1. Chan MP, Andea AA, Harms PW, et al. Genomic copy number analysis of a spectrum of blue nevi identifies recurrent aberrations of entire chromosomal arms in melanoma ex blue nevus. *Mod Pathol.* 2016;29(3):227-239.
2. Costa S, Byrne M, Pissaloux D, et al. Melanomas Associated With Blue Nevi or Mimicking Cellular Blue Nevi: Clinical, Pathologic, and Molecular Study of 11 Cases Displaying a High Frequency of GNA11 Mutations, BAP1 Expression Loss, and a Predilection for the Scalp. *The American journal of surgical pathology.* 2016;40(3):368-377.
3. Dai J, Tetzlaff MT, Schuchter LM, Elder DE, Elenitsas R. Histopathologic and mutational analysis of a case of blue nevus-like melanoma. *Journal of cutaneous pathology.* 2016;43(9):776-780.
4. Gerami P, Pouryazdanparast P, Vemula S, Bastian BC. Molecular analysis of a case of nevus of ota showing progressive evolution to melanoma with intermediate stages resembling cellular blue nevus. *Am J Dermatopathol.* 2010;32(3):301-305.
5. Held L, Metzler G, Eigentler TK, et al. Recurrent nodules in a periauricular plaque-type blue nevus with fatal outcome. *Journal of cutaneous pathology.* 2012;39(12):1088-1093.
6. Maize JC, Jr., McCalmont TH, Carlson JA, Busam KJ, Kutzner H, Bastian BC. Genomic analysis of blue nevi and related dermal melanocytic proliferations. *The American journal of surgical pathology.* 2005;29(9):1214-1220.
7. North JP, Yeh I, McCalmont TH, LeBoit PE. Melanoma ex blue nevus: two cases resembling large plaque-type blue nevus with subcutaneous cellular nodules. *Journal of cutaneous pathology.* 2012;39(12):1094-1099.
8. Yeh I, Fang Y, Busam KJ. Melanoma arising in a large plaque-type blue nevus with subcutaneous cellular nodules. *The American journal of surgical pathology.* 2012;36(8):1258-1263.
9. Shoo BA, Sagebiel RW, Kashani-Sabet M. Discordance in the histopathologic diagnosis of melanoma at a melanoma referral center. *Journal of the American Academy of Dermatology.* 2010;62(5):751-756.
10. Abdel-Rahman MH, Pilarski R, Cebulla CM, et al. Germline BAP1 mutation predisposes to uveal melanoma, lung adenocarcinoma, meningioma, and other cancers. *J Med Genet.* 2011;48(12):856-859.
11. Harbour JW, Onken MD, Roberson ED, et al. Frequent mutation of BAP1 in metastasizing uveal melanomas. *Science.* 2010;330(6009):1410-1413.
12. Testa JR, Cheung M, Pei J, et al. Germline BAP1 mutations predispose to malignant mesothelioma. *Nature genetics.* 2011;43(10):1022-1025.
13. Wiesner T, Murali R, Fried I, et al. A distinct subset of atypical Spitz tumors is characterized by BRAF mutation and loss of BAP1 expression. *The American journal of surgical pathology.* 2012;36(6):818-830.
14. Wiesner T, Obenauf AC, Murali R, et al. Germline mutations in BAP1 predispose to melanocytic tumors. *Nature genetics.* 2011;43(10):1018-1021.
15. Bastian BC, LeBoit PE, Hamm H, Brocker EB, Pinkel D. Chromosomal gains and losses in primary cutaneous melanomas detected by comparative genomic hybridization. *Cancer Res.* 1998;58(10):2170-2175.
16. Bastian BC, LeBoit PE, Pinkel D. Mutations and copy number increase of HRAS in Spitz nevi with distinctive histopathological features. *Am J Pathol.* 2000;157(3):967-972.
17. Bastian BC, Olshen AB, LeBoit PE, Pinkel D. Classifying melanocytic tumors based on DNA copy number changes. *Am J Pathol.* 2003;163(5):1765-1770.

18. Bastian BC, Wesselmann U, Pinkel D, Leboit PE. Molecular cytogenetic analysis of Spitz nevi shows clear differences to melanoma. *J Invest Dermatol.* 1999;113(6):1065-1069.
19. Bastian BC, Xiong J, Frieden IJ, et al. Genetic changes in neoplasms arising in congenital melanocytic nevi: differences between nodular proliferations and melanomas. *Am J Pathol.* 2002;161(4):1163-1169.
20. Boi S, Tebaldi T, Re A, et al. Increased frequency of minimal homozygous deletions is associated with poor prognosis in primary malignant melanoma patients. *Genes Chromosomes Cancer.* 2014;53(6):487-496.
21. Chandler WM, Rowe LR, Florell SR, Jahromi MS, Schiffman JD, South ST. Differentiation of malignant melanoma from benign nevus using a novel genomic microarray with low specimen requirements. *Arch Pathol Lab Med.* 2012;136(8):947-955.
22. Gast A, Scherer D, Chen B, et al. Somatic alterations in the melanoma genome: a high-resolution array-based comparative genomic hybridization study. *Genes Chromosomes Cancer.* 2010;49(8):733-745.
23. Jonsson G, Dahl C, Staaf J, et al. Genomic profiling of malignant melanoma using tiling-resolution arrayCGH. *Oncogene.* 2007;26(32):4738-4748.
24. Kaufmann WK, Carson CC, Omolo B, et al. Mechanisms of chromosomal instability in melanoma. *Environ Mol Mutagen.* 2014;55(6):457-471.
25. Stark M, Hayward N. Genome-wide loss of heterozygosity and copy number analysis in melanoma using high-density single-nucleotide polymorphism arrays. *Cancer Res.* 2007;67(6):2632-2642.
26. Wiesner T, Kutzner H, Cerroni L, Mihm MC, Jr., Busam KJ, Murali R. Genomic aberrations in spitzoid melanocytic tumours and their implications for diagnosis, prognosis and therapy. *Pathology.* 2016;48(2):113-131.
27. Yeh I, de la Fouchardiere A, Pissaloux D, et al. Clinical, histopathologic, and genomic features of Spitz tumors with ALK fusions. *The American journal of surgical pathology.* 2015;39(5):581-591.