

Exam No	Sex	BCOR mut=1; BCOR wt=0	Material sent	Func	Gene	ExonicFunc	AChange	Start	End	Ref	Obs	mutation_load	reads_total	reads_variant	Age	Morphology	Cytogenetics	IPSS	IPSS-R
10-012914	Male	1	BM	exonic	BCOR	frameshift insertion	NM_001123384:c.4172_4173insGGAGA:p.T1391fs	39921492	39921492	-	TCT	0.470394737	304	143	81	RARS-T	normal karyotype	low	low
09-011649	Male	1	BM	exonic	BCOR	frameshift insertion	NM_001123384:c.3367_3368insAA:p.D1123fs	39923670	39923670	-	TT	0.74611399	193	144	61.9	RCMD	normal karyotype	intermediate 1	intermediate
09-002790	Male	1	BM	exonic;s	BCOR	stopgain SNV	NM_001123384:c.G4663T:p.E1555X	39913509	39913509	C	A	0.707317073	779	551	62.9	RAEB-1	normal karyotype	intermediate 1	intermediate
10-022162	Male	1	BM	exonic	BCOR	stopgain SNV	NM_001123384:c.C3433T:p.R1145X	39923604	39923604	G	A	0.96835443	158	153	80.8	RAEB-1	normal karyotype	intermediate 1	intermediate
09-026043	Male	1	BM	exonic	BCOR	stopgain SNV	NM_001123384:c.C3436T:p.R1146X	39923601	39923601	G	A	0.75	240	180	67.8	RAEB-1	normal karyotype	intermediate 1	high
09-025370	Male	1	BM	exonic	BCOR	frameshift deletion	NM_001123383:c.811delT:p.S271fs	39933788	39933788	A	-	0.459821429	448	206	75.2	RAEB-2	normal karyotype	intermediate 2	high
10-024978	Male	1	BM	exonic	BCOR	frameshift insertion	NM_001123383:c.1655_1656insG:p.A552fs	39932944	39932944	-	C	0.829424307	469	389	69.6	RAEB-2	normal karyotype	intermediate 2	very high
05-002208	Male	1	BM	exonic	BCOR	frameshift deletion	NM_001123384:c.3248delC:p.P1083fs	39923789	39923789	G	-	0.752124646	706	531	83.6	RAEB-2	normal karyotype	not known	not known
07-008922	Male	1	BM	exonic	BCOR	frameshift deletion	NM_001123384:c.3819delA:p.K1273fs	39922197	39922197	T	-	0.132882883	444	59	81.2	RCMD	normal karyotype	intermediate 1	low
08-018374	Male	1	BM	exonic	BCOR	nonsynonymous SNV	NM_001123384:c.G4234A:p.E1412K	39921430	39921430	C	T	0.111111111	153	17	69.3	RAEB-1	normal karyotype	intermediate 1	low
09-005669	Male	1	BM	exonic	BCOR	frameshift deletion	NM_001123384:c.4888_4897del:p.1630_1633del	39911577	39911586	GAA	-	0.602006689	299	180	58	RAEB-1	normal karyotype	intermediate 1	high
08-005827	Male	1	BM	exonic	BCOR	frameshift insertion	NM_001123384:c.4568_4569insA:p.M1523fs	39914638	39914638	-	T	0.206395349	344	71	61.3	RAEB-2	normal karyotype	intermediate 2	high
09-008756	Male	0	BM												60.6	RAEB-1	normal karyotype	intermediate 1	intermediate
09-011674	Male	0	BM												67.5	RCMD	normal karyotype	intermediate 1	intermediate
08-021733	Male	0	BM												80.9	RAEB-1	normal karyotype	intermediate 1	intermediate
07-000867	Male	0	BM												75.6	RAEB-2	normal karyotype	intermediate 2	high
08-009253	Male	0	BM												83.8	RAEB-2	normal karyotype	intermediate 2	intermediate
12-006306	Male	0	BM												83.1	RARS-T	normal karyotype	low	low
09-004647	Male	0	BM												67.5	RAEB-1	normal karyotype	intermediate 1	high
12-019861	Male	0	BM												72.5	RAEB-2	normal karyotype	intermediate 2	very high
07-002826	Male	0	BM												80.8	RCMD	normal karyotype	low	very low
11-009704	Male	0	BM												81	RARS-T	normal karyotype	low	very low
08-008723	Male	0	BM												68.3	RA	normal karyotype	low	low
08-025142	Male	0	BM												83.8	RCMD	normal karyotype	low	low
07-001244	Male	0	BM												62.4	RAEB-1	normal karyotype	intermediate 1	low
08-006600	Male	0	BM												80.7	RAEB-1	normal karyotype	intermediate 1	low
08-007100	Male	0	BM												70.1	RAEB-1	normal karyotype	intermediate 1	low
08-019074	Male	0	BM												68.3	RAEB-1	normal karyotype	intermediate 1	low
09-005067	Male	0	BM												62.5	RAEB-1	normal karyotype	intermediate 1	low
09-013930	Male	0	BM												63.2	RCMD	normal karyotype	intermediate 1	low
10-018512	Male	0	BM												58.8	RAEB-1	normal karyotype	intermediate 1	low

11-031738	Male	0	BM													69.7	RAEB-1	normal karyotype	intermediate 1	low	
05-001144	Male	0	BM													27.6	RA	normal karyotype	intermediate 1	intermediate	
11-026392	Male	0	BM													70.8	RAEB-2	normal karyotype	intermediate 1	intermediate	
07-011921	Male	0	BM													81.7	RAEB-2	normal karyotype	intermediate 2	high	
09-002844	Male	0	BM													62.6	RAEB-2	normal karyotype	intermediate 2	high	
08-017576	Male	0	BM													61.8	RAEB-2	normal karyotype	not known	not known	
09-000188	Male	0	BM													75.3	RAEB-2	normal karyotype	not known	not known	
10-027628	Female	1	BM	exonic	BCOR	nonsynonymous SNV	NM_001123383:c.G1294A:p.V432I	39933305	39933305	C	T	0.444444444	1611	716	74.8	RCMD-RS	normal karyotype	low	low	very low	
10-008924	Female	1	BM	splicing	BCOR	splice-site mutation	NM_001123384:E10	39921389	39921391	CAC	-	0.207920792	202	42	76.1	RARS	normal karyotype	low	low	low	
11-002758	Female	1	BM	exonic	BCOR	stopgain SNV	NM_001123383:c.C1118A:p.S373X	39933481	39933481	G	T	0.458398744	637	292	72.1	RCMD-RS	normal karyotype	low	low	low	
12-029317	Female	1	BM	exonic	BCOR	nonsynonymous SNV	NM_001123384:c.A4085G:p.Q1362R	39921579	39921579	T	C	0.443762781	489	217	70.7	RCMD-RS	normal karyotype	low	low	low	
09-007410	Female	1	BM	exonic	BCOR	stopgain SNV	NM_001123384:c.G3382T:p.E1128X	39923655	39923655	C	A	0.439285714	280	123	68.5	RCMD-RS	normal karyotype	low	low	intermediate	
09-011104	Female	1	BM	exonic	BCOR	stopgain SNV	NM_001123384:c.C3433T:p.R1145X	39923604	39923604	G	A	0.405919662	473	192	86.9	RAEB-2	normal karyotype	intermediate 1	intermediate	intermediate	
09-018857	Female	1	BM	exonic	BCOR	frameshift insertion	NM_001123383:c.2751_2752insTTTA:p.T917fs	39931848	39931848	-	TAA	0.221078134	1651	365	70.8	RAEB-1	normal karyotype	intermediate 1	intermediate	intermediate	
09-018165	Female	1	BM	exonic;s	BCOR	stopgain SNV	NM_001123384:c.3448_3449insTA:p.G1150_L1151delinsX	39923589	39923589	-	TA	0.391061453	537	210	59.4	RAEB-1	normal karyotype	intermediate 1	intermediate	high	
10-034166	Female	1	BM	splicing	BCOR	splice-site mutation	NM_001123384:E14	39913137	39913137	A	G	0.365384615	780	285	74.9	RAEB-2	normal karyotype	intermediate 2	intermediate	high	
11-030207	Female	1	BM	exonic	BCOR	stopgain SNV	NM_001123383:c.A634T:p.K212X	39933965	39933965	T	A	0.161574314	1931	312	85.1	RAEB-2	normal karyotype	intermediate 2	intermediate	high	
09-002017	Female	1	BM	exonic	BCOR	frameshift insertion	NM_001123383:c.916_917insT:p.H306fs	39933683	39933683	-	A	0.40776699	103	42	70.7	RAEB-2	normal karyotype	intermediate 2	intermediate	very high	
12-003018	Female	0	BM													70.4	RCMD-RS	normal karyotype	low	low	very low
09-025638	Female	0	BM													67.6	RCMD-RS	normal karyotype	low	low	low
10-036519	Female	0	BM													76.3	RARS	normal karyotype	low	low	low
11-026186	Female	0	BM													74.7	RCMD-RS	normal karyotype	low	low	low
11-029005	Female	0	BM													78.3	RCMD-RS	normal karyotype	low	low	low
11-032640	Female	0	BM													65.3	RCMD-RS	normal karyotype	low	low	low
12-013743	Female	0	BM													74.9	RCMD-RS	normal karyotype	low	low	low
10-026107	Female	0	BM													75.8	RCMD-RS	normal karyotype	intermediate 1	intermediate	low
11-032285	Female	0	BM													72.8	RAEB-1	normal karyotype	intermediate 1	intermediate	low
06-012269	Female	0	BM													79.5	RAEB-1	normal karyotype	intermediate 1	intermediate	intermediate
07-007280	Female	0	BM													73.8	RAEB-1	normal karyotype	intermediate 1	intermediate	intermediate
08-004706	Female	0	BM													60.6	RAEB-1	normal karyotype	intermediate 1	intermediate	intermediate
08-006581	Female	0	BM													56.4	RAEB-1	normal karyotype	intermediate 1	intermediate	intermediate
08-018593	Female	0	BM													71.9	RAEB-1	normal karyotype	intermediate 1	intermediate	intermediate
09-010719	Female	0	BM													73.5	RAEB-1	normal karyotype	intermediate 1	intermediate	intermediate
10-011271	Female	0	BM													75.8	RAEB-2	normal karyotype	intermediate 1	intermediate	intermediate
10-031376	Female	0	BM													74	RAEB-2	normal karyotype	intermediate 2	intermediate	intermediate

07-012062	Female	0	BM													71.7	RAEB-2	normal karyotype	intermediate 2	high
08-003790	Female	0	BM													83.2	RAEB-2	normal karyotype	intermediate 2	high
09-025284	Female	0	BM													89.1	RAEB-2	normal karyotype	intermediate 2	high
10-017048	Female	0	BM													85.7	RAEB-2	normal karyotype	intermediate 2	high
10-022875	Female	0	BM													87.4	RAEB-2	normal karyotype	intermediate 2	high
07-013982	Female	0	BM													76.3	RARS	normal karyotype	not known	not known
11-014829	Female	0	BM													74.4	RCMD-RS	normal karyotype	not known	not known