

HOTAIR

HOTAIR (for HOX transcript antisense RNA) is a human gene located on [chromosome 12](#). It is the first example of an RNA expressed on one chromosome that has been found to influence transcription on another chromosome.

It has been considered as a negative prognostic factor in liver, colon, and laryngeal squamous cancer patients, and identified as a critical [glioma biomarker](#) for tumor grade, molecular subtype, diagnosis, and prognosis. ^{1) 2) 3)}

HOTAIR was reported to promote [glioblastoma](#) (GBM) [cell cycle](#) by regulating a predominant [PRC2](#) complex component [EZH2](#) ⁴⁾.

Knockdown of HOTAIR was found to exert a glioma-suppressive function by regulating the [miR 326/FGF1](#)- signaling pathway in vitro and in vivo, indicating the HOTAIRmiR-326-FGF1 axis as a potential therapeutic strategy for glioma treatment ⁵⁾.

Knockdown of HOTAIR can also increase permeability of the blood-tumor barrier (BTB) by reducing tight junction-related proteins in glioma microvascular endothelial cells via the miR148b-3p/USF1 pathway, facilitating the delivery of antineoplastic drugs ⁶⁾.

In addition, HOTAIR is a direct target of [bromodomain](#) and extraterminal (BET) domain proteins in GBM ⁷⁾ BET proteins are functional requisites for GBM cell growth and malignancy; ⁸⁾ thus, targeting HOTAIR may block BET protein-induced malignancy of GBM and overcome resistance of GBM to BET bromodomain inhibitors (BBIs), which show broad anticancer effects.

¹⁾

Zhang, J.X., Han, L., Bao, Z.S., Wang, Y.Y., Chen, L.Y., Yan, W., Yu, S.Z., Pu, P.Y., Liu, N., You, Y.P., et al.; Chinese Glioma Cooperative Group (2013). HOTAIR, a cell cycle-associated Long non-coding RNA and a strong predictor of survival, is preferentially expressed in classical and mesenchymal glioma. *Neuro-oncol.* 15, 1595–1603.

²⁾

Shen, J., Hodges, T.R., Song, R., Gong, Y., Calin, G.A., Heimberger, A.B., and Zhao, H. (2018). Serum HOTAIR and GAS5 levels as predictors of survival in patients with glioblastoma. *Mol. Carcinog.* 57, 137–141.

³⁾

Tan, S.K., Pastori, C., Penas, C., Komotar, R.J., Ivan, M.E., Wahlestedt, C., and Ayad, N.G. (2018). Serum Long non-coding RNA HOTAIR as a novel diagnostic and prognostic biomarker in glioblastoma multiforme. *Mol. Cancer* 17, 74.

⁴⁾

Zhang, K., Sun, X., Zhou, X., Han, L., Chen, L., Shi, Z., Zhang, A., Ye, M., Wang, Q., Liu, C., et al. (2015). Long non-coding RNA HOTAIR promotes glioblastoma cell cycle progression in an EZH2 dependent manner. *Oncotarget* 6, 537–546.

⁵⁾

Ke, J., Yao, Y.L., Zheng, J., Wang, P., Liu, Y.H., Ma, J., Li, Z., Liu, X.B., Li, Z.Q., Wang, Z.H., and Xue, Y.X. (2015). Knockdown of long non-coding RNA HOTAIR inhibits malignant biological behaviors of human glioma cells via modulation of miR-326. *Oncotarget* 6, 21934–21949.

⁶⁾

Sa, L., Li, Y., Zhao, L., Liu, Y., Wang, P., Liu, L., Li, Z., Ma, J., Cai, H., and Xue, Y. (2017). The Role of HOTAIR/miR-148b-3p/USF1 on Regulating the Permeability of BTB. *Front. Mol. Neurosci.* 10, 194.

⁷⁾

Pastori, C., Kapranov, P., Penas, C., Peschansky, V., Volmar, C.H., Sarkaria, J.N., Bregy, A., Komotar, R., St Laurent, G., Ayad, N.G., and Wahlestedt, C. (2015). The Bromodomain protein BRD4 controls HOTAIR, a Long non-coding RNA essential for glioblastoma proliferation. Proc. Natl. Acad. Sci. USA 112, 8326–8331.

⁸⁾

Xu, L., Chen, Y., Mayakonda, A., Koh, L., Chong, Y.K., Buckley, D.L., Sandanaraj, E., Lim, S.W., Lin, R.Y., Ke, X.Y., et al. (2018). Targetable BET proteins- and E2F1- dependent transcriptional program maintains the malignancy of glioblastoma. Proc. Natl. Acad. Sci. USA 115, E5086–E5095.

From:

<https://neurosurgerywiki.com/wiki/> - **Neurosurgery Wiki**

Permanent link:

<https://neurosurgerywiki.com/wiki/doku.php?id=hotair>

Last update: **2024/06/07 02:54**

