

MEDICINE of THE HIGHEST ORDER



Secondary Acute Lymphoblastic Leukemia in a Patient with Presumptive Weaver Syndrome

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EZH2 overexpression PRC2 BBP43 EZH2 SET FISTOR F	
Figure 2: Mechanism of PRC2 dependent methylation in altering gene expression	
Biological Functions	
Promotion of autophagy and apoptosis	
Promotion of DNA damage repair	
and biological functions of EZH2	

Weaver Syndrome is a rare overgrowth disorder, but case reports have indicated an increased malignancy risk, especially of solid tumors. In general, overgrowth disorders are known to be commonly associated with an increased malignancy risk in childhood. In our review of the contemporary literature, only two cases were identified of patients with Weaver Syndrome developing ALL, both in childhood. If a germline mutation is confirmed in this patient, this case raises many further questions. These questions include: 1) Is this patient's current ALL clone related to the childhood ALL clone? 2) Are patients with germline EZH2 mutations at risk for

the development of secondary ALL? Somatic EZH2 mutations in leukemia are typically associated with poor overall survival outcomes and chemotherapy response. Further follow-up of this case is of great interest.

1.	Basel-Vanagaite L. Acute lympho
2.	Cohen AS, Yap DB, Lewis ME, et a
	307.
3.	Gibson WT, Hood RL, Zhan SH, et

- 2020;10(3):94. doi:10.3390/jpm100300
- phenotype. Am J Med Genet A. 2013; 161(12): 2972-2980.

We wish to thank the family of our case patient for their support. Verbal consent was obtained, and images included with family permission.

Discussion

recurrent de novo ALL?

3) In this case, do the TET2 and/or DNMT3A mutations interact with a germline EZH2 mutation to contribute to

Selected References

olastic leukemia in Weaver syndrome. *Am J Med Genet A*. 2010 Feb;152A(2):383-6. doi: 10.1002/ajmg.a.33244. PMID: 2010167 al. Weaver Syndrome-Associated EZH2 Protein Variants Show Impaired Histone Methyltransferase Function In Vitro. Hum Mutat. 2016;37(3):301

al. Mutations in EZH2 cause Weaver syndrome. Am J Hum Genet. 2012;90(1):110-118. doi:10.1016/j.ajhg.2011.11.018 Hoermann G, Greiner G, Griesmacher A, Valent P. Clonal Hematopoiesis of Indeterminate Potential: A Multidisciplinary Challenge in Personalized Hematology. J Pers Med.

Imagawa E, Higashimoto K, Sakai Y, et al. Mutations in genes encoding polycomb repressive complex 2 subunits cause Weaver syndrome. Hum Mutat. 2017; 38(6): 637-648 Tatton-Brown K, Hanks S, Ruark E, Zachariou A, Duarte SDV, Ramsay E, et al. Germline mutations in the oncogene EZH2 cause Weaver syndrome and increased human height. Oncotarget. 2011; 2 1127–1133. 10.18632/oncotarget.385 Tatton-Brown K, Murray A, Hanks S, et al. Weaver syndrome and EZH2 mutations: Clarifying the clinical

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