



The World's Congress of Surgery



Genotype phenotype correlation in RET positive Medullary Thyroid Cancer: A single institution experience

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Introduction

- Demographics, presentation and prognosis differ between MTC subtypes based on RET mutation
- Screening is important for early diagnosis and prophylactic thyroidectomy

Conclusion

- 35.5% Hereditary MTC 634 most common mutation (65.7%)
- Occurrence of pheochromocytoma and HPT in 634 consistent with literature
- 804 mutation had higher basal calcitonin and nodal recurrence

Methodology

- Retrospective study- single institution
- January 2002 December 2022

	Results							
MTC n=234/5246 (4.46%)	70	Demographics	634 n=46 (65.7%)	804 n=12 (17.1%)	618 n=4 (5.7%)	918 n=4 (5.7%)	790 n=2 (2.9%)	620 n=2 (2.9%)
RET Tested	(35.5%)	Median age (yrs)(range)	27 (9-68)	43 (22-64)	37 (18-39)	16.5 (8-37)	43 (29-57)	37 p=.03 (33-41)
197 /234 (84.2%)		Gender(M:F)	20:26	6:6	2:2	2:2	0:2	2:0
Sporadic	Hereditar	ry				DET 63	21	

Results

Objective

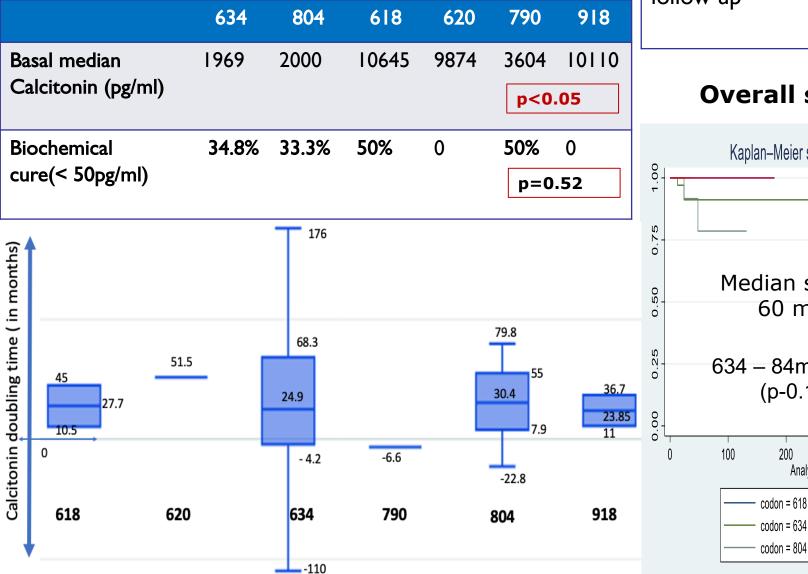
To analyze mutation profile in RET positive MTCs for differences in demographics, presentation and outcomes

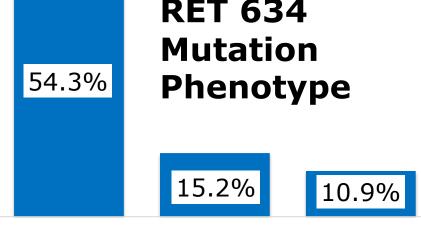
Presentation

- Goitre with LN most common presentation
- 21.7% (10/46) 643 mutation and 25% (1/4) 918 mutation presented with pheochromocytoma
- 33/64 individuals of 37 index families screened tested positive for RET mutation
- 5 underwent prophylactic thyroidectomy (634-2,804-2,618-1)

Pheochromocytoma

	Incidence	Synchronous	MTC before Pheo	Pheo before MTC
634	54.3% n=25/46	68% (mean age 35.7yrs)	28% (mean duration 15.6 yrs)	4% (duration 10 yrs)
918	25% n=1/4	100% (37 yrs)	-	-





HPT PHEO CLA

HPT seen only in 634 mutation

Outcomes	634	804	618	620	790	918
Nodal mets at presentation	50% (23)	66.6% (8)	50% (2)	50% (1)	50% (1)	75% (3)
Metastasis at presentation	17.4% (8)	8.3% (I)	25% (1)	-	-	-
Nodal recurrence	7.4% (8)	25% (3)	25% (1)	50% (1)	p=0.021 634 vs 804	25% (1)
Metastasis on follow up	8.6% (4)	8.3% (1)	-	50% (1)	-	-

Overall survival

Kaplan-Meier survival estimates Median survival 60 months 634 – 84m vs 804 -48m (p-0.15) 200 300 400 500 Analysis time codon = 618 · codon = 620 codon = 634 codon = 790

codon = 918

Disease free survival

