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Explore Recent Scientific Contributions Using Invivoscribe's Products

Key opinion leaders across the world have recently published papers that discuss the advantages of using highly sensitive next-generation sequencing (NGS) for Minimal Residual Disease (MRD) testing.

Key Topics Discussed:

- MRD testing for Multiple Myeloma, CLL & AML with NGS
- Transitioning from Capillary Electrophoresis to NGS
- Advantages of using NGS versus Flow Cytometry



Ig Gene Clonality Analysis Using Next-Generation Sequencing for Improved Minimal Residual Disease Detection with Significant Prognostic Value in Multiple Myeloma Patients

Ha, | et al. (2022) The Journal of Molecular Diagnostics. 24(1):48-56.

Routine Evaluation of Minimal Residual Disease in Myeloma Using Next-Generation Sequencing Clonality Testing

Ho, C et al. (2021) The Journal of Molecular Diagnostics. 23(2):181-199.





NGS Analysis of Clonality and Minimal Residual Disease in a Patient with Concurrent Richter's Transformation and CLL/SLL

Kadkol, SS et al. (2021) Case Reports in Hematology. 2021:9740281.

Immunoglobulin Gene Rearrangement in Koreans with Multiple Myeloma: Clonality Assessment and Repertoire Analysis Using Next-Generation Sequencing Kim, M et al. (2021) *PLoS One.* 16(6):e0253541.





Validation of a Next-Generation Sequencing-Based T-Cell Receptor Gamma Gene Rearrangement Diagnostic Assay: Transitioning from Capillary Electrophoresis to Next-Generation Sequencing

Ho, C C et al. (2021) The Journal of Molecular Diagnostics. 23(7):805-815.

Key Publications:

Simple Deep Sequencing-Based Post-Remission MRD Surveillance Predicts Clinical Relapse in B-ALL

Cheng, S et al. (2018) Journal of Hematology & Oncology. 11(1):105.





Targeted Deep Sequencing Reveals Clinically Relevant Subclonal IgHV Rearrangements in Chronic Lymphocytic Leukemia

Stamatopoulos, B et al. (2017) Leukemia. 31(4):837-845.

A Next-generation Sequencing-Based Assay for Minimal Residual Disease Assessment in AML: Patients with FLT3-ITD Mutations

Levis, M J et al. (2018) Blood Advances. 2(8):825-831.





Establishment of Immunoglobulin Heavy (IGH) Chain Clonality Testing by Next-Generation Sequencing for Routine Characterization of B-Cell and Plasma Cell Neoplasms

Arcila, M E et al. (2018) The Journal of Molecular Diagnostics. 21(2):330-342.

Standardized Minimal Residual Disease Detection by Next-Generation Sequencing in Multiple Myeloma

Yao, Q et al. (2019) Frontiers in Oncology. 9:449.



None of the claims in the publications have been validated by Invivoscribe or reviewed by a regulatory authority



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About Invivoscribe

If you are interested in more information on how to implement NGS LymphoTrack® Assays for clonality and MRD testing on the Illumina® or Thermo Fisher® platforms used in your laboratory, please contact

inquiry@invivoscribe.com



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