Detection of genomic aberrations in molecularly defined Burkitt's lymphoma by array-based, high resolution, single nucleotide polymorphism analysis

René Scholtysik,[1] Markus Kreuz,[2] Wolfram Klapper,[3] Birgit Burkhardt,[4] Alfred C. Feller,[5] Michael Hummel,[6] Markus Loeffler,[2] Maciej Rosolowski,[2] Carsten Schwaenen,[7] Rainer Spang,[8] Harald Stein,[6] Christoph Thorns,[5] Lorenz Trümper,[9] Inga Vater,[10] Swen Wessendorf,[7] Thorsten Zenz,[7] Reiner Siebert,[10] and Ralf Küppers,[1] for the "Molecular Mechanisms in Malignant Lymphomas" Network Project of the Deutsche Krebshilfe

[1]Institute of Cell Biology (Cancer Research), Medical School, University of Duisburg-Essen, Essen; [2]Institute of Medical Informatics, Statistics and Epidemiology (IMISE), University of Leipzig, Leipzig; [3]Department of Pathology, Hematopathology Section and Lymph Node Registry, University Hospital Schleswig-Holstein, Campus Kiel/Christian-Albrechts-University, Kiel; [4]NHL-BFM Study Center, Department of Pediatric Hematology and Oncology, Justus-Liebig-University, Giessen; [5]Institute of Pathology, University of Lübeck, Lübeck; [6]Institute of Pathology, Charité, Campus Benjamin Franklin, Berlin; [7]Internal Medicine III, University Hospital of Ulm, Ulm; [8]Institute of Functional Genomics, University of Regensburg, Regensburg; [9]Department of Hematology/Oncology, University Hospital Göttingen, Göttingen and [10]Institute of Human Genetics, Christian-Albrechts University Kiel & University Hospital Schleswig-Holstein, Campus Kiel, Kiel, Germany

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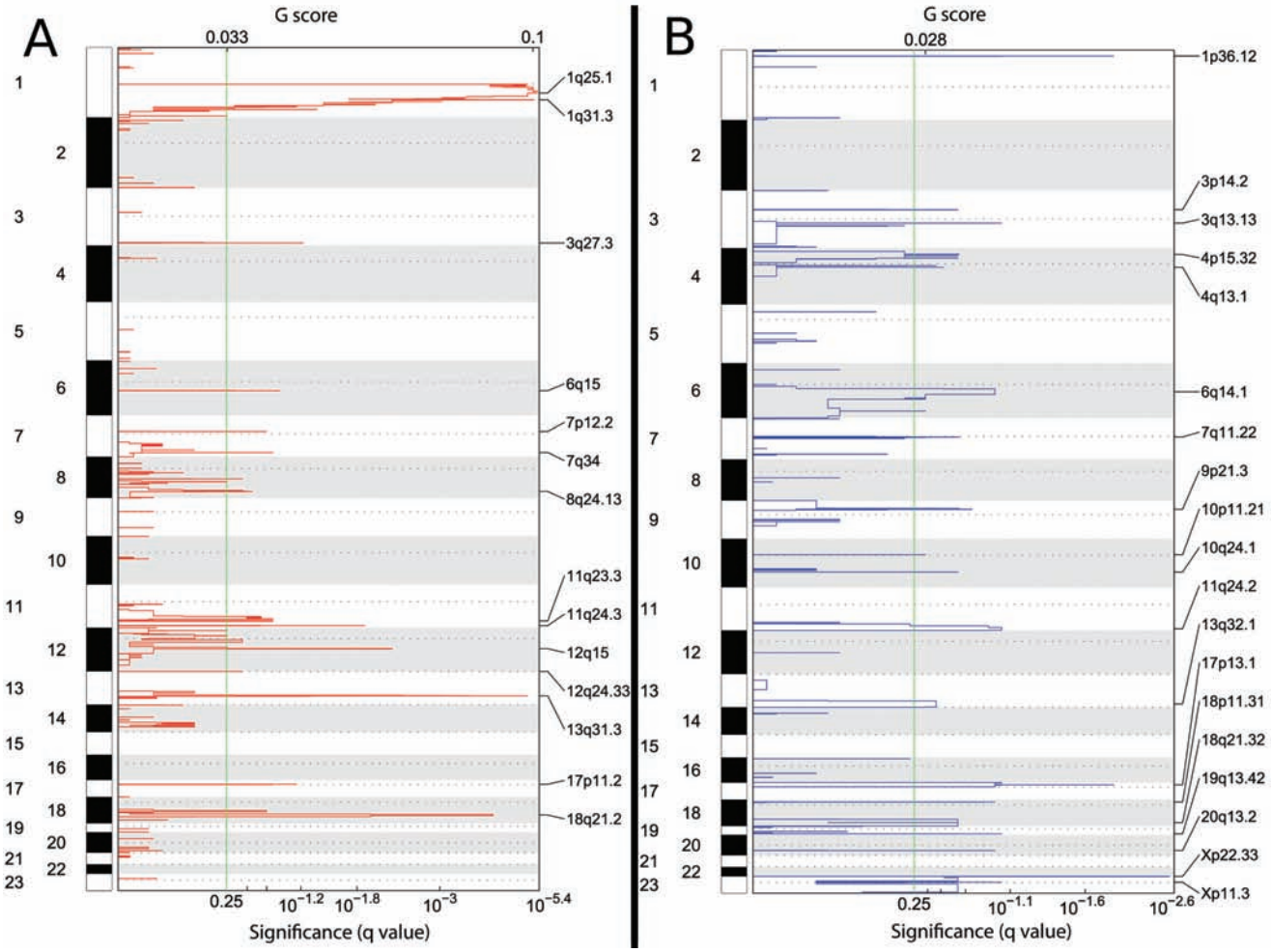
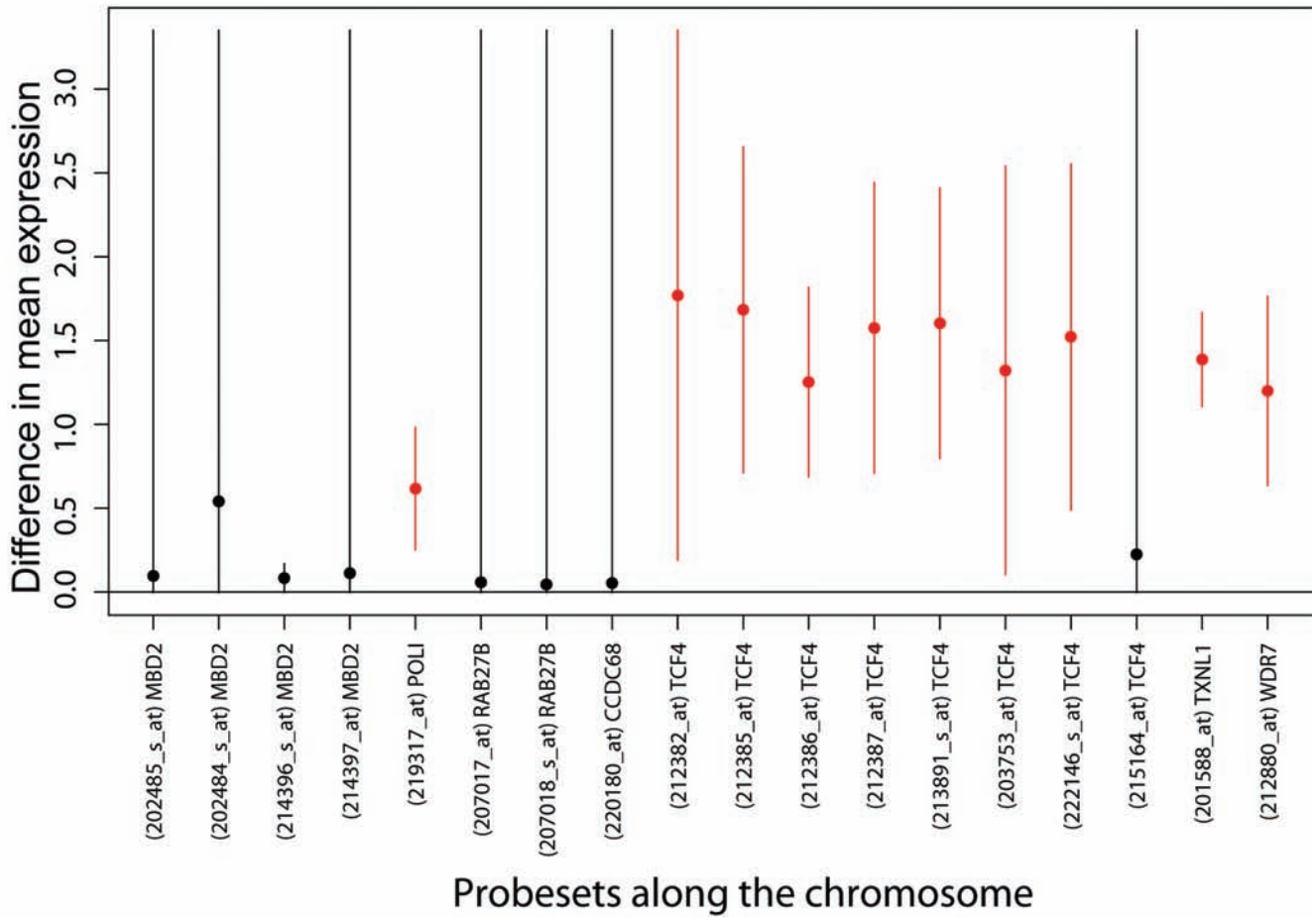


Online Supplementary Figure S1. Recurrence of gains (A) and losses (B) as given by GISTIC. y-axis: genomic position with chromosomes marked; G-score: significance score provided by the GISTIC algorithm; q-value: statistical significance, corrected for multiple testing. Cytobands are displayed on the right margin to better localize spikes of significance. When comparing the frequencies with those in Figure 1, it is important to note that the GISTIC algorithm also considers the extent of copy number changes.



Online Supplementary Figure S2. Differential gene expression for the gained region 18q21.2. Simultaneous, two-sided 95% bootstrap confidence intervals for the differences in expression between cases with and without gain on 18q21.2. Some of the intervals were truncated to fit the picture. The estimates for the differences in expression are denoted by bold dots. Their positive values reflect increased expression in the cases showing a gain. Significant probe-sets are depicted in red. One of them is *POLI*.

Online Supplementary Table S1. High-level amplifications.

Array	Copy number	Chromosome	Start base	Stop base	Start SNP	Stop SNP	Start base_extended	Stop base_extended	Molecular diagnosis	GCB.ABC	Length	Percent_CNP
MPI-078	≥5	11	117026813	118271578	157866	158053	117008604	118321366	mBL	GCB	267	0.00
MPI-078	≥5	12	40211576	46865076	163705	164296	40185579	46868499	mBL	GCB	1263	14.01
MPI-078	≥5	14	76103995	78099690	184753	185035	76056803	78134215	mBL	GCB	466	31.76
MPI-078	≥5	14	80108557	81754501	185199	185297	80033917	81769270	mBL	GCB	265	9.43
MPI-078	≥5	14	86628568	89463608	185629	185886	86596899	89467100	mBL	GCB	534	7.87
MPI-095	≥5	10	1030138	1106294	132940	132945	1001958	1115769	mBL	GCB	9	0.00
MPI-095	≥5	12	67281049	68369481	165990	166104	67278530	68393146	mBL	GCB	215	0.00
MPI-095	≥5	13	94349071	96913193	178097	178282	94338710	96975940	mBL	GCB	449	0.67
MPI-018	≥5	1	149777411	151724330	21707	22025	149770061	151758544	mBL	GCB	319	7.52
MPI-018	≥5	1	155175515	156116877	22517	22742	155151979	156150021	mBL	GCB	226	4.42
MPI-018	≥5	1	156313411	180997834	22774	27939	156301926	181026453	mBL	GCB	5166	20.81
MPI-018	≥5	10	61560943	62676104	300479	300722	61547712	62700755	mBL	GCB	244	0.00
MPI-018	≥5	10	63258917	63653400	300843	300911	63250873	63655389	mBL	GCB	69	0.00
MPI-018	≥5	10	64475241	64888399	301128	301180	64459413	64904559	mBL	GCB	53	15.09
MPI-018	≥5	17	19109505	19426327	434721	434758	18857962	19455348	mBL	GCB	38	28.95
MPI-018	≥5	17	19504358	20161920	434767	434851	19500060	20403550	mBL	GCB	85	3.53
MPI-018	≥5	21	26741469	28129476	479307	479671	26736201	28137133	mBL	GCB	365	18.08
MPI-018	≥5	21	28390206	29161249	479710	479917	28375852	29161798	mBL	GCB	208	1.44
MPI-048	≥5	13	90776963	90832063	381231	381243	90766136	90843580	mBL	GCB	13	0.00
MPI-048	≥5	18	49306894	55355975	452186	453503	49301604	55369820	mBL	GCB	1318	11.00
MPI-071	≥5	18	46033349	46192632	451420	451456	46018071	46199252	mBL	GCB	37	16.22
MPI-071	≥5	18	49019835	55250645	452135	453451	49015687	55260238	mBL	GCB	1317	11.01
MPI-089	≥5	1	238329580	238336668	39139	39144	238321944	238338426	mBL	GCB	6	0.00
MPI-171	≥5	1	166223758	166317208	25130	25180	166220861	166345649	mBL	GCB	51	0.00
MPI-171	≥5	1	171855554	171859428	26157	26163	171851736	171863619	mBL	GCB	7	0.00
MPI-532	≥5	11	65110065	67367795	328207	328378	65015493	67383143	mBL	GCB	172	37.79
MPI-532	≥5	11	71403860	73401270	328958	329147	71399055	73405476	mBL	GCB	190	4.74
MPI-532	≥5	11	106685152	108051787	336044	336264	106658439	108075704	mBL	GCB	221	45.25
MPI-532	≥5	11	109944451	111421334	336579	336893	109909301	111443258	mBL	GCB	315	0.00
MPI-532	≥5	11	112740629	113348687	337226	337350	112720334	113364827	mBL	GCB	125	0.00
MPI-532	≥5	18	41463382	44010072	450484	450980	41456402	44041120	mBL	GCB	497	5.23
MPI-532	≥5	18	50013288	53161776	452341	452949	49999395	53164072	mBL	GCB	609	10.18

Online Supplementary Table S2. Homozygous deletions.

Array	Copy number	Chromosome	Start base	Stop base	Start SNP	Stop SNP	Start base_ extended	Stop base_ extended	Molecular diagnosis	GCB.ABC	Length	Percent_ CNP
MPI-006	0	7	38096725	38158081	100968	100979	38011359	38176607	mBL	GCB	15	100.00
MPI-006	0	9	20018452	21850428	124293	124450	19990277	21880326	mBL	GCB	412	11.65
MPI-006	0	9	21999960	22389693	124462	124496	21990770	22402948	mBL	GCB	83	0.00
MPI-006	0	14	105746660	106109648	187524	187534	105553453	106142730	mBL	GCB	21	100.00
MPI-016	0	22	21388199	21546762	230290	230295	21276793	21576156	mBL	GCB	8	100.00
MPI-095	0	22	21163978	21546762	230280	230295	21105150	21576156	mBL	GCB	25	100.00
MPI-001	0	22	21405412	21546762	484677	484681	21389135	21576156	mBL	GCB	5	100.00
MPI-003	0	14	105149735	105170875	402004	402008	105071276	105238706	mBL	GCB	5	100.00
MPI-018	0	10	98422661	98547438	307777	307795	98407920	98587392	mBL	GCB	19	0.00
MPI-018	0	17	51518046	51522852	439078	439086	51512358	51532981	mBL	GCB	9	100.00
MPI-018	0	22	21463635	21580110	484678	484683	21405412	21582785	mBL	GCB	6	100.00
MPI-032	0	1	23476031	23690889	3701	3732	23457834	23714049	mBL	GCB	32	0.00
MPI-048	0	22	20711383	21546762	484566	484681	20711025	21576156	mBL	GCB	116	100.00
MPI-048	0	X	638171	851166	490077	490108	636217	890087	mBL	GCB	32	100.00
MPI-071	0	14	105547042	105835970	402012	402019	105399872	105988336	mBL	GCB	8	100.00
MPI-081	0	8	12264620	12286403	240783	240788	12040126	12446597	mBL	GCB	6	100.00
MPI-082	0	22	21163978	21546762	484657	484681	21105150	21576156	mBL	GCB	25	100.00
MPI-082	0	X	47033985	47142737	494155	494163	46975369	47150006	mBL	GCB	9	0.00
MPI-085	0	14	105547042	105835970	402012	402019	105399872	105988336	mBL	GCB	8	100.00
MPI-101	0	6	29979615	30015940	186477	186487	29957598	30023128	mBL	GCB	11	100.00
MPI-101	0	9	43847424	44108554	273923	273927	43618057	64197638	mBL	GCB	5	100.00
MPI-114	0	6	79036117	79083405	194803	194807	79019345	79109332	mBL	GCB	5	100.00
MPI-114	0	X	22730841	22745968	492173	492178	22718153	22793958	mBL	GCB	6	100.00
MPI-171	0	3	75535509	75627862	96855	96861	75476955	75647561	mBL	GCB	7	100.00
MPI-528	0	18	62058576	62062341	454841	454846	62053230	62063122	mBL	GCB	6	100.00
MPI-528	0	22	21031117	21546762	484646	484681	21020620	21576156	mBL	GCB	36	100.00
MPI-529	0	1	23538204	23657473	3719	3726	23535004	23673235	mBL	GCB	8	0.00
MPI-529	0	22	21105150	21546762	484656	484681	21083648	21576156	mBL	GCB	26	100.00
MPI-532	0	6	29984938	30015940	186479	186487	29980936	30023128	mBL	GCB	9	100.00
MPI-532	0	22	21083648	21546762	484655	484681	21058712	21576156	mBL	GCB	27	100.00

Online Supplementary Table S3. SEE EXCEL FILE

Online Supplementary Table S4. SEE EXCEL FILE

Online Supplementary Table S5. Gene dosage for the miRNA supercluster 17-92 on chromosome 13q31.3.

miRNA	Mean expression with gain (A)	Mean expression (B) with normal copy number	ΔCT: columns A-B	*P* value (Mann-Whitney-U Test)	Adjusted *P* value (Bonferroni)
17-3p	30.67	31.71	-1.05	0.13	1
17-5p	27.90	29	-1.10	0.049	0.39
18a-probe 1	30.79	30.96	-0.178	0.92	1
18a-probe 2	29.75	30.29	-0.544	0.63	1
19a	27.31	28.44	-1.12	0.38	1
19b	24.73	26.15	-1.41	0.085	0.68
20a	24.53	25.95	-1.42	0.049	0.39
92	25.00	25.44	-0.437	0.63	1

Expression values are normalised CTvalues (see the Design and Methods section), thus a lower value means higher expression in the cases with gains.