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- 3) Design & Methods. The following authors were responsible for specific investigations (please detail):
- Cristina Gattazzo and Renato Zambello were responsible for the study design;
- Cristina Gattazzo, Mirna Balsamo, Elisa Scquizzagto, Antonella Teramo and Elena Vendrame were responsible for blood samples processing;
- Anna Cabrelle was responsible for flow cytometry assay;
- Cristina Gattazzo was responsible for DNA and RNA purification, TOPO TA cloning, DNA methylation assay;
- Elisa Scquizzato, Antonella Teramo and Elena Vendrame were responsible for PCR investigation (semi-quantitative and RT-PCR);
- Marta Miorin was responsible of plasmid sequencing.
- 4) **Results**. The following authors were responsible for specific portions of the results, including figures and tables (please indicate the person responsible for each figure and each table):
- Cristina Gattazzo, Marta Miorin were responsible for methylation assay, KIR sequencing, data collection, analysis of data and Figure 2, 3 and 4;

- Antonella Teramo, Elisa Scquizzato and Elena Vendrame were responsible for KIR genotyping analysis, KIR and DNMTs expression analysis, Figure 1 and Table 2;
- Anna Cabrelle was responsible for flow cytometry analysis and cell sorting;
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- Renato Zambello and Gianpietro Semenzato were responsible of data analysis.
- 5) Writing the manuscript. The following authors were responsible for writing the manuscript:
- Cristina Gattazzo and Renato Zambello were responsible for preparation and writing the manuscript;
- Gianpietro Semenzato was responsible for reviewing the manuscript.
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