

First results from the RETgistry:

A global consortium for the study of resistance to RET inhibitors in *RET*-altered tumors



Alissa J. Cooper^{1*}, Alexander Drilon², Julia Rotow³, Stephen V. Liu⁴, Oliver Gautschi⁵, Katherine E. Smith⁶, Dae-Ho Lee⁷, Misako Nagasaka⁸, Herbert H.F. Loong⁹, Nathan Pennell¹⁰, Jyoti D. Patel¹¹, Martin Früh¹², Benjamin Solomon¹³, Georg Pall¹⁴, Natalie Uy¹⁵, Jonathan W. Riess¹⁶, Lori Wirth¹, Beow Y. Yeap¹, Justin F. Gainor¹, Jessica J. Lin^{1*}

1. Massachusetts General Hospital Cancer Center, Boston, MA 2. Memorial Sloan Kettering Cancer Center, New York, NY 3. Dana-Farber Cancer Institute, Boston, MA 4. Georgetown University, Washington, DC 5. Cantonal Hospital, Lucerne, Switzerland 6. Mayo Clinic, Rochester, MN 7. Asan Medical Center, Seoul, South Korea 8. University of California-Irvine, Orange, CA 9. The Chinese University of Hong Kong, Hong Kong 10. Cleveland Clinic, Cleveland, OH 11. Northwestern University, Evanston, IL 12. Kantonsspital, St.Gallen, Switzerland 13. Peter MacCallum Cancer Center, Melbourne, Australia 14. Tirol Kliniken, Innsbruck, Austria 15. University of Washington, Seattle, WA 16. University of California-Davis, Davis, CA. *Emails: acooper@mgh.harvard.edu, jilin1@partners.org

BACKGROUND

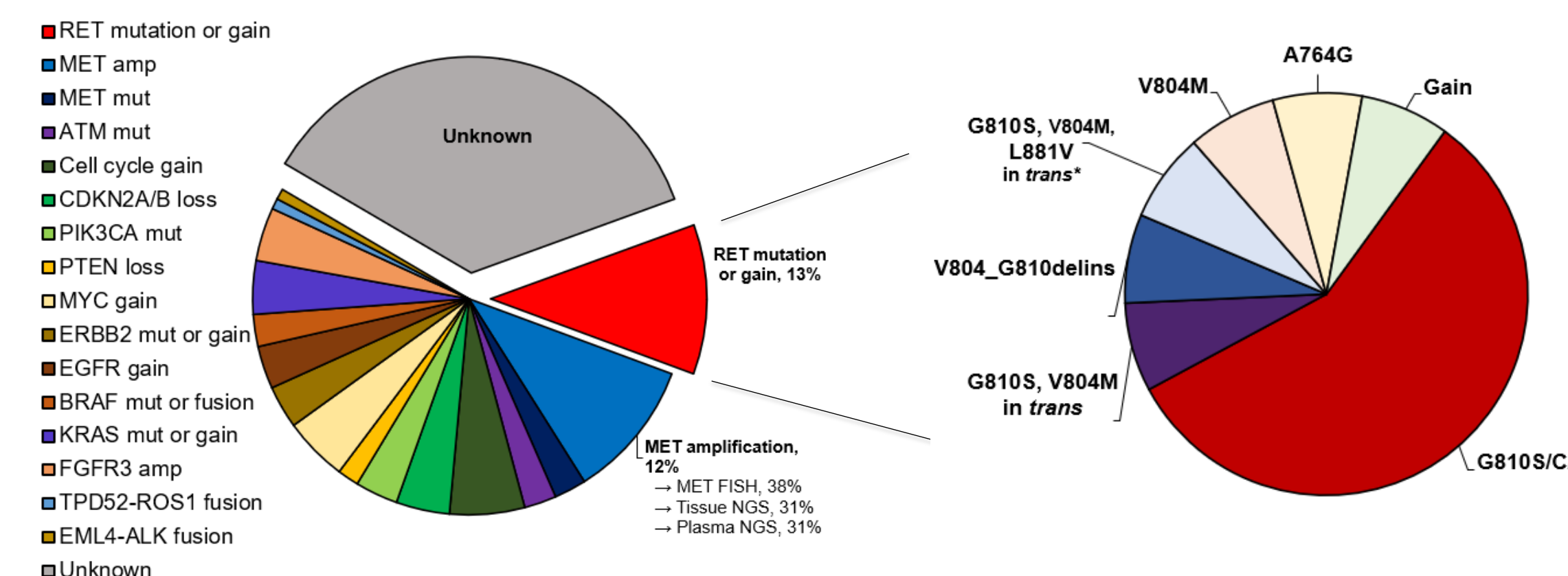
- Rearranged during transfection (*RET*) gene alterations are the oncogenic driver in diverse tumor types. *RET*-selective tyrosine kinase inhibitors (TKIs) selpercatinib and pralsetinib are effective, but acquired drug resistance remains a challenge.
- Here, we report the initial results from the RETgistry, an international consortium aimed at elucidating mechanisms of resistance to *RET* TKIs across *RET*-altered solid tumors.

METHODS

- This was a retrospective analysis performed across 16 institutions.
- Patients (pts) were eligible if they had an advanced solid tumor harboring an oncogenic *RET* alteration, received ≥ 1 *RET* TKI with disease progression, and underwent resistant tumor or liquid biopsy for next-generation sequencing (NGS).

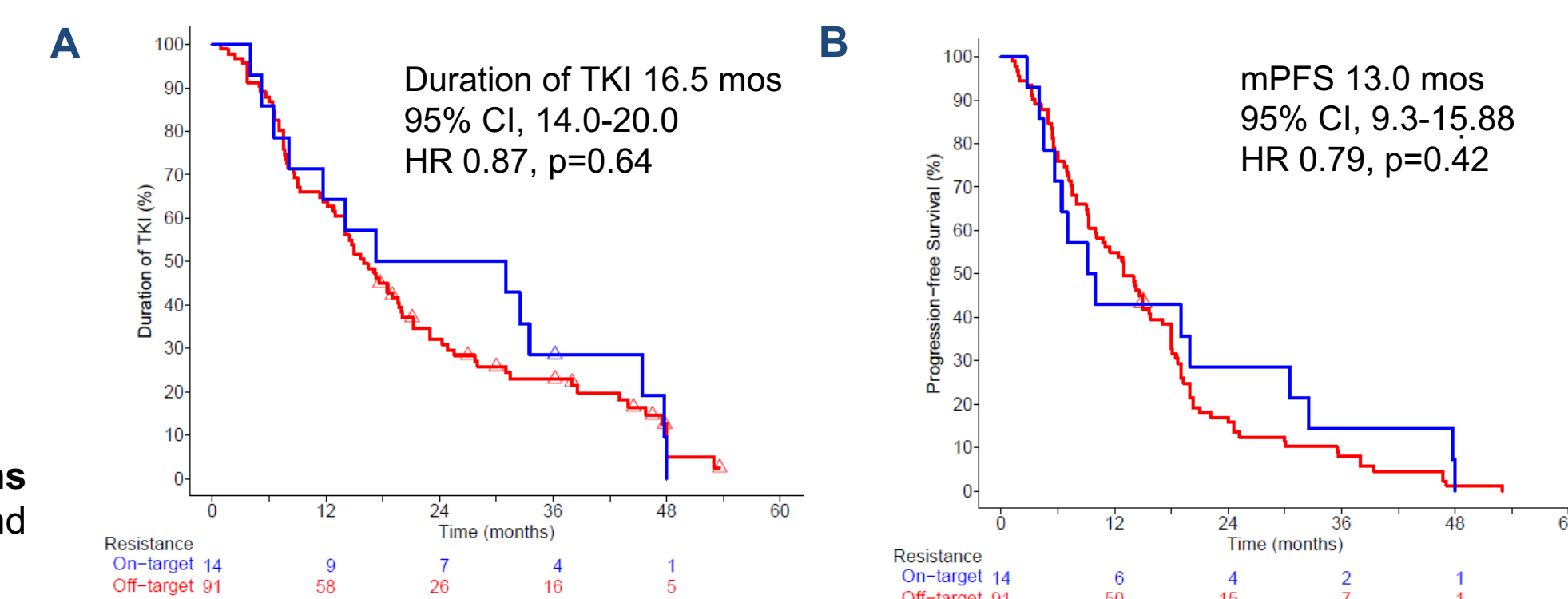
RESULTS

- 105 time-distinct biopsies were included in analysis, obtained from 89 pts with progression on a *RET*-selective TKI (Fig. 1). 97% of samples had baseline NGS.
- Acquired *RET* mutations were detected in 13% (G810X, in 10%) (Fig. 2, 3).
- Potential off-target resistance gene alterations identified in 46 cases (44%) included *MET* amplification (12%), *BRAF* V600E or fusion (3%), *KRAS* gain or mutation (5%), *ERBB2* amplification (2%), *EGFR* amplification (3%), *ROS1* fusion (1%), *ALK* fusion (1%), and activating *PIK3CA* mutation or *PTEN* loss (5%) (Fig. 2, 4).



RESULTS

- On-target mutations were detected in 15% of thyroid cancer samples and 12.9% of NSCLC samples; off-target alterations, in 40% of thyroid cancer and 44.7% of NSCLC samples.
- Median duration of *RET* TKI preceding biopsies [first-line, n=34 (32%); second-line, n=42 (40%); third-/greater-line, n=29 (28%)] was 16.5 months (mos) (95% CI, 14.0-20.0).
- Median PFS was 13.0 mos (95% CI, 9.3-15.88).
- The duration of TKI therapy (HR 0.87, p=0.64) or PFS (HR 0.79, p=0.42) (Fig. 5) did not differ according to the presence of on-target vs off-target resistance.



CONCLUSIONS

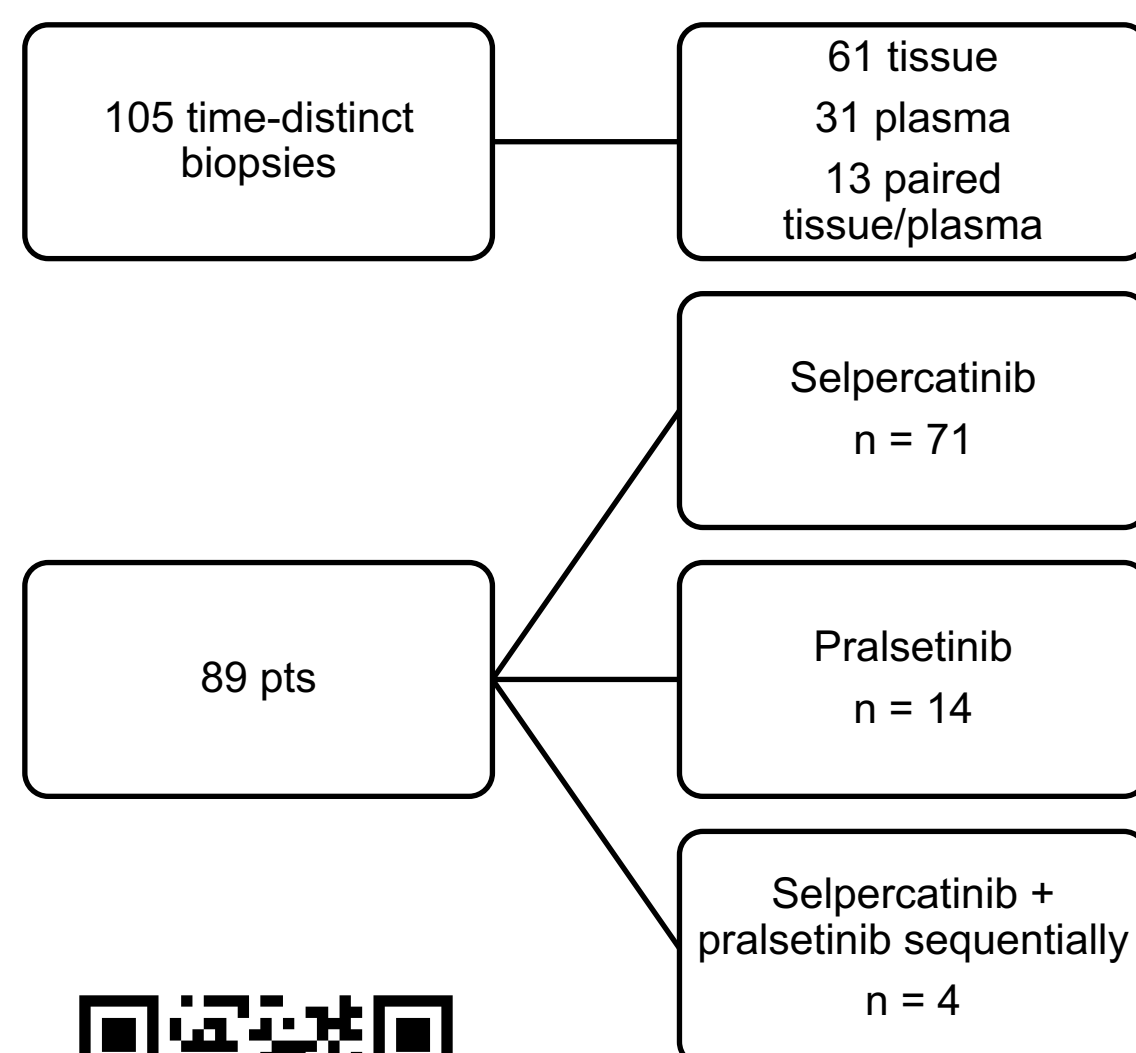
- On-target resistance to *RET* inhibition due to acquired *RET* mutations was less common than off-target resistance, identified in 13% of the analyzed cases.
- Several potentially actionable off-target gene alterations were detected in post-treatment biopsies, (e.g., *MET* amplification, *BRAF* or *KRAS* alterations, *ALK* or *ROS1* fusion).
- Further studies are warranted to enable the development of strategies to address resistance in pts with *RET*-altered tumors.

Acknowledgements: We thank the patients and investigators for their contributions, and The Happy Lungs Project for their support.

Table 1. Characteristics of enrolled pts.

Characteristic	n (%), n=89
Age at diagnosis, median (range)	58 (21-86)
Male	42 (47)
Never or light smoker	79 (89)
Tumor type	n=89
Non-small cell lung cancer (NSCLC)	73 (82)
Medullary thyroid cancer (MTC)	13 (15)
Papillary thyroid cancer	2 (2)
Anaplastic thyroid cancer	1 (1)
NSCLC, RET fusion	n=73
KIF5B-RET	51 (70)
CDC6-RET	15 (21)
Other or unknown	7 (10)
MTC, RET mutation	n=13
M918T	7 (54)
Other	6 (46)
RET TKI line of therapy	n=105
First-line	34 (32)
Second-line	42 (40)
Third- or greater-line	29 (28)
Resistant biopsy obtained, median (range)	15.4 mos (1.8-58.8)
Baseline co-mutations, pre-RET TKI	n=93
TP53	27 (29)
CDKN2A/B loss	12 (13)

Fig. 1. Summary of samples collected from the enrolled pts.



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Fig. 2. Putative on- and off-target resistance mechanisms detected in post-RET TKI biopsies. The diagnostic method used for *MET* amplification detection is listed.

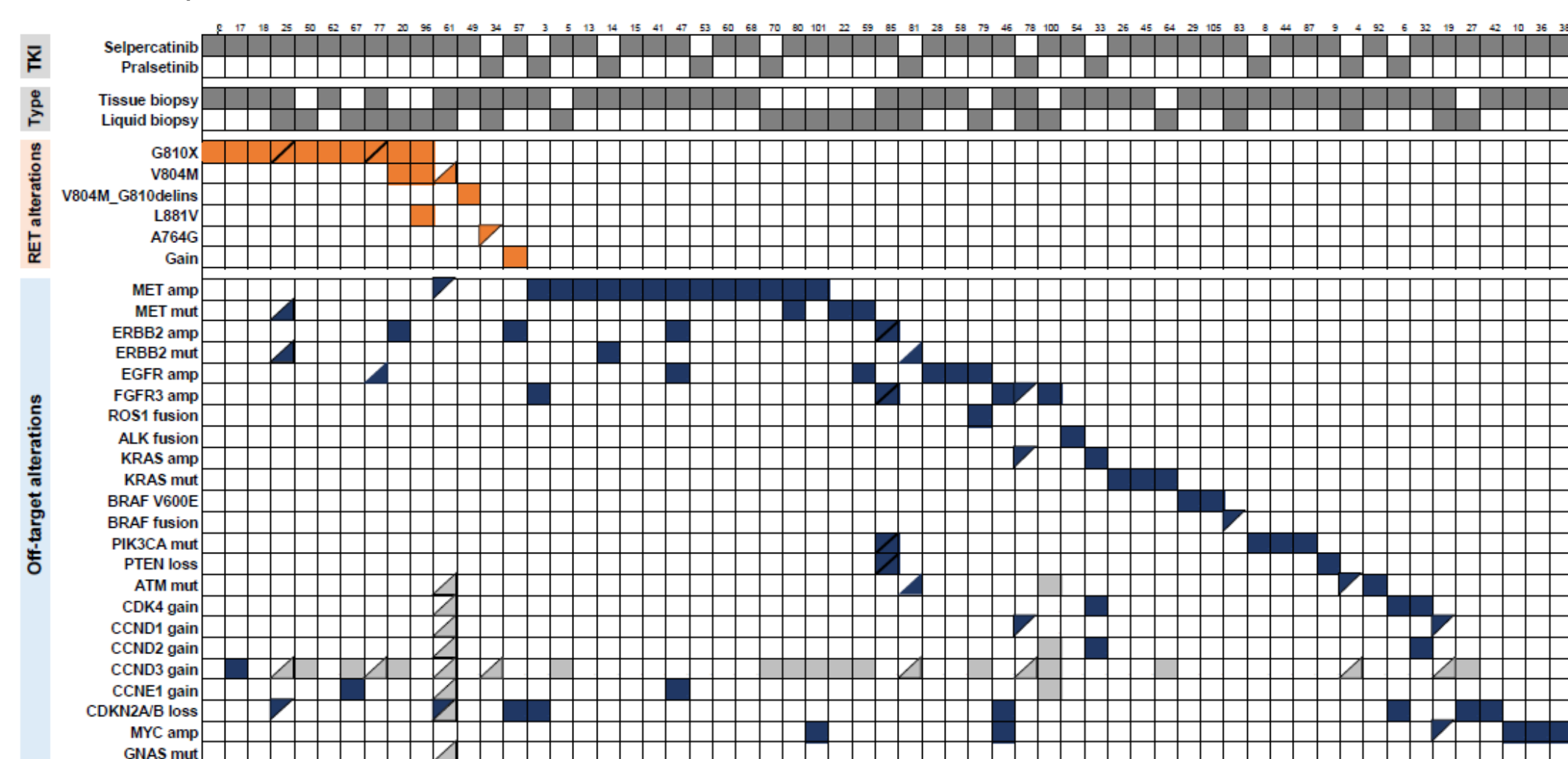


Fig. 4. Heatmap depicting gene alterations detected in post-RET-selective TKI biopsies (n=60/105) with putative on- or off-target resistance mechanisms, as in Fig. 2). Light gray = samples in which the NGS assay did not probe the indicated genes. For samples with paired tissue/liquid biopsies (n=10), findings are shown using the upper (tissue) and lower (liquid) triangle, respectively.