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617.ACUTE MYELOID LEUKEMIAS: BIOMARKERS, MOLECULAR MARKERS AND MINIMAL RESIDUAL DISEASE IN DIAGNOSIS AND PROGNOSIS**Establishment of an NPM1 Mutation Copy Number Estimator for Xpert® NPM1 Mutation Test**Mengying(Mona) Liu¹, Mengying Liu¹, Yuanyuan Liu, PhD¹, Huilin Wei, MD PhD², Tran Tran¹, Lin Yuan, MS², Gwo-Jen Day¹¹Cepheid, Sunnyvale, CA²Hematology Oncology R&D, Cepheid, Sunnyvale, CA

Objectives: The nucleophosmin (*NPM1*) is the most mutated gene (~30%) in Acute Myeloid Leukemia (AML)¹. Three *NPM1* mutations (type A, B, and D) represent ~84% in *NPM1*-mutated AML cases while other uncommon subtypes occupy ~16%². Xpert® *NPM1* mutation, an automated cartridge-based test for measuring *NPM1* mutation transcript levels (type A, B and D), is standardized to quantify the amount of mutated *NPM1* relative to *ABL1* control gene based on delta Ct in peripheral blood³. Since mutated *NPM1* level is crucial for risk assessment, medication selection, and ongoing therapeutic monitoring in AML^{4,5}, it is important to obtain the *NPM1* mutation copy number (CN). The aim of this work is to develop *NPM1* mutation CN estimator and to compare %*NPM1* mutation/*ABL1* reporting between delta Ct-based and CN-based methods.

Methods: Five levels of *NPM1* mutations (A, B, D) and *ABL1* IVT-RNA panels as well as two lots of Xpert® *NPM1* mutation tests were used to generate standard curves for CN and %CN reporting. The cell lysates from cell lines carrying either *NPM1* mutation A, B, or D and AML clinical samples containing *NPM1* mutations were examined to evaluate the CN and %CN between two lots of the Xpert® *NPM1* mutation tests and to compare the delta Ct-based and CN-based methods for reporting %*NPM1* mutation/*ABL1*.

Results: Linearity was demonstrated in Ct vs CN input for *NPM1* mutation and *ABL1* with R² above 0.96 for Lot1 and Lot2. Less than 3-fold difference was exhibited for CN and %CN across two lots of Xpert® *NPM1* mutation test. Less than 3-fold difference was observed in %*NPM1* mutation/*ABL1* reporting between delta Ct-based and CN-based approaches.

Conclusion: An *NPM1* mutation copy number estimator for Xpert® *NPM1* mutation test was established, which will provide diagnostic and prognostic values for *NPM1*-mutated AML patients.

References:

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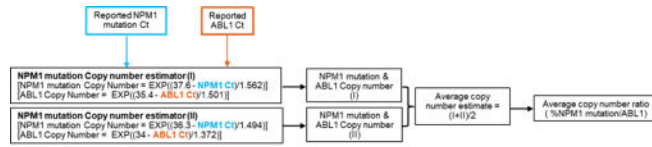


Figure 1: Two sets of NPM1 mutation copy number estimator for **Xpert®** NPM1 mutation test, which will provide diagnostic and prognostic values for NPM1-mutated AML patients. Enter reported NPM1 mutation Ct and ABL1 Ct into the formulas (I) and (II) to calculate the copy number. Average copy number of NPM1 mutation and ABL1 from both formulas will be utilized in obtaining averaged %NPM1 mutation/ABL1.

Figure 1

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