Chromosome Analysis, Bone Marrow with Reflex to Genomic Microarray

Patient: [redacted] DOB: [redacted] Age: 17 Gender: M **Patient Identifiers:** [redacted] **Visit Number (FIN):** [redacted]	Client: [redacted] Physician: [redacted]	ARUP Test Code: 2007130 Collection Date: 03/20/2018 Received in lab: 03/21/2018 Completion Date: 03/29/2018

Interpretation

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 Specimen received

 Specimen type:       Bone Marrow (Right)
 Reason for referral: B-Cell neoplasm; Burkitt Lymphoma, DLBCL,
B-ALL
 Test performed:      Chromosome Analysis


 Laboratory analysis

 Number of cells counted:    25
 Number of cells analyzed:   25
 Number of cells karyotyped: 18
 ISCN Band level:            400
 Banding Method:             G-Banding

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 Chromosome Results: 47,XY,t(8;14)(q24;q32),+mar[1]/46,XY[24]
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 Diagnostic Impression:
Two cell lines were detected in multiple cultures from this
patient. One cell line showed a translocation between
chromosomes 8 and 14, and gain of a marker chromosome in 1/25
(4%) cells.

The remaining 24/25 (96%) cells showed a normal male chromosome
complement.

t(8;14) and marker was observed in only a single cell. In
general, a single cell abnormality does not define an abnormal
clone and is interpreted as in-vitro artifact. However, in this
case, the presence of t(8;14) and marker was observed in a
concurrent chromosome study from a Left side specimen (ARUP
accession 18079118552). Therefore, this single cell is presumed
to be part of an abnormal clonal population based on concurrent
chromosome studies.

The (8;14)(q24;q32) translocation resulting in IGH/MYC
rearrangement is a recurrent abnormality observed in lymphoid
disorders, most frequently in non-Hodgkin lymphomas, especially
Burkitt lymphoma. Clinical correlation is recommended.

Since this analysis revealed an abnormal result, the cytogenomic
microarray analysis, which was ordered as a reflex study, will
not be performed unless we are notified otherwise. Please
contact ARUP Client Services at 1-800-242-2787 if you want to
pursue the microarray analysis.

NOTE: FISH P ALL and FISH ALYMP were performed on this sample
and reported under ARUP accessions 18079118697 and 18079118698
respectively. FISH results were NORMAL. With low level
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mosaicism, the sensitivity of the technique that detects mosaicism may vary. The culture technique may have enriched the abnormal cell line in chromosome analysis.

This result has been reviewed and approved by Xinjie Xu, PhD, FACMG
Electronic Signature

NOTE: Only the Chromosome Analysis, Bone Marrow results are reported on this enhanced report. When the result of Cytogenetics Result is either "no growth" or "normal," Cytogenomic SNP Microarray - Oncology is performed. If the cytogenomic microarray test was performed, those results can be accessed via a patient report or electronic medical records system after the genomic microarray has been completed.

Test developed and characteristics determined by ARUP Laboratories. See Compliance Statement B: aruplab.com/CS

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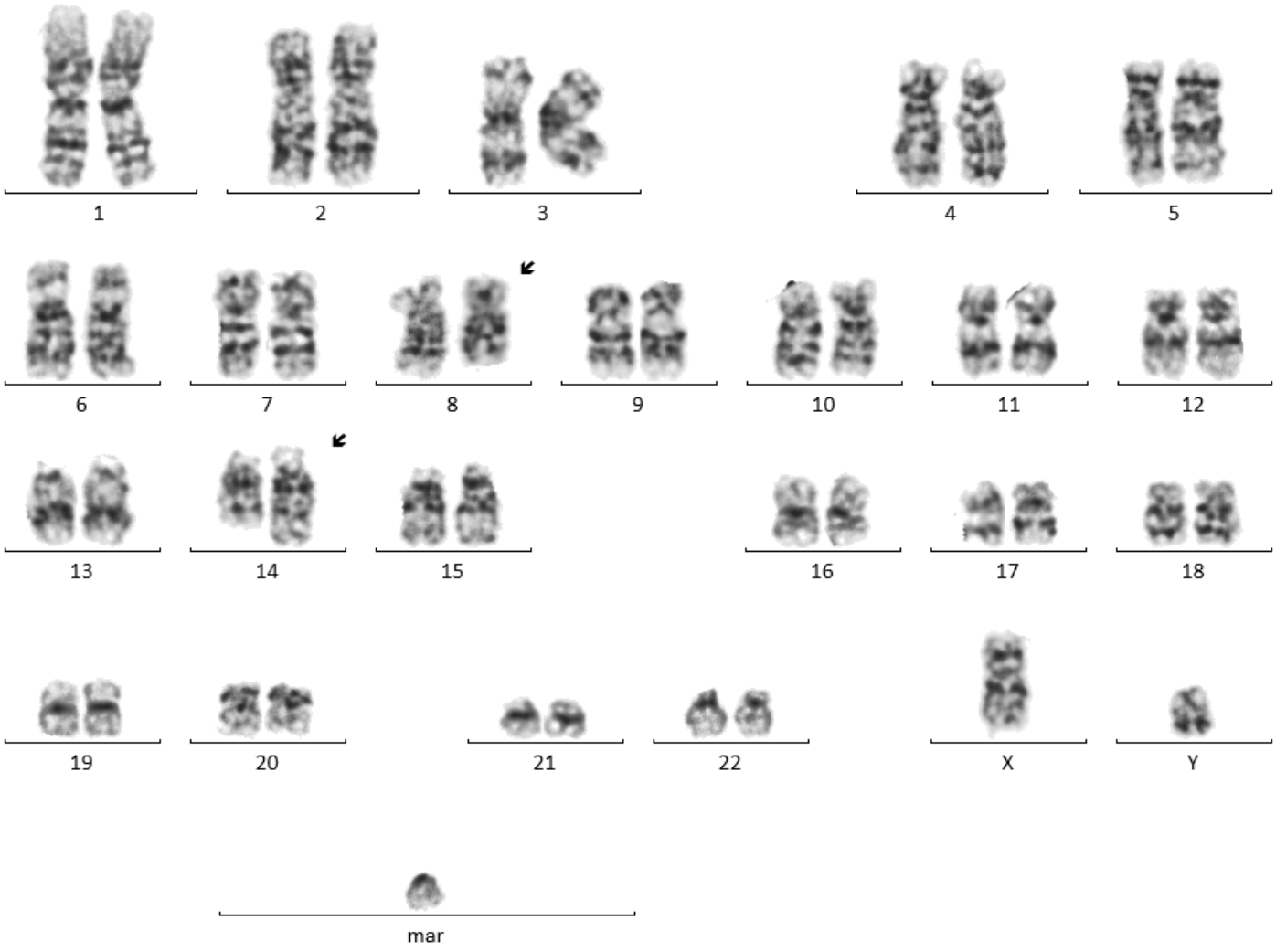
Patient: [redacted] | **Date of Birth:** [redacted] | **Gender:** M | **Physician:** [redacted]
Patient Identifiers: [redacted] | **Visit Number (FIN):** [redacted]

Slide ID: 0064



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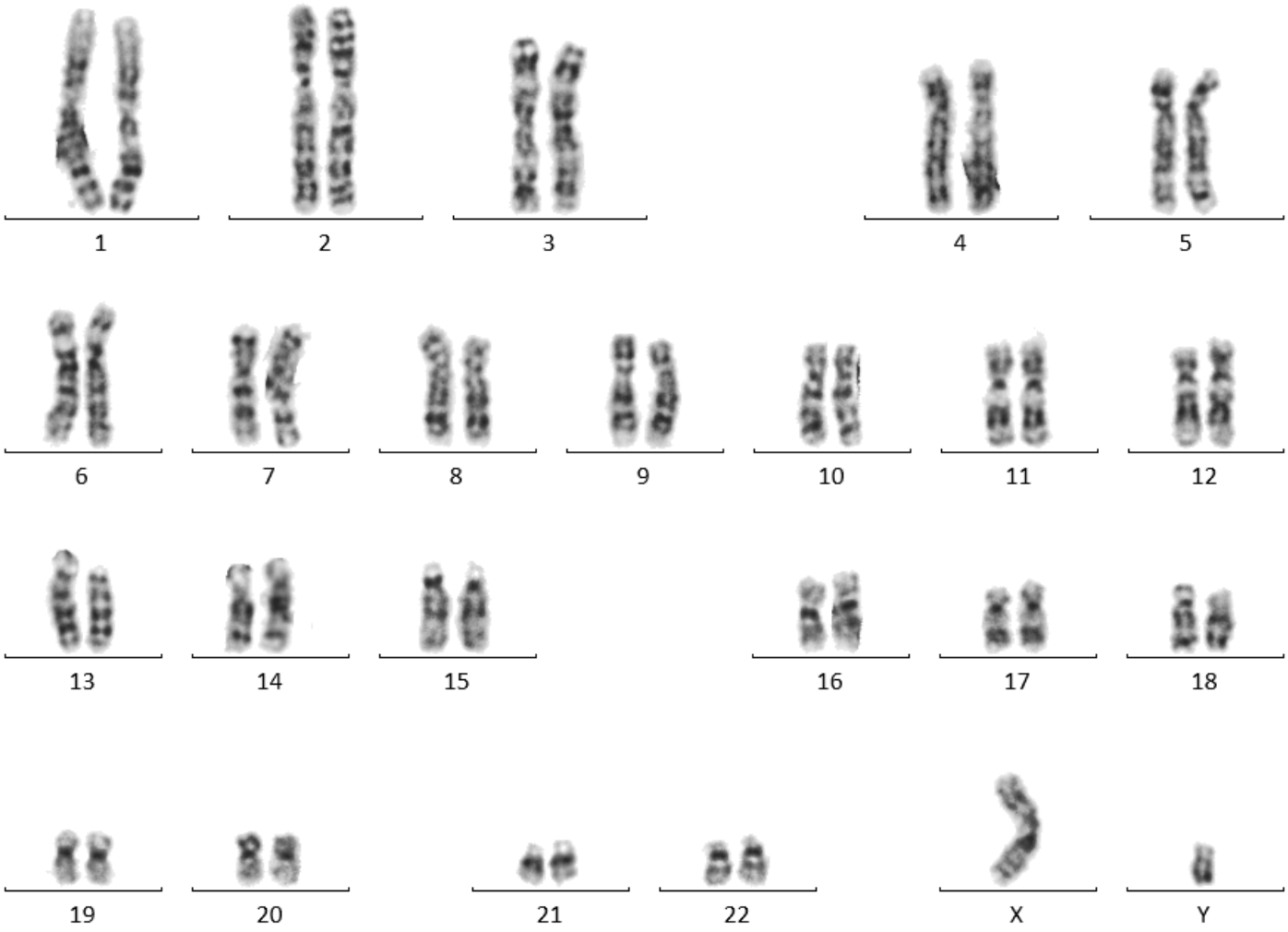
Patient Identifiers: [redacted] | **Visit Number (FIN):** [redacted]

Slide ID: 0090



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