
Expression profiling of adult acute lymphoblastic leukemia identifies a *BCR-ABL1*-like subgroup characterized by high non-response and relapse rates

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SUPPLEMENTAL DATA for

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Content

Supplemental Table S1. Characteristics and outcome of HOVON patients who were included in the molecular analyses compared with the total study cohort

Supplemental Table S2. Reverse transcription PCR primers used for the detection of tyrosine kinase fusion genes. Design as reported by Roberts *et al.*, New Engl J Med 2014

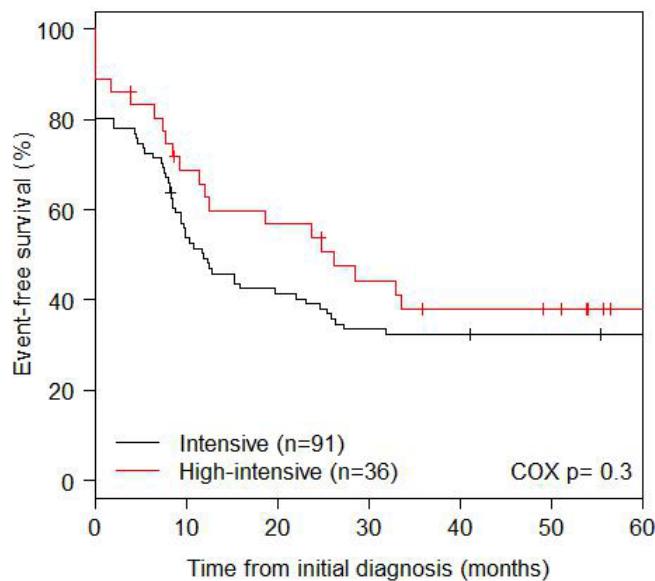
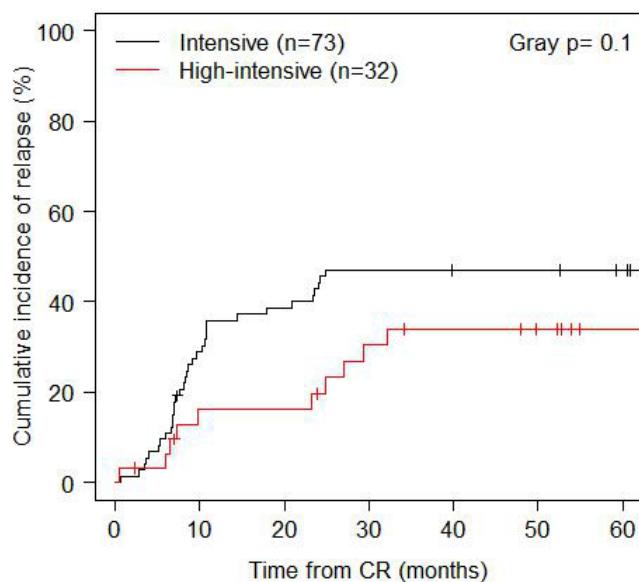
Supplemental Figure S1. Clinical outcome for High-intensive and Intensive treatment protocols

Supplemental Table S1. Characteristics and outcome of HOVON patients who were included in the molecular analyses compared with the total study cohort

	Included in BCR-ABL1-like paper				Total
	no [# and column %]	yes			
Total	305 100%	102 100%			407 100%
Trial code					
HO18	96 31%	38 37%			134 33%
HO37	141 46%	41 40%			182 45%
HO70	28 9%	8 8%			36 9%
HO71	40 13%	15 15%			55 14%
Sex					
M	169 55%	46 45%			215 53%
F	136 45%	56 55%			192 47%
Age					
Median	36	36			36
Range	15-69	16-68			15-69
WBC [x 10⁹/l]					
Median	7.1	14.1			8.8
Range	0.4-878	0.7-505			0.4-878
WBC: <30 vs. >=30 x 10E9/l					
<30	240 79%	68 67%			308 76%
>=30	65 21%	34 33%			99 24%
CR after cycle 1					
no	83 27%	35 34%			118 29%
yes	222 73%	67 66%			289 71%
Risk based on WBC, (cyto)genetics and CR1					
standard	162 53%	38 37%			200 49%
high	143 47%	64 63%			207 51%
CR (on protocol) achieved					
no	38 12%	19 19%			57 14%
yes	267 88%	83 81%			350 86%
Event free survival					
median	15 mo	13 mo			15 mo
at 10 years	31%	30%			31%
Disease free survival from CR					
median	17 mo	24 mo			18 mo
at 10 years	35%	37%			35%
Overall survival					
median	23 mo	23 mo			23 mo
at 10 years	35%	34%			35%

Supplemental Table S2. Reverse transcription PCR primers used for the detection of tyrosine kinase fusion genes. Design as reported by Roberts *et al.*, New Engl J Med 2014

Gene fusion	Forward Primer ID	Forward sequence 5' to 3'	Reverse Primer ID	Reverse sequence 5' to 3'
ETV6-ABL1	ETV6_exon5_F4	ggagaataatcaactgcccagcgtcct	ABL1_exon4_R2	gccaccgtcaggctgtattctcc
NUP214-ABL1	NUP_exon20_F5	cagtggccctggaggaaaacccagt	ABL1_exon4_R2	gccaccgtcaggctgtattctcc
RANBP2-ABL1	RANBP2_exon16_F2	tggttcttgcgaaatgcagattca	ABL1_exon4_R2	gccaccgtcaggctgtattctcc
RCSD1-ABL1	RCSD1_exon3_F3	cagccagtaaaccacccgaaggaa	ABL1_exon4_R2	gccaccgtcaggctgtattctcc
SNX2-ABL1	SNX2_exon3_F1	cggAACCTTCTCCTGcagtacacc	ABL1_exon4_R2	gccaccgtcaggctgtattctcc
ZMZ1-ABL1	ZMZ1_exon17_F2	aggaccggcagatgaacaccaactg	ABL1_exon4_R2	gccaccgtcaggctgtattctcc
PAG1-ABL2	PAG1_exon7_F1	ccttcaggagaaggaaaggggggagg	ABL2_exon6_R3	gtcggttatggggacacaccatag
RCSD1-ABL2	RCSD1_exon3_F3	cagccagtaaaccacccgaaggaa	ABL2_exon6_R3	gtcggttatggggacacaccatag
ZC3HAV1-ABL2	ZC3HAV1_exon11_F2	tgtcagagatccatcacctacatcca	ABL2_exon6_R3	gtcggttatggggacacaccatag
SSBP2-CSF1R	SSBP2_exon12_F2	tcatgccttagtccagcagattcaacca	CSF1R_exon14_R2	tggctcatatcttcagctggaca
MYH9-CSF1R	MYH9_exon1_F2	gcgggaaggcggcgaggag	CSF1R_exon14_R2	tggctcatatcttcagctggaca
ETV6-NTRK3	ETV6_exon5_F4	ggagaataatcaactgcccagcgtcct	NTRK3_exon15_R1	atcttgtcttggtcggctgaggt
ATF7IP-JAK2	ATF7IP_exon12_F2	aaccctatacaaccagcacggctct	JAK2_exon20_R4	tgttgtcatgtctgtagggatttcagga
BCR-JAK2	BCR_exon1_F1	tgcccataaggcggccacggcact	JAK2_exon20_R4	tgttgtcatgtctgtagggatttcagga
EBF1-JAK2	EBF_exon14_F2	cacgagcatgaacggatacggctct	JAK2_exon20_R4	tgttgtcatgtctgtagggatttcagga
ETV6-JAK2	ETV6_exon3_F1	atggcaaagtctctgtctgac	JAK2_exon20_R4	tgttgtcatgtctgtagggatttcagga
PAX5-JAK2	PAX5_exon3_F2	acaatgacaccgtgcctagctcag	JAK2_exon20_R4	tgttgtcatgtctgtagggatttcagga
PPFIBP1-JAK2	PPFIBP1_exon10_F1	tgcaggatgaaaggaaagggttga	JAK2_exon20_R4	tgttgtcatgtctgtagggatttcagga
SSBP2-JAK2	SSBP2_exon7_F1	ggcacttggaggtgtcccaggaaagt	JAK2_exon20_R4	tgttgtcatgtctgtagggatttcagga
STRN3-JAK2	STRN3_exon7_F3	tgaaggaggctggagaagcacggagt	JAK2_exon20_R4	tgttgtcatgtctgtagggatttcagga
TPR-JAK2	TPR_exon38_F1	tggaaatgccttccaaagaagtga	JAK2_exon20_R4	tgttgtcatgtctgtagggatttcagga
TERF2-JAK2	TERF2_exon8_F	agacttgggtggaagaggatgaactgt	JAK2_exon20_R4	tgttgtcatgtctgtagggatttcagga
EBF1-PDGFRB	EBF1_exon11_F	ccaccatcgattatggttccagaggt	PDGFRB_exon13_R2	tttcatgtggcctgagaatggctc
TNIP1-PDGFRB	TNIP1_exon13_F2	aagcactgagcatccaaacc	PDGFRB_exon13_R2	tttcatgtggcctgagaatggctc
ZEB2-PDGFRB	ZEB2_exon9_F2	cccaacttggaggcgtactttgctga	PDGFRB_exon13_R2	tttcatgtggcctgagaatggctc
SSBP2-PDGFRB	SSBP2_exon7_F1	ggcacttggaggtgtcccaggaaagt	PDGFRB_exon13_R2	tttcatgtggcctgagaatggctc
SSBP2-PDGFRB	SSBP2_exon12_F2	tcatgccttagtccagcagattcaacca	PDGFRB_exon13_R2	tttcatgtggcctgagaatggctc
MYB-TYK2	MYB_exon6_F1	tcaggctccgcctacagtcaactc	TYK2_exon19_R2	tccgggttacagtcaagacgtcag
MYH9-IL2RB	MYH9_exon1_F2	gcgggaaggcggcgaggag	IL2RB_exon10_R2	caccctggccatctgtctcacaccaa
IQGAP2-TSLP	IQGAP2_exon1_F2	agatcttggcgctagggaaat	TSLP_exon1_2_R1	tggctatattcacagcacaaaataa

A**B**

Supplemental Figure S1. Clinical outcome for High-intensive and Intensive treatment protocols

Patients had been included in one of four consecutive clinical trials of the Dutch-Belgium HOVON study group or had been treated accordingly in intensive (HOVON-18 or HOVON-37, n=91, 1993-2005) or pediatric-inspired high-intensive (HOVON-70 or HOVON-71, n=36, 2005-2009) protocols.

(A) Event-free survival curves for High-intensive and Intensive treatment protocols. The events were no CR on protocol, relapse, and death. (B) Cumulative incidence of relapse curves from time of CR for High-intensive and Intensive treatment protocols. For CIR analyses, relapse was taken as event, and death as a competing event.