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Risk Assessment With Ultra-Low-Pass Whole-Genome Sequencing of Cell-Free DNA for Large B-Cell Lymphoma. Zhao D, Lang N, Liu T, Shelton V, Rangel-Patiño J, Gong IY, Hong M, Travas A, Murad V, Alrekhais I, Metser U, Prica A, Kukreti V, Bhella S, Vijenthira A, Crump M, Kuruvilla J, Arruda A, Minden MD, Lam B, Kridel R.**JCO Precis Oncol**. 2025 Jul;9:e2500200. doi: 10.1200/PO-25-00200. Epub 2025 Jul 9.PMID: 40632977

Single-cell Transcriptional Atlas of Human Hematopoiesis Reveals Genetic and Hierarchy-Based Determinants of Aberrant AML Differentiation. Zeng AGX, Iacobucci I, Shah S, Mitchell A, Wong G, Bansal S, Chen D, Gao Q, Kim H, Kennedy JA, Arruda A, Minden MD, Haferlach T, Mullighan CG, Dick JE.**Blood Cancer Discov.** 2025 Jul 1;6(4):307-324. doi: 10.1158/2643-3230.BCD-24-0342.PMID: 40294241

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