# **110P ESMO 2022 Characteristics and survival** outcomes of patients with **RET** fusion-positive (**RET**-fp) solid tumours receiving **non-RET** inhibitor therapy in a real-world setting

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# BACKGROUND

- *RET* gene fusions are oncogenic drivers in multiple tumour types.<sup>1,2</sup>
- Current standards of care for patients with *RET*-fp solid tumours have limited efficacy and/or significant off-target toxicity;<sup>2,3</sup> there is a need for efficacious precision therapies that selectively target RET alterations.
- The genomic landscape and natural history of patients with RET-fp solid tumours is unknown
- Using real-world data, we described the clinical characteristics and survival outcomes of patients with RET-fp metastatic solid tumours who received non-selective RET inhibitor therapy, and assessed the prognostic value of RET fusions in solid tumours.

# **O** METHODS

### **Study design**

- Clinical characteristics and survival outcomes for RET inhibitor-naïve patients with metastatic solid tumours were collected from the nationwide (US-based) de-identified Flatiron Health/Foundation Medicine clinico-genomic database (FH-FMI CGDB; version April 2022)
- The de-identified data originated from ~280 cancer clinics in the US (~800 care sites); retrospective longitudinal clinical data were derived from electronic health records, comprising patient-level structured and unstructured data, curated via technology-enabled abstraction, and were linked to genomic data derived from Foundation Medicine comprehensive genomic profiling (CGP) tests in the FH-FMI CGDB by de-identified, deterministic matching.<sup>4</sup>
- Eligible patients had:
- $\geq 1$  documented clinical visit in the Flatiron Health network between 1 January 2011 and 31 March 2022, and underwent ≥1 DNA Foundation Medicine CGP test prior to 1 April 2022
- A visit gap ≤90 days after initial diagnosis.
- Patients with a diagnosis of RET-fp non-small cell lung cancer were excluded.
- *RET* positivity was defined by the presence of a fusion with a predicted known/likely functional status.
- A *RET* wild-type (*RET*-WT) status was determined when the CGP test was unable to detect qualifying RET fusions.

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### Matching of *RET*-WT and *RET*-fp patients

- in a 4:1 ratio, using two models:
- as the index date.

#### **Objectives**

- covariate-matched RET-WT cohorts.
- patterns and genomic alterations
- were assessed.

# **RESULTS**

#### **Patient characteristics**

- comprised 104 patients (Table 1).

#### **Table 1. Baseline characteristics**

#### Characteristic

#### Mean age, years (SD)

**Sex**,\* n (%) Female Male

**Race**, n (%) Asian / Black or African Amer Hispanic or Latino / White

Other / Missing

**ECOG PS**<sup>†</sup>, n (%)

≥2

#### Missing

Prior lines of therapy, n (%)

≥2

Missing

**Practice type**, n (%) Academic Community

Mean time from initial diagno CGP report date, months (SD)

\*Data missing for one patient in the non-matched *RET*-WT cohort. <sup>†</sup>Closest ECOG PS 30 days prior to 7 days after CGP report date. CGP, comprehensive genomic profiling; ECOG PS, Eastern Cooperative Oncology Group performance status; RET-fp, RET fusion-positive; RET-WT, RET wild type.

#### Disclosures

Pilar Garrido reports an advisory board role for Abbvie, Amgen, AstraZeneca, Bayer, GlaxoSmithKline, Janssen, Lilly, MSD, Novartis, Pfizer, Roche, Takeda and Sanofi; Speaker role for AstraZeneca, Janssen, MSD, Medscape, Novartis, Pfizer, Roche, Takeda and TouchIme. For co-author disclosures please refer to the abstract: https://oncologypro.esmo.org/

• Mahalanobis distance matching was used to match *RET*-WT with *RET*-fp patients

- In both models, patients were matched using preselected covariates: age, gender, race, tumour type, practice type, Eastern Cooperative Oncology Group (ECOG) performance status, year of CGP, time from initial diagnosis to CGP report date and number of treatment lines prior to the CGP report date

In model 1, the CGP report date was used as the index date; in model 2, the date of metastatic diagnosis (after adjustment for immortal time bias) was used

• The primary objective was to evaluate overall survival (OS) in the RET-fp and

Secondary objectives included descriptions of patient characteristics, treatment

- Tumour mutational burden (TMB), microsatellite instability (MSI) and oncogenic alterations in ALK, BRAF, ERBB2, EGFR, NTRK, ROS1, MET and KRAS

• Overall, there were 7,220 eligible patients with *RET*-WT solid tumours.

The RET-fp cohort comprised 26 patients and the matched RET-WT cohort

	<i>RET</i> -fp (N=26)	Matched <i>RET</i> -WT (N=104)	Non-matched <i>RET</i> -WT (N=7,116)
	65.3 (10.3)	61.9 (12.0)	64.5 (10.1)
	11 (42.3)	44 (42.3)	4,033 (56.7)
	15 (57.7)	60 (57.7)	3,082 (43.3)
ican	0 / 1 (3.8)	0 / 4 (3.8)	173 (2.4) / 615 (8.6)
	0 / 21 (80.8)	0 / 91 (87.5)	15 (0.2) / 4,597 (64.6)
	3 (11.5) / 1 (3.8)	9 (8.7) / 0	1,120 (15.7) / 596 (8.4)
	7 (26.9)	26 (25.0)	1,656 (23.3)
	7 (26.9)	34 (32.7)	2,339 (32.9)
	1 (3.8)	4 (3.8)	869 (12.2)
	11 (38.5)	40 (38.5)	2,252 (31.6)
	2 (7.7)	8 (7.7)	807 (11.3)
	6 (32.1)	24 (32.1)	2,580 (26.3)
	7 (26.9)	28 (26.9)	1,932 (27.2)
	11 (42.3)	44 (42.3)	1,797 (25.3)
	4 (15.4)	15 (14.4)	999 (14.0)
	22 (84.6)	89 (85.6)	6,117 (86.0)
sis to	8.1 (10.7)	7.9 (8.6)	14.5 (22.9)

• In total, nine distinct *RET*-fp tumour types were identified; the most frequent were colorectal (34.6%), pancreatic (15.4%) and thyroid (15.4%) cancers (**Figure 1**).

• Of nine different *RET* fusion partners detected, the most common were NCOA4 (46.2%), CCDC6 (23.1%) and ERC1 (7.7%)

## Figure 1. Tumour types in the *RET*-fp cohort (N=26)



Patients with RET-fp (NSCLC excluded). \*Neuroendocrine tumours included one gastrointestinal tumour and two unspecified anatomical locations. CUP, cancer of unknown primary; NSCLC, non-small cell lung cancer; RET-fp, RET fusion-positive; SCLC, small cell lung cancer.

#### TMB and MSI

- In the *RET*-fp cohort, 65.4% of patients had a low TMB status (7.7% were TMB-high) and 50.0% had a low MSI status (3.8% were MSI-high) (Table 2)
- One patient in the RET-fp cohort had high MSI and an ERBB2 amplification
- No other assessed oncogenic co-alterations were identified in this cohort.
- Most patients in the matched *RET*-WT cohort also had low levels of TMB (84.6%) and MSI (82.7%) (Table 2)
- Patients in this cohort had an equal rate of *ERBB2* (3.8%), and a higher rate of BRAF (5.8%) and KRAS (37.5%) alterations than patients in the RET-fp cohort.

Co-occurring biomarkers and molecular characteristics*	<i>RET</i> -fp (N=26)	Matched <i>RET</i> -WT (N=104)	Non-matched <i>RET</i> -WT (N=7,116)		
TMB status, n (%) High (≥20 mut/Mb) Medium (<20, ≥5.7 mut/Mb) Low (<5.7 mut/Mb) Missing	2 (7.7) 5 (19.2) 17 (65.4) 2 (7.7)	0 16 (15.4) 88 (84.6) 0	215 (3.0) 1,323 (18.6) 5,578 (78.4) 0		
<b>MSI-high</b> , n (%) Yes No Unknown / Missing	1 (3.8) 13 (50.0) 12 (46.1)	0 86 (82.7) 18 (17.3)	119 (1.7) 6,021 (84.6) 976 (13.7)		
Oncogenic alteration, n (%) ALK BRAF ERBB2 EGFR MET KRAS NTRK ROS1	0 0 1 (3.8) 0 0 0 0	0 6 (5.8) 4 (3.8) 0 0 39 (37.5) 0 0	13 (0.2) 389 (5.5) 300 (4.2) 53 (0.7) 9 (0.1) 2,622 (36.8) 13 (0.2) 11 (0.2)		

\*Variants of known/likely functional status. MSI, microsatellite instability; mut/Mb, mutations per megabase; RET-fp. RET fusion-positive; RET-WT, RET wild type; TMB, tumour mutational burden.

#### Survival outcomes

- When the CGP report date was used as the index date (model 1; Figure 2A):
- Median OS was 6.0 months (95% CI 1.6–9.9) in the RET-fp cohort and 9.7 months (95% CI 6.3–11.7) in the matched *RET*-WT cohort
- The hazard ratio (HR) was 1.8 (95% CI 1.1–3.1).
- When the initial diagnosis date was used as the index date (model 2; Figure 2B):
- Median OS was 6.9 months (95% CI 1.6–9.6) in the *RET*-fp cohort and 11.2 months (95% CI 7.7–16.9) in the matched RET-WT cohort
- The HR was 2.2 (95% CI 1.3–3.7).

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### Table 2. Biomarkers and molecular characteristics

Figure 2. OS using A) CGP report date as the index date (model 1); B) initial diagnosis date as the index date\* (model 2)





#### References



1. Kato, et al. Clin Cancer Res 2017 2. Belli, et al. Clin Cancer Res 2020 3. Drilon, et al. Nat Rev Clin Oncol 2018 4. Singal, et al. JAMA 2019

\*Corrected for immortal time bias.

CGP, comprehensive genomic profiling; OS, overall survival; RET-fp, RET fusion-positive; RET-WT, RET wild type

# **O** CONCLUSIONS

- Despite the small sample size, patients with *RET*-fp solid tumours had a shorter median OS than matched patients with RET-WT tumours and may have an increased risk of death.
- Oncogenic co-alterations were infrequent in patients with *RET-*fp tumours, which suggests that *RET* fusions are the primary oncogenic drivers in these tumours.
- Our data highlight the need for effective RET inhibitors that could improve the survival of patients with solid tumours harbouring **RET** fusions.

# SUMMARY



Median OS: RET-fp vs matched RET-WT Model 1: 6.0 vs 9.7 months Model 2: 6.9 vs 11.2 months





Low frequency of genetic co-alterations in patients with **RET**-fp solid tumours



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