

# KRAS Mutations and Prognostic Implications in Appendiceal Cancers

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An NCI-Designated Cancer Center

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### INTRODUCTION

- Despite shared embryonic origin, appendiceal cancers (AC) have distinct clinical and molecular features compared to colorectal cancers (CRC).
- Detection of mutant KRAS has been associated with worse survival in CRC, whereas in AC the prognostic significance of mutant KRAS (55-65% prevalence) has yet to be characterized.

#### **METHODS AND STATISTICS**

- AC tissues from 891 individual patients underwent DNA (592, NextSeq, or WES, NovaSeq) and WTS (NovaSeq) sequencing at Caris Life Sciences (Phoenix, AZ). Low-grade appendiceal mucinous neoplasms were excluded.
- Chi-square, Fishers-exact, and Mann Whitney U tests were used to determine statistical significance and were adjusted for multiple hypothesis testing (p<0.05).
- Overall survival was calculated using insurance claims of AC patients from time of sample collection to last contact.
- Hazard ratios (HRs) were calculated using the Cox proportional hazards model (log-rank test).
- Multivariate regression analysis was performed on age, sex, histology and KRAS<sup>mut</sup>. vs KRAS<sup>wt</sup>.

#### RESULTS

- AC histological subtypes comprised mucinous adenocarcinoma (44.9%), signet ring cell carcinoma (13.4%), goblet cell carcinoma (23.8%) and adenocarcinoma- not otherwise specified (NOS, 18%).
- Among all AC specimens, KRAS was the most common mutation (49 %).
- KRAS<sup>mut</sup> vs KRAS<sup>wt</sup> tumors were more frequently associated with *TP53*<sup>mut</sup> (48.6% vs. 36.8%, respectively; p<0.01) and GNAS<sup>mut</sup> (42.8% vs. 5.7%, respectively; p<0.00001).
- Most frequent co-mutant gene with *KRAS* was TP53 (61.2%) in adenocarcinoma-NOS, and GNAS (46.9%) in mucinous adenocarcinomas.
- KRAS G12D is the most common mutation.
- Median OS among KRAS<sup>mut</sup> vs KRAS<sup>wt</sup> was 35.0 vs. 24.1 months, respectively (HR=0.65, 95% CI: 0.54-0.80, p<0.0001).</li>
- KRAS<sup>mut</sup> was associated with a trend towards improved survival in mucinous adenocarcinomas (HR 0.61; 95% CI:0.43-0.86, p0.005) but not with signet ring cell (HR 1.20; 95% CI:0.66-2.20, p 0.53) or goblet cell carcinoma (HR 1.17; 95% CI: 0.58-2.34, p 0.648).
- On multivariate analysis KRAS<sup>mut</sup> was an independent prognostic factor for improved survival among all AC (HR 0.63, 95% CI:0.51-0.80, p 0.0001).
- Notably, GNAS<sup>mut</sup> was associated with improved survival (HR 0.57, 95% CI:0.47-0.70, p<0.00001), while TP53<sup>mut</sup> was associated with poorer survival (HR 1.58, 95% CI:1.35-1.86, p<0.00001) among all AC.

## CONCLUSIONS

- KRAS was one of most frequently mutated genes in ACs. In contrast to CRC, KRAS<sup>mut</sup> was associated with significantly improved survival.
- This observed survival advantage remained consistent in the histologic subgroup of mucinous adenocarcinoma, but not in goblet and signet ring cell cancers.
- Prospective trials evaluating survival advantage of KRAS mutations and its implications in choosing future targeted therapies should be performed as well as analysis on response to current therapy.

#### **Table 1:** Baseline characteristics

	All cancers	Mucinous Adenocarcinoma	Signet Ring Cell Carcinoma	Goblet   Neuroendocrine	Adenocarcinoma- NOS	
Total cases	891	400	119	212	160	
Median age [range]	62 [22 – 90]	61 [29 – 90]	63 [40 – 88]	62 [22 – 90]	60 [28 – 85]	
	N (%)	N (%)	N (%)	N (%)	N (%)	Chi-square (p-value)
Male Gender	458 (51.4)	180 (45.0)	62 (52.1)	86 (40.6)	96 (60)	16.12 (0.001)
Race						
White	493/639 (73.8)	209/291 (71.8)	73/90 (81.1)	132/172 (76.7)	79/115 (68.7)	
African American	97/639 (14.5)	41/291 (14.1)	13/90 (14.4)	16/172 (9.3)	27/115 (23.5)	20.29
Asian Pacific Islander	32/639 (4.8)	15/291 (5.1)	2/90 (2.2)	12/172 (7.0)	3/115 (2.6)	(0.016)
Others	46/639 (6.9)	26/291 (8.9)	2/90 (2.2)	12/172 (7.0)	6/115 (5.2)	
Ethnicity						
Hispanic	89/641 (13.2)	53/300 (17.7)	10/90 (11.1)	11/161 (6.8)	15/117 (13.0)	11.18
Non-Hispanic	579/641 (86.7)	247/300 (82.3)	80/90 (88.9)	150/161 (93.2)	102/117 (87.0)	(0.011)

<sup>\*</sup> Race and ethnicity data is only available for 70% of patients

Figure 1: Frequency of KRAS mutation subtypes



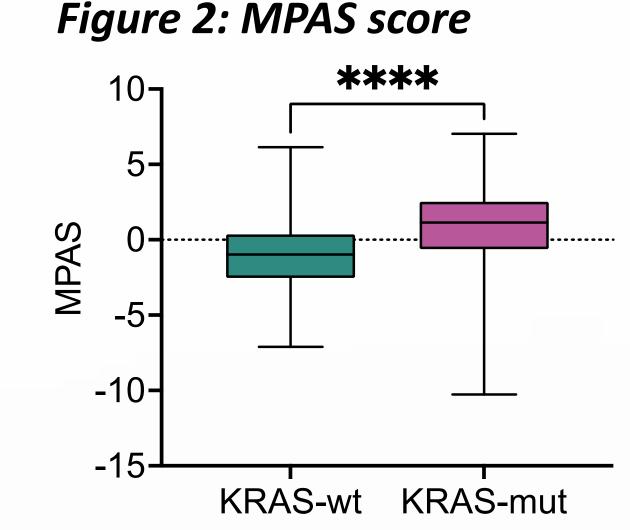


Figure 3: Molecular profiling: KRAS mutant vs. wild type

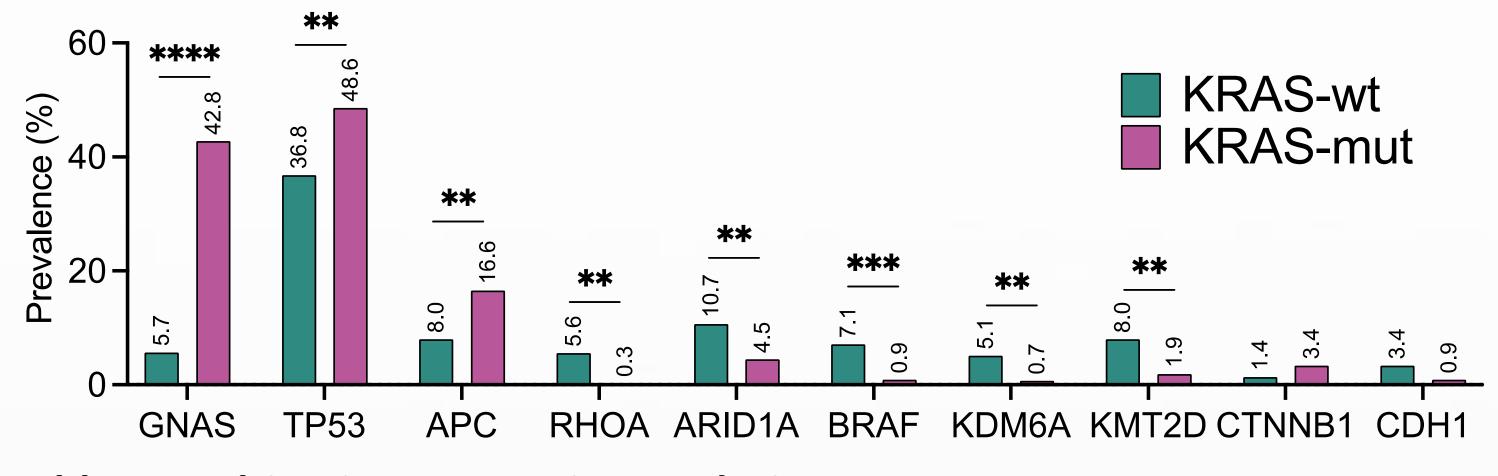


Table 2: Multivariate regression analysis

	All cancers		Mucinous Adenocarcinoma		
	HR (95% CI)	P*	HR (95% CI)	P*	
Age Group					
<50 years	Reference		Reference		
≥ 50 years	1.30 (1.01 – 1.67)	0.042	1.40 (0.94 – 2.09)	0.092	
Sex					
Male	Reference		Reference		
Female	0.96 (0.79 – 1.16)	0.700	0.78 (0.57 – 1.07)	0.121	
Race					
White	Reference		Reference		
African American	0.62 (0.39 – 1.12)	0.11	1.07 (0.47 – 2.43)	0.878	
Asian or Pacific islander	0.75 (0.53 – 1.06)	0.104	0.80(0.45 - 1.44)	0.460	
Others	0.70 (0.44 – 1.11)	0.131	1.05 (0.54 – 2.06)	0.878	
Mutation Profile	Reference - wild type				
KRAS -mutant	0.63 (0.51 – 0.80)	0.0001	0.64 (0.43 - 0.96)	0.031	
TP53 -mutant	1.6 (1.3 – 1.96)	0.0	1.84 (1.33 – 2.56)	0.0002	
BRAF -mutant	0.8 (0.49 – 1.32)	0.389	0.58 (0.23 – 1.50)	0.263	
APC -mutant	1.02 (0.75 – 1.39)	0.877	1.13 (0.65 – 1.98)	0.659	
GNAS -mutant	0.84 (0.63 – 1.11)	0.220	0.83 (0.58 – 1.18)	0.291	

Figure 4: Survival analysis - KRAS mutant vs wild type

