





203-787-1717 203-901-1289



Patient: Jane A. Doe

**DOB/Gender:** 10/10/61 (58 yrs) - Female

Patient ID/MRN: 123456 Date Collected: 04/09/2019



Case#/Status: X19-00352 - Final

Report Category: **Neoplastic** 



Provider: John Doe, M.D.

Hematology Oncology Associates

Tel: 800-123-4567 Fax: 800-765-4321



Lymphoplasmacytic lymphoma (see comment)



### COMMENT

The overall findings are consistent with involvement by a LOW GRADE NON-HODGKIN B-CELL LYMPHOMA. Although the plasma cell component is morphologically limited and a clonal population of plasma cells was not definitively detected by flow cytometry or immunohistochemistry, in the setting of a monoclonal IgM of approximately 2 g/dL, the presence of a MYD88 mutation is most consistent with involvement by a LYMPHOPLASMACYTIC LYMPHOMA. Correlation with clinical and laboratory findings is advised.

All myeloid and lymphoid neoplasms are now classified and named in accordance with the newly revised 2017 version of the WHO Classification of Tumours of Haematopoietic and Lymphoid Tissues.



# **COMPONENT DIAGNOSES**

Biopsy: Non-Hodgkin B-cell lymphoma (see comment)

Aspirate: Maturing trilineage hematopoiesis with frequent small lymphocytes Flow Cytometry: Suspicious for a non-Hodgkin B-cell lymphoproliferative disorder (see

comment)

Normal female karyotype Karyotyping:

No Clonal Abnormalities Detected with probes specific for recurrent FISH:

abnormalities in Plasma Cell Myeloma (see interpretation for probes

tested)

Molecular: MYD88 mutation(s) detected: P.L265P (see comment)





### **CLINICAL DATA**

Monoclonal paraproteinemia.

Received CBC, reported on 4/9/2019: WBC 6.3; RBC 4.36; HGB 12.9; HCT 39.6; MCV 91; MCH 29.7; MCHC 32.6; RDW 14.4; PLT 213; MPV 7.1; LYM 37.4%; MON 10.1%; NEU 'NP'; EOS 'NP'; BAS 'NP' (NP = not provided)

**Electronically Signed By:** Frank Bauer, MD (04/18/19 17:39)





Bone marrow, core & clot biopsies: non-Hodgkin B-cell lymphoma (see comment)



The findings are consistent with involvement by a NON-HODGKIN B-CELL LYMPHOMA with a differential that includes a marginal zone lymphoma and lymphoplasmacytic lymphoma. Correlation with clinical and laboratory findings is advised.



# MICROSCOPIC DESCRIPTION

Marrow Cellularity: Mildly hypercellular (60%)

Infiltrate: Approximately 30% of the cellularity is comprised of an interstitial infiltrate of

lymphocytes that are small sized with round nuclei, condensed chromatin, indistinct nucleoli and

small amounts of cytoplasm occurring singly and in one large aggregate.

Myeloid Maturation: Normal Erythroid Maturation: Normal

Myeloid/Erythroid Ratio: Mildly increased

Megakaryocytes: Normal in number and morphology

Granulomas: Not identified

Iron Stain: No stainable iron is seen in this decalcified specimen

Marrow Reticulin: Mild increase in fibrosis is noted in association with the lymphoid aggregate

Marrow Trabeculae: Normal

Clot preparation: Similar findings to the core biopsy

PAS / Giemsa: Examined

Special Stains: Giemsa, Iron, PAS, Reticulin

Immunostains: CD20, CD79a and Pax-5 highlight numerous lymphocytes. An aggregate of CD3 positive T cells is seen. CD138 positive plasma cells account for less than 5% of the marrow

cellularity and are polyclonal for immunoglobulin kappa and lambda light chains.

#### **Additional Studies:**

| Stain                    | Result                            |
|--------------------------|-----------------------------------|
| CD3 (MRQ-39)             | See microscopic description above |
| CD20 (L26)               | See microscopic description above |
| CD79a (SP18)             | See microscopic description above |
| CD138/syndecan-1 (B-A38) | See microscopic description above |
| Kappa (L1C1)             | See microscopic description above |
| Lambda (Lamb14)          | See microscopic description above |
| PAX5 (SP34)              | See microscopic description above |

**Electronically Signed By:** Frank Bauer, MD (04/12/19 17:37)

#### **GROSS DESCRIPTION:**

- 1. The specimen is received in formalin labeled with the patient's initials and requisition number, and consists of 1 piece of bone marrow core measuring 1.0 x 0.2 x 0.2 cm. The specimen is submitted in 1 cassette after decalcification.
- 2. The specimen is received in formalin labeled with patient's initials and requisition number, and consists of 1 piece of bone marrow clot measuring  $0.5 \times 1.4 \times 1.4$

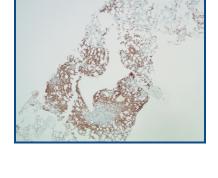
Disclaimer: The adequacy of staining is verified by the appropriate positive and negative controls. The reagents used for these assays are analyte specific reagents (ASR). Their performance characteristics have been validated by Precipio, Inc., New Haven, CT. They have not been reviewed by the FDA. The FDA has deemed that such approval is unwarranted. These assays are for clinical use and should not be viewed as experimental or "research use only".



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Patient: Jane A. Doe

Case #: X19-00352



CD79a





Bone marrow, aspirate: Maturing trilineage hematopoiesis with frequent small lymphocytes



The marrow aspirate smear is spicular and cellular with maturing trilineage hematopoiesis and scattered small lymphocytes. Megakaryocytes are normal in number and morphology. The myeloid: erythroid (M:E) ratio is approximately 8:1. Erythroid maturation is normal. Myeloid maturation is normal. No increase in immature cells is noted. Scant, focal iron is seen without ring sideroblasts on iron stain of the marrow aspirate.

Number of cells counted: 222

| Cell Type        | Percent | Ref. Range    |
|------------------|---------|---------------|
| Blasts           | 0%↓     | 0.3 - 3.0 %   |
| Immature myeloid | 9%↓     | 12.0 - 21.0 % |
| Mature myeloid   | 63 % 🕇  | 35.0 - 55.0 % |
| Eosinophils      | 3 %     | 1.0 - 3.0 %   |
| Basophils        | 0 %     | 0.0 - 1.0 %   |
| Lymphocytes      | 16 % 🕇  | 10.0 - 15.0 % |
| Plasma cells     | 1%      | 0.0 - 1.0 %   |
| Monocytes        | 0 %     | 0.0 - 1.0 %   |
| Erythroid        | 8%↓     | 15.0 - 25.0 % |
| M:E ratio        | 8:1 🕇   | 2 - 4:1       |

**Electronically Signed By:** Frank Bauer, MD (04/12/19 17:37)







Bone marrow, aspirate: Suspicious for a non-Hodgkin B-cell lymphoproliferative disorder (see comment)



# **COMMENT**

The differential diagnosis includes lymphoplasmacytic lymphoma and a marginal zone lymphoma. Most cases of chronic lymphocytic lymphoma, mantle cell lymphoma and follicular lymphoma will express either CD5 or CD10. Correlation with the concurrent bone marrow core and aspirate morphology and cytogenetic findings is advised.

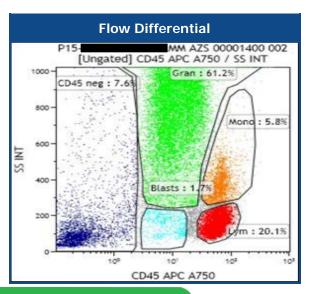


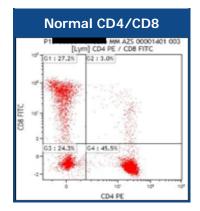
# INTERPRETATION

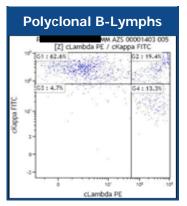
The lymphocytes (20%) include 17% B-cells with an excess of immunoglobulin kappa to lambda light chains (5:1), but are negative for CD5 and CD10. Seventy-four percent (74%) of the lymphocytes are mature T-cells with a normal CD4/CD8 ratio, and 10% natural killer (NK) cells. Less than 1% of the cellularity are plasma cells. Although excess cytoplasmic kappa light chain is detected, no cohesive population of cells is observed.

### **RESULT**

Analysis Time: 4/12/19 13:04 Viability: 98% (Normal > 80%) Specimen: BM, Lavender-top tube







### Flow Cytometry Differential

| Lymphocytes:   | 20% |
|----------------|-----|
| Monocytes:     | 6%  |
| Granulocytes:  | 61% |
| Plasma Cells:  | <1% |
| Blasts:        | 2%  |
| nRBC & Debris: | 8%  |





|     | Plasma Cells   |  |
|-----|--|--|
| %   | Marker   | %  |
| 82  | CD19   | 93   |
| 74  | CD56   | 17   |
| 48  | сКарра   | 54   |
| 68  | cLambda  | 4  |
| 81  | IgA  | 19   |
| 25  | IgG  | 47   |
| 2   | IgM  | 87   |
| 17  |  |  |
| 17  |  |  |
| 13  |  |  |
| 100 |  |  |
| 10  |  |  |
| 13  |  |  |
| 6   |  |  |
|     | 82<br>74<br>48<br>68<br>81<br>25<br>2<br>17<br>17<br>13<br>100<br>10 | %       Marker         82       CD19         74       CD56         48       cKappa         68       cLambda         81       IgA         25       IgG         2       IgM         17       17         13       100         10       13 |

**Electronically Signed By:** Frank Bauer, MD (04/12/19 17:38)

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Bone marrow, aspirate: Normal female karyotype



KARYOTYPE "ISCN": 46,XX[20]; Normal Female Karyotype

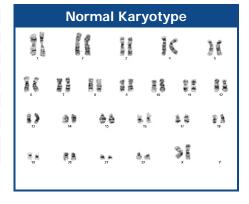
Conventional cytogenetic analysis shows a female karyotype with no evidence of an acquired clonal abnormality. This does not exclude the possibility of an abnormality that cannot be detected at the chromosomal level or exists at a low residual level.

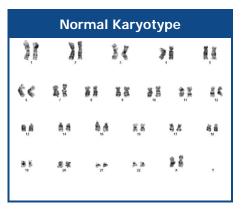
Interpretation of this specimen's cytogenetic results should be made in conjunction with morphologic, immunophenotypic, and clinical findings. The results of this analysis do not exclude the possibility of genetic alterations below the band-resolution of this test or abnormalities due to other etiologies.

FISH studies are recommended and much more sensitive than G-band analysis for cases of plasma cell neoplasia because plasma cells have a very low proliferation rate in culture. FISH studies are recommended for clinically-significant abnormalities. (NCCN Guidelines, ver 2.2013, Multiple Myeloma, National Comprehensive Cancer Network, nccn.org)

### **Analysis**

| Cells Counted:    | 20                         |
|-------------------|----------------------------|
| Cells Analyzed:   | 20                         |
| Cells Imaged:     | 3                          |
| Cells Karyotyped: | 3                          |
| Band Level:       | 450                        |
| Banding Type:     | G-Banding                  |
| Indication:       | Monoclonal paraproteinemia |
|                   |                            |





**Electronically Signed By:** Frank Bauer, MD (04/18/19 16:40)







Bone marrow, aspirate: No Clonal Abnormalities Detected with probes specific for recurrent abnormalities in Plasma Cell Myeloma (see interpretation for probes tested).

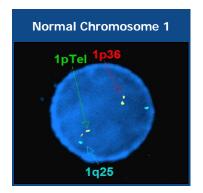


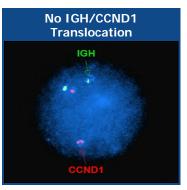
FISH "ISCN": nuc ish (p58x2,1q25x2)[200],(CEP9x2)[200],(CCND1x2,IGHx2)[200],(D13S319x2,13q34x2)[200],(p53x2)[200]

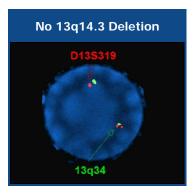
Fluorescence in situ hybridization (FISH) with a panel of probes specific for detection of recurring chromosome abnormalities in plasma dysplasia was performed on uncultured bone marrow cells.

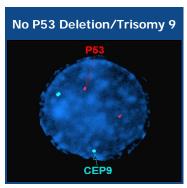
The regions/loci represented in these probe mixes were:

- 1. IGH/CCND1 dual color, dual translocation probes to 11q13 & 14q32 regions respectively, reveal a hybridization pattern within normal limits in 200 analyzed nuclei.
- 2. P53 (17p13.1), used to detect deletion/copy number abnormalities of chromosome 17p, reveals a hybridization pattern within normal limits in 200 analyzed nuclei.
- 3. 1p36 Microdeletion Region Probe LSI p58 (1p36) /TelVysion 1p/LSI 1q25, used to detect copy number abnormalities of chromosome 1, reveal a hybridization pattern within normal limits in 200 analyzed nuclei.
- 4. CEP9 (centromere probe to chromosome 9), used to detect copy number abnormalities of chromosome 9, reveals a hybridization pattern within normal limits in 200 analyzed nuclei.
- 5. D13S319 (13q14.3) and 13q34, used to detect copy number abnormalities/deletion of chromosome 13, reveal a hybridization pattern within normal limits in 200 analyzed nuclei.









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Bone marrow, aspirate: MYD88 mutation(S) detected: P.L265P (see comment)



MYD88 mutation is the most frequent genomic abnormality in diffuse large cell lymphoma (DLBCL) activated B-cell-like (ABC) subtype, detected in 40% of cases. MYD88 is rarely mutated in the germinal center B-cell-like (GCB) DLBL, therefore, it can be used to differentiate between the two subtypes. MYD88 mutation is detected in approximately 90% of cases of Waldenstrom macroglobulinemia/lymphoplasmacytic lymphoma. MYD88 mutation analysis can be a useful prognostic tool for patients with IgM-MGUS since the L265P mutation is associated with a higher risk of disease progression and a greater disease burden. MYD88 mutation has also been reported to be common (40%) in central nervous system lymphoma.

#### **METHODOLOGY:**

Total nucleic acid was extracted from patient's plasma, PB/BM cells or paraffin-embedded tissues (FFPE). Bi-directional Sanger sequencing of exon 5 of MYD88 was performed, including the L265P mutation hot spot. This is a sequencing-based assay which has a typical sensitivity of 10-15% for detecting MYD88 mutations in a wild-type background. Various factors including quantity and quality of nucleic acid, sample preparation, and sample age can affect assay performance.

#### **REFERENCES:**

1. Trøen G, Warsame A, Delabie J. CD79B and MYD88 Mutations in Splenic Marginal Zone Lymphoma. ISRN Oncol. 2013;2013:252318.

2. Xu L, et al. MYD88 L265P in Waldenstrom's macroblobulinemia, IgM monoclonal gammopathy, and other B-cell lymphoproliferative disorders using conventional and quantitative allele-specific PCR. Blood. 2013;121(11):2051-2058.

The performance characteristics of this test have been determined by the laboratory. This test has not been approved by the FDA. The FDA has determined such clearance or approval is not necessary. The laboratory is CLIA certified to perform high complexity clinical testing.

**Electronically Signed By:** Frank Bauer, MD (04/18/19 16:36)

Patient: Jane A. Doe

**Received:** 04/09/19 11:04

**Case #**: X19-00352

**Reported:** 04/18/19 17:43



**Received Information:** 2 Formalin containers, 10 smears, 2 green-top tubes, 1 lavender-top tube