

AUTHOR CONTRIBUTION FORM

Title	Transcriptome analysis reveals molecular profiles associated with evolving steps of monoclonal gammopathies
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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.):

L.L.C. and N.C.G. were responsible for the design of the study, supervised all genomic analysis and prepared the initial version of the paper. E.F. carried out the microarray hibridization. L.A.C and F.S.B accomplished the bioinformatic analysis. M.E.S. performed the qRT-PCR analysis. J.M.H.R., J.J.L, J.B., A.O., M.L.M., J.H and A.I.T. are clinicians who contributed to sample collection. R.G.S. participated in the design of the study and supervised the statistical analysis. M.V.M, and J.F.S.M gave the final approval of the version to be published.