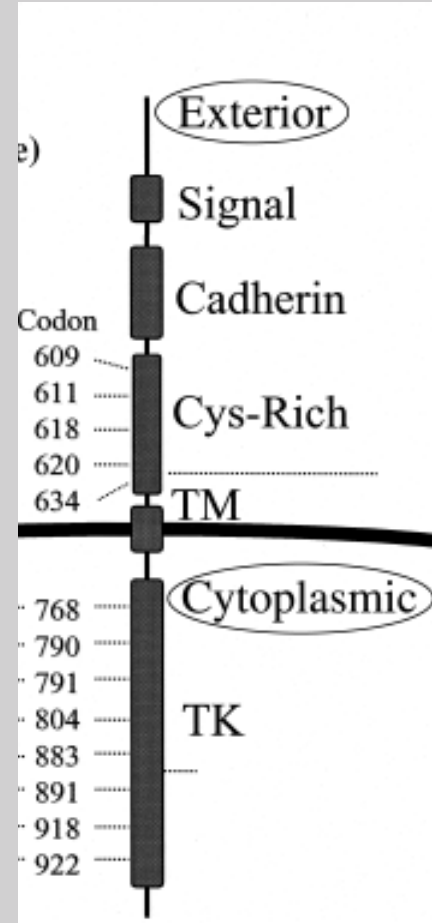
Familial MTC: genetics and risk stratification

RET → transmembrane receptor

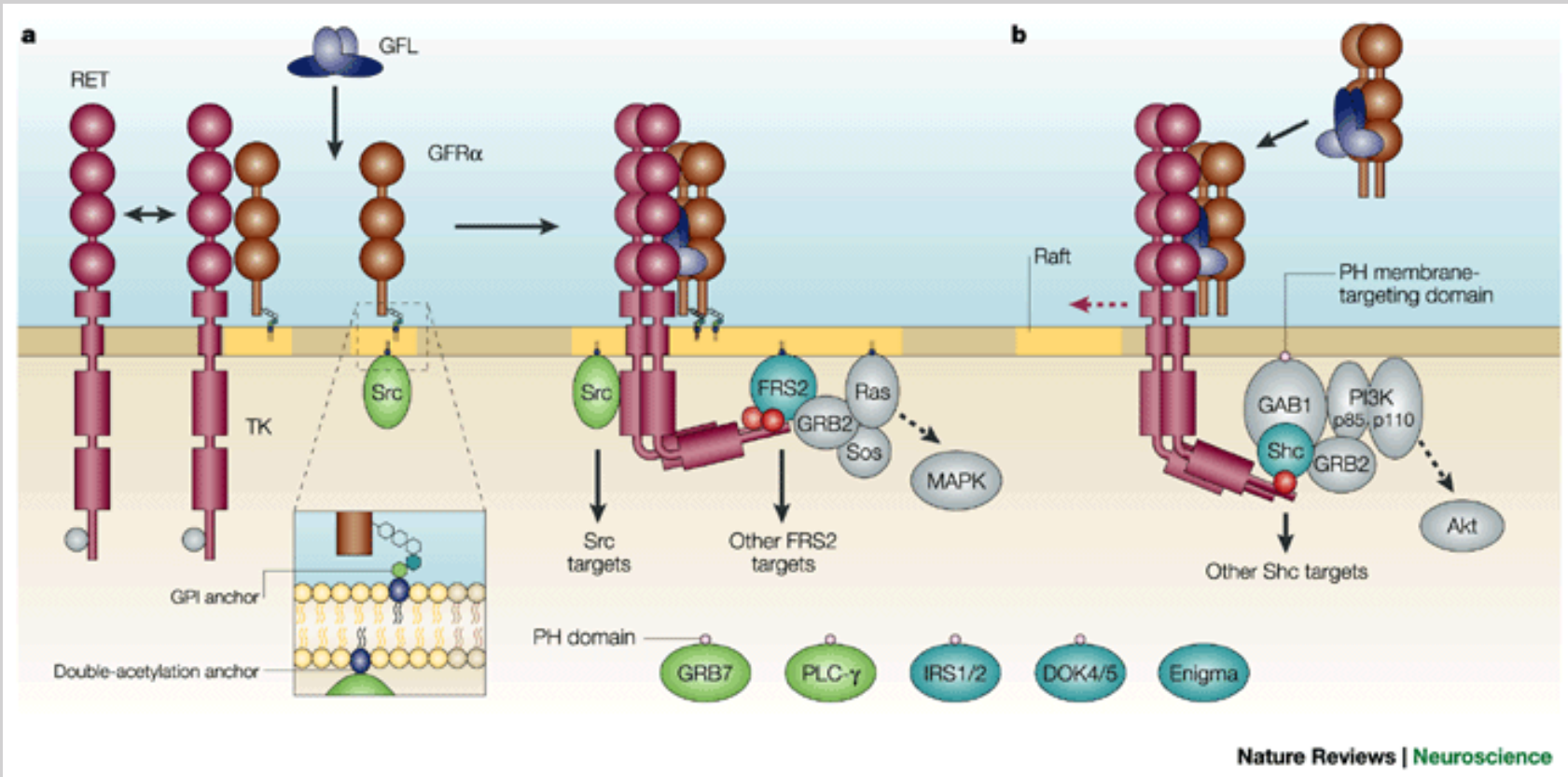
extracellular domain

transmembrane domain

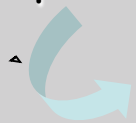
tyrosine kinase domain



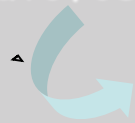
Familial MTC: genetics and risk stratification



In the presence of GDNF the receptor complex is activated

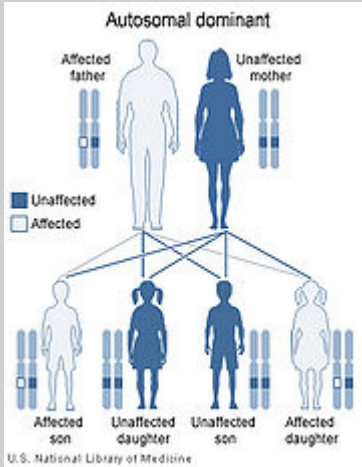


autofosforilazione



signal transduction pathway activation

Familial MTC: genetics and risk stratification



RET mutations

constitutive supraphysiological
activation of the RET receptor
tyrosine kinase



cell hyperstimulation

thyroid C cells ◀

▶
adrenal medullary cells

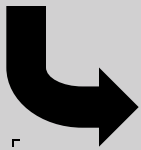
▶
parathyroid chief cells

Familial MTC: genetics and risk stratification

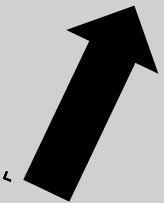
'Gain-of-function' mutations

cysteine-rich extracellular domains

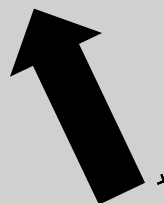
High transforming ability



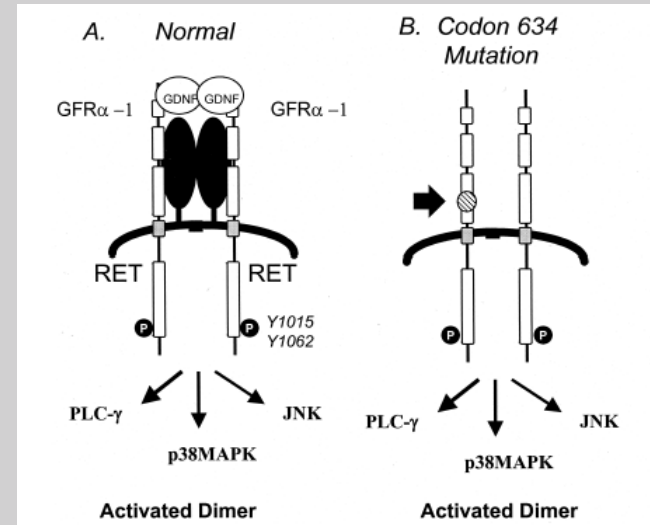
ligand-independent dimerization and cross-phosphorylation of mutant RET receptor proteins



addition of one more cysteine residue in codons 533, 606 or 631



loss of a cysteine residue irrespective of the amino acid substituting for cysteine



Familial MTC: genetics and risk stratification

'Gain-of-function' mutations

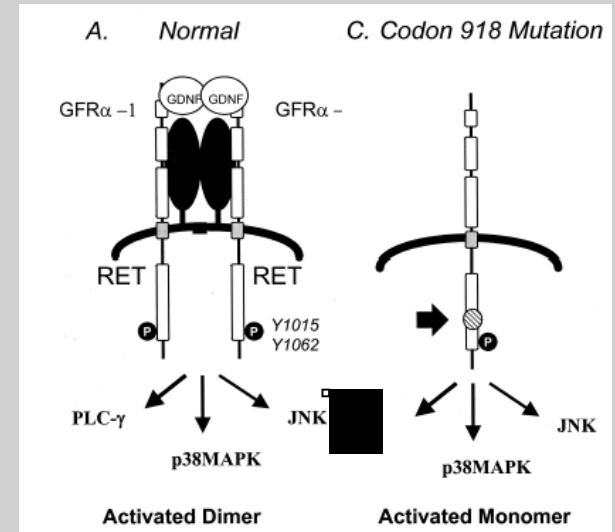
intracellular tyrosine kinase domain
codons 768, 790, 791, 804 and 891

facilitate the access of
ATP to its binding site

preferential binding of
intracellular substrate

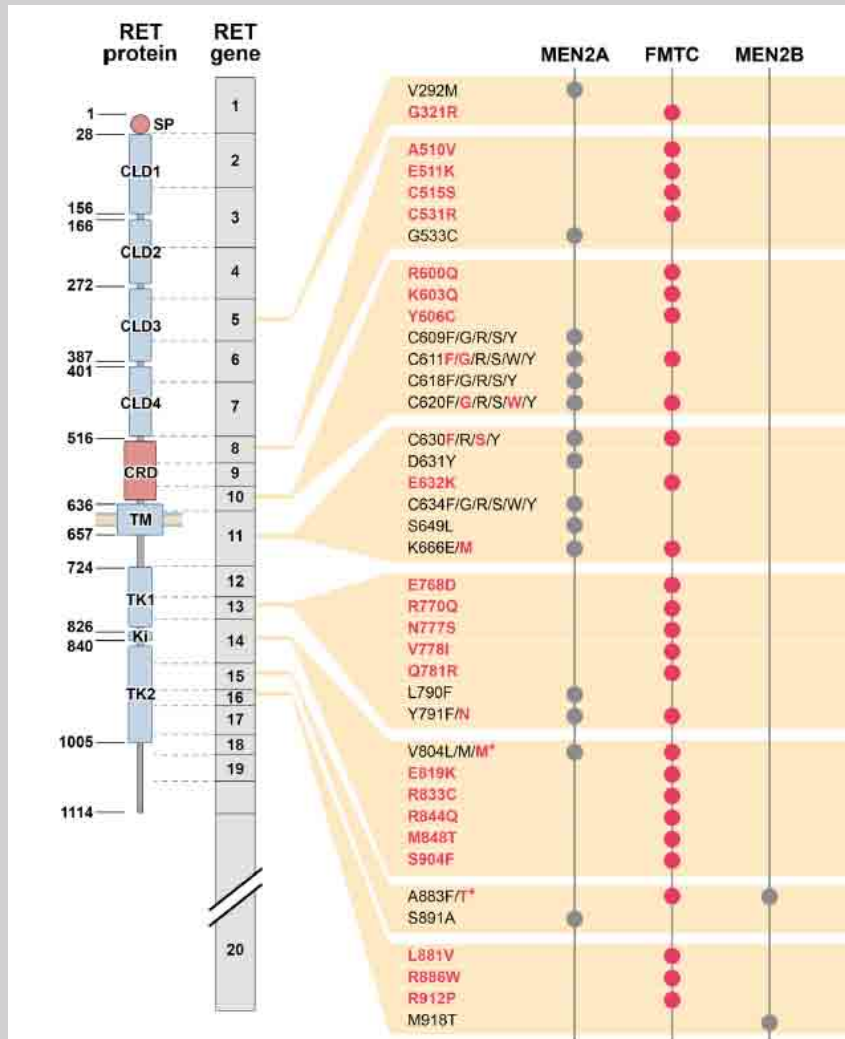
intracellular catalytic core
codon 918

Very high
transforming ability



Mutations in the intracellular kinase domain cause
RET autophosphorylation, independently of
dimerization and of the ligand

GENOTYPE-PHENOTYPE CORRELATION



Mutations in codons 609, 611, 618, 620 (exon 10) co-segregate with Hirshprung Disease and MEN2 syndromes

Mulligan et al. 1994 Hum Mol Genetics 3:2163
 Mulligan et al. 1994 Nat Genetics 6: 70
 Romeo et al. 1998 J Int Med 243: 515
 Sijmons et al. 1998 Gut 43: 542
 Nishikawa et al. 2003 Eur J Hum Genetics 11:364
 Pasini et al. 2002 Surgery 131: 373

Familial MTC: genetics and risk stratification

RET mutations

Table 2. Germline mutations of the *RET* proto-oncogene in MEN-2A (24)

| Affected codon | Exon | Clinical syndrome | Percentage of all MEN-2 mutations |
|----------------|------|------------------------------|-----------------------------------|
| 609 | 10 | MEN-2A/FMTC | 0-1 |
| 611 | 10 | MEN-2A/FMTC | 2-3 |
| 618 | 10 | MEN-2A/FMTC | 3-5 |
| 620 | 10 | MEN-2A/FMTC | 6-8 |
| 630 | 11 | FMTC | <0.1 |
| 634 | 11 | MEN-2A | 80-90 |
| 768 | 13 | FMTC | 0-1 |
| 790 | 13 | MEN-2A/ FMTC | <0.1 |
| 791 | 13 | FMTC | <0.1 |
| 804 | 14 | FMTC (age of onset variable) | 0-1 |
| 883 | 15 | MEN-2B | |
| 891 | 15 | FMTC | 0-1 |
| 918 | 16 | MEN-2B | 10-20 |
| 920 | | | |
| 922 | | Sporadic/MEN-2B | |

MEN-2A: multiple endocrine neoplasia type 2, FMTC: familial medullary thyroid carcinoma



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Table 3 Distribution of *RET* missense germline mutations in Continental Europe (500 RET families)^a. Owing to rounding, not all numbers add up.

| ATA class | <i>RET</i> mutation | Germany, Halle 1994–2012 ^b | Italy, multicenter (18) | France, multicenter (19, 20) | Total |
|-----------|---------------------|--|----------------------------|---------------------------------|--------------|
| | | <i>n</i> (%) | <i>n</i> (%) | <i>n</i> (%) | <i>n</i> (%) |
| D | M918T | 32 (16) | 17 (9) | 3 (3) | 52 (10.4) |
| D | A883F | 0 | 0 | 0 | 0 |
| C | C634R/G/F/S/W/Y | 73 (36) | 52 (26) | 46 (47) | 171 (34.2) |
| B | C630R/F/S/Y | 1 (0.5) | 4 (2) | 0 | 5 (1.0) |
| B | C620R/G/F/S/W/Y | 14 (7) | 9 (5) | 12 (12) | 35 (7.0) |
| B | C618R/G/F/S/Y | 11 (5) | 15 (8) | 6 (6) | 32 (6.4) |
| B | C611R/G/F/S/W/Y | 6 (3) | 1 (1) | 1 (1) | 8 (1.6) |
| B | C609R/F/S/Y | 1 (0.5) | 6 (3) | 1 (1) | 8 (1.6) |
| A | G533C | 0 | 0 | 0 | 0 |
| A | E768D | 2 (1) | 9 (5) | 2 (2) | 13 (2.6) |
| A | L790F | 26 (13) | 8 (4) | 4 (4) | 38 (7.6) |
| A | Y791F | 14 (7) | 2 (1) | 0 | 16 (3.2) |
| A | V804L/M | 19 (9) | 52 (26) | 15 (15) | 86 (17.2) |
| A | S891A | 6 (3) | 23 (12) | 7 (7) | 36 (7.2) |
| Total | Any | 205 (100) | 198 (100) | 97 (100) | 500 (100) |

ATA, American Thyroid Association; RET, rearranged during transfection.

^aConsidering series with a minimum of 30 European RET families only that specified familial RET prevalence.

^bUpdated from reference (17) (141 RET families).





Familial MTC: genetics and risk stratification

Risk associated to RET mutations

Table 3. American Thyroid Association (ATA) risk level and timing of prophylactic thyroidectomy in multiple endocrine neoplasia type 2 (MEN-2A)

| ATA risk level | Age of prophylactic surgery |
|--|--|
| Level A (codons 768, 790, 791, 804, and 891) | Consider operative resection before age 5 years May delay operative resection if: <ul style="list-style-type: none"> ▪ Normal annual serum calcitonin and ▪ Normal annual neck ultrasound (no lesions >5 mm and no concerning adenopathy) and ▪ Less aggressive family history and ▪ Family preference |
| Level B (codons 609, 611, 618, 620, and 630) | Consider operative resection before age 5 years May delay operative resection if: <ul style="list-style-type: none"> ▪ Normal annual serum calcitonin and ▪ Normal annual neck ultrasound (no lesions >5 mm and no concerning adenopathy) and ▪ Less aggressive family history and ▪ Family preference |
| Level C (codon 634) | Before 5 years of age |
| Level D (codon 883, 918) Tandem mutation (804-805, 804-806, 804-904) | First month of life |





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Familial MTC: genetics and risk stratification

DNA-based screening

Following the discovery of RET mutations in the germline of patients with MEN type 2, DNA-based analysis of the RET proto-oncogene was rapidly integrated into the routine clinical armamentarium.

Machens et al. 2009 J Intern Med 266: 114

Cost-effective identification of affected family members

Gilchrist et al. Clin Genet 2004; 66: 349–53

legal and ethical importance
indicating the need for prophylactic
thyroidectomy

gold standard of care

Rosenthal et al Thyroid 2005; 15: 140–5

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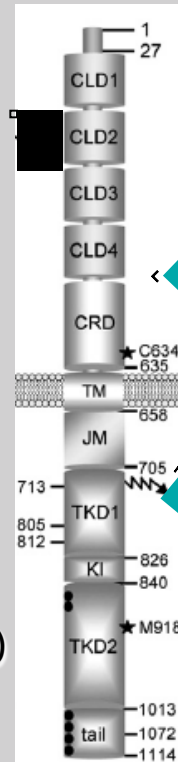


Familial MTC: genetics and risk stratification

Hereditary MTC

Germline RET mutations

- ▶ 95% of MEN2A kindreds
Hirschsprung disease
Lichen amyloidosis
- ▶ 88% of FMTC kindreds
- ▶ >95% of MEN 2B kindreds (codon 918)



Prospective family screening

| MUTATED CODONS | EXONS | PHENOTYPE | NO |
|--------------------------|----------------|-------------------------------------|----|
| 609 611 618 620 | 10 | FMTC FMTC FMTC-MEN 2A FMTC | |
| 630 634 | 11 | FMTC FMTC-MEN 2A | |
| 768 790 | 13 13 | FMTC FMTC | |
| 804 848 | 14 14 | FMTC FMTC | |
| 883 891 904 | 15 15 15 | FMTC FMTC FMTC | |
| 918 | 16 | MEN 2B | |

Elisei et al. 2007 J Clin Endocrinol Metab 92:4725-9

Castellone et al. 2008 Endocrinol Metab Clin North Am 37:363-74, viii





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Familial MTC: genetics and risk stratification

Hereditary MTC

- ▶ 6-7% of “sporadic” MTC carry a germline RET mutation

Elisei et al. 2007 J Clin Endocrinol Metab 92:4725-9

Wohllk et al. 1996 J Clin Endocrinol Metab 81: 3740-3745

RET genetic testing should be encouraged
in all newly diagnosed MTC patients

The National Comprehensive Cancer Network
Clinical Practice Guidelines in Oncology
2009

biochemical screening
for MEN2

hyperparathyroidism

pheochromocytoma

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Familial MTC: genetics and risk stratification

How is genetic analysis performed?

1. Patient referral for MTC or family history of MTC
2. History - evaluate family history
3. Clinical examination
4. Informed consent signature
5. Blood withdrawal (no fasting needed)
6. Sample sent to the Lab



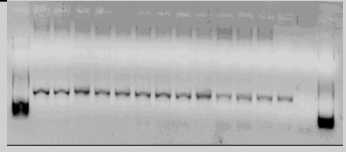
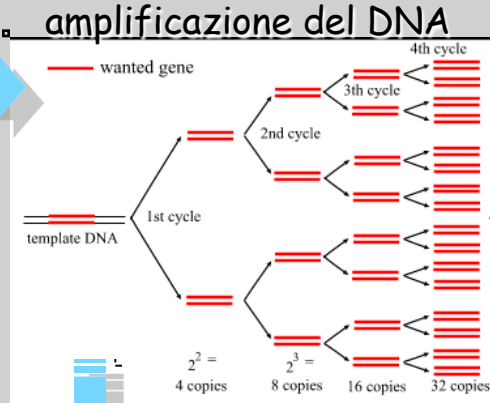
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DNA
isolation



denaturation

GENOMIC DNA DIRECT SEQUENCING

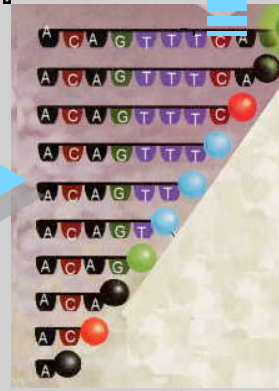
Capillary electrophoresis



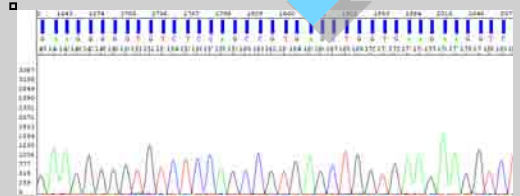
purification



purification



Sequencing reaction



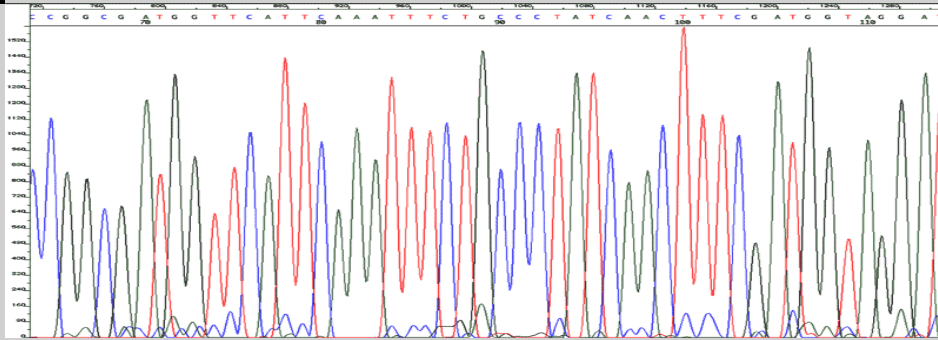
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Electropherogram analysis
by a Technician
by the Physician in charge

Comparison with the normal sequence
→ any SNP?

DIAGNOSIS

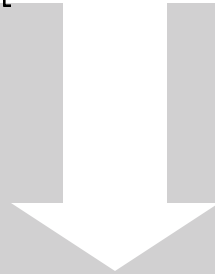




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Familial MTC: genetics and risk stratification

Genetic screening in proband and in first degree relatives is fundamental



High likelihood of developing MTC during lifespan

consider prophylactic thyroidectomy

follow up



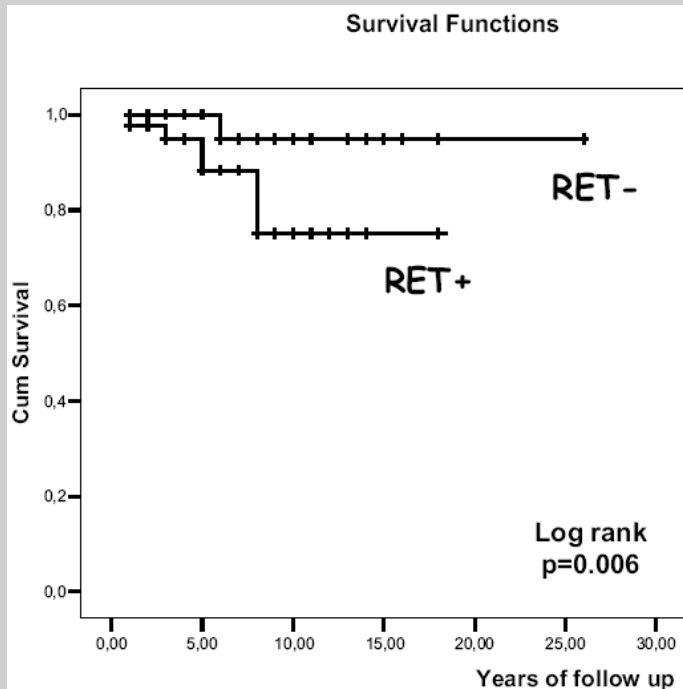
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Sporadic MTC

40-50% of sporadic MTC display somatic RET mutations

Romei et al. 1996 J Clin Endocrinol Metab 81:1619–1622
Schilling et al. 2001 Int J Cancer 95:62–66

Zedenius et al. 1995 J Clin Endocrinol Metab 80:3088–3090
Zedenius et al. 1998 Cancer Detect Prev 22:544–548



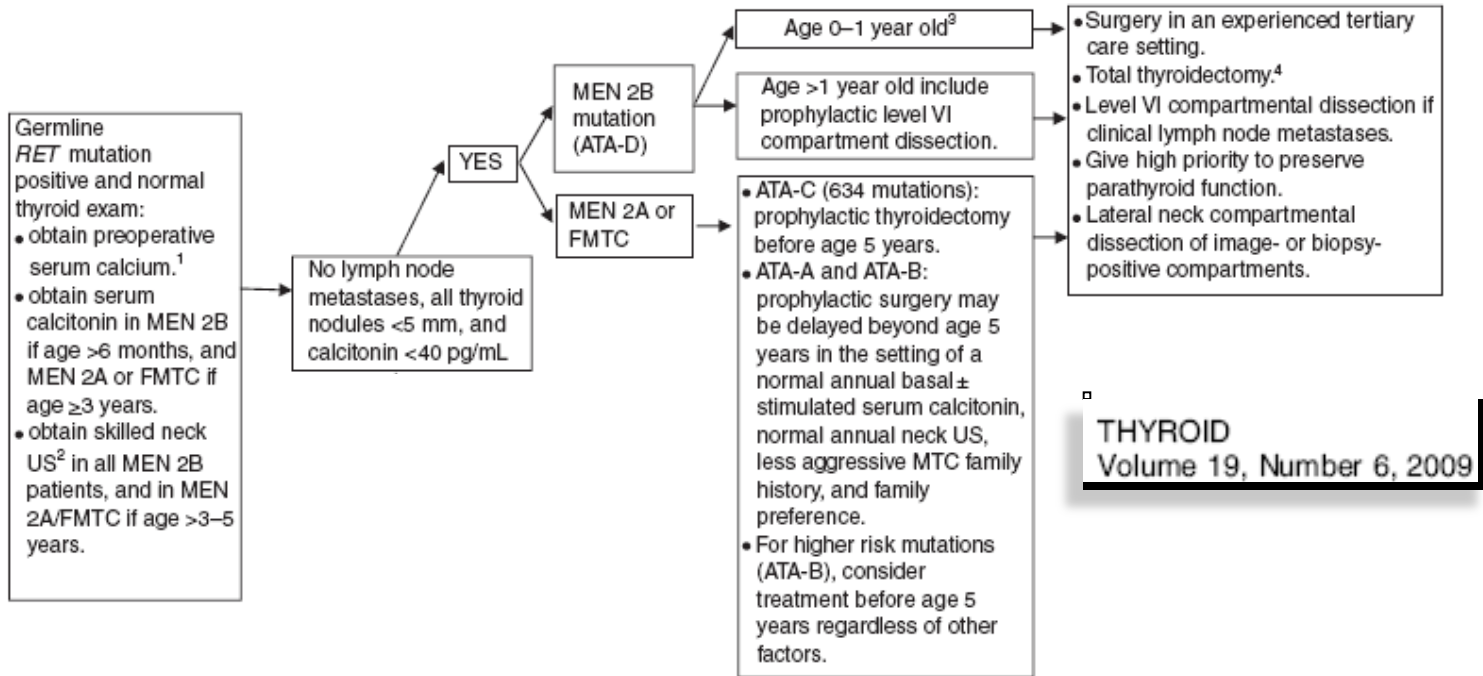
- Somatic RET mutations correlate with
- presence of lymph node metastases at diagnosis
 - worse outcome
 - disease persistence after surgery
 - lower survival rate

bad prognostic factor

Elisei et al. 2008 J Clin Endocrinol Metab 93:682-7

Familial MTC: genetics and risk stratification

Medullary Thyroid Cancer: Management Guidelines of the American Thyroid Association



¹Treat hyperparathyroidism with 4 gland resection and autograft to heterotopic site, or subtotal parathyroidectomy. Consider cryopreservation. PHEO preoperative screening should begin by age 8 years for MEN 2B and mutated *RET* codons 634 and 630; otherwise by age 20 years for other *RET* mutations.

²Neck US to include the superior mediastinum and central and lateral neck compartments.

³Insufficient data to recommend routine prophylactic level VI compartment dissection.

⁴Parathyroid glands resected or devascularized should be autografted in the neck in *RET*-negative, MEN 2B, and FMTC patients, while MEN 2A glands should be autografted to a heterotopic site.

FIG. 1. Initial diagnosis and therapy of pre-clinical disease.



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RET mutation analysis is a fundamental step in
the diagnostic work-up in
medullary thyroid carcinoma patients





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Familial MTC: genetics and risk stratification

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THANK YOU!

University of Ferrara
Department of Medical Sciences
Section of Endocrinology



Ettore degli Uberti