

**Manuscript no. HAEMATOL2010/026831 entitled "Detection of genomic aberrations in molecularly defined Burkitt lymphoma by array-based high resolution single nucleotide polymorphism analysis"**

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**Information about the contributions of each person named as having participated in the study**

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2) Authors who participated in the **conception of the study**: René Scholtysik, Markus Kreuz, Lorenz Trümper, Reiner Siebert, Ralf Küppers

3) **Design & Methods**. The following authors were responsible for specific investigations:

- René Scholtysik was responsible for generating and evaluating SNP chip data and integrating the various data sets.
- Markus Kreuz, Markus Loeffler, Maciej Rosolowski and Rainer Spang were responsible for the bioinformatic evaluation of the data
- Wolfram Klapper, Harald Stein and Alfred C. Feller were responsible for the pathological evaluation of the cases
- Birgit Burkhardt was responsible for diagnosing paediatric Burkitt's lymphoma cases and providing their clinical data
- Alfred C. Feller and Christoph Thorns were responsible for generating the miRNA expression data
- Michael Hummel and Harald Stein were responsible for generating the gene expression data
- Lorenz Trümper was responsible for providing clinical data and the DNA and RNA of the samples.
- Inga Vater and Reiner Siebert were responsible for generating molecular cytogenetic data
- Swen Wessendorf and Carsten Schwaenen were responsible for generating comparative genomic hybridization data
- Thorsten Zenz was responsible for generating TP53 mutation data
- Ralf Küppers was responsible for supervision of the project and integrating and evaluating of the various data sets.

4) **Results.** The following authors were responsible for specific portions of the results, including figures and tables:

- René Scholtysik was responsible for generating and evaluating SNP chip data and integrating the various data sets (incl. all Tables).
- Alfred C. Feller and Christoph Thorns were responsible for generating the miRNA expression data (Supplementary Figure S5).
- Michael Hummel and Harald Stein were responsible for generating the gene expression data (Figure 2)
- Wolfram Klapper, Harald Stein and Alfred C. Feller were responsible for the pathological evaluation of the cases.
- Markus Kreuz, Markus Loeffler, Maciej Rosolowski and Rainer Spang were responsible for the bioinformatic evaluation of the data (all figures and Tables).
- Swen Wessendorf and Carsten Schwaenen were responsible for generating comparative genomic hybridization data.
- Thorsten Zenz was responsible for generating TP53 mutation data.
- Inga Vater and Reiner Siebert were responsible for generating molecular cytogenetic data.
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- Lorenz Trümper was responsible for providing clinical data and the DNA and RNA of the samples.
- Ralf Küppers was responsible for supervision of the project and integrating and evaluating of the various data sets.

5) **Writing the manuscript.** The following authors were responsible for writing the manuscript: René Scholtysik, Markus Kreuz, Reiner Siebert and Ralf Küppers were responsible for writing of the manuscript.

6) **Contributors Listed in Acknowledgments:** We thank Ludger Klein-Hitpaß for generating the SNP-chip profiles and Claudia Becher for support with the FISH analyses. This work was supported by the Deutsche Krebshilfe through grant 107532.