



## Chronic Lymphocytic Leukemia (CLL) Pre-Infusion Data

**Registry Use Only**

Sequence Number:

Date Received:

CIBMTR Center Number: \_\_\_\_\_

CIBMTR Research ID: \_\_\_\_\_

Event date: \_\_ \_\_ \_\_ \_\_ / \_\_ \_\_ / \_\_ \_\_  
                  YYYY    MM    DD

HCT type (check all that apply):  Autologous     Allogeneic, unrelated     Allogeneic, related

Product type (check all that apply):

Bone marrow     PBSC     Single cord blood unit     Multiple cord blood units     Other product. Specify: \_\_\_\_\_

**Subsequent Transplant or Cellular Therapy**

**If this is a report of a second or subsequent transplant or cellular therapy for the same disease subtype and this baseline disease insert has not been completed for the previous transplant or cellular therapy (e.g. patient was on TED track for the prior HCT, prior HCT was autologous with no consent, prior cellular therapy was not reported to the CIBMTR), begin the form at question one.**

**If this is a report of a second or subsequent transplant or cellular therapy for a different disease, begin the form at question one.**

Is this the report of a second or subsequent transplant or cellular therapy for the same disease?

- Yes - **Go to questions 149**
- No - **Go to question 1**

**Disease Assessment at Diagnosis**

1. What was the date of diagnosis: \_\_ \_\_ \_\_ \_\_ / \_\_ \_\_ / \_\_ \_\_  
YYYY MM DD

2. Was documentation submitted to the CIBMTR (e.g. pathology report used for diagnosis)?  Yes  No

3. Did a histologic transformation occur at any time after CLL diagnosis?

- Yes →
- No

4. Date of transformation: \_\_ \_\_ \_\_ \_\_ / \_\_ \_\_ / \_\_ \_\_  
YYYY MM DD

5. Specify the disease classification after transformation

Diffuse large B-cell lymphoma (Richter syndrome) – **Also complete CIBMTR form 2018 - LYM**

Other disease classification →

6. Specify other disease classification: \_\_\_\_\_

7. Was documentation submitted to the CIBMTR (e.g. pathology report at transformation)?  Yes  No

**Autoimmune disorder(s) at diagnosis:**

8. Immune hemolytic anemia  Yes  No  Unknown
9. Immune thrombocytopenia  Yes  No  Unknown

10. Other

- Yes →
- No
- Unknown

11. Specify other autoimmune disorder: \_\_\_\_\_

12. Rai stage (at diagnosis)

- Known →
- Unknown

13. What was the Rai stage? (at diagnosis)

Stage 0 —Low risk — lymphocytosis (> 15,000 x 10<sup>9</sup>/L) in blood or bone marrow only without lymphadenopathy, hepatosplenomegaly, anemia or thrombocytopenia

Stage 1 - Intermediate risk — lymphocytosis plus enlarged lymph nodes (lymphadenopathy) without hepatosplenomegaly, anemia or thrombocytopenia

Stage II - Intermediate risk — lymphocytosis plus enlarged liver or spleen with or without lymphadenopathy

Stage III - High risk — lymphocytosis plus anemia (Hgb < 11.0 g/dL) with or without enlarged liver, spleen, or lymph nodes

Stage IV - High risk — lymphocytosis plus thrombocytopenia (platelet count < 100 x 10<sup>9</sup>/L) with or without anemia or enlarged liver, spleen, or lymph nodes

14. Binet stage (at diagnosis)

- Known →  
 Unknown

15. What was the Binet stage? (at diagnosis) (Five lymphoid bearing areas are possible: axillary, cervical, inguino-femoral, liver, and spleen.)

- Stage A — two or fewer lymphoid bearing areas enlarged, without anemia or thrombocytopenia  
 Stage B — three or more lymphoid bearing areas enlarged, without anemia or thrombocytopenia  
 Stage C — presence of anemia (Hgb < 10.0 g/dL) or thrombocytopenia (platelet count < 100 x 10<sup>9</sup>/L)

16. Were systemic symptoms (B symptoms) present (unexplained fever > 38° C ; or night sweats; unexplained weight loss of > 10% of body weight in six months before diagnosis)?

- Yes  No  Unknown

17. Was extranodal disease present?

- Yes →  
 No

**Specify site(s) of disease:**

18. Central nervous system (CNS)  Yes  No  
 19. Lung  Yes  No  
 20. Other site  
 Yes → 21. Specify other site: \_\_\_\_\_  
 No

**Laboratory Studies at Diagnosis**

22. WBC:

- Known →  
 Unknown

23. \_\_\_\_\_ • \_\_\_\_\_  x 10<sup>9</sup>/L (x 10<sup>3</sup>/mm<sup>3</sup>)  x 10<sup>6</sup>/L

24. Hemoglobin: (untransfused)

- Known →  
 Unknown

25. \_\_\_\_\_ • \_\_\_\_\_  g/dL  g/L  mmol/L

26. Platelets: (untransfused)

- Known →  
 Unknown

27. \_\_\_\_\_  x 10<sup>9</sup>/L (x 10<sup>3</sup>/mm<sup>3</sup>)  x 10<sup>6</sup>/L

28. Lymphocytes:

- Known →  
 Unknown

29. \_\_\_\_\_ %

30. Prolymphocytes:

- Known →  
 Unknown

31. \_\_\_\_\_ %

32. LDH:

Known →

Unknown

33. \_\_\_\_\_ • \_\_\_\_\_  U/L   $\mu$ kat/L

34. Upper limit of normal for LDH: \_\_\_\_\_ • \_\_\_\_\_  U/L   $\mu$ kat/L

35. Serum  $\beta$ 2 microglobulin:

Known →

Unknown

36. \_\_\_\_\_ • \_\_\_\_\_   $\mu$ g/dL  mg/L  nmol/L

37. Upper limit of normal for serum  $\beta$ 2 microglobulin: \_\_\_\_\_ • \_\_\_\_\_  
  $\mu$ g/dL  mg/L  nmol/L

38. Lymphocytes in bone marrow

Known →

Unknown

39. \_\_\_\_\_ %

40. Leukemia cell type: (may be determined at any time after diagnosis)  B-cell  T-cell  Unknown

41. Were tests for molecular markers performed (e.g. PCR)?

Yes →

No

Unknown

42. Date sample collected: \_\_\_\_ / \_\_\_\_ / \_\_\_\_  
 YYYYY MM DD

43. Immunoglobulin heavy chain variable (IGHV) mutation

Positive →

Negative →

Not done

44. Specify method used:

ASO IGHV RQ-PCR

Consensus IGHV PCR

Consensus IGHV PCR using HTS

Nested ASO IGHV PCR

Other method → 45. Specify other method: \_\_\_\_\_

46. NOTCH 1 mutation  Positive  Negative  Not done

47. P53 mutation  Positive  Negative  Not done

48. SF3B1 mutation  Positive  Negative  Not done

49. Other molecular marker

Positive →

Negative →

Not done

50. Specify other molecular marker: \_\_\_\_\_

51. Was documentation submitted to the CIBMTR?  Yes  No

**Immunophenotype: (may be determined at any time after diagnosis)**

52. Was flow cytometry (immunophenotyping) performed?

- Yes →
- No
- Unknown

53. CD5+	<input type="checkbox"/> Positive	<input type="checkbox"/> Negative	<input type="checkbox"/> Not done
54. CD19+	<input type="checkbox"/> Positive	<input type="checkbox"/> Negative	<input type="checkbox"/> Not done
55. CD20+	<input type="checkbox"/> Positive	<input type="checkbox"/> Negative	<input type="checkbox"/> Not done
56. CD23+	<input type="checkbox"/> Positive	<input type="checkbox"/> Negative	<input type="checkbox"/> Not done
57. CD38+	<input type="checkbox"/> Positive → <input type="checkbox"/> Negative <input type="checkbox"/> Not done		
58. Specify percent positivity: <input type="checkbox"/> ≥30% positivity <input type="checkbox"/> <30% positivity			
59. Slg	<input type="checkbox"/> Positive	<input type="checkbox"/> Negative	<input type="checkbox"/> Not done
60. ZAP-70 - mutated	<input type="checkbox"/> Positive	<input type="checkbox"/> Negative	<input type="checkbox"/> Not done

61. Were cytogenetics tested (karyotyping or FISH)?

- Yes →
- No
- Unknown

62. Results of tests:

- Abnormalities identified →
- No evaluable metaphases
- No abnormalities

**Specify cytogenetic abnormalities identified at diagnosis:**

**Trisomy**

63. +12  Yes     No

**Translocation**

64. t(11;14)  Yes     No

65. Any other translocation of 14  Yes     No

**Deletion**

66. del(11q) / 11q-  Yes     No

67. del(13q) / 13q-  Yes     No

68. del(17p) / 17p-  Yes     No

**Other**

69. Chromosome 6 abnormalities  Yes     No

70. Chromosome 8 abnormalities  Yes     No

71. Other abnormality

Yes → 72. Specify other abnormality: \_\_\_\_\_

No

73. Was documentation submitted to the CIBMTR (e.g. cytogenetic or FISH report)?

Yes     No

**Pre-HCT or Pre-Infusion Therapy**

74. Was therapy given?

- Yes →  
 No  
 Unknown

**Line of Therapy**

75. Systemic therapy:

- Yes →  
 No

76. Date therapy started

- Known →  
 Unknown

77. Date started: \_\_\_/\_\_\_/\_\_\_  
YYYY MM DD

78. Date therapy stopped

- Known →  
 Unknown

79. Date stopped: \_\_\_/\_\_\_/\_\_\_  
YYYY MM DD

80. Number of cycles

- Known →  
 Unknown

81. Number of cycles: \_\_\_\_\_

- |                                       |                              |                             |
|---------------------------------------|------------------------------|-----------------------------|
| 82. Alemtuzumab (Campath)             | <input type="checkbox"/> Yes | <input type="checkbox"/> No |
| 83. Bendamustine                      | <input type="checkbox"/> Yes | <input type="checkbox"/> No |
| 84. Chlorambucil (Leukeran)           | <input type="checkbox"/> Yes | <input type="checkbox"/> No |
| 85. Cladribine (2-CdA, Leustatin)     | <input type="checkbox"/> Yes | <input type="checkbox"/> No |
| 86. Corticosteroids                   | <input type="checkbox"/> Yes | <input type="checkbox"/> No |
| 87. Cyclophosphamide (Cytoxan)        | <input type="checkbox"/> Yes | <input type="checkbox"/> No |
| 88. Cytarabine (Ara-C)                | <input type="checkbox"/> Yes | <input type="checkbox"/> No |
| 89. Doxorubicin (Adriamycin)          | <input type="checkbox"/> Yes | <input type="checkbox"/> No |
| 90. Etoposide (VP-16, VePesid)        | <input type="checkbox"/> Yes | <input type="checkbox"/> No |
| 91. Fludarabine (Fludara)             | <input type="checkbox"/> Yes | <input type="checkbox"/> No |
| 92. Gemcitabine (Gemzar)              | <input type="checkbox"/> Yes | <input type="checkbox"/> No |
| 93. Ibrutinib (Imbruvica)             | <input type="checkbox"/> Yes | <input type="checkbox"/> No |
| 94. Idelalisib (Zydelig)              | <input type="checkbox"/> Yes | <input type="checkbox"/> No |
| 95. Ifosfamide (Ifex)                 | <input type="checkbox"/> Yes | <input type="checkbox"/> No |
| 96. Lenalidomide (Revlimid)           | <input type="checkbox"/> Yes | <input type="checkbox"/> No |
| 97. Nelarabine                        | <input type="checkbox"/> Yes | <input type="checkbox"/> No |
| 98. Nitrogen mustard (mustine)        | <input type="checkbox"/> Yes | <input type="checkbox"/> No |
| 99. Obinutuzumab                      | <input type="checkbox"/> Yes | <input type="checkbox"/> No |
| 100. Oblimersen                       | <input type="checkbox"/> Yes | <input type="checkbox"/> No |
| 101. Ofatumumab (Arzerra, HuMax-CD20) | <input type="checkbox"/> Yes | <input type="checkbox"/> No |
| 102. Pentostatin (Nipent)             | <input type="checkbox"/> Yes | <input type="checkbox"/> No |
| 103. Rituximab (anti-CD20, Rituxan)   | <input type="checkbox"/> Yes | <input type="checkbox"/> No |
| 104. Venetoclax                       | <input type="checkbox"/> Yes | <input type="checkbox"/> No |
| 105. Vincristine (VCR, Oncovin)       | <input type="checkbox"/> Yes | <input type="checkbox"/> No |
| 106. Other systemic therapy           |                              |                             |

- Yes →  
 No

107. Specify other systemic therapy: \_\_\_\_\_



123. Date assessed: \_\_ \_\_ \_\_ \_\_ / \_\_ \_\_ / \_\_ \_\_  
YYYY MM DD

124. Were tests for molecular markers performed (e.g. PCR)?

- Yes →
- No
- Unknown

125. Date sample collected: \_\_ \_\_ \_\_ \_\_ / \_\_ \_\_ / \_\_ \_\_  
YYYY MM DD

126. Immunoglobulin heavy chain variable (IGHV) mutation

- Positive →
- Negative →
- Not done

127. Specify method used:

- ASO IGHV RQ-PCR
- Consensus IGHV PCR
- Consensus IGHV PCR using HTS
- Nested ASO IGHV PCR
- Other method →

128. Specify other method:

\_\_\_\_\_

129. NOTCH 1 mutation

- Positive
- Negative
- Not done

130. P53 mutation

- Positive
- Negative
- Not done

131. SF3B1 mutation

- Positive
- Negative
- Not done

132. Other molecular marker

- Positive →
- Negative →
- Not done

133. Specify other molecular marker:

\_\_\_\_\_

134. Was the disease status assessed via flow cytometry (minimum 4-color flow) (immunophenotyping)?

- Yes →
- No

135. Date sample collected: \_\_ \_\_ \_\_ \_\_ / \_\_ \_\_ / \_\_ \_\_  
YYYY MM DD

136. Was disease detected?  Yes  No



137. Was the disease status assessed via cytogenetic testing (karyotyping or FISH)?  
 Yes →  
 No

138. Was the disease status assessed via FISH?  
 Yes →  
 No

139. Date sample collected:  
 \_\_ \_\_ / \_\_ \_\_ / \_\_ \_\_  
 YYYY MM DD

140. Was disease detected?  
 Yes  No

141. Was the disease status assessed via karyotyping?  
 Yes →  
 No

142. Date sample collected:  
 \_\_ \_\_ / \_\_ \_\_ / \_\_ \_\_  
 YYYY MM DD

143. Was disease detected?  
 Yes  No

144. Was the disease status assessed by clinical / hematologic assessment?  
 Yes →  
 No

145. Date assessed: \_\_ \_\_ / \_\_ \_\_ / \_\_ \_\_  
 YYYY MM DD

146. Was disease detected?  Yes  No

147. Did disease relapse/progress following this line of therapy?  
 Yes →  
 No

148. Date of relapse/progression:  
 \_\_ \_\_ / \_\_ \_\_ / \_\_ \_\_  
 YYYY MM DD

Copy questions 75-148 if needed for multiple lines of therapy.

**Disease Assessment at Last Evaluation Prior to the Start of the Preparative Regimen / Infusion**

149. Did the recipient have known nodal involvement?  
 Yes →  
 No

150. Specify the size of the largest nodal mass: \_\_\_\_ cm x \_\_\_\_ cm

151. Was extranodal disease present?  
 Yes →  
 No

Specify site(s) of involvement:

152. Central nervous system (CNS)  Yes  No

153. Lung  Yes  No

<p>154. Other site  <input type="checkbox"/> Yes →  <input type="checkbox"/> No</p>	<p>155. Specify other site: _____</p>
<p>156. Polymphocytes  <input type="checkbox"/> Known →  <input type="checkbox"/> Unknown</p>	<p>157. _____ %</p>
<p>158. Serum β2 microglobulin:  <input type="checkbox"/> Known →  <input type="checkbox"/> Unknown</p>	<p>159. _____ • _____ <input type="checkbox"/> μg/dL <input type="checkbox"/> mg/L <input type="checkbox"/> nmol/L</p> <p>160. Upper limit of normal for serum β2 microglobulin: _____ • _____  <input type="checkbox"/> μg/dL <input type="checkbox"/> mg/L <input type="checkbox"/> nmol/L</p>
<p>161. Lymphocytes in bone marrow:  <input type="checkbox"/> Known →  <input type="checkbox"/> Unknown</p>	<p>162. _____ %</p>
<p>163. Were tests for molecular markers performed (e.g. PCR)?  <input type="checkbox"/> Yes →  <input type="checkbox"/> No  <input type="checkbox"/> Unknown</p>	<p>164. Date sample collected: __ __ / __ __ / __ __                  YYYY MM DD</p> <p>165. Immunoglobulin heavy chain variable (IGHV) mutation  <input type="checkbox"/> Positive →  <input type="checkbox"/> Negative  <input type="checkbox"/> Not done</p> <div style="border: 1px solid black; padding: 5px; margin: 5px 0;"> <p>166. Specify method used:  <input type="checkbox"/> ASO IGHV RQ-PCR  <input type="checkbox"/> Consensus IGHV PCR  <input type="checkbox"/> Consensus IGHV PCR using HTS  <input type="checkbox"/> Nested ASO IGHV PCR  <input type="checkbox"/> Other method →</p> </div> <div style="border: 1px solid black; padding: 5px; margin: 5px 0;"> <p>167. Specify other method:                  _____</p> </div> <p>168. NOTCH 1 mutation <input type="checkbox"/> Positive <input type="checkbox"/> Negative <input type="checkbox"/> Not done</p> <p>169. P53 mutation <input type="checkbox"/> Positive <input type="checkbox"/> Negative <input type="checkbox"/> Not done</p> <p>170. SF3B1 mutation <input type="checkbox"/> Positive <input type="checkbox"/> Negative <input type="checkbox"/> Not done</p> <p>171. Other molecular marker  <input type="checkbox"/> Positive →  <input type="checkbox"/> Negative →  <input type="checkbox"/> Not done</p> <div style="border: 1px solid black; padding: 5px; margin: 5px 0;"> <p>172. Specify other molecular marker: _____</p> </div> <p>173. Was documentation submitted to the CIBMTR? <input type="checkbox"/> Yes <input type="checkbox"/> No</p>

174. Was the disease status assessed via flow cytometry (minimum 4-color flow) (immunophenotyping)?

- Yes  
 No

175. Date sample collected: \_\_\_ / \_\_\_ / \_\_\_  
 YYYY MM DD

176. Was disease detected?  Yes  No

177. Were cytogenetics tested (karyotyping or FISH)?

- Yes  
 No  
 Unknown

178. Results of tests:

- Abnormalities identified  
 No evaluable metaphases  
 No abnormalities

**Specify cytogenetic abnormalities detected at last evaluation prior to the start of the preparative regimen / infusion:**

**Trisomy**

179. +12  Yes  No

**Translocation**

180. t(11;14)  Yes  No

181. Any other translocation of 14  Yes  No

**Deletion**

182. del(11q) / 11q-  Yes  No

183. del(13q) / 13q-  Yes  No

184. del(17p) / 17p-  Yes  No

**Other**

185. Chromosome 6 abnormalities  Yes  No

186. Chromosome 8 abnormalities  Yes  No

187. Other abnormality

- Yes → 188. Specify other abnormality:  
 No \_\_\_\_\_

189. Was the disease status assessed by clinical / hematologic assessment?

- Yes  
 No

190. Date assessed: \_\_\_ / \_\_\_ / \_\_\_  
 YYYY MM DD

191. Was disease detected?  Yes  No

**Disease Status at the Last Evaluation Prior to the Start of the Preparative Regimen / Infusion**

192. What was the disease status?

- Complete remission (CR) — no lymphadenopathy; no organomegaly; neutrophils  $\geq 1.5 \times 10^9/L$ ; platelets  $> 100 \times 10^9/L$ ; hemoglobin  $> 11.0$  g/dL; lymphocytes  $< 4 \times 10^9/L$ ; bone marrow  $< 30\%$  lymphocytes; absence of constitutional symptoms - **Go to question 193**
- Partial remission (PR) —  $\geq 50\%$  decrease in peripheral blood lymphocyte count from pretreatment value;  $\geq 50\%$  reduction in lymphadenopathy if present pretreatment;  $\geq 50\%$  reduction in liver and spleen size if enlarged pretreatment; one or more of the following: neutrophils  $\geq 1.5 \times 10^9/L$  or 50% improvement over baseline, platelets  $> 100 \times 10^9/L$  or 50% improvement over baseline, hemoglobin  $> 11.0$  g/dL or 50% improvement over baseline - **Go to question 193**
- Stable disease (SD) — no change; not complete remission, partial remission, nor progressive disease - **Go to question 193**
- Progressive disease (Prog) — one or more of the following:  $\geq 50\%$  increase in the sum of the products of  $\geq 2$  lymph nodes ( $\geq 1$  node must be  $\geq 2$  cm) or new nodes;  $\geq 50\%$  increase in liver or spleen size, or new hepatomegaly or splenomegaly;  $\geq 50\%$  increase in absolute lymphocyte count to  $\geq 5 \times 10^9/L$ ; transformation to a more aggressive histology - **Go to question 193**
- Untreated — no chemotherapy given in the 6 months prior to HCT - **Go to question 193**
- Not assessed - **Go to First Name**

193. Date assessed: \_\_\_/\_\_\_/\_\_\_  
 YYY Y / MM / DD

First Name: \_\_\_\_\_

Last Name: \_\_\_\_\_

E-mail address: \_\_\_\_\_

Date: \_\_\_/\_\_\_/\_\_\_  
 YYY Y / MM / DD