

Supplementary Table S2. Data of DNA copy number normalized to Log2 of chromosomes 14, 17, and 22. The values in yellow represent zones of the chromosome where we found a gain in T-ALL and B-ALL samples. Chr: chromosome, Start: starting position, End: finishing position, GSM: sample identifier.

T-ALL Samples											
Chr	Start	End	GSM1963457	GSM1963456	GSM1963455	GSM1963454	GSM1963453	GSM1963452	GSM1963451	GSM1963450	GSM1963449
chr14	21444377	21444436	1.81E-01	-2.79E-02	5.12E-01	-1.35E-01	2.27E-01	1.52E-01	-5.04E-01	2.85E-01	2.99E-01
chr14	21646729	21646788	1.87E-01	4.59E-01	-1.75E+00	-3.99E-01	-4.43E-01	-2.28E-01	7.91E-02	4.11E-01	1.02E-01
B-ALL Samples											
Chr	Start	End	GSM1963407	GSM1963406	GSM1963405	GSM1963404	GSM1963403	GSM1963402	GSM1963401	GSM1963400	GSM1963399
chr14	21444377	21444436	-4.46E-01	1.89E-01	7.96E-02	2.49E-01	4.30E-01	-1.22E-01	-4.25E-02	-4.44E-01	4.81E-01
chr14	21646729	21646788	-2.74E-01	1.84E-01	7.18E-02	3.05E-01	5.66E-01	9.03E-02	3.54E-03	-3.85E-01	2.78E-01
T-ALL Samples											
Chr	Start	End	GSM1963457	GSM1963456	GSM1963455	GSM1963454	GSM1963453	GSM1963452	GSM1963451	GSM1963450	GSM1963449
chr17	41755952	41756011	-4.75E-01	3.78E-02	3.73E-01	3.19E-01	-1.43E-02	4.11E-01	6.96E-01	1.57E-01	1.56E-01
chr17	41788252	41788311	6.75E-02	-1.44E-01	7.47E-01	3.61E-01	1.76E-02	-1.60E-01	2.81E-01	-2.25E-01	5.06E-01
chr17	41819694	41819753	-4.96E-01	-1.87E-01	7.00E-01	3.95E-01	-1.26E-01	5.49E-02	1.95E-01	-3.12E-01	2.57E-01
chr17	41854798	41854857	-5.44E-01	-6.43E-03	5.36E-01	5.25E-01	3.81E-01	1.64E-01	1.58E-01	-5.98E-01	-2.68E-01
chr17	41876006	41876065	-4.54E-01	-3.51E-01	4.91E-01	3.46E-01	2.72E-01	7.99E-02	8.81E-02	-6.14E-01	-4.35E-02
chr17	41958730	41958789	-5.01E-01	-2.22E-02	7.21E-02	2.60E-01	1.01E-01	4.05E-01	6.29E-01	-5.55E-03	3.65E-01
chr17	41999354	41999413	-3.01E-01	-2.59E-01	5.92E-01	4.25E-01	7.19E-02	2.35E-01	6.03E-01	9.03E-02	1.13E-01
chr17	42005133	42005192	-4.48E-01	9.57E-02	5.07E-01	6.10E-01	1.72E-01	-1.97E-01	2.21E-01	-3.51E-01	1.38E-01
B-ALL Samples											
Chr	Start	End	GSM1963407	GSM1963406	GSM1963405	GSM1963404	GSM1963403	GSM1963402	GSM1963401	GSM1963400	GSM1963399
chr17	41755952	41756011	1.55E-01	-8.37E-02	-4.26E-01	-3.83E-01	-2.01E-01	1.59E-04	-6.93E-01	-4.73E-01	6.89E-01
chr17	41788252	41788311	-1.03E-02	6.16E-01	-1.17E-02	-2.09E-01	-1.83E-01	2.79E-01	-2.55E-01	-7.88E-04	4.83E-01
chr17	41819694	41819753	-3.70E-01	2.68E-01	-2.53E-01	7.31E-02	-2.95E-01	-5.47E-01	-6.98E-01	-6.97E-01	9.96E-01
chr17	41854798	41854857	-5.97E-01	6.92E-02	-5.82E-01	-4.24E-01	-5.48E-01	-6.81E-01	-8.59E-01	-6.52E-01	1.14E+00
chr17	41876006	41876065	-7.22E-01	3.88E-01	-6.13E-01	5.58E-03	-2.76E-01	-7.16E-01	-6.07E-01	-3.28E-01	1.17E+00
chr17	41958730	41958789	8.66E-02	-1.38E-02	-1.79E-01	-4.45E-01	-5.39E-01	-1.57E-01	-5.68E-01	-6.78E-01	4.61E-01

chr17	41999354	41999413	3.17E-01	2.21E-01	-4.66E-01	-4.39E-01	-4.95E-01	-4.38E-01	-6.51E-01	-2.05E-02	7.85E-01
chr17	42005133	42005192	-2.77E-02	4.13E-01	-4.67E-01	-2.91E-01	-6.24E-01	-3.66E-01	-3.44E-01	-5.14E-01	5.91E-01
T-ALL Samples											
Chr	Start	End	GSM1963457	GSM1963456	GSM1963455	GSM1963454	GSM1963453	GSM1963452	GSM1963451	GSM1963450	GSM1963449
chr22	20026630	20026689	-2.16E-01	-8.50E-01	-1.96E+00	-9.70E-01	1.14E-01	1.09E-01	1.22E-01	-5.38E-01	2.08E-01
chr22	20037371	20037430	-7.03E-01	-3.99E-01	-8.94E-01	-3.78E-01	-7.05E-02	8.93E-01	3.28E-01	-7.28E-01	1.65E-01
B-ALL Samples											
Chr	Start	End	GSM1963407	GSM1963406	GSM1963405	GSM1963404	GSM1963403	GSM1963402	GSM1963401	GSM1963400	GSM1963399
chr22	20026630	20026689	-1.02E-01	-3.15E-01	-4.36E-01	2.37E-01	4.23E-01	5.49E-01	8.45E-02	-8.78E-02	-5.45E-01
chr22	20037371	20037430	1.87E-01	-1.16E-01	-7.11E-01	-7.12E-02	1.02E+00	3.05E-01	-4.44E-01	-6.37E-01	-2.53E-01