

Positive Quality Intervention: Selpercatinib (Retevmo®) Genomic Testing Management

Description: This PQI is developed to provide guidance to genomic testing with respect to selpercatinib.

Background: *RET*-altered cancers include both *RET* fusions and *RET* mutations. Both alterations involve activating *RET* signaling pathways that promote unwanted cell proliferation in cancers. NCCN guidelines for NSCLC include a Category 2A recommendation for *RET* testing as part of broad molecular profiling in routine clinical practice. In multiple guidelines, *RET* testing is considered as part of a larger initial panel or secondary single analyte test following negative results for other genetic variants such as EGFR, ALK, and ROS1. Molecular testing within *RET*-mutated medullary thyroid cancer (MTC) is applicable as approximately 50% of patients with sporadic MTC have somatic *RET* mutations. In American Thyroid Association, NCCN, and ESMO guidelines, *RET* testing should be considered within the MTC space. Next generation sequencing (NGS) analyzes DNA and/or RNA when detecting *RET*. This method requires a small amount of tissue for multiplex testing for many common and rare cancer-related biomarkers. Tissue testing is often considered as *RET* alteration may not be found in the blood through liquid biopsy and up to 30% of *RET* alterations can be missed if only ctDNA is tested. There are multiple testing methods for *RET* that will help determine patient eligibility for selpercatinib, noting that indicated tumor types are associated with specific alterations (Review Supplemental Information section for approved indications).

<i>RET</i> alteration to test	Associated tumor type(s)
RET-fusion	NSCLC, Thyroid
RET-mutation	MTC

PQI Process:

- Consider the following preferred testing methods when planning for *RET* genomic testing:
 - Perform NGS when applicable
 - Account for 2-4 weeks for test completion
 - Both DNA and RNA-based NGS testing methods are appropriate and care team should discuss the general advantages and disadvantages of both
 - RNA-based NGS is able to reveal unbiased fusion information and there are no intron coverage issues ^{1,2}
 - Reverse Transcription-PCR
 - Quick and relatively inexpensive; test completion with 1-2 days
 - PCR testing is designed predominantly for fusions and *RET* fusion frequency is underestimated
 - FISH
 - High rate of false positive/false negative
 - Should only be considered in rare circumstances (ex. if NGS or RT-PCR are not available)

Patient Centered Activities:

- Provide education to patients regarding genetic testing and what to expect
- Provide testing schedule to patient
- Refer to [Selpercatinib \(Retevmo®\) Management](#) PQI and [Oral Chemotherapy Education \(OCE\)](#) Sheet
- Patient Assistance: [NCODA Financial Assistance Tool](#)

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Supplementary Information:

Selpercatinib is a kinase inhibitor indicated for the treatment of:

- Adult patients with metastatic *RET* fusion-positive non-small cell lung cancer (NSCLC)*
- Adult and pediatric patients 12 years of age and older with advanced or metastatic *RET*-mutant medullary thyroid cancer (MTC) who require systemic therapy*
- Adult and pediatric patients 12 years of age and older with advanced or metastatic *RET* fusion-positive thyroid cancer who require systemic therapy and who are radioactive iodine-refractory (if radioactive iodine is appropriate)*

*Approved under accelerated approval based on overall response rate and duration of response³

References:

1. Garinet S, Laurent-Puig P, Blons H, Oudart JB. Current and future molecular testing in NSCLC, what can we expect from new sequencing technologies? *J Clin Med*.
2. Drilon A, Wang L, Arcila ME, et al. Broad, hybrid capture-based next-generation sequencing identifies actionable genomic alterations in lung adenocarcinomas otherwise negative for such alterations by other genomic testing approaches. *Clin Cancer Res*. 2015;21(16):3631-3639
3. [RETEVMO® \[package insert\]. Lilly USA, LLC. Indianapolis, IN.](#)

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