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Supplementary Materials and Methods

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Supplemental Tables

Supplemental Table 1: UDS results

UPI	Gender Age*	Sample date	Therapy	BCR-ABL1/ ABL1 (%IS)	T315I (%)	Additional TKI resistance mutations, mutation load (%)							Phase ¹	
1	f 58,8	ID: Dec-05; therapy: im												
		Sep-06	im	60.351	0	F359C; 0								BP
		Oct-06	ni	62.838	1	F359C; 0								BP
		Nov-06	ni	85.153	84	F359C; 8								BP
2	f 84,4	ID: Jul-01; therapy: HU, im												
		Jan-07	HU	98.645	0	G250E; 0 E255K; 0 E255V; 0 F359V; 0 Y253F; 0 M351T; 0 H396R; 0								BP
		Jun-07	HU	34.445	1	G250E; 26 E255K; 6 E255V; 2 F359V; 2 Y253F; 14 M351T; 11 H396R; 24								BP
		Aug-08	da	32.870	99	G250E; 0 E255K; 0 E255V; 0 F359V; 0 Y253F; 0 M351T; 0 H396R; 0								N.A.
3	m 67,6	ID: Jul-00; therapy: HU, im												
		Oct-07	im	50.194	0									N.A.
		Aug-08	im	21.501	1									N.A.
		Dec-08	da	33.486	9									N.A.
		Feb-09	da	19.626	27									N.A.
4	m 68,9	ID: Jul-04; therapy: HU, im												
		Aug-07	im	62.013	0	M244V; 5 F359V; 0								CHR
		Jan-08	im	34.743	1	M244V; 95 F359V; 2								CP
		Apr-08	im	47.587	11	M244V; 86 F359V; 2								CHR
		Nov-08	im	33.507	40	M244V; 51 F359V; 4								CP
5	m 59,6	ID: Feb-09; therapy: FLAG-Ida, im, da no MMR												
		Aug-11	da	5.533	0	G250E; 3 F317L; 25 F317L; 17 M244V; 14 F317L; 12 F359I; 2								N.A.
		Sep-11	da	1.222	1	G250E; 3 F317L; 27 F317L; 12 M244V; 8 F317L; 12 F359I; 2								N.A.
		Oct-11	da	3.359	1	G250E; 0 F317L; 35 F317L; 16 M244V; 6 F317L; 20 F359I; 0								N.A.
		Jan-12	da	7.099	87	G250E; 0 F317L; 7 F317L; 0 M244V; 0 F317L; 1 F359I; 0								N.A.
6	m 18,8	ID: Jun-11; therapy: im no MMR												
		Nov-11	im	4.696	0									N.A.
		Feb-12	im	55.383	1									AP
		May-12	im	51.966	99									N.A.
7	m 33,5	ID: Jul-12; therapy: im; da no MMR												
		Apr-13	da	41.933	0	M244V; 0								N.A.
		Apr-13	da	10.960	1	M244V; 0								CHR
		Jun-13	da	16.480	4	M244V; 0								N.A.
		Jul-13	da	6.423	16	M244V; 2								N.A.

8	f 23,3	ID: Feb-04; therapy: alloSCT, im											
		Jan-11	da	12.698	0	E255K; 0							BP
		Mar-11	im	3.858	2	E255K; 23							N.A.
		Jun-11	da	38.019	57	E255K; 33							N.A.
9	f 56,1	ID: Nov-06; therapy: im											
		Dec-07	da	27.037	2	F317L; 0	M244V; 2					BP	
		Feb-08	im	4.374	82	F317L; 7	M244V; 4					CP	
10	m 57,2	ID: Feb-12; therapy: da no MMR											
		May-12	da	15.529	0							CP	
		Aug-12	da	2.168	3							N.A.	
		Jan-13	da	0.700	30							N.A.	
11	m 45,6	ID: Feb-07; therapy: HU, im											
		Jul-07	im	2.231	4	E459K; 1						N.A.	
		Oct-07	im	19.529	62	E459K; 7						CHR	
12	f 72,7	Mar-10	ID	5.378	0	M244V; 0						BP	
		Sep-10	im	1.968	4	M244V; 0						CHR	
		Nov-10	im	6.986	54	M244V; 10						N.A.	
		ID: Sep-07; therapy: im, HU											
13	f 44,8	May-11	da	8.522	0	M244V; 0	F359V; 84	E355G; 7				N.A.	
		May-11	da	2.239	4	M244V; 5	F359V; 67	E355G; 3				N.A.	
		Feb-12	da	0.610	100	M244V; 0	F359V; 0	E355G; 0				N.A.	
14	f 63,9	ID: before Nov-04, therapy: im											
		Jul-06	im	61.908	0	E255K; 0	M244V; 1					N.A.	
		Aug-06	im	36.130	5	E255K; 0	M244V; 0					N.A.	
		Apr-07	im	68.947	34	E255K; 5	M244V; 3					N.A.	
15	m 69,9	ID: Dec-07; therapy: im CCyR: Apr-08											
		Apr-08	im	0.118	0	E255V; 0	Y253H; 0	L248V; 0				AP	
		Jul-08	im	9.969	5	E255V; 3	Y253H; 41	L248V; 22				AP	
		Nov-08	da	66.324	99	E255V; 0	Y253H; 0	L248V; 0				BP	
16	m 61,3	Feb-12	ID	62.688	0	E255K; 0	E255V; 0	F359C; 0	E355G; 0			CP	
		May-12	im	50.435	7	E255K; 63	E255V; 2	F359C; 10	E355G; 8			BP	
		Jul-12	da	44.527	89	E255K; 7	E255V; 0	F359C; 0	E355G; 0			BP	
17	m 69,8	ID: Nov-00; therapy: HU, im											
		Jun-06	im	37.327	8	G250E; 49	F359C; 8					CP	
		Oct-06	im	58.829	9	G250E; 71	F359C; 12					CP	
		Jan-07	im	58.751	9	G250E; 76	F359C; 5					CP	
		Feb-08	im	40.385	4	G250E; 94	F359C; 0					CP	
		Sep-08	im	30.392	99	G250E; 0	F359C; 0					CP	

18	f 52,1	ID: therapy: Feb-02; HU, IFN, im, da											
		Jul-07	da	17.407	0								N.A.
		Feb-08	da	6.456	10								N.A.
		May-08	da	7.972	4								N.A.
		Jun-08	da	30.130	94							N.A.	
19	f 62,0	ID: May-09; therapy: im, ni											
		Feb-13	alloSCT	13.318	0	G250E; 0	E255K; 0	E255V; 0	F317L; 100	F486S; 0			N.A.
		May-13	alloSCT	2.380	11	G250E; 1	E255K; 9	E255V; 7	F317L; 88	F486S; 2			N.A.
		May-13	alloSCT	33.063	51	G250E; 7	E255K; 6	E255V; 30	F317L; 53	F486S; 7			N.A.
20	m 61,0	Sep-11	ID	45.975	0							CP	
		Jan-12	im	27.036	14							CP	
		Apr-12	im	49.833	91							CP	
21	m 55,7	ID: Apr-08; therapy: im no MMR											
		Aug-08	im	7.863	0	E255V; 0						N.A.	
		Apr-09	im	74.007	27	E255V; 2						AP	
22	m 42,4	ID: Nov-01; therapy: HU, alloSCT, IFN											
		Feb-10	im	53.805	0							BP	
		Jun-10	da	65.635	30							N.A.	
23	m 61,4	ID: Sep-12; therapy: im; da no MMR											
		Feb-14	da	15.362	0							N.A.	
		May-14	da	18.336	30							N.A.	
24	m 81,3	Nov-11	ID	72.252	0	E255K; 0	E255V; 0	E459K; 0	Y253F; 0			BP	
		Jan-12	im	52.970	31	E255K; 8	E255V; 2	E459K; 15	Y253F; 3			BP	
25	m 69,3	ID: Jun-11, therapy: im											
		Jan-14	ni	26.360	0	F359V; 64	Y253H; 3	E450K; 6				N.A.	
		Aug-14	da	38.323	40	F359V; 0	Y253H; 0	E450K; 0				CP	
26	f 61,4	ID: Jul-96; therapy: im											
		Jul-09	im	42.570	0	E459K; 3						N.A.	
		Nov-09	da	45.103	46	E459K; 0						BP	
27	m 65,2	ID: Jul-06; therapy: im											
		Mar-12	ni	2.794	0							N.A.	
		Jun-12	ni	2.834	46							N.A.	
28	m 41,1	ID: Feb-12; therapy: ni no MMR											
		May-12	ni	7.444	0							N.A.	
		Aug-12	ni	66.459	49							BP	

29	m 58,6	ID: Jan-08; therapy: im no MMR											
		Sep-08	im	43.194	0	F359V; 20	Y253H; 2						BP
		Dec-08	da	48.440	55	F359V; 0	Y253H; 0						N.A.
30	m 52,1	ID: Oct-12; therapy: HU, ni no MMR											
		Mar-13	ni	43.170	0	E255K; 4	E255V; 3	F359V; 79					N.A.
		May-13	da	28.060	56	E255K; 9	E255V; 0	F359V; 30					BP
31	m 71,6	ID: Dec-08; therapy: da, im, cytarabin, HU no MMR											
		Nov-13	ni	39.294	0	F359V; 0	E255K; 0						CP
		Jan-14	ni	47.174	64	F359V; 20	E255K; 13						CHR
32	m 59,5	Dec-08	ID	51.109	0	M244V; 0							CP
		Aug-09	im	58.749	68	M244V; 29							CHR
33	m 45,8	ID: Dec-10; therapy: im no MMR											
		Nov-11	im	59.422	0	F317I; 0	F317L; 0	Y253H;100					BP
		Feb-12	im	29.683	76	F317I; 15	F317L; 7	Y253H;100					BP
34	m 85,3	ID: Aug-98; therapy: HU; im, da, ni											
		Jan-12	ni	46.585	0	E255K; 2	F317L; 0	V299L; 18					CP
		Mar-13	ni	53.703	88	E255K; 1	F317L; 7	V299L; 0					CP
35	m 62,3	ID: Sep-10; therapy: da no MMR											
		Apr-11	da	33.591	0								BP
		Jun-11	da	30.393	94								BP
36	m 51,1	ID: Apr-13, therapy: im no MMR											
		Sep-13	im	37.657	0	Q252H; 10							BP
		Oct-13	da	40.200	96	Q252H; 2							BP
37	f 61,3	ID: Jul-06; therapy: im MMR (Jan-07) CCyR (Mar-08)											
		Nov-07	im	47.754	0	E255V; 99							BP
		Apr-08	da	5.379	100	E255V; 99							CP
38	f 38,8	ID: Sept-07; therapy cytarabine -daunorubicin, im, alloSCT, da											
		Oct-08	ni	28.947	0								CHR
		Dec-08	ni	0.046	100								BP
39	f 60,5	ID: Oct-01; therapy: im											
		Mar-09	im	N.A.	0								BP
		Aug-09	da	58.644	100								BP

40	m 72,0	ID: Apr-10, therapy: im no MMR										
		May-10	im	50.260	0							BP
		Jul-10	im	28.353	100							BP

Supplemental Table 1: For patients of cohort 1, given are sampling dates, patients' characteristic (gender, age, date of initial diagnosis, disease phase), mutation loads and *BCR-ABL1/ABL1* (%IS). In case molecular and/or cytogenetic response data was available since initial diagnosis, achievement or absence of treatment milestones is given.

Abbreviations: UPI indicates unique patient identifier; TKI, tyrosine kinase domain; IS, international scale; N.A., not analyzed; f, female; m, male; ID: initial diagnosis; im, imatinib; da, dasatinib; ni, nilotinib; HU, Hydroxyurea, IFN; Interferon alpha; alloSCT, allogeneic stem cell transplantation, IDA-FLAG, idarubicin, fludarabine, cytarabine, G-CSF; CHR, complete hematologic response; CP, chronic phase; AP, accelerated phase; BP, blast phase (definitions according to World Health Organization¹); MMR, major molecular response; CCyR: complete cytogenetic response.

*Age at the time point when T315I had reached the Sanger Sequencing detection limit (end of monitoring period)

Supplemental Table 2: Cytogenetic and molecular cytogenetic results

UPI	Sample date	Additional cytogenetic aberrations affecting the <i>BCR-ABL1</i> sequence ¹ ; ISCN formula	Results FISH
1	Sep-06	no	<i>BCR-ABL+</i>
1	Oct-06	N.A.	<i>BCR-ABL+</i>
1	Nov-06	no	<i>BCR-ABL+</i>
2	Jan-07	yes; 47,XX,t(9;22)(q34;q11),+der(22)t(9;22)(q34;q11)	<i>BCR-ABL+</i> ; duplication of colocalization signal ²
2	Jun-07	No	<i>BCR-ABL+</i>
2	Aug-08	N.A.	N.A.
3	Oct-07	N.A.	<i>BCR-ABL+</i>
3	Aug-08	N.A.	N.A.
3	Dec-08	N.A.	N.A.
3	Feb-09	N.A.	N.A.
4	Aug-07	no	<i>BCR-ABL+</i>
4	Jan-08	no	<i>BCR-ABL+</i>
4	Apr-08	no	<i>BCR-ABL+</i>
4	Nov-08	no	<i>BCR-ABL+</i>
5	Aug-11	N.A.	N.A.
5	Sep-11	N.A.	N.A.
5	Oct-11	N.A.	N.A.
5	Jan-12	N.A.	N.A.
6	Nov-11	N.A.	N.A.
6	Feb-12	yes; 43-44,XY,der(1)t(1;5)(p34;q21)t(1;7)(q21;q31),-3,der(5)t(5;17)(q11;q21),der(6)del(6)(p11)del(6)(q22),-7,t(9;22;12)(q34;q11;q22),der(9)t(6;9)(?q22;p11),der(15)t(3;15)(q13;q25),-17,der(19)t(1;19)(q41;p11), +der(22)t(9;22)(q34;q11)	<i>BCR-ABL+</i> ; duplication of colocalization signal ²
6	May-12	N.A.	N.A.
7	Apr-13	N.A.	N.A.
7	Apr-13	no	<i>BCR-ABL+</i>
7	Jun-13	N.A.	N.A.
7	Jul-13	N.A.	N.A.
8	Jan-11	N.A.	N.A.
8	Mar-11	N.A.	N.A.
8	Jun-11	N.A.	N.A.
9	Dec-07	yes; 46,XX,der(5)t(5;7)(q22;q22),-7,t(9;22)(q34;q11),+der(22)t(9;22)(q34;q11)	<i>BCR-ABL+</i> ; duplication of colocalization signal ²
9	Feb-08	N.A.	<i>BCR-ABL+</i>
10	May-12	N.A.	<i>BCR-ABL+</i>
10	Aug-12	N.A.	N.A.
10	Jan-13	N.A.	N.A.
11	Jul-07	N.A.	<i>BCR-ABL+</i>

11	Oct-07	no	<i>BCR-ABL+</i>
12	Mar-10	no	<i>BCR-ABL+</i>
12	Sep-10	no	N.A.
12	Nov-10	N.A.	N.A.
13	May-11	N.A.	N.A.
13	May-11	N.A.	N.A.
13	Feb-12	N.A.	N.A.
14	Jul-06	N.A.	<i>BCR-ABL+</i> ; duplication of colocalization signal ²
14	Aug-06	N.A.	<i>BCR-ABL+</i> ; duplication of colocalization signal ²
14	Apr-07	yes; 47,XX,t(9;22)(q34;q11), +der(22)t(9;22)(q34;q11)	<i>BCR-ABL+</i> ; duplication of colocalization signal ²
15	Apr-08	No	<i>BCR-ABL-</i>
15	Jul-08	No	<i>BCR-ABL+</i>
15	Nov-08	N.A.	N.A.
16	Feb-12	N.A.	<i>BCR-ABL+</i>
16	May-12	no	N.A.
16	Jul-12	N.A.	N.A.
17	Jun-06	no	<i>BCR-ABL+</i>
17	Oct-06	no	<i>BCR-ABL+</i>
17	Jan-07	no	<i>BCR-ABL+</i>
17	Feb-08	no	<i>BCR-ABL+</i>
17	Sep-08	no	<i>BCR-ABL+</i>
18	Jul-07	N.A.	N.A.
18	Feb-08	N.A.	N.A.
18	May-08	N.A.	N.A.
18	Jun-08	no	<i>BCR-ABL+</i>
19	Feb-13	N.A.	N.A.
19	May-13	N.A.	N.A.
19	May-13	N.A.	N.A.
20	Sep-11	no	<i>BCR-ABL+</i>
20	Jan-12	N.A.	N.A.
20	Apr-12	N.A.	N.A.
21	Aug-08	N.A.	<i>BCR-ABL+</i>
21	Apr-09	yes; 47,XY,t(9;22)(q34;q11),dup(17)(q23q25), +der(22)t(9;22)(q34;q11)	<i>BCR-ABL+</i> ; duplication of colocalization signal ²
22	Feb-10	yes; 48,XY,+8,t(9;22)(q34;q11), +der(22)t(9;22)(q34;q11)	<i>BCR-ABL+</i> ; duplication of colocalization signal ²
22	Jun-10	yes; 46,XY,t(9;22)(q34;q11), +der(22)t(9;22)(q34;q11)	<i>BCR-ABL+</i> ; duplication of colocalization signal ²
23	Feb-14	N.A.	N.A.
23	May-14	N.A.	N.A.
24	Nov-11	no	<i>BCR-ABL+</i>
24	Jan-12	N.A.	N.A.

25	Jan-14	N.A.	N.A.
25	Aug-14	N.A.	N.A.
26	Jul-09	N.A.	N.A.
26	Nov-09	yes; 47,XX,-7,+8,+8,der(8;12)(p10;q10), t(9;22)(q34;q11),i(17)(q10), +der(22)t(8;22)(q11;q11)t(8;12)(q24;p11)t(12;22)(p13;q11)t(9;22)(q34;q11)	<i>BCR-ABL+</i> ; duplication of colocalization signal ²
27	Mar-12	N.A.	N.A.
27	Jun-12	N.A.	N.A.
28	May-12	N.A.	N.A.
28	Aug-12	yes; 46,XY,dic(7;9)(p12;p12),t(9;22)(q34;q11), +der(22)t(9;22)(q34;q11)	<i>BCR-ABL+</i> ; duplication of colocalization signal ²
29	Sep-08	N.A.	N.A.
29	Dec-08	no	<i>BCR-ABL+</i> ; duplication of colocalization signal ²
30	Mar-13	N.A.	<i>BCR-ABL+</i>
30	May-13	no	N.A.
31	Nov-13	N.A.	N.A.
31	Jan-14	N.A.	N.A.
32	Dec-08	no	<i>BCR-ABL+</i>
32	Aug-09	no	<i>BCR-ABL+</i>
33	Nov-11	no	<i>BCR-ABL+</i>
33	Feb-12	no	<i>BCR-ABL+</i>
34	Jan-12	N.A.	N.A.
34	Mar-13	N.A.	N.A.
35	Apr-11	yes; 48,XY,t(9;22)(q34;q11),+21, +der(22)t(9;22)(q34;q11)	N.A.
35	Jun-11	no	N.A.
36	Sep-13	no	<i>BCR-ABL+</i>
36	Oct-13	N.A.	N.A.
37	Nov-07	no	<i>BCR-ABL+</i>
37	Apr-08	no	<i>BCR-ABL+</i>
38	Oct-08	no	<i>BCR-ABL+</i>
38	Dec-08	no	<i>BCR-ABL-</i>
39	Mar-09	no	<i>BCR-ABL+</i>
39	Aug-09	N.A.	N.A.
40	May-10	no	<i>BCR-ABL+</i>
40	Jul-10	N.A.	<i>BCR-ABL+</i>

Supplemental Table 2: For patients of cohort 1, given are additional cytogenetic aberrations.

Abbreviations: N.A. indicates not analyzed; FISH, Fluorescence in situ hybridization; ISCN: International System of Cytogenetic Nomenclature

¹Listed are only cytogenetic aberrations in addition to the Philadelphia chromosome affecting the *BCR-ABL1* sequence. Patients with other chromosomal aberrations are listed in the category “no”. For 14/42 patients of cohort 2 cytogenetic or molecular cytogenetic data was available. No aberration affecting the *BCR-ABL1* copy number was detected.

²duplication of colocalization signal indicates duplication of Philadelphia chromosome

Supplemental Table 3: Primer sequences

454_1st PCR-F	GAGCAGCAGAAGAAGTGTTCAGA
454_1st PCR-R	CTTGGAGTGAGGCATCTCAG
454_Seq1-F	TGTCTATGGTGTGTCCCCCA
454_Seq2-F	TGCTGTACATGGCCACTCAG
454_Seq1-R	ACTTGTTGTAGGCCAGGCTC
454_Seq2-R	CCTGCAGCAAGGTA CT CACA

Supplemental Table 3: Primers given in table were used to generate amplicons for 454 XL+ sequencing.

Supplemental Table 4: PCR amplification

<i>BCR-ABL1</i> fusion transcript amplification (1st PCR):		
95 °C	5 min	
95 °C	30 sec	x29
60 °C	30 sec	
72 °C	3 min 30 sec	
72 °C	7 min	
Sequencing amplicon amplification (nested PCR)		
95 °C	5 min	
95 °C	30 sec	x29
60 °C	30 sec	
72 °C	50 sec	
72 °C	7 min	

Supplemental Table 4: PCR conditions to generate amplicons for 454 XL+ sequencing are given.

Supplemental Table 5: Dilution experiment

		mutation load				
	mixing ratio	Q252H	E255K	T315I	F359V	E499E
Sample A	90	2%		97%		
Sample B	10		14%	64%	21%	100%
calculated mutation load		2%	1%	93%	2%	10%

	mixing ratio	Q252H	E255K	T315I	F359V	E499E
Sample A	99	2%		97%		
Sample B	1		14%	64%	21%	100%
calculated mutation load		2%	0%*	97%	0%*	1%

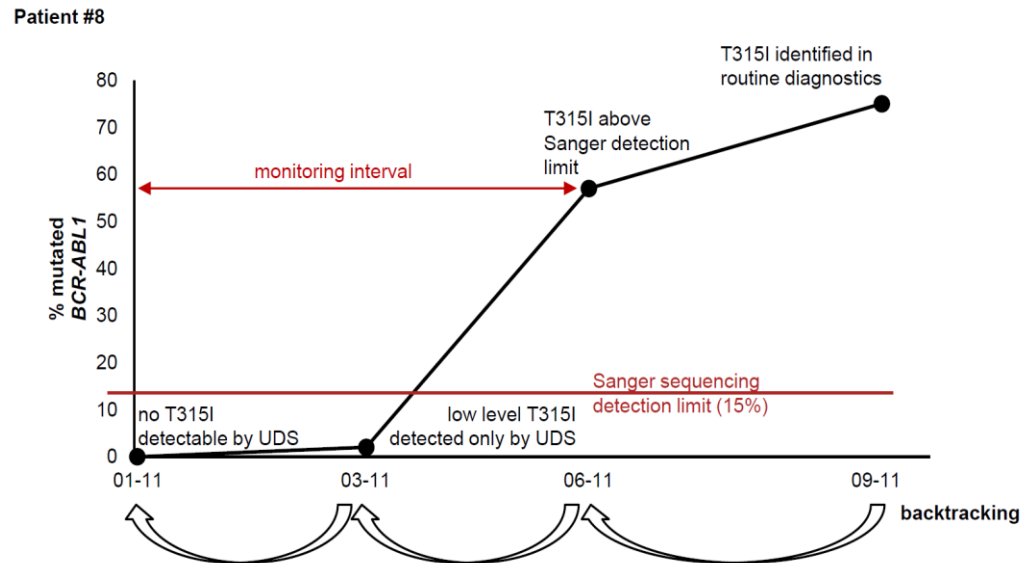
	mixing ratio	M244V	E255K	T315I
Sample C	1	2%		61%
Sample D	2		99%	
calculated mutation load		1%	66%	20%

Supplemental Table 5: Samples were mixed in the indicated ratio to simulate mutations with low and defined loads.

*Mutation loads below 1% are not called.

Supplemental Figures

Figure 1



Supplemental Figure 1: Patient #8 illustrates our study design. From the initial time point when T315I was identified during routine diagnostics, we backtracked earlier samples until no T315I mutation was detectable. We selected the sample from June 2011 as endpoint of the monitoring interval, because the T315I mutation load had reached the Sanger sequencing detection limit, and excluded the sample from September 2011.

Figure 2



Supplemental Figure 2: Individual sequencing reads (33 of 2562) of patient #33 (s. Figure 5D) are shown. The c.944C>T (T315I), the c.949T>A (F317I) and the c.951T>A (F317L) mutated bases are highlighted in yellow.

References

1. Baccharani M, Deininger MW, Rosti G, et al. European LeukemiaNet recommendations for the management of chronic myeloid leukemia: 2013. *Blood*. 2013;122(6):872-884.