Manuscript no. HAEMATOL/2010/021865 entitled "MICA polymorphism identified by whole genome array associated with NKG2D-mediated cytotoxicity in T-cell large granular lymphocyte leukemia"

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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: Viny, Clemente, Maciejewski

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

   • Viny was responsible for acquisition of biologic tissue, handling of genomic material, analysis of array data, confirmatory studies for SNP identification
   • Clemente was responsible for development of cytotoxic killing assay and for flow cytometric studies
   • Ishwaran was responsible for construction of statistical model for SNP identification through Random Forests analysis and modification for this application
   • Askar and Zhang were responsible for Methodology of allelic discrimination SSOP for MICA
   • Maciejewski was responsible for mentroship and guidance from the inception of this project

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

   • Viny is responsible for array data, PCR data, analysis of allele data, and all figures and tables
   • Clemente is responsible for flow cytometry data including killing assay
   • Jasek is responsible for sequencing data
   • Askar and Zhang are responsible for SSOP MIC allele data
   • Ishwaran is responsible for statistical analysis of SNP array data
   • Nowacki is responsible for statistical analysis of killing assay data
   • Maciejewski is responsible for oversight of all results

5) Writing the manuscript. The following authors were responsible for writing the manuscript:

   • Viny was responsible for writing of this manuscript in its entirety
   • All other authors provided critical review
6) Contributors Listed in Acknowledgments:

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