

Belgian expert consensus for tumor-agnostic treatment of NTRK gene fusion-driven solid tumors with larotrectinib

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ABSTRACT

Fusions of NTRK (neurotrophic tyrosine receptor kinase) genes with 5' partner genes can result in the expression of chimeric proteins that drive oncogenesis through ligand-independent kinase activation. Despite variable frequencies of NTRK fusions in different tumor types, the fact that they are common to a wide range of cancers raises the possibility of developing tumor-agnostic treatments specifically targeting NTRK fusion products, irrespective of tumor type.

The first-generation Trk (tropomyosin receptor kinase) inhibitor, larotrectinib, was the first tumor-agnostic treatment of NTRK fusion-positive cancers in adults and children, to be approved in the European Union.

This consensus, developed by a Belgian multidisciplinary expert panel, aims to highlight the unmet medical need associated to NTRK fusion-driven cancer treatment and, based on current knowledge of NTRK fusions and larotrectinib treatment outcome and safety, provide comprehensive guidance to oncologists regarding NTRK fusion-driven cancer diagnostics and the best use of larotrectinib in real-world clinical settings.

1. NTRK (neurotrophic tyrosine receptor) genes: function and oncogenic fusions

The NTRK genes (*NTRK1*, *NTRK2*, and *NTRK3*) encode the tropomyosin receptor kinase (Trk) family of proteins (respectively TrkA, TrkB, and TrkC), which play a critical role in the modulation of various biological processes and cellular functions (Amatu et al., 2019). Due to intra- or inter-chromosomal rearrangements, fusions of NTRK genes with a wide range of 5' partner genes can occur, leading to the expression of chimeric proteins that drive oncogenesis through constitutive (i. e., ligand-independent) kinase activation (Cocco et al., 2018; Kojadinovic et al., 2021). As a result, NTRK gene fusions act as primary

oncogenic drivers for a broad range of adult and pediatric cancers, including rare tumors frequently associated with NTRK gene fusions and more common tumors where NTRK gene fusions are rarer (Cocco et al., 2018; Kojadinovic et al., 2021; Bourgeois et al., 2000; El Demellawy et al., 2016; Skalova et al., 2010; Tognon et al., 2002; Stransky et al., 2014) and represent an important clinically actionable target (Fig. 1). Moreover, these fusions generally do not co-occur with other known oncogenic driver mutations (Rosen et al., 2020). Detection of NTRK gene fusions can be done by prescreening with immunohistochemistry (IHC) and, in the case of positive IHC results, subsequent confirmation by fluorescence in-situ hybridization (FISH), reverse transcriptase polymerase chain reaction (RT-PCR), or DNA/RNA next-generation

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sequencing (NGS) (Solomon et al., 2019).

2. The case for larotrectinib: a pan-Trk inhibitor

Larotrectinib is a highly-selective, pan-Trk inhibitor that potently blocks the adenosine triphosphate binding site of TrkA, TrkB, and TrkC, with half maximal inhibitory concentration (IC₅₀) values in the low nanomolar range (5–11 nM) (European Medicines Agency, 2021a; Federman and McDermott, 2019).

To date, larotrectinib has been evaluated in three clinical trials: an adult phase I trial (NCT02122913) (Drilon et al., 2018), a pediatric phase I/II trial (SCOUT; NCT02637687) (Laetsch et al., 2018), and an

adult/adolescent phase II basket trial (NAVIGATE; NCT02576431) (Drilon et al., 2018; Drilon et al., 2016). In a pooled analysis of 175 NTRK gene fusion-positive patients from these three studies, larotrectinib was shown to induce an early and durable response. Time to response ranged from 0.9 to 6.6 months, with a median time to response of 1.8 months; 12-month and 24-month durations of response (DOR) were 81 % and 66 %, respectively with an objective response rate (ORR) of 78 %, independent of tumor histology, age, and/or NTRK gene fusion status (McDermott et al., 2020). Larotrectinib has also been shown to cross the blood/brain barrier (Drilon et al., 2019; Ziegler et al., 2018). Analysis of a subset of 24 NTRK gene fusion-positive patients with primary central nervous system tumors showed that larotrectinib was

Adult tumor types

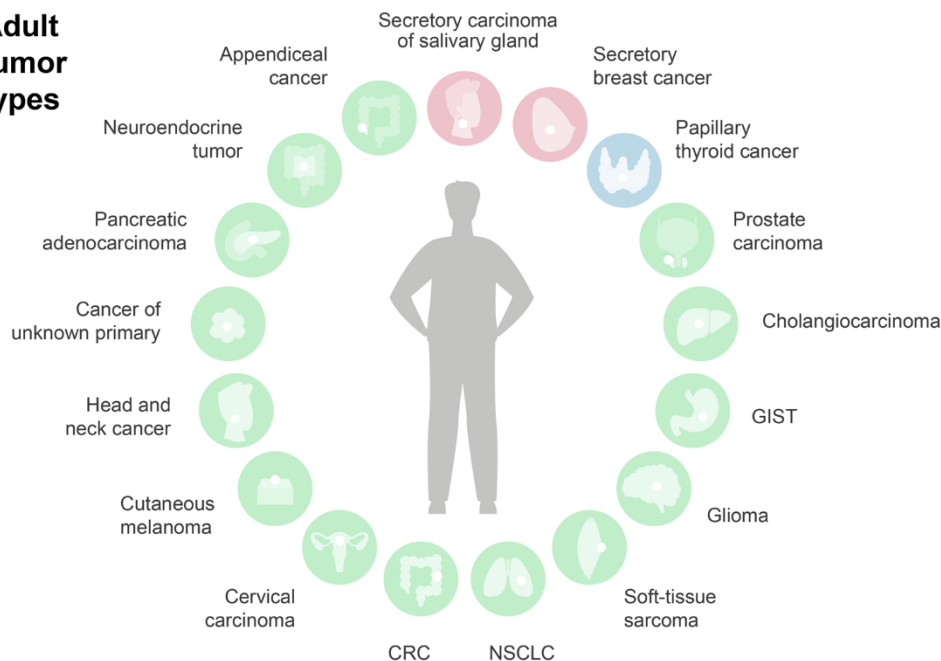
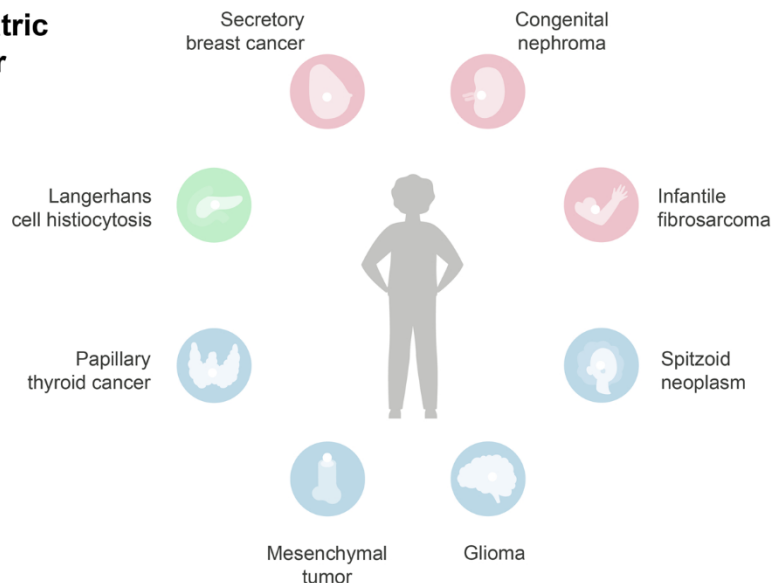


Fig. 1. Prevalence of NTRK gene fusions according to tumor type.


For ranges of NTRK gene fusion prevalence associated to each cancer type and corresponding references, please refer to table T1 of supplementary materials. CRC, colorectal cancer; GIST, gastrointestinal stromal tumors; NSCLC, non-small cell lung cancer; NTRK, neurotrophic tyrosine receptor kinase. This list of tumor types is non-exhaustive.

Pediatric tumor types



NTRK Fusion frequency

High (>90%) Medium (5-40%) Low (<5%)

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active in the central nervous system, with a disease control rate at ≥ 24 weeks of 63 % (Perreault et al., 2020). For larotrectinib efficacy data by tumor type, please refer to Table T2.

Response rates in treatment-naïve patients appeared to be higher than in patients with prior lines of treatment (ORR = 91 % for naïve patients vs. ORR ≥ 70 % for non-naïve patients). However, in light of the high response rates observed for patients with prior lines of treatment and since no cross-resistance between larotrectinib and other treatments was evidenced, it is clear that larotrectinib is also beneficial to previously treated patients (Drilon et al., 2020). In addition, a rapid, sustained, and clinically meaningful improvement of quality of life (QoL) was reported for both adult and pediatric patients with NTRK gene fusion-driven cancers treated with larotrectinib, with improvements in QoL scoring observed for 91 % of adults and 67 % of children who initially reported suboptimal QoL (Kummar et al., 2020).

Overall, larotrectinib is well tolerated. In a pooled safety analysis of 279 patients from the three studies mentioned above (McDermott et al., 2020), the majority (>90 %) of adverse events were grade 1 and 2; the most common grade 3/4 treatment-related adverse events were increased levels of alanine aminotransferase (4 %), decreased neutrophil count (2 %), and anemia (3 %). Dose reductions due to treatment-related adverse events were reported in 11 % of NTRK gene fusion-positive patients and discontinuations in only 2 % of all patients. No treatment-related deaths were reported (McDermott et al., 2020).

In addition to these studies, a global, prospective, multi-cohort, non-interventional phase IV study (NCT04142437, ON-TRK) is currently recruiting patients with the aim of collecting real-world efficacy and safety data for larotrectinib (ClinicalTrials.gov, 2021).

While randomized controlled trials (RCTs) are in general the gold standard to evaluate the efficacy and safety of a given treatment versus the current standard of care (SoC), such study designs are difficult to apply to tumors driven by biomarkers that rarely occur in cancers. Basket trial designs are a valid alternative and are typically preferred to RCTs for the assessment of cancer treatments, such as larotrectinib, that target specific oncogenic genomic alterations rather than specific tumor histologies.

In addition, the growth modulation index (GMI), a metric which uses the patients as their own control, has also been proposed to assess investigational drugs targeting disease for which RCT data are unavailable (Italiano et al., 2020a). GMI is defined as the ratio of progression-free survival (PFS) on the current line of therapy over time to progression on the last prior line of therapy. A drug is deemed to display meaningful clinical activity if the calculated GMI is ≥ 1.33 (Von Hoff, 1998). In an analysis of 122 eligible patients with NTRK gene fusion-positive cancers, 63 % of adult patients and 82 % of pediatric patients had a GMI ≥ 1.33 , suggesting that PFS was improved following treatment with larotrectinib compared to the previous line of treatment (Italiano et al., 2020a; Italiano et al., 2020b). The GMI presents several limitations such as potential differences in how PFS and time to disease progression are assessed for different treatment lines or the fact that it cannot be used to assess front-line treatments. However, in addition to the high ORR observed for larotrectinib, the results obtained using the GMI support the benefits of larotrectinib for the treatment of patients with NTRK gene fusion-positive cancers.

Based on the results of the three aforementioned clinical trials, the use of larotrectinib as tumor-agnostic treatment for both adults and children with NTRK gene fusion-positive solid tumors was approved by the Food and Drug Administration (FDA) in 2018 (Food and Drug Administration, 2018) and by the European Medicines Agency (EMA) in 2019 (European Medicines Agency, 2019). It was the second drug to be approved with tumor agnostic indications by the FDA, and the first by the EMA. According to the EMA, larotrectinib monotherapy is indicated for the treatment of adult and pediatric patients with solid tumors that display a NTRK gene fusion (i) who have a disease that is locally advanced, metastatic, or where surgical resection is likely to result in unsatisfactory treatment options

(European Medicines Agency, 2021a). Larotrectinib is available in oral liquid (20 mg/mL) and capsule (25 and 100 mg) formulations; the recommended dosage is two 100 mg doses a day for adults and two doses of 100 mg/m² of body surface a day for children (with a maximum dose of 100 mg) (European Medicines Agency, 2021a). Besides larotrectinib, another Trk inhibitor, entrectinib, has been approved by the FDA and EMA for use as a monotherapy for the treatment of adult and pediatric patients of at least 12 years of age with NTRK gene fusion-positive solid tumors, and for the treatment of adult patients with ROS1-positive, advanced non-small cell lung cancer (NSCLC) not previously treated with ROS1 inhibitors (Food and Drug Administration, 2021; European Medicines Agency, 2021b). Further discussion relative to entrectinib or other Trk-inhibitors is beyond the scope of the present consensus.

In addition to the fact that NTRK gene fusions act as primary oncogenic drivers, recently also trends suggest the existence of an association between NTRK gene fusions and poor prognosis (Bazhenova et al., 2020), further underlining the importance of early detection and treatment of NTRK gene fusion-driven cancers.

3. Tailoring NTRK gene fusion-driven cancer diagnostics and treatment

The present consensus was developed by a Belgian multidisciplinary expert panel composed of 10 Health Care Professionals specialized in adult and pediatric oncology, and/or pathology. All members of the panel participated in the preparation, review, and finalization of the document, and declare having worked according to their best medical judgement. In addition, though Bayer was involved in the coordination of the manuscript's development, the members also declare having worked freely and without undue influence from external entities. In this document, we, the authors, aim to provide guidance to physicians regarding NTRK gene fusion-driven cancer diagnostics and the use of larotrectinib in clinical settings.

The conclusions of this consensus with regards to NTRK gene fusion-driven cancer diagnostics and treatment, as well as a non-exhaustive list of cancer types to which these may apply, are summarized in Fig. 2. The different categories [AH_{SoC}, AH_{not}, AL_{SoC-T+}, AL_{SoC-T}, AL_{not}, LocA, and Loc] are also referred to in the text below, whenever relevant.

Of note, this consensus was built on the basis of currently available data and knowledge regarding NTRK gene fusion-driven cancers and Trk inhibitors as well as the available standard of care for tumor types. As precision oncology is a rapidly evolving field, the recommendations put forth here may have to be updated in the future should new information come to light.

3.1. Diagnostic approach for NTRK gene fusion-driven cancers

We agree that, ideally, all locally advanced and metastatic solid tumors [AH_{SoC}, AH_{not}, AL_{SoC-T+}, AL_{SoC-T}, AL_{not}, LocA] should be systematically tested for NTRK gene fusions (eventually via pre-screening with IHC followed or directly by confirmatory testing by RNA-NGS, RT-PCR, or FISH) in parallel to other actionable oncogenic drivers. Molecular testing should be conducted at the time of diagnosis, as recommended by the Belgian Personalized Medicine Commission (ComPerMed, 2021).

However, for tumor types that already undergo broad genomic testing via DNA and RNA-NGS at the time of diagnosis [AL_{SoC-T+}] (e.g., advanced lung adenocarcinoma and squamous cell), we suggest the inclusion of NTRK gene fusion testing in the RNA-NGS testing panel. In particular for small biopsies (e.g., small bronchial biopsies in lung cancer patient), DNA-NGS and/or RNA-NGS testing should be prioritized over IHC as NGS analyses deliver optimum results with minimum sample requirements, and the inclusion of NTRK gene fusion testing in the testing panel would entail little to no additional cost or burden of work.

While testing all locally advanced and metastatic solid tumors in



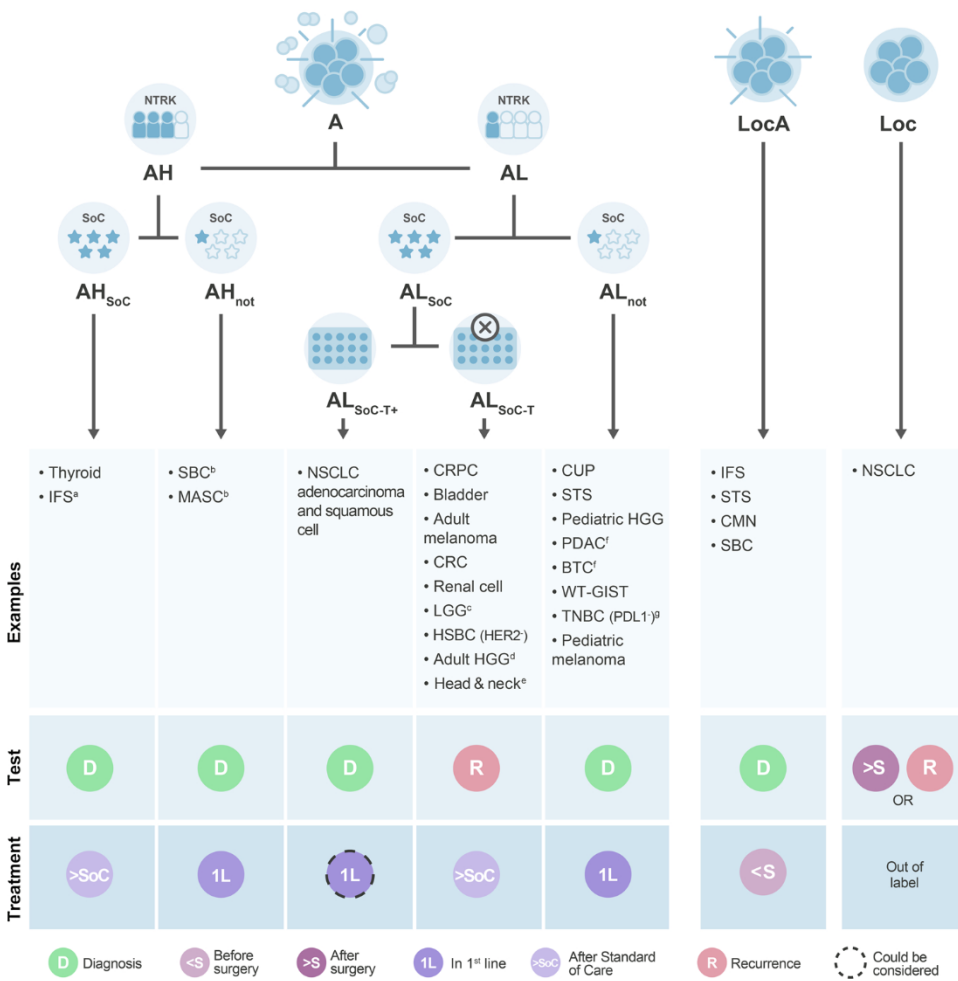


Fig. 2. Algorithm for the timing of NTRK gene fusion testing and larotrectinib treatment for given cancer types according to disease advancement, NTRK gene fusion prevalence, and availability of a satisfactory standard of care.

A, advanced (unresectable/metastatic) solid tumors; **LocA**, locally advanced solid tumors; **Loc**, localized solid tumors (in which clinically actionable information can be useful in later disease stage); **AH**, A tumors with high prevalence of NTRK gene fusions; **AL**, A tumors with low prevalence of NTRK gene fusions; **AH_{SoC}**, AH tumors with satisfactory SoC available; **AH_{not}**, AH tumors with non-satisfactory SoC; **AL_{SoC}**, AL tumors with satisfactory SoC available; **AL_{not}**, AL tumors with non-satisfactory SoC; **AL_{SoC-T+}**, AL_{SoC} tumors on which broad molecular testing is carried out in routine; **AL_{SoC-T}**, AL_{SoC} tumors on which broad molecular testing is not carried out in routine.

For the present consensus, a satisfactory SoC is defined as a SoC significantly improving patient outcome as determined by efficacy (ORR, OS), duration of response (DOR, PFS), time to response, safety, and QoL. The lists of cancer types used as examples for each category are non-exhaustive; tumor types should be assessed on a case-by-case basis to determine if they belong to a given category. ^a For IFS patients, it is reasonable to consider larotrectinib as 1L treatment in case of severe illness and/or toxicity of SoC. ^b Testing for NTRK gene fusions is needed for proper diagnosis of this specific tumor type. ^c With the exception of IDH-WT astrocytoma which is considered as a HGG. ^d For adult glioblastoma, testing should be done at diagnosis and larotrectinib treatment could be considered after the 1L SoC, i.e., chemoradiotherapy. ^e With the exception of salivary gland tumors, for which NTRK fusion testing should be done at diagnosis and larotrectinib treatment considered as 1L. ^f For aggressive forms of PDAC and BTC, larotrectinib could be

considered after 1 L SoC. ^g PDL1-positive subtypes should be tested at recurrence after 1 L SoC and larotrectinib treatment could be considered in 2 L. 1 L, first line treatment; 2 L, second line treatment; BTC, Biliary tract carcinoma; CMN, congenital mesoblastic nephroma; CRC, colorectal cancer; CRPC, castration-resistant prostate cancer; CUP, cancers of unknown primary site; DOR, duration of response; HGG, high grade glioma; HER2, human epidermal growth factor receptor 2; HSBC (HER2⁻), hormone-sensitive HER2-negative breast cancer; IDH-WT, isocitrate dehydrogenase-wild type; IFS, infantile fibrosarcoma; LGG, low grade glioma; NSCLC, non-small cell lung cancer; ORR, objective response rate; OS, overall survival; PDAC, pancreatic ductal adenocarcinoma; PDL1, programmed death-ligand 1; PFS, progression-free survival; SBC, secretory breast cancer; SCC, squamous cell carcinoma; SoC, standard of care; STS, soft tissue sarcoma; TNBC (PDL1⁻), PDL1-negative triple-negative breast cancer; WT-GIST, wild type gastrointestinal stromal tumors; QoL, quality of life.

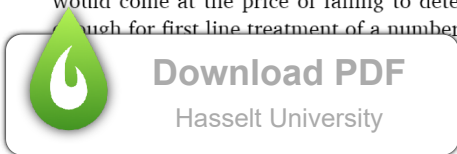
parallel would without doubt enable early detection of clinically actionable oncogenic drivers and generally improve first line treatment choices, several obstacles associated to the reality of clinical settings, such as limitations in testing capacities and in the number of tests that can be carried out on some tumor tissue samples, can impede the testing process. Therefore, we suggest an alternative testing strategy, namely that **systematic NTRK gene fusion testing** should be done at diagnosis, at the very least in rare tumor types or tumor types with unsatisfactory treatment options, such as undifferentiated and anaplastic thyroid cancer, high grade glioma (HGG), pancreatic cancer, cholangiocarcinoma, microsatellite instability-high (MSI-H) colorectal cancer (CRC), salivary gland carcinoma, soft tissue sarcoma (STS) and pediatric cancers, etc. **For other tumor types**, we propose carrying out **NTRK gene fusion testing serially**, after testing for other more common clinically actionable oncogenic drivers. However, such an approach would come at the price of failing to detect NTRK gene fusions early enough for first line treatment of a number of patients.

recurrence in which clinically actionable information can be useful in later disease stage (e.g., HGG, surgical or locoregional stage lung adenocarcinoma, etc.), testing for NTRK gene fusions could be done immediately after surgery or, alternatively, at first disease recurrence or tumor progression.

Finally, for tumor types with a very high prevalence of NTRK gene fusions (>80%), such as secretory breast cancer (SBC), infantile fibrosarcoma (IFS), mammary analogue of the salivary gland carcinoma (MASC), and congenital mesoblastic nephroma (CMN), NTRK gene fusion testing is required for complete, reliable diagnosis. For such cancer types, testing should be carried out at diagnosis using confirmatory testing methods (RNA-NGS, RT-PCR, or FISH) directly rather than an initial IHC prescreening followed by subsequent confirmatory testing.

Additionally, we would also like to highlight the continuous research being performed to facilitate NTRK gene fusion testing further, e.g. by adding plasma next-generation sequencing of circulating tumour DNA (ctDNA) to tissue-based testing the detection rate of oncogenic drivers is

[Loc] with high risk of



increased. Rolfo et al. demonstrated that ctDNA NGS is a rapid, non-invasive screening method for this rare genomic target that may improve identification of patients who can benefit from TRK-targeted therapy (Rolfo et al., 2021).

3.2. Larotrectinib treatment in NTRK gene fusion positive cancers

To define the place of larotrectinib in the treatment sequence for patients with NTRK gene fusion-driven tumors, we propose a categorization based on the degree of **disease advancement** (unresectable/metastatic vs. locally advanced), the **prevalence of NTRK gene fusions**, and, most importantly, the non-satisfactory outcome for the **SoC** of given cancer types.

For the present consensus, tumor types with unmet medical need are defined as cancers lacking a SoC that significantly improves patient outcome, assessed with regards to treatment efficacy (ORR, PFS, overall survival [OS]), DOR, time to response, safety profile, and patient QoL.

The criterion referring to the prevalence of NTRK gene fusions was specifically included in the categorization to highlight the importance of the testing and treatment not only for tumor types with high frequencies of NTRK gene fusions, but also for tumors with lower prevalence of these genetic alterations.

3.2.1. Advanced solid tumors with non-satisfactory SoC

For patients with **advanced** (unresectable, metastatic) solid tumors with **unmet medical need** [AH_{not}, AL_{not}], larotrectinib should be considered as a **first line** treatment regardless of the prevalence of NTRK gene fusions. Examples of tumor types that fit into this category include salivary gland carcinoma, cancers of unknown primary site, most STS (especially rare sub-types), pediatric HGG, pancreatic ductal adenocarcinoma, cholangiocarcinoma, wild-type gastrointestinal stromal tumors (GIST), triple negative breast cancer (with the exception of programmed death-ligand 1 [PDL1] positive subtypes), and pediatric melanoma. Please note that the previous list is non-exhaustive; The treating physician should evaluate on case by case basis if certain tumor types provide his patient adequately satisfactory SoC or not.

Salivary gland carcinoma, such as MASC (known for its very high prevalence of NTRK gene fusions) are considered not to have a satisfactory SoC because response to chemotherapy is limited and presents high toxicity. In these tumor types, NTRK gene fusion testing should be performed at diagnosis and, for NTRK gene fusion-positive cases, larotrectinib should be considered as first line treatment, particularly in absence of other targets such as AR (androgen receptor). This contrasts to other head and neck cancers (mainly squamous cell and lymphoma subtypes), for which a satisfactory SoC is available (see Section 3.2.2 'Advanced solid tumors with satisfactory SoC' below for further information).

Similarly, chemotherapeutic SoC for **cancers of unknown primary site** are currently non satisfactory and these cancer types should be tested for NTRK gene fusions prior to the first line of treatment to keep larotrectinib as an early treatment option.

While a SoC currently exists for **STS**, response rates suggest it does not significantly improve patient outcome. In addition, most sarcoma subtypes are not very sensitive to chemotherapy. NTRK gene fusion testing should therefore be carried out at early stages (in particular for rare sub-types of STS) in order for larotrectinib to be potentially used as a first line treatment.

With regards to **pediatric HGG**, due to the absence of standard treatment guidelines and the fact that the most common therapeutic approaches (surgery, radiotherapy, and chemotherapy) are rarely curative, early testing for NTRK gene fusion and first line treatment with larotrectinib appears of particular interest, in particular for younger children with non-brainstem disease, for which NTRK gene fusion prevalence is notoriously high (Wu et al., 2014). Such an approach was the subject of a case report, where in regression of ETV6-NTRK3

line treatment with larotrectinib (Alharbi et al., 2020). This contrasts with **adult HGG**, for which radiotherapy, generally in combination with temozolomide or other chemotherapy is considered to be a satisfactory first line SoC (please refer to Section 3.2.2 'Advanced solid tumors with satisfactory SoC' below for further information).

NTRK gene fusion testing should be performed at diagnosis for **pancreatic ductal adenocarcinoma** (ideally after excluding the presence of KRAS mutations) and **cholangiocarcinoma**, and, in NTRK gene fusion-positive cases, larotrectinib could be considered as a first line treatment, in particular in young and fit patients. However, as these tumor types are notoriously aggressive, should the treating physician anticipate delays in obtaining results, chemotherapeutic treatment may be initiated while testing is still being carried out. In such cases, larotrectinib may be considered as a second line treatment following this initial round of chemotherapy treatment or, if there is a lack of response or excessive toxicity of chemotherapy, first line treatment may be shifted towards larotrectinib. In addition, if there are indications that disease is progressing despite larotrectinib treatment administration, we advise not waiting three months before re-assessing treatment options.

As **wild-type subtypes of GIST** show relatively poor response to chemotherapy and other treatments, NTRK gene fusion testing should ideally be carried out at diagnosis, and, if the presence of NTRK gene fusions is highlighted, larotrectinib treatment should be considered as a first line treatment.

For **PDL1-negative triple negative breast cancer and breast cancer of the secretory type** (known for its very high prevalence of NTRK gene fusions), we advise NTRK gene fusions to be tested at diagnosis. Larotrectinib can be considered as first line treatment in NTRK gene fusion-positive cases. For PDL1-positive triple negative breast cancer subtypes, NTRK gene fusion testing should be done at disease recurrence or after failure of first line immune-oncology treatment and larotrectinib could be considered as a subsequent line of treatment in positive cases.

Finally, due to its rarity and the subsequent lack of research and satisfactory treatment options, **pediatric melanoma** should also be tested at diagnosis to keep larotrectinib as a potential first line treatment. This contrasts with adult melanoma, for which immune-oncology treatment and BRAF/MEK-inhibitors are considered as a satisfactory first line SoC (see Section 3.2.2 'Advanced solid tumors with satisfactory SoC' below for further information).

3.2.2. Advanced solid tumors with satisfactory SoC

For patients with **advanced** (unresectable, metastatic) solid tumors for which there is a satisfactory **SoC** [AH_{SoC}, AL_{SoC}], larotrectinib treatment should be considered in a **second or later line** (after failure of SoC). Such a strategy may, for example, be applied to metastatic castration-resistant prostate cancer, bladder cancer, renal cell carcinoma, thyroid cancer, SCC and lymphoma subtypes of the head & neck, IFS, adult melanoma, adult HGG, CRC, advanced NSCLC adenocarcinoma and squamous cell subtypes, low grade glioma (with the exception of isocitrate dehydrogenase [IDH] wild type astrocytoma), and hormone-sensitive human epidermal growth factor receptor 2 (HER2) negative breast cancer. Please note that the previous list is non-exhaustive; tumor types should be assessed on a case-by-case basis to determine if they belong to this category.

For **prostate cancer**, specifically **metastatic castration-resistant disease**, NTRK gene fusion testing should be carried out immediately after the first line treatment due to the fact that a high proportion (~40%) of patients have rapidly progressive disease despite treatment with the SoC. This early assessment of the NTRK gene fusion status would enable the use of larotrectinib as a second line treatment for NTRK gene fusion-positive cases. The same approach could be applied to **bladder cancer and renal cell cancer**, in particular if the patient does not respond to treatment with checkpoint inhibitors.

As the NTRK gene fusion-driven variants of **thyroid cancer** appear to be particularly aggressive and also display good responses to Trk

inhibition (Chu et al., 2020), larotrectinib should be administered after failure of radioactive iodine treatment in order to limit the burden and long-term adverse effects of repeated exposure to radioactive treatment, which implies that testing should ideally occur at diagnosis in order to have an immediate next line option in case of relapse.

Head and neck cancers of squamous cell and lymphoma subtype should ideally already be tested at time of diagnosis of metastatic disease, however larotrectinib treatment could be considered after first line SoC. As discussed above, these recommendations do not apply to salivary gland carcinoma (please refer to Section 3.2.1 ‘Advanced solid tumors with non-satisfactory SoC’ above for more information).

Chemotherapy is currently the recommended first line treatment for IFS. However, as argued in a 2020 international consensus, an exception to this guideline should be made for IFS patients with metastatic disease or who require quick treatment response (Orbach et al., 2020). In these specific cases and cases where conventional treatment might result in severe adverse events, it is reasonable to consider larotrectinib as a first line treatment, which implies that testing should ideally occur at diagnosis. In light of the high prevalence of NTRK gene fusions driving IFS (>95%), testing for NTRK fusions is even required to properly diagnose this specific tumor type (Cocco et al., 2018).

For **adult melanoma** (regardless of PDL1 status), NTRK gene fusion testing should be done in all unresectable or metastatic patients for whom other actionable drivers, in particular *BRAF* and *NRAS* mutations, have not been detected, and larotrectinib may be considered after first line immune-oncology treatment. As discussed above, these recommendations do not apply to pediatric melanoma (please refer to Section 3.2.1 ‘Advanced solid tumors with non-satisfactory SoC’ above for more information).

Adult HGG patients should ideally already be tested for NTRK gene fusions at diagnosis, due to the high chances of recurrence. This would enable the use of larotrectinib as a second line treatment for NTRK gene fusion-positive cases, after first line radiotherapy – chemotherapy treatment. As discussed above, these recommendations do not apply to pediatric HGG (please refer to Section 3.2.1 ‘Advanced solid tumors with non-satisfactory SoC’ above for more information).

For **CRC**, NTRK gene fusion testing should ideally be done at diagnosis, however, due to the fact that there is little concordance between the MSI-H/mismatch repair-deficient (MMRd) status and NTRK gene fusion status, testing is likely to be carried out after the first line of treatment. In patients with MSI-H/MMRd tumors undergoing first line pembrolizumab treatment, testing for NTRK gene fusions should be done after this first line so that larotrectinib can be considered for the second line of treatment. In patients with microsatellite stable (MSS)/ mismatch repair (MMR) proficient tumors, NTRK gene fusion testing should be done in wild type patients (*RAS* & *BRAF*-negative) following failure of first line chemotherapy combinations, and larotrectinib could be considered as a subsequent line of treatment. As described in the previous section, if NTRK gene fusion testing is included as part of a broad genomic testing panel (duo DNA & RNA NGS) carried out at the time of diagnosis of metastatic CRC or advanced NSCLC adenocarcinoma and squamous cell subtypes, larotrectinib can only be considered a first line treatment option if the treating physician considers the available treatment option not adequately satisfactory. The treating physician should evaluate on case by case basis if certain tumor types provide his/her patient adequately satisfactory SoC or not (Planchard et al., 2018).

Hormone-sensitive HER2 negative breast cancer patients should be tested for NTRK gene fusions after other treatment options (hormone-based therapy, biological agents, and first line chemotherapy) have been exhausted, and larotrectinib could thereafter be considered for treatment of NTRK gene fusion-positive cases.

Patients with **low grade glioma (with exception of IDH-wild type astrocytoma)** generally have a good prognosis, and current SoC (resection, concurrent chemotherapy and radiotherapy) provides durable control of the disease while being relatively well tolerated. Therefore, testing should be carried

out after disease relapse or if the disease progresses to a HGG subtype (please refer to Section 3.2.1 ‘Advanced solid tumors with non-satisfactory SoC’ above for testing and treatment recommendations for HGG), and, for NTRK gene fusion-positive disease, larotrectinib can be considered as second line treatment. For **IDH-wild type astrocytoma**, which is generally considered and treated as a HGG (please refer to Section 3.2.1 ‘Advanced solid tumors with non-satisfactory SoC’ above), early testing and use of larotrectinib as a second line treatment for NTRK gene fusion-positive disease is preferable.

For cancers where the treating physician considers the available SoC not as adequately satisfactory, the use of larotrectinib as a first line treatment could be considered in order to alleviate the risk of treatment attrition associated with sequential lines of therapy. Examples of this adapted strategy could include **advanced NSCLC adenocarcinoma** or **metastatic CRC patients** with of a positive result following the NTRK gene fusion testing carried out at the time of diagnosis.

3.2.3. Locally advanced solid tumours

For patients with **locally advanced** tumors [LocA] where surgery can result in severe morbidity, larotrectinib could be considered as **neoadjuvant** therapy.

Such a recommendation was previously made by the 2020 international consensus for locally advanced cases of IFS in order to avoid potentially debilitating surgery (Orbach et al., 2020). The successful use of larotrectinib as neoadjuvant therapy in an STS patient was previously reported in Belgium, with the patient achieving a partial response enabling curative limb-sparing surgery (Percy et al., 2021). Similarly, neoadjuvant larotrectinib treatment may also be considered for **congenital mesoblastic nephroma** (CMN; which the present consensus posits may in fact be a renal form of IFS) if the tumor cannot be surgically removed (DuBois et al., 2018), though such cases are rare. **Secretory breast cancer** (both adult and pediatric forms) is another tumor type that is included in this category, with the aim of improving chances of successful resectability during later surgical procedures.

4. Conclusions and perspectives

The benefits of larotrectinib as a tumor-agnostic treatment for adult and pediatric patients with NTRK gene fusion-driven cancers are currently supported by the recent efficacy data, rapid and durable clinical responses, improvements in QoL, and favorable safety profile of the drug. Despite the overall rarity of NTRK gene fusions, the availability of a targeted treatment, added to the fact that gene fusions-positive cancer patients have a relatively poor prognosis, highlights the importance of early testing for this clinically actionable target and subsequent treatment of NTRK gene fusion-positive patients.

While screening of all metastatic and locally advanced patients for NTRK gene fusions should be considered with the aim of providing adequate treatment to target patients as early as indicated, the reality of clinical settings requires some degree of compromise. In this consensus, considering the limitations and obstacles of real-world clinical settings, we propose a categorization of cancer types based on disease advancement, NTRK gene fusion prevalence, and unmet medical need with regards to SoC and, for each category, provide concrete guidance regarding the timing of NTRK gene fusion testing and place of larotrectinib in the treatment algorithm.

Author contributions

Authors contributed equally to the development of the present consensus.

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Bayer SA-NV Belgium set up two expert meetings to discuss the NTRK fusion-driven cancer diagnostics and the best use of larotrectinib



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Declaration of Competing Interest

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

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Appendix A. Supplementary data

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