

AUTHOR CONTRIBUTION FORM

Title	The analysis of clonal diversity and therapy responses using STAT3 mutations as a molecular marker in large granular lymphocytic leukemia
First author	Satu Mustjoki

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Responsible author (author responsible for the integrity of the work as a whole)

Name: Satu Mustjoki

Institute: Hematology Research Unit Helsinki, Department of Hematology, University of Helsinki and Helsinki University Central Hospital Cancer Center, Helsinki, Finland

e-mail: satu.mustjoki@helsinki.fi

Author contributions

Please describe the contributions of each author, indicating who was responsible for each part of the study and the preparation of the manuscript (collection of data, experiments, data analysis, generation of figures, interpretation of data, preparation of the text, etc.):

H.L.M.R designed the study, coordinated the project, performed sequence analysis, did laboratory studies, analyzed the data and wrote the paper. S.M. designed the study, coordinated the project, analyzed the data and wrote the paper. T.O., M.J.C., and A.J. provided patient samples and patient data, participated in the laboratory studies and contributed to write the paper. S.L., P.E., T.L., D.E.H, and E.I.A. performed sequence analysis. S.A.U.Z. and J.M.L.M designed and performed bioinformatics analysis. T.P.L, J.P.M., and K.P. participated in the study design, data analysis and contributed to write the paper. All authors read and approved the final manuscript.