



Disease Classification

CIBMTR Use Only
Sequence Number:

Date Received:

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CIBMTR Research ID: _____
Event date: ____ - ____ - ____

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- AML with t(8;21); (q22; q22.1); RUNX1-RUNX1T1 (281)
- AML with inv(16)(p13.1;1q22) or t(16;16)(p13.1; q22); CBFB-MYH11 (282)
- APL with PML-RARA (283)
- AML with BCR-ABL1 (provisional entity) (3)
- AML with mutated NPM1 (4)
- AML with biallelic mutations of CEBPA (297)
- AML with mutated RUNX1 (provisional entity) (298)
- AML with 11q23 (MLL) abnormalities (i.e., t(4;11), t(6;11), t(9;11), t(11;19)) (284)
- AML with myelodysplasia – related changes (285)
- Therapy related AML (t-AML) (9)

AML, not otherwise specified

- AML, not otherwise specified (280)
- AML, minimally differentiated (286)
- AML without maturation (287)
- AML with maturation (288)
- Acute myelomonocytic leukemia (289)
- Acute monoblastic / acute monocytic leukemia (290)
- Acute erythroid leukemia (erythroid / myeloid and pure erythroleukemia) (291)
- Acute megakaryoblastic leukemia (292)
- Acute basophilic leukemia (293)
- Acute panmyelosis with myelofibrosis (294)
- Myeloid sarcoma (295)
- Myeloid leukemia associated with Down syndrome (299)

4. Did AML transform from MDS or MPN?

- Yes – **Also complete MDS Disease Classification questions**
- No

5. Is the disease (AML) therapy related?

- Yes
- No
- Unknown

6. Did the recipient have a predisposing condition?

- Yes - **Go to question 7**
- No - **Go to question 9**
- Unknown - **Go to question 9**

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7. Specify condition:

- Bloom syndrome - **Go to question 9**
- Down syndrome - **Go to question 9**
- Fanconi anemia - **Also complete CIBMTR Form 2029 - Go to question 9**
- Dyskeratosis congenita - **Go to question 9**
- Other condition - **Go to question 8**

8. Specify other condition: _____

Labs at diagnosis

9. Were cytogenetics tested (karyotyping or FISH)? (at diagnosis)

- Yes - **Go to question 10**
 No - **Go to question 21**
 Unknown - **Go to question 21**

10. Were cytogenetics tested via FISH?

- Yes – **Go to question 11**
 No - **Go to question 15**

11. Results of tests:

- Abnormalities identified – **Go to question 12**
 No abnormalities - **Go to question 15**

Specify cytogenetic abnormalities identified at diagnosis:

12. Specify number of distinct cytogenetic abnormalities:

- One (1)
 Two (2)
 Three (3)
 Four or more (4 or more)

13. Specify abnormalities (check all that apply)

- 5
 -7
 -17
 -18
 -X
 -Y
 +4
 +8
 +11
 +13
 +14
 +21
 +22
 t(3;3)

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- t(6;9)
- t(8;21)
- t(9;11)
- t(9;22)
- t(15;17) and variants
- t(16;16)
- del(3q) / 3q-
- del(5q) / 5q-
- del(7q) / 7q-
- del(9q) / 9q-
- del(11q) / 11q-
- del(16q) / 16q-
- del(17q) / 17q-
- del(20q) / 20q-
- del(21q) / 21q-
- inv(3)
- inv(16)
- (11q23) any abnormality
- 12p any abnormality
- Other abnormality - **Go to question 14**

14. Specify other abnormality: _____

15. Were cytogenetics tested via karyotyping?

- Yes – **Go to question 16**
- No - **Go to question 20**

16. Results of tests:

- Abnormalities identified – **Go to question 17**
- No evaluable metaphases - **Go to question 21**
- No abnormalities - **Go to question 21**

Specify cytogenetic abnormalities identified at diagnosis:

17. Specify number of distinct cytogenetic abnormalities:

CIBMTR Center Number: _____

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- One (1)
- Two (2)
- Three (3)
- Four or more (4 or more)

18. Specify abnormalities: (check all that apply)

- 5
- 7
- 17
- 18
- X
- Y
- +4
- +8
- +11
- +13
- +14
- +21
- +22
- t(3;3)
- t(6;9)
- t(8;21)
- t(9;11)
- t(9;22)
- t(15;17) and variants
- t(16;16)
- del(3q) / 3q-
- del(5q) / 5q-
- del(7q) / 7q-
- del(9q) / 9q-
- del(11q) / 11q-
- del(16q) / 16q-
- del(17q) / 17q-
- del(20q) / 20q-
- del(21q) / 21q-
- inv(3)
- inv(16)

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- (11q23) any abnormality
- 12p any abnormality
- Other abnormality - **Go to question 19**

19. Specify other abnormality: _____

20. Was documentation submitted to the CIBMTR? (e.g. cytogenetic or FISH report)
- Yes
 - No

21. Were tests for molecular markers performed (e.g. PCR, NGS)? (at diagnosis)
- Yes – **Go to question 22**
 - No – **Go to question 34**
 - Unknown – **Go to question 34**

Specify molecular markers identified at diagnosis:

22. CEBPA
- Positive – **Go to question 23**
 - Negative - **Go to question 24**
 - Not done - **Go to question 24**

23. Specify CEBPA mutation
- Biallelic (homozygous)
 - Monoallelic (heterozygous)
 - Unknown

24. FLT3 – D835 point mutation
- Positive
 - Negative
 - Not done

25. FLT3 – ITD mutation
- Positive- **Go to question 26**
 - Negative- **Go to question 26**
 - Not done- **Go to question 27**

26. FLT3 – ITD allelic ratio

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Known - **Go to question 27**

Unknown - **Go to question 28**

27. Specify FLT3 - ITD allelic ratio: ____ ▪ ____

28. IDH1

Positive

Negative

Not done

29. IDH2

Positive

Negative

Not done

30. KIT

Positive

Negative

Not done

31. NPM1

Positive

Negative

Not done

32. Other molecular marker

Positive- **Go to question 33**

Negative- **Go to question 33**

Not done- **Go to question 34**

33. Specify other molecular marker: _____

Copy and complete questions 32-33 for multiple molecular markers

Labs between diagnosis and last evaluation:

34. Were cytogenetics tested (karyotyping or FISH)? (between diagnosis and last evaluation)

Yes - **Go to question 35**

No - **Go to question 46**

Unknown - **Go to question 46**

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CIBMTR Research ID: _____

35. Were cytogenetics tested via FISH?

Yes – **Go to question 36**

No - **Go to question 40**

36. Results of tests:

Abnormalities identified – **Go to question 37**

No abnormalities - **Go to question 40**

Specify cytogenetic abnormalities identified between diagnosis and last evaluation:

37. Specify number of distinct cytogenetic abnormalities:

One (1)

Two (2)

Three (3)

Four or more (4 or more)

38. Specify abnormalities (check all that apply)

-5

-7

-17

-18

-X

-Y

+4

+8

+11

+13

+14

+21

+22

t(3;3)

t(6;9)

t(8;21)

t(9;11)

t(9;22)

t(15;17) and variants

t(16;16)

del(3q) / 3q-

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- del(5q) / 5q-
- del(7q) / 7q-
- del(9q) / 9q-
- del(11q) / 11q-
- del(16q) / 16q-
- del(17q) / 17q-
- del(20q) / 20q-
- del(21q) / 21q-
- inv(3)
- inv(16)
- (11q23) any abnormality
- 12p any abnormality
- Other abnormality - **Go to question 39**

39. Specify other abnormality: _____

40. Were cytogenetics tested via karyotyping?

Yes – **Go to question 41**

No - **Go to question 45**

41. Results of tests:

Abnormalities identified – **Go to question 42**

No evaluable metaphases - **Go to question 46**

No abnormalities - **Go to question 46**

Specify cytogenetic abnormalities identified between diagnosis and last evaluation:

42. Specify number of distinct cytogenetic abnormalities:

One (1)

Two (2)

Three (3)

Four or more (4 or more)

43. Specify abnormalities (check all that apply)

-5

-7

-17

-18

-X

CIBMTR Center Number: _____

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- Y
- +4
- +8
- +11
- +13
- +14
- +21
- +22
- t(3;3)
- t(6;9)
- t(8;21)
- t(9;11)
- t(9;22)
- t(15;17) and variants
- t(16;16)
- del(3q) / 3q-
- del(5q) / 5q-
- del(7q) / 7q-
- del(9q) / 9q-
- del(11q) / 11q-
- del(16q) / 16q-
- del(17q) / 17q-
- del(20q) / 20q-
- del(21q) / 21q-
- inv(3)
- inv(16)
- (11q23) any abnormality
- 12p any abnormality
- Other abnormality - **Go to question 44**

44. Specify other abnormality: _____

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

- Yes
- No

46. Were tests for molecular markers performed (e.g. PCR, NGS)? (between diagnosis and last evaluation)

Yes – **Go to question 47**

No – **Go to question 59**

Unknown – **Go to question 59**

Specify molecular markers identified between diagnosis and last evaluation:

47. CEBPA

Positive – **Go to question 48**

Negative - **Go to question 49**

Not done - **Go to question 49**

48. Specify CEBPA mutation

Biallelic (homozygous)

Monoallelic (heterozygous)

Unknown

49. FLT3 – D835 point mutation

Positive

Negative

Not done

50. FLT3 – ITD mutation

Positive - **Go to question 51**

Negative - **Go to question 53**

Not done - **Go to question 53**

51. FLT3 – ITD allelic ratio

Known - **Go to question 52**

Unknown - **Go to question 53**

52. Specify FLT3 - ITD allelic ratio: ____ ▪ ____

53. IDH1

Positive

Negative

Not done

54. IDH2

CIBMTR Center Number: _____

CIBMTR Research ID: _____

- Positive
- Negative
- Not done

55. KIT

- Positive
- Negative
- Not done

56. NPM1

- Positive
- Negative
- Not done

57. Other molecular marker

- Positive- **Go to question 58**
- Negative- **Go to question 58**
- Not done- **Go to question 59**

58. Specify other molecular marker: _____

Copy and complete questions 57-58 to report multiple other molecular markers

Labs at last evaluation:

59. Were cytogenetics tested (karyotyping or FISH)? (at last evaluation)

- Yes - **Go to question 60**
- No - **Go to question 71**
- Unknown - **Go to question 71**

60. Were cytogenetics tested via FISH?

- Yes – **Go to question 61**
- No - **Go to question 65**

61. Results of tests:

- Abnormalities identified – **Go to question 62**
- No abnormalities - **Go to question 65**

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Specify cytogenetic abnormalities identified at last evaluation:

62. Specify number of distinct cytogenetic abnormalities:

- One (1)
- Two (2)
- Three (3)
- Four or more (4 or more)

63. Specify abnormalities (check all that apply)

- 5
- 7
- 17
- 18
- X
- Y
- +4
- +8
- +11
- +13
- +14
- +21
- +22
- t(3;3)
- t(6;9)
- t(8;21)
- t(9;11)
- t(9;22)
- t(15;17) and variants
- t(16;16)
- del(3q) / 3q-
- del(5q) / 5q-
- del(7q) / 7q-
- del(9q) / 9q-
- del(11q) / 11q-
- del(16q) / 16q-
- del(17q) / 17q-
- del(20q) / 20q-

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- del(21q) / 21q-
- inv(3)
- inv(16)
- (11q23) any abnormality
- 12p any abnormality
- Other abnormality - **Go to question 64**

64. Specify other abnormality: _____

65. Were cytogenetics tested via karyotyping?

- Yes – **Go to question 66**
- No - **Go to question 71**

66. Results of tests:

- Abnormalities identified – **Go to question 67**
- No evaluable metaphases - **Go to question 71**
- No abnormalities - **Go to question 71**

Specify cytogenetic abnormalities identified at last evaluation:

67. Specify number of distinct cytogenetic abnormalities:

- One (1)
- Two (2)
- Three (3)
- Four or more (4 or more)

68. Specify abnormalities (check all that apply)

- 5
- 7
- 17
- 18
- X
- Y
- +4
- +8
- +11
- +13

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- +14
- +21
- +22
- t(3;3)
- t(6;9)
- t(8;21)
- t(9;11)
- t(9;22)
- t(15;17) and variants
- t(16;16)
- del(3q) / 3q-
- del(5q) / 5q-
- del(7q) / 7q-
- del(9q) / 9q-
- del(11q) / 11q-
- del(16q) / 16q-
- del(17q) / 17q-
- del(20q) / 20q-
- del(21q) / 21q-
- inv(3)
- inv(16)
- (11q23) any abnormality
- 12p any abnormality
- Other abnormality - **Go to question 69**

69. Specify other abnormality: _____

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

- Yes
- No

71. Were tests for molecular markers performed (e.g. PCR, NGS)? (at last evaluation)

- Yes - **Go to question 72**
- No - **Go to question 84**
- Unknown - **Go to question 84**

Specify molecular markers identified at last evaluation:

72. CEBPA
- Positive – **Go to question 73**
 - Negative - **Go to question 74**
 - Not done - **Go to question 74**
73. Specify CEBPA mutation
- Biallelic (homozygous)
 - Monoallelic (heterozygous)
 - Unknown
74. FLT3 – D835 point mutation
- Positive
 - Negative
 - Not done
75. FLT3 – ITD mutation
- Positive - **Go to question 76**
 - Negative - **Go to question 78**
 - Not done - **Go to question 78**
76. FLT3 – ITD allelic ratio
- Known - **Go to question 77**
 - Unknown - **Go to question 78**
77. Specify FLT3 - ITD allelic ratio: ____ ▪ ____
78. IDH1
- Positive
 - Negative
 - Not done
79. IDH2
- Positive
 - Negative
 - Not done
80. KIT
- Positive

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Negative

Not done

81. NPM1

Positive

Negative

Not done

82. Other molecular marker

Positive- **Go to question 83**

Negative- **Go to question 83**

Not done- **Go to question 84**

83. Specify other molecular marker: _____

Copy and complete questions 82-83 to report multiple other molecular markers

CNS Leukemia

84. Did the recipient have central nervous system leukemia at any time prior to the start of the preparative regimen / infusion?

Yes

No

Unknown

Status at transplantation:

85. What was the disease status (based on hematological test results)?

Primary induction failure – **Go to question 89**

1st complete remission (no previous bone marrow or extramedullary relapse) (include CRi)– **Go to question 86**

2nd complete remission – **Go to question 86**

≥ 3rd complete remission – **Go to question 86**

1st relapse – **Go to question 88**

2nd relapse – **Go to question 88**

≥ 3rd relapse – **Go to question 88**

No treatment – **Go to question 89**

86. How many cycles of induction therapy were required to achieve 1st complete remission? (includes CRi)

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1

2

≥ 3

87. Was the recipient in remission by flow cytometry?

Yes – **Go to question 89**

No – **Go to question 89**

Unknown – **Go to question 89**

Not applicable – **Go to question 89**

88. Date of most recent relapse: _____

YYYY

MM

DD

89. Date assessed: _____ - **Go to signature line**

YYYY

MM

DD

Acute Lymphoblastic Leukemia (ALL)

90. Specify ALL classification:

B-lymphoblastic leukemia / lymphoma

- B-lymphoblastic leukemia / lymphoma, NOS (B-cell ALL, NOS) (191)
- B-lymphoblastic leukemia / lymphoma with t(9;22)(q34.1;q11.2); BCR-ABL1 (192)
- B-lymphoblastic leukemia / lymphoma with t(v;11q23.3); KMT2A rearranged (193)
- B-lymphoblastic leukemia / lymphoma with t(1;19)(q23;p13.3); TCF3-PBX1 (194)
- B-lymphoblastic leukemia / lymphoma with t(12;21)(p13.2;q22.1); ETV6-RUNX1 (195)
- B-lymphoblastic leukemia / lymphoma with t(5;14)(q31.1;q32.3); IL3-IGH (81)
- B-lymphoblastic leukemia / lymphoma with Hyperdiploidy (51-65 chromosomes) (82)
- B-lymphoblastic leukemia / lymphoma with Hypodiploidy (<45 chromosomes) (83)
- B-lymphoblastic leukemia / lymphoma, BCR-ABL1-like (provisional entity) (94)
- B-lymphoblastic leukemia / lymphoma, with iAMP21 (provisional entity) (95)

T-cell lymphoblastic leukemia / lymphoma

- Early T-cell precursor lymphoblastic leukemia (provisional entity) (96)
- Natural killer (NK)- cell lymphoblastic leukemia / lymphoma (provisional entity) (97)

91. Did the recipient have a predisposing condition?

Yes - **Go to question 92**

No - **Go to question 94**

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Unknown - **Go to question 94**

92. Specify condition:

Aplastic anemia - **Go to question 94** Also complete CIBMTR Form 2028 — APL

Bloom syndrome - **Go to question 94**

Down syndrome - **Go to question 94**

Fanconi anemia - **Go to question 924** Also complete CIBMTR Form 2029 — FAN

Other condition - **Go to question 93**

93. Specify other condition:

94. Were tyrosine kinase inhibitors given for therapy at any time prior to start of the preparative regimen / infusion?
(e.g. imatinib mesylate, dasatinib, etc.)

Yes

No

Laboratory studies at diagnosis:

95. Were cytogenetics tested (karyotyping or FISH)? (at diagnosis)

Yes - **Go to question 96**

No - **Go to question 106**

Unknown - **Go to question 106**

96. Were cytogenetics tested via FISH? (at diagnosis)

Yes - **Go to question 97**

No - **Go to question 101**

97. Results of tests: (at diagnosis)

Abnormalities identified - **Go to question 98**

No abnormalities - **Go to question 101**

Specify cytogenetic abnormalities identified:

98. Specify number of distinct cytogenetic abnormalities:

One (1)

Two (2)

Three (3)

Four or more (4 or more)

99. Specify abnormalities: (check all that apply)

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- 7
- +4
- +8
- +17
- +21
- t(1;19)
- t(2;8)
- t(4;11)
- t(5;14)
- t(8;14)
- t(8;22)
- t(9;22)
- t(10;14)
- t(11;14)
- t(12;21)
- del(6q) / 6q-
- del(9p) / 9p-
- del(12p) / 12p-
- add(14q)
- (11q23) any abnormality
- 9p any abnormality
- 12p any abnormality
- Hyperdiploid (> 50)
- Hypodiploid (< 45)
- iAMP21
- Other abnormality - **Go to question 100**

100. Specify other abnormality:

101. Were cytogenetics tested via karyotyping? (at diagnosis)

Yes - **Go to question 102**

No - **Go to question 107**

102. Results of tests: (at diagnosis)

Abnormalities identified - **Go to question 103**

No evaluable metaphases - **Go to question 107**

No abnormalities - **Go to question 107**

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Specify cytogenetic abnormalities identified:

103. Specify number of distinct cytogenetic abnormalities:

- One (1)
- Two (2)
- Three (3)
- Four or more (4 or more)

104. Specify abnormalities: (check all that apply)

- 7
- +4
- +8
- +17
- +21
- t(1;19)
- t(2;8)
- t(4;11)
- t(5;14)
- t(8;14)
- t(8;22)
- t(9;22)
- t(10;14)
- t(11;14)
- t(12;21)
- del(6q) / 6q-
- del(9p) / 9p-
- del(12p) / 12p-
- add(14q)
- (11q23) any abnormality
- 9p any abnormality
- 12p any abnormality
- Hyperdiploid (> 50)
- Hypodiploid (< 45)
- iAMP21
- Other abnormality – **Go to question 105**

105. Specify other abnormality: _____

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

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Yes

No

107. Were tests for molecular markers performed (e.g. PCR, NGS)? (at diagnosis)

Yes – **Go to question 108**

No – **Go to question 113**

Unknown – **Go to question 113**

Specify molecular markers identified at diagnosis:

108. BCR / ABL

Positive

Negative

Not done

109. TEL-AML / AML1

Positive

Negative

Not done

110. Other molecular marker

Positive – **Go to question 111**

Negative – **Go to question 111**

Not done – **Go to question 112**

111. Specify other molecular marker:

Copy and complete questions 110-111 for additional molecular markers

Laboratory studies between diagnosis and last evaluation:

112. Were cytogenetics tested (karyotyping or FISH)? (between diagnosis and last evaluation)

Yes - **Go to question 113**

No - **Go to question 124**

Unknown - **Go to question 124**

113. Were cytogenetics tested via FISH? (between diagnosis and the last evaluation)

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Yes - **Go to question 114**

No - **Go to question 118**

114. Results of tests: (between diagnosis and the last evaluation)

Abnormalities identified - **Go to question 115**

No abnormalities - **Go to question 118**

Specify cytogenetic abnormalities identified:

115. Specify number of distinct cytogenetic abnormalities:

One (1)

Two (2)

Three (3)

Four or more (4 or more)

116. Specify abnormalities: (check all that apply)

-7

+4

+8

+17

+21

t(1;19)

t(2;8)

t(4;11)

t(5;14)

t(8;14)

t(8;22)

t(9;22)

t(10;14)

t(11;14)

t(12;21)

del(6q) / 6q-

del(9p) / 9p-

del(12p) / 12p-

add(14q)

(11q23) any abnormality

9p any abnormality

12p any abnormality

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- Hyperdiploid (> 50)
- Hypodiploid (< 45)
- iAMP21
- Other abnormality – **Go to question 117**

117. Specify other abnormality:

118. Were cytogenetics tested via karyotyping? (between diagnosis and the last evaluation)

Yes - **Go to question 119**

No - **Go to question 124**

119. Results of tests: (between diagnosis and the last evaluation)

- Abnormalities identified - **Go to question 120**
- No evaluable metaphases - **Go to question 124**
- No abnormalities - **Go to question 124**

Specify cytogenetic abnormalities identified:

120. Specify number of distinct cytogenetic abnormalities:

- One (1)
- Two (2)
- Three (3)
- Four or more (4 or more)

121. Specify abnormalities: (check all that apply)

- 7
- +4
- +8
- +17
- +21
- t(1;19)
- t(2;8)
- t(4;11)
- t(5;14)
- t(8;14)
- t(8;22)
- t(9;22)
- t(10;14)
- t(11;14)

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CIBMTR Research ID: _____

- t(12;21)
- del(6q) / 6q-
- del(9p) / 9p-
- del(12p) / 12p-
- add(14q)
- (11q23) any abnormality
- 9p any abnormality
- 12p any abnormality
- Hyperdiploid (> 50)
- Hypodiploid (< 45)
- iAMP21
- Other abnormality – **Go to question 122**

122. Specify other abnormality:

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

- Yes
- No

124. Were tests for molecular markers performed (e.g. PCR, NGS)? (between diagnosis and last evaluation)

- Yes – **Go to question 125**
- No – **Go to question 129**
- Unknown – **Go to question 129**

Specify molecular markers identified between diagnosis and last evaluation:

125. BCR / ABL

- Positive
- Negative
- Not done

126. TEL-AML / AML1

- Positive
- Negative
- Not done

127. Other molecular marker

- Positive – **Go to question 128**
- Negative – **Go to question 128**

Not done – **Go to question 129**

128. Specify other molecular marker:

Copy and complete questions 127-128 for additional molecular markers

Laboratory studies at last evaluation:

129. Were cytogenetics tested (karyotyping or FISH)? (at last evaluation)

Yes - **Go to question 130**

No - **Go to question 141**

Unknown - **Go to question 141**

130. Were cytogenetics tested via FISH?

Yes - **Go to question 131**

No - **Go to question 135**

131. Results of tests:

Abnormalities identified - **Go to question 132**

No abnormalities - **Go to question 135**

Specify cytogenetic abnormalities identified at last evaluation:

132. Specify number of distinct cytogenetic abnormalities:

One (1)

Two (2)

Three (3)

Four or more (4 or more)

133. Specify abnormalities: (check all that apply)

-7

+4

+8

+17

+21

t(1;19)

t(2;8)

t(4;11)

t(5;14)

t(8;14)

CIBMTR Center Number: _____

CIBMTR Research ID: _____

- t(8;22)
- t(9;22)
- t(10;14)
- t(11;14)
- t(12;21)
- del(6q) / 6q-
- del(9p) / 9p-
- del(12p) / 12p-
- add(14q)
- (11q23) any abnormality
- 9p any abnormality
- 12p any abnormality
- Hyperdiploid (> 50)
- Hypodiploid (< 45)
- iAMP21
- Other abnormality – **Go to question 134**

134. Specify other abnormality:

135. Were cytogenetics tested via karyotyping? (at last evaluation)

Yes - **Go to question 136**

No - **Go to question 141**

136. Results of tests:

Abnormalities identified - **Go to question 137**

No evaluable metaphases - **Go to question 141**

No abnormalities - **Go to question 141**

Specify cytogenetic abnormalities identified at last evaluation:

137. Specify number of distinct cytogenetic abnormalities:

- One (1)
- Two (2)
- Three (3)
- Four or more (4 or more)

138. Specify abnormalities: (check all that apply)

-7

+4

CIBMTR Center Number: _____

CIBMTR Research ID: _____

- +8
- +17
- +21
- t(1;19)
- t(2;8)
- t(4;11)
- t(5;14)
- t(8;14)
- t(8;22)
- t(9;22)
- t(10;14)
- t(11;14)
- t(12;21)
- del(6q) / 6q-
- del(9p) / 9p-
- del(12p) / 12p-
- add(14q)
- (11q23) any abnormality
- 9p any abnormality
- 12p any abnormality
- Hyperdiploid (> 50)
- Hypodiploid (< 45)
- iAMP21
- Other abnormality – **Go to question 139**

139. Specify other abnormality:

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

- Yes
- No

141. Were tests for molecular markers performed (e.g. PCR, NGS)? (at last evaluation)

- Yes – **Go to question 142**
- No – **Go to question 146**
- Unknown – **Go to question 146**

Specify molecular markers identified at last evaluation:

CIBMTR Center Number: _____

CIBMTR Research ID: _____

142. BCR / ABL

- Positive
- Negative
- Not done

143. TEL-AML / AML1

- Positive
- Negative
- Not done

144. Other molecular marker

- Positive – **Go to question 145**
- Negative – **Go to question 145**
- Not done – **Go to question 146**

145. Specify other molecular marker:

Copy and complete questions 144-145 for additional molecular markers

CNS Leukemia

146. Did the recipient have central nervous system leukemia at any time prior to the start of the preparative regimen / infusion?

- Yes
- No
- Unknown

Status at transplantation:

147. What was the disease status (based on hematological test results)?

- Primary induction failure – **Go to question 151**
- 1st complete remission (no previous marrow or extramedullary relapse)(include CRi) – **Go to question 148**
- 2nd complete remission – **Go to question 148**
- ≥ 3rd complete remission – **Go to question 148**
- 1st relapse – **Go to question 150**
- 2nd relapse – **Go to question 150**
- ≥ 3rd relapse – **Go to question 150**
- No treatment – **Go to question 151**

CIBMTR Center Number: _____

CIBMTR Research ID: _____

- Essential thrombocythemia (includes primary thrombocytosis, idiopathic thrombocytosis, hemorrhagic thrombocythemia) (58)
- Polycythemia vera (PCV) (57)
- Primary myelofibrosis (includes chronic idiopathic myelofibrosis (CIMF), angiogenic myeloid metaplasia (AMM), myelofibrosis/sclerosis with myeloid metaplasia (MMM), idiopathic myelofibrosis) (167)
- Mastocytosis**
- Myeloproliferative neoplasm (MPN), unclassifiable (60)
- Myeloid/lymphoid neoplasms with PDGFRA rearrangement**
- Myeloid/lymphoid neoplasms with PDGFRB rearrangement**
- Myeloid/lymphoid neoplasms with FGFR1 rearrangement**
- Myeloid/lymphoid neoplasms with PCM1-JAK2-**
- Chronic myelomonocytic leukemia (CMML) (54)
- Juvenile myelomonocytic leukemia (JMML/JCML) (no evidence of Ph¹ or BCR/ABL) (36) – **Go to question 212**
- Atypical chronic myeloid leukemia (aCML), BCR-ABL1- – Go to question 265**
- MDS / MPN with ring sideroblasts and thrombocytosis (MDS / MPN-RS-T)**
- Myelodysplastic / myeloproliferative neoplasm, unclassifiable (69)

168. Was the disease (MDS/MPN) therapy related?

- Yes
- No
- Unknown

169. Did the recipient have a predisposing condition?

- Yes – **Go to question 168**
- No – **Go to question 170**
- Unknown – **Go to question 170**

170. Specify condition:

- Aplastic anemia – **Go to question 172**
- Bloom syndrome – **Go to question 172**
- Down syndrome – **Go to question 172**
- Fanconi anemia – **Go to question 172**
- Other condition – **Go to question 171**

171. Specify other condition:

Laboratory studies at diagnosis of MDS:

CIBMTR Center Number: _____

CIBMTR Research ID: _____

172. WBC

Known

Unknown

173. _____ • _____ x 10⁹/L (x 10³/mm³)
 x 10⁶/L

174. Hemoglobin

Known

Unknown

175. _____ • _____ g/dL
 g/L
 mmol/L

176. Was RBC transfused ≤ 30 days before date of test?

Yes

No

177. Platelets

Known

Unknown

178. _____ x 10⁹/L (x 10³/mm³)
 x 10⁶/L

179. Were platelets transfused ≤ 7 days before date of test?

Yes

No

180. Neutrophils

Known

Unknown

181. _____%

182. Blasts in bone marrow

Known

Unknown

CIBMTR Center Number: _____

CIBMTR Research ID: _____

183. _____ %

184. Were cytogenetics tested (karyotyping or FISH)?

Yes – **Go to question 185**

No – **Go to question 212**

Unknown – **Go to question 212**

185. Results of tests:

Abnormalities identified – **Go to question 186**

No evaluable metaphases – **Go to question 212**

No abnormalities – **Go to question 212**

Specify abnormalities identified at diagnosis:

186. Specify number of distinct cytogenetic abnormalities:

One (1)

Two (2)

Three (3)

Four or more (4 or more)

Monosomy

187. -5

Yes

No

188. -7

Yes

No

189. -13

Yes

No

190. -20

Yes

No

191. -Y

Yes

CIBMTR Center Number: _____

CIBMTR Research ID: _____

No

Trisomy

192. +8

Yes

No

193. +19

Yes

No

Translocation

194. t(1;3)

Yes

No

195. t(2;11)

Yes

No

196. t(3;3)

Yes

No

197. t(3;21)

Yes

No

198. t(6;9)

Yes

No

199. t(11;16)

Yes

No

Deletion

200. del(3q) / 3q-

CIBMTR Center Number: _____

CIBMTR Research ID: _____

Yes

No

201. del(5q) / 5q-

Yes –

No

202. del(7q) / 7q-

Yes

No

203. del(9q) / 9q-

Yes

No

204. del(11q) / 11q-

Yes

No

205. del(12p) / 12p-

Yes

No

206. del(13q) / 13q-

Yes

No

207. del(20q) / 20q-

Yes

No

Inversion

208. inv(3)

Yes

No

Other

209. i17q

CIBMTR Center Number: _____

CIBMTR Research ID: _____

Yes

No

210. Other abnormality

Yes – **Go to question 211**

No – **Go to question 212**

211. Specify other abnormality:

212. Did the recipient progress or transform to a different MDS / MPN subtype between diagnosis and the start of the preparative regimen?

Yes – **Go to question 213**

No – **Go to question 216**

213. Specify the MDS / MPN subtype after transformation:

Refractory cytopenia with unilineage dysplasia (RCUD) (includes refractory anemia (RA)) (51) – **Go to question 214**

Refractory anemia with ringed sideroblasts (RARS) (55) – **Go to question 214**

Refractory anemia with excess blasts-1 (RAEB-1) (61) – **Go to question 214**

Refractory anemia with excess blasts-2 (RAEB-2) (62) – **Go to question 214**

Refractory cytopenia with multilineage dysplasia (RCMD) (64) – **Go to question 214**

Childhood myelodysplastic syndrome (Refractory cytopenia of childhood (RCC)) (68) – **Go to question 214**

Myelodysplastic syndrome with isolated del(5q) (5q- syndrome) (66) – **Go to question 214**

Myelodysplastic syndrome (MDS), unclassifiable (50) – **Go to question 214**

Chronic neutrophilic leukemia (165) – **Go to question 214**

Chronic eosinophilic leukemia, NOS (166) – **Go to question 214**

Essential thrombocythemia (includes primary thrombocytosis, idiopathic thrombocytosis, hemorrhagic thrombocythemia) (58) – **Go to question 214**

Polycythemia vera (PCV) (57) – **Go to question 214**

Primary myelofibrosis (includes chronic idiopathic myelofibrosis (CIMF), angiogenic myeloid metaplasia (AMM), myelofibrosis/sclerosis with myeloid metaplasia (MMM), idiopathic myelofibrosis) (167) – **Go to question 214**

Mastocytosis

Myeloproliferative neoplasm (MPN), unclassifiable (60) – **Go to question 214**

Myeloid/lymphoid neoplasms with PDGFRA rearrangement

Myeloid/lymphoid neoplasms with PDGFRB rearrangement

Myeloid/lymphoid neoplasms with FGFR1 rearrangement

Myeloid/lymphoid neoplasms with PCM1-JAK2-

CIBMTR Center Number: _____ CIBMTR Research ID: _____

Chronic myelomonocytic leukemia (CMML) (54) – **Go to question 214**

Atypical chronic myeloid leukemia (aCML), BCR-ABL1- – **Go to question 265**

MDS / MPN with ring sideroblasts and thrombocytosis (MDS / MPN-RS-T)

Myelodysplastic / myeloproliferative neoplasm, unclassifiable (69) – **Go to question 214**

Transformed to AML (70) – **Go to question 215**

214. Specify the date of the most recent transformation: _____ - _____ - _____ - **Go to question 216**

215. Date of MDS diagnosis: _____ - _____ - _____ - **Go to signature line**

Laboratory studies at last evaluation prior to the start of the preparative regimen:

216. WBC

Known

Unknown

217. _____ • _____ x 10⁹/L (x 10³/mm³)
 x 10⁶/L

218. Hemoglobin

Known

Unknown

219. _____ • _____ g/dL
 g/L
 mmol/L

220. Was RBC transfused ≤ 30 days before date of test?

Yes

No

221. Platelets

Known

Unknown

222. _____ x 10⁹/L (x 10³/mm³)
 x 10⁶/L

CIBMTR Center Number: _____ CIBMTR Research ID: _____

223. Were platelets transfused \leq 7 days before date of test?

Yes

No

224. Neutrophils

Known

Unknown

225. _____ %

226. Blasts in bone marrow

Known

Unknown

227. _____ %

228. Were cytogenetics tested (karyotyping or FISH)?

Yes – **Go to question 229**

No – **Go to question 256**

Unknown – **Go to question 256**

229. Results of tests:

Abnormalities identified – **Go to question 230**

No evaluable metaphases – **Go to question 256**

No abnormalities – **Go to question 256**

Specify cytogenetic abnormalities identified at last evaluation prior to the start of the preparative regimen:

230. Specify number of distinct cytogenetic abnormalities:

One (1)

Two (2)

Three (3)

Four or more (4 or more)

Monosomy

CIBMTR Center Number: _____

CIBMTR Research ID: _____

Yes

No

232. -7

Yes

No

233. -13

Yes

No

234. -20

Yes

No

235. -Y

Yes

No

Trisomy

236. +8

Yes

No

237. +19

Yes

No

Translocation

238. t(1;3)

Yes

No

239. t(2;11)

Yes

No

240. t(3;3)

CIBMTR Center Number: _____

CIBMTR Research ID: _____

Yes

No

241. t(3;21)

Yes

No

242. t(6;9)

Yes

No

243. t(11;16)

Yes

No

Deletion

244. del(3q) / 3q-

Yes

No

245. del(5q) / 5q-

Yes

No

246. del(7q) / 7q-

Yes

No

247. del(9q) / 9q-

Yes

No

248. del(11q) / 11q-

Yes

No

249. del(12p) / 12p-

Yes

No

CIBMTR Center Number: _____

CIBMTR Research ID: _____

250. del(13q) / 13q-

Yes

No

251. del(20q) / 20q-

Yes

No

Inversion

252. inv(3)

Yes

No

Other

253. i17q

Yes

No

254. Other abnormality

Yes – **Go to question 255**

No – **Go to question 256**

255. Specify other abnormality:

Status at transplantation:

256. What was the disease status?

- Complete remission (CR) – requires all of the following, maintained for ≥ 4 weeks: * bone marrow evaluation: $< 5\%$ myeloblasts with normal maturation of all cell lines * peripheral blood evaluation: hemoglobin ≥ 11 g/dL untransfused and without erythropoietin support; ANC $\geq 1000/\text{mm}^3$ without myeloid growth factor support; platelets $\geq 100 \times 10^9/\text{L}$ without thrombopoietic support; 0% blasts - **Go to question 260**
- Hematologic improvement (HI) – requires one measurement of the following, maintained for ≥ 8 weeks without ongoing cytotoxic therapy; specify which cell line was measured to determine HI response: * HI-E – hemoglobin increase of ≥ 1.5 g/dL untransfused; for RBC transfusions performed for Hgb ≤ 9.0 , reduction in RBC units transfused in 8 weeks by ≥ 4 units compared to the pre-treatment transfusion number in 8 weeks * HI-P – for pre-treatment platelet count of $> 20 \times 10^9/\text{L}$, platelet absolute increase of $\geq 30 \times 10^9/\text{L}$; for pre-treatment platelet count of $< 20 \times 10^9/\text{L}$, platelet absolute increase of $\geq 20 \times 10^9/\text{L}$ and $\geq 100\%$ from pre-treatment level * HI-N – neutrophil count increase of $\geq 100\%$ from pre-treatment level and an absolute increase of $\geq 500/\text{mm}^3$ - **Go to question 257**
- No response (NR) / stable disease (SD) – does not meet the criteria for at least HI, but no evidence of disease progression - **Go to question 260**

Hodgkin and Non-Hodgkin Lymphoma

268. Specify the lymphoma histology: (at transplant)

Hodgkin Lymphoma Codes

- Hodgkin lymphoma, not otherwise specified (150)
- Lymphocyte depleted (154)
- Lymphocyte-rich (151)
- Mixed cellularity (153)
- Nodular lymphocyte predominant Hodgkin lymphoma (155)
- Nodular sclerosis (152)

Non-Hodgkin Lymphoma Codes

B-cell Neoplasms

- ALK+ large B-cell lymphoma (1833)
- B-cell lymphoma, unclassifiable, with features intermediate between DLBCL and classical Hodgkin lymphoma (149)
- Burkitt lymphoma (111)
- Burkitt-like lymphoma with 11q aberration (1834)
- Diffuse large B-cell Lymphoma (cell of origin unknown) (107)
- Diffuse, large B-cell lymphoma- Germinal center B-cell type (1820) – Go to question 270
- Diffuse, large B-cell lymphoma- Activated B-cell type (non-GCB) (1821) – Go to question 270
- DLBCL associated with chronic inflammation (1825)
- Duodenal-type follicular lymphoma (1815)
- EBV+ DLBCL, NOS (1823)
- EBV+ mucocutaneous ulcer (1824)
- Extranodal marginal zone B-cell lymphoma of mucosal associated lymphoid tissue type (MALT) (122)
- Follicular, predominantly small cleaved cell (Grade I follicle center lymphoma) (102)
- Follicular, mixed, small cleaved and large cell (Grade II follicle center lymphoma) (103)
- Follicular, predominantly large cell (Grade IIIA follicle center lymphoma) (162)
- Follicular, predominantly large cell (Grade IIIB follicle center lymphoma) (163)
- Follicular, predominantly large cell (Grade IIIA vs IIIB not specified) (1814)
- Follicular (grade unknown) (164)
- HHV8+ DLBCL, NOS (1826)
- High-grade B-cell lymphoma, with MYC and BCL2 and/or BCL6 rearrangements (1831)
- High-grade B-cell lymphoma, NOS (1830)
- Intravascular large B-cell lymphoma (136)
- Large B-cell lymphoma with IRF4 rearrangement (1832)
- Lymphomatoid granulomatosis (1835)
- Mantle cell lymphoma (115)
- Nodal marginal zone B-cell lymphoma (± monocytoid B-cells) (123)
- Pediatric nodal marginal zone lymphoma (1813)
- Pediatric-type follicular lymphoma (1816)
- Plasmablastic lymphoma (1836)
- Primary cutaneous follicle center lymphoma (1817)
- Primary cutaneous DLBCL, leg type (1822)
- Primary diffuse, large B-cell lymphoma of the CNS (118)
- Primary effusion lymphoma (138)

CIBMTR Center Number: _____

CIBMTR Research ID: _____

- Primary mediastinal (thymic) large B-cell lymphoma (125)
- Splenic marginal zone B-cell lymphoma (124)
- Splenic B-cell lymphoma/leukemia, unclassifiable (1811)
- Splenic diffuse red pulp small B-cell lymphoma (1812)
- T-cell / histiocytic rich large B-cell lymphoma (120)
- Waldenstrom macroglobulinemia / Lymphoplasmacytic lymphoma (173)
- Other B-cell lymphoma (129) – **Go to question 269**

T-cell and NK-cell Neoplasms

- Adult T-cell lymphoma / leukemia (HTLV1 associated) (134)
- Aggressive NK-cell leukemia (27)
- Angioimmunoblastic T-cell lymphoma (131)
- Anaplastic large-cell lymphoma (ALCL), ALK positive (143)
- Anaplastic large-cell lymphoma (ALCL), ALK negative (144)
- Breast implant-associated anaplastic large-cell lymphoma (1861)
- Chronic lymphoproliferative disorder of NK cells (1856)
- Extranodal NK / T-cell lymphoma, nasal type (137)
- Enteropathy-type T-cell lymphoma (133)
- Follicular T-cell lymphoma (1859)
- Hepatosplenic T-cell lymphoma (145)
- Indolent T-cell lymphoproliferative disorder of the GI tract (1858)
- Monomorphic epitheliotropic intestinal T-cell lymphoma (1857)
- Mucosis fungoides (141)
- Nodal peripheral T-cell lymphoma with TFH phenotype (1860)
- Peripheral T-cell lymphoma (PTCL), NOS (130)
- Primary cutaneous $\gamma\delta$ T-cell lymphoma (1851)
- Primary cutaneous CD8+ aggressive epidermotropic cytotoxic T-cell lymphoma (1852)
- Primary cutaneous acral CD8+ T-cell lymphoma (1853)
- Primary cutaneous CD4+ small/medium T-cell lymphoproliferative disorder (1854)
- Primary cutaneous CD30+ T-cell lymphoproliferative disorders [Primary cutaneous anaplastic large-cell lymphoma (C-ALCL), lymphoid papulosis] (147)
- Sezary syndrome (142)
- Subcutaneous panniculitis-like T-cell lymphoma (146)
- Systemic EBV+ T-cell lymphoma of childhood (1855)
- T-cell large granular lymphocytic leukemia (126)
- Other T-cell / NK-cell lymphoma (139) **Go to question 269**

Posttransplant lymphoproliferative disorders (PTLD)

- Classical Hodgkin lymphoma PTLD (1876)
- Florid follicular hyperplasia PTLD (1873)
- Infectious mononucleosis PTLD (1872)
- Monomorphic PTLD (B- and T-/NK-cell types) (1875)
- Plasmacytic hyperplasia PTLD (1871)
- Polymorphic PTLD (1874)

269. Specify other lymphoma histology: _____ – **Go to question 271**

270. Assignment of DLBCL (germinal center B-cell type vs. Activated B-cell type) subtype was based on:

- Immunohistochemistry (e.g. Han's algorithm)

CIBMTR Center Number: _____

CIBMTR Research ID: _____

Gene expression profile

Unknown method

271. Is the lymphoma histology reported at transplant a transformation from CLL?

Yes – **Go to question 272**

No - **Go to question 273**

272. Was any 17p abnormality detected?

Yes– **Go to question 277**

No– **Go to question 277**

273. Is the lymphoma histology reported at transplant a transformation from a different lymphoma histology? (Not CLL)

Yes – **Go to question 274**

No – **Go to question 277**

274. Specify the original lymphoma histology: (prior to transformation) same option list as Q268

275. Specify other lymphoma histology: _____

276. Date of original lymphoma diagnosis: _____ - _____ - _____ (report the date of diagnosis of original lymphoma subtype)

277. Was a PET (or PET/CT) scan performed? (at last evaluation prior to the start of the preparative regimen / infusion)

Yes – **Go to question 278**

No – **Go to question 283**

278. Was the PET (or PET/CT) scan positive for lymphoma involvement at any disease site?

Yes

No

279. Date of PET scan

Known– **Go to question 280**

Unknown – **Go to question 281**

280. Date of PET (or PET/CT) scan: _____

YYYY MM DD

281. Deauville (five-point) score of the PET (or PET/CT) scan

Known – **Go to question 282**

Unknown – **Go to question 283**

282. Scale

- 1- no uptake or no residual uptake
- 2- slight uptake, but below blood pool (mediastinum)
- 3- uptake above mediastinal, but below or equal to uptake in the liver
- 4- uptake slightly to moderately higher than liver
- 5- markedly increased uptake or any new lesion

Status at transplantation / infusion:

283. What was the disease status?

- Disease untreated– **Go to question 285**
- PIF res - Primary induction failure – resistant: NEVER in COMPLETE remission but with stable or progressive disease on treatment. – **Go to question 284**
- PIF sen / PR1 - Primary induction failure – sensitive: NEVER in COMPLETE remission but with partial remission on treatment. – **Go to question 284**
- PIF unk - Primary induction failure – sensitivity unknown– **Go to question 284**
- CR1 - 1st complete remission: no bone marrow or extramedullary relapse prior to transplant– **Go to question 284**
- CR2 - 2nd complete remission– **Go to question 284**
- CR3+ - 3rd or subsequent complete remission– **Go to question 284**
- REL1 unt - 1st relapse – untreated; includes either bone marrow or extramedullary relapse– **Go to question 284**
- REL1 res - 1st relapse – resistant: stable or progressive disease with treatment– **Go to question 284**
- REL1 sen - 1st relapse – sensitive: partial remission (if complete remission was achieved, classify as CR2) – **Go to question 284**
- REL1 unk - 1st relapse – sensitivity unknown– **Go to question 284**
- REL2 unt - 2nd relapse – untreated: includes either bone marrow or extramedullary relapse– **Go to question 284**
- REL2 res - 2nd relapse – resistant: stable or progressive disease with treatment– **Go to question 284**
- REL2 sen - 2nd relapse – sensitive: partial remission (if complete remission achieved, classify as CR3+)– **Go to question 284**
- REL2 unk - 2nd relapse – sensitivity unknown– **Go to question 284**
- REL3+ unt - 3rd or subsequent relapse – untreated; includes either bone marrow or extramedullary relapse– **Go to question 284**
- REL3+ res - 3rd or subsequent relapse – resistant: stable or progressive disease with treatment– **Go to question 284**
- REL3+ sen - 3rd or subsequent relapse – sensitive: partial remission (if complete remission achieved, classify as CR3+)– **Go to question 284**
- REL3+ unk - 3rd relapse or greater – sensitivity unknown– **Go to question 284**

CIBMTR Center Number: _____ CIBMTR Research ID: _____

284. Total number of lines of therapy received: (at any time between diagnosis and HCT / infusion)

1 line

2 lines

3+ lines

285. Date assessed: _____ - **Go to signature line**

Multiple Myeloma / Plasma Cell Disorder (PCD)

287. Specify the multiple myeloma/plasma cell disorder (PCD) classification:

- Multiple myeloma-IgG (181) - **Go to questions 289**
- Multiple myeloma-IgA (182) - **Go to questions 289**
- Multiple myeloma-IgD (183) - **Go to questions 289**
- Multiple myeloma-IgE (184) - **Go to questions 289**
- Multiple myeloma-IgM (not Waldenstrom macroglobulinemia) (185) - **Go to questions 289**
- Multiple myeloma-light chain only (186) - **Go to questions 289**
- Multiple myeloma-non-secretory (187) - **Go to questions 290**
- Plasma cell leukemia (172) - **Go to question 295**
- Solitary plasmacytoma (no evidence of myeloma) (175) - **Go to question 295**
- Amyloidosis (174) - **Go to question 295**
- Osteosclerotic myeloma / POEMS syndrome (176) - **Go to questions 295**
- Light chain deposition disease (177) - **Go to questions 295**
- Other plasma cell disorder (179) - **Go to question 288**

288. Specify other plasma cell disorder: _____ - **Go to question 295**

289. Light chain

kappa

lambda

290. What was the Durie-Salmon staging (at diagnosis)?

- Stage I (All of the following: Hgb > 10g/dL; serum calcium normal or <10.5 mg/dL; bone x-ray normal bone structure (scale 0), or solitary bone plasmacytoma only; low M-component production rates IgG < 5g/dL, IgA < 3g/dL; urine light chain M-component on electrophoresis <4g/24h) – **Go to questions 291**
- Stage II (Fitting neither Stage I or Stage III) – **Go to questions 291**

CIBMTR Center Number: _____ CIBMTR Research ID: _____

Stage III (One of more of the following: Hgb < 8.5 g/dL; serum calcium > 12 mg/dL; advanced lytic bone lesions (scale 3); high M-component production rates IgG >7g/dL, IgA > 5g/dL; Bence Jones protein >12g/24h) – **Go to questions 291**

Unknown – **Go to questions 292**

291. What was the Durie-Salmon sub classification (at diagnosis)?

A - relatively normal renal function (serum creatinine < 2.0 mg/dL)

B - abnormal renal function (serum creatinine ≥ 2.0 mg/dL)

I.S.S.:

292. Serum β 2-microglobulin: _____ • _____ μ g/dL
 mg/L
 nmol/L

293. Serum albumin: _____ • _____ g/dL
 g/L

294. Stage

1 (β ₂-mic < 3.5, S. albumin ≥ 3.5)

2 (not fitting stage 1 or 3)

3 (β ₂-mic ≥ 5.5; S. albumin —)

295. Were cytogenetics tested (karyotyping or FISH)?

Yes – **Go to questions 296**

No – **Go to question 317**

Unknown – **Go to question 317**

296. Results of tests:

Abnormalities identified – **Go to question 297**

No evaluable metaphases – **Go to question 317**

No abnormalities – **Go to question 317**

Specify cytogenetic abnormalities identified at any time prior to the start of the preparative regimen:

Trisomy

297. +3

Yes

CIBMTR Center Number: _____

CIBMTR Research ID: _____

No

298. +5

Yes

No

299. +7

Yes

No

300. +9

Yes

No

301. +11

Yes

No

302. +15

Yes

No

303. +19

Yes

No

Translocation

304. t(4;14)

Yes

No

305. t(6;14)

Yes

No

306. t(11;14)

Yes

No

CIBMTR Center Number: _____

CIBMTR Research ID: _____

307. t(14;16)

Yes

No

308. t(14;20)

Yes

No

Deletion

309. del (13)/13q-

Yes

No

310. del (17)/17p-

Yes

No

Other

311. Hyperdiploid (>50)

Yes

No

312. Hypodiploid (<46)

Yes

No

313. Any abnormality at 1q

Yes

No

314. Any abnormality at 1p

Yes

No

315. Other abnormality

Yes

No

316. Specify other abnormality: _____

Status at transplantation:

317. What was the disease status?

Stringent complete remission (sCR). - CR as defined, plus: normal free light chain ratio, and absence of clonal cells in the bone marrow by immunohistochemistry or immunofluorescence (confirmation with repeat bone marrow biopsy not needed). (Presence and/or absence of clonal cells is based upon the κ/λ ratio. An abnormal κ/λ ratio by immunohistochemistry and/or immunofluorescence requires a minimum of 100 plasma cells for analysis. An abnormal ratio reflecting the presence of an abnormal clone is κ/λ of $> 4:1$ or $< 1:2$.) sCR requires two consecutive assessments made at any time before the institution of any new therapy, and no known evidence of progressive or new bone lesions if radiographic studies were performed; radiographic studies are not required to satisfy sCR requirements. - **Go to question 318**

Complete remission (CR) — negative immunofixation on serum and urine samples, and disappearance of any soft tissue plasmacytomas, and $< 5\%$ plasma cells in the bone marrow (confirmation with repeat bone marrow biopsy not needed). CR requires two consecutive assessments made at any time before the institution of any new therapy, and no known evidence of progressive or new bone lesions if radiographic studies were performed; radiographic studies are not required to satisfy CR requirements. - **Go to question 318**

Near complete remission (nCR) — serum and urine M-protein detectable by immunoelectrophoresis (IFE), but not on electrophoresis (negative SPEP & UPEP); $< 5\%$ plasma cells in bone marrow. nCR requires two consecutive assessments made at any time before the initiation of any new therapy, and no known evidence of progressive or new bone lesions if radiographic studies were performed; radiographic studies are not required to satisfy nCR requirements. - **Go to question 318**

Very good partial remission (VGPR) — serum and urine M-protein detectable by immunofixation but not on electrophoresis, or $\geq 90\%$ reduction in serum M-protein and urine M-protein level < 100 mg/24 hours. VGPR requires two consecutive assessments made at any time before the institution of any new therapy, and no known evidence of progressive or new bone lesions if radiographic studies were performed; radiographic studies are not required to satisfy VGPR requirements. - **Go to question 318**

Partial remission (PR) — $\geq 50\%$ reduction in serum M-protein, and reduction in 24-hour urinary M-protein by $\geq 90\%$ or to < 200 mg/24 hours. If the serum and urine M-protein are unmeasurable (i.e., do not meet any of the following criteria: • serum M-protein ≥ 1 g/dL. Urine M-protein ≥ 200 mg/24 hours • serum free light chain assay shows involved level ≥ 10 mg/dL, provided serum free light chain ratio is abnormal), a $\geq 50\%$ decrease in the difference between involved and uninvolved free light chain levels is required in place of the M-protein criteria. If serum and urine M-protein are unmeasurable, and serum free light assay is also unmeasurable, a $\geq 50\%$ reduction in plasma cells is required in place of M-protein, provided the baseline bone marrow plasma cell percentage was $\geq 30\%$. In addition to the above listed criteria, a $\geq 50\%$ reduction in the size of soft tissue plasmacytomas is also required, if present at baseline. PR requires two consecutive assessments made at any time before the institution of any new therapy, and no known evidence of progressive or new bone lesions if radiographic studies were performed; radiographic studies are not required to satisfy PR requirements. - **Go to question 318**

Stable disease (SD) — not meeting the criteria for CR, VGPR, PR or PD. SD requires two consecutive assessments made at any time before the institution of any new therapy, and no known evidence of progressive or new bone lesions if radiographic studies were performed; radiographic studies are not required to satisfy SD requirements. - **Go to question 318**

Progressive disease (PD) — requires any one or more of the following: Increase of $\geq 25\%$ from baseline in: serum M-component and/or (absolute increase ≥ 0.5 g/dL) (for progressive disease, serum M-component increases of ≥ 1 g/dL are sufficient to define relapse if the starting M-component is ≥ 5 g/dL). Urine M-component and/or (absolute increase ≥ 200 mg/24 hours) for recipients without measurable serum and urine M-protein levels: the difference between involved and uninvolved free light chain levels (absolute increase > 10

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mg/dL). Bone marrow plasma cell percentage (absolute percentage $\geq 10\%$) (relapse from CR has a 5% cutoff vs. 10% for other categories of relapse) definite development of new bone lesions or soft tissue plasmacytomas, or definite increase in the size of any existing bone lesions or soft tissue plasmacytomas. Development of hypercalcemia (corrected serum calcium > 11.5 mg/dL or 2.65 mmol) that can be attributed solely to the plasma cell proliferative disorder PD requires two consecutive assessments made at any time before classification as disease progression, and/or the institution of any new therapy - **Go to question 318**

Relapse from CR (Rel) (untreated) — requires one or more of the following: reappearance of serum or urine M-protein by immunofixation or electrophoresis development of $\geq 5\%$ plasma cells in the bone marrow (relapse from CR has a 5% cutoff vs. 10% for other categories of relapse) appearance of any other sign of progression (e.g., new plasmacytoma, lytic bone lesion, hypercalcemia) Rel requires two consecutive assessments made at any time before classification as relapse, and/or the institution of any new therapy. – **Go to question 318**

Unknown – **Go to signature line**

Not applicable (Amyloidosis with no evidence of myeloma) – **Go to signature line**

318. Date assessed: _____ - **Go to signature line**

YYYY

MM

DD

Solid Tumors

319. Specify the solid tumor classification:

- Breast cancer (250)
- Lung, small cell (202)
- Lung, non-small cell (203)
- Lung, not otherwise specified (230)
- Germ cell tumor, extragonadal (225)
- Testicular (210)
- Ovarian (epithelial) (214)
- Bone sarcoma (excluding Ewing family tumors) (273)
- Ewing family tumors of bone (including PNET) (275)
- Ewing family tumors, extraosseous (including PNET) (276)
- Fibrosarcoma (244)
- Hemangiosarcoma (246)
- Leiomyosarcoma (242)
- Liposarcoma (243)
- Lymphangio sarcoma (247)
- Neurogenic sarcoma (248)
- Rhabdomyosarcoma (232)
- Synovial sarcoma (245)
- Soft tissue sarcoma (excluding Ewing family tumors) (274)

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- Central nervous system tumor, including CNS PNET (220)
- Medulloblastoma (226)
- Neuroblastoma (222)
- Head / neck (201)
- Mediastinal neoplasm (204)
- Colorectal (228)
- Gastric (229)
- Pancreatic (206)
- Hepatobiliary (207)
- Prostate (209)
- External genitalia (211)
- Cervical (212)
- Uterine (213)
- Vaginal (215)
- Melanoma (219)
- Wilm tumor (221)
- Retinoblastoma (223)
- Thymoma (231)
- Renal cell (208)
- Other solid tumor (269) – **Go to question 320**
- Solid tumor, not otherwise specified (200)

320. Specify other solid tumor: _____ - **Go to signature line**

Severe Aplastic Anemia

321. Specify the severe aplastic anemia classification:

- Acquired severe aplastic anemia, not otherwise specified (301)
- Acquired SAA secondary to hepatitis (302)
- Acquired SAA secondary to toxin / other drug (303)
- Acquired amegakaryocytosis (not congenital) (304)
- Acquired pure red cell aplasia (not congenital) (306)
- Dyskeratosis congenita (307)
- Other acquired cytopenic syndrome (309) – **Go to question 322**

322. Specify other acquired cytopenic syndrome: _____ - **Go to signature line**

Inherited Abnormalities of Erythrocyte Differentiation or Function

323. Specify the inherited abnormalities of erythrocyte differentiation or function classification:

- Paroxysmal nocturnal hemoglobinuria (PNH) (56)
- Shwachman-Diamond (305)
- Diamond-Blackfan anemia (pure red cell aplasia) (312)
- Other constitutional anemia (319) – **Go to question 324**
- Fanconi anemia (311) (If the recipient developed MDS or AML, indicate MDS or AML as the primary disease).
- Sickle thalassemia (355)
- Sickle cell disease (356)
- Beta thalassemia major (357)
- Other hemoglobinopathy (359) – **Go to question 325**

324. Specify other constitutional anemia: _____

325. Specify other hemoglobinopathy: _____

- **Go to signature line**

Disorders of the Immune System

326. Specify disorder of immune system classification:

- Adenosine deaminase (ADA) deficiency / severe combined immunodeficiency (SCID) (401)
- Absence of T and B cells SCID (402)
- Absence of T, normal B cell SCID (403)
- Omenn syndrome (404)
- Reticular dysgenesis (405)
- Bare lymphocyte syndrome (406)
- Other SCID (419) – **Go to question 327**
- SCID, not otherwise specified (410)
- Ataxia telangiectasia (451)
- HIV infection (452)
- DiGeorge anomaly (454)
- Common variable immunodeficiency (457)

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- Leukocyte adhesion deficiencies, including GP180, CD-18, LFA and WBC adhesion deficiencies (459)
- Kostmann agranulocytosis (congenital neutropenia) (460)
- Neutrophil actin deficiency (461)
- Cartilage-hair hypoplasia (462)
- CD40 ligand deficiency (464)
- Other immunodeficiencies (479) – **Go to question 328**
- Immune deficiency, not otherwise specified (400)
- Chediak-Higashi syndrome (456)
- Griscelli syndrome type 2 (465)
- Hermansky-Pudlak syndrome type 2 (466)
- Chronic granulomatous disease (455)
- Wiskott-Aldrich syndrome (453)
- X-linked lymphoproliferative syndrome (458)

327. Specify other SCID: _____ - **Go to signature line**

328. Specify other immunodeficiency: _____ - **Go to signature line**

Inherited Abnormalities of Platelets

329. Specify inherited abnormalities of platelets classification:

- Congenital amegakaryocytosis / congenital thrombocytopenia (501)
- Glanzmann thrombasthenia (502)
- Other inherited platelet abnormality (509) – **Go to question 330**

330. Specify other inherited platelet abnormality: _____ - **Go to signature line**

Inherited Disorders of Metabolism

331. Specify inherited disorders of metabolism classification:

- Osteopetrosis (malignant infantile osteopetrosis) (521)

Leukodystrophies

- Metachromatic leukodystrophy (MLD) (542)
- Adrenoleukodystrophy (ALD) (543)
- Krabbe disease (globoid leukodystrophy) (544)

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- Lesch-Nyhan (HGPRT deficiency) (522)
- Neuronal ceroid lipofuscinosis (Batten disease) (523)

Mucopolysaccharidoses

- Hurler syndrome (IH) (531)
- Scheie syndrome (IS) (532)
- Hunter syndrome (II) (533)
- Sanfilippo (III) (534)
- Morquio (IV) (535)
- Maroteaux-Lamy (VI) (536)
- β -glucuronidase deficiency (VII) (537)
- Mucopolysaccharidosis (V) (538)
- Mucopolysaccharidosis, not otherwise specified (530)

Mucolipidoses

- Gaucher disease (541)
- Niemann-Pick disease (545)
- I-cell disease (546)
- Wolman disease (547)
- Glucose storage disease (548)
- Mucolipidoses, not otherwise specified (540)

Polysaccharide hydrolase abnormalities

- Aspartyl glucosaminidase (561)
- Fucosidosis (562)
- Mannosidosis (563)
- Polysaccharide hydrolase abnormality, not otherwise specified (560)
- Other inherited metabolic disorder (529) – **Go to question 332**
- Inherited metabolic disorder, not otherwise specified (520)

332. Specify other inherited metabolic disorder: _____ - **Go to signature line**

Histiocytic disorders

333. Specify histiocytic disorder classification:

- Hemophagocytic lymphohistiocytosis (HLH) (571)
- Langerhans cell histiocytosis (histiocytosis-X) (572)
- Hemophagocytosis (reactive or viral associated) (573)
- Malignant histiocytosis (574)

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- Other histiocytic disorder (579) – **Go to question 334**
- Histiocytic disorder, not otherwise specified (570)

334. Specify other histiocytic disorder: _____ - **Go to signature line**

Autoimmune Diseases

335. Specify autoimmune disease classification:

Arthritis

- Rheumatoid arthritis (603)
- Psoriatic arthritis / psoriasis (604)
- Juvenile idiopathic arthritis (JIA): systemic (Stills disease) (640)
- Juvenile idiopathic arthritis (JIA): oligoarticular (641)
- Juvenile idiopathic arthritis (JIA): polyarticular (642)
- Juvenile idiopathic arthritis (JIA): other (643) **Go to question 337**
- Other arthritis (633) – **Go to question 336**

Multiple sclerosis

- Multiple sclerosis (602)

Connective tissue diseases

- Systemic sclerosis (scleroderma) (607)
- Systemic lupus erythematosus (SLE) (605)
- Sjögren syndrome (608)
- Polymyositis / dermatomyositis (606)
- Antiphospholipid syndrome (614)
- Other connective tissue disease (634) – **Go to question 338**

Vasculitis

- Wegener granulomatosis (610)
- Classical polyarteritis nodosa (631)
- Microscopic polyarteritis nodosa (632)
- Churg-Strauss (635)
- Giant cell arteritis (636)
- Takayasu (637)
- Behcet syndrome (638)
- Overlap necrotizing arteritis (639)
- Other vasculitis (611) – **Go to question 339**

Other neurological autoimmune diseases

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- Myasthenia gravis (601)
- Other autoimmune neurological disorder (644) – **Go to question 340**

Hematological autoimmune diseases

- Idiopathic thrombocytopenic purpura (ITP) (645)
- Hemolytic anemia (646)
- Evan syndrome (647)
- Other autoimmune cytopenia (648) – **Go to question 341**

Bowel diseases

- Crohn's disease (649)
- Ulcerative colitis (650)
- Other autoimmune bowel disorder (651) – **Go to question 342**

336. Specify other arthritis: _____

337. Specify other juvenile idiopathic arthritis (JIA): _____

338. Specify other connective tissue disease: _____

339. Specify other vasculitis: _____

340. Specify other autoimmune neurological disorder: _____

341. Specify other autoimmune cytopenia: _____

342. Specify other autoimmune bowel disorder: _____

- Go to signature line

Other Disease

343. Specify other disease: _____

First Name: _____

Last Name: _____

E-mail address: _____

CIBMTR Center Number: _____ CIBMTR Research ID: _____

Date: _____
 YYYY MM DD