

Sinonasal tumor

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During the past few years, there has been an expansion in our understanding of [gene fusions](#) and [translocations](#) involved in [cancer](#) of the [sinonasal tract](#).

Larkin et al. review the downstream biologic effects, clinical characteristics, and pathologic features of these tumors. The molecular consequences and neo-antigens resulting from these chromosomal [aberrations](#) are considered and targets for current and future [clinical trials](#) discussed.

Recent findings: Several new, clinically relevant, chromosomal aberrations have been discovered and evaluated to varying degrees in [sinonasal](#) tumors including DEK-AFF2, BRD4::NUT, ADCK4::NUMBL, and ETV6::NTRK3. Sinonasal malignancies demonstrate a diverse genetic landscape and varying clinical courses. Recent studies illustrate that gene fusions and translocations may play a role in [carcinogenesis](#) in certain sinonasal tumor subtypes and may be used to develop new biomarker-driven and patient-centered treatments ¹⁾.

[Sinonasal undifferentiated carcinoma](#)

see [Sinonasal tract schwannoma](#).

[Human papillomavirus-related multiphenotypic sinonasal carcinoma](#)

¹⁾

Larkin R, Hermesen MA, London NR Jr. Translocations and Gene Fusions in Sinonasal Malignancies. Curr Oncol Rep. 2023 Feb 8. doi: 10.1007/s11912-023-01364-x. Epub ahead of print. PMID: 36753024.

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