Massive parallel sequencing unveils homologous recombination deficiency in follicular dendritic cell sarcoma

Luisa Lorenzi,[1] Torsten Haferlach,[2] Luigi Mori,[3] Matteo Simbeni,[1] Wencke Walter,[2] Piera Balzarini,[1] Manja Meggendorfer,[2] Claudia Döring,[4] Silvia Lonardi,[1] Mattia Bugatti,[1] Claudio Agostinelli,[5] Jay Mehta,[6] Anita Borges,[7] Abbas Agaimy,[8] Ingrid Simonitsch-Klupp,[9] José Cabeçadas,[10] Elias Campo,[11] Stefano Aldo Pileri,[12] Fabio Facchetti[1] Martin Leo Hansmann[13] and Sylvia Hartmann[4]

[1]Pathology Unit, ASST Spedali Civili di Brescia, Department of Molecular and Translational Medicine, University of Brescia, Brescia, Italy; [2]MLL Munich Leukemia Laboratory, Munich, Germany; [3]Laboratory of Molecular Medicine, Department of Clinical and Experimental Science, University of Brescia, Brescia, Italy; [4]Dr Senckenberg Institute of Pathology, Goethe University, Frankfurt, Germany; [5]Department of Medical and Surgical Sciences, University of Bologna, Bologna, Italy; [6]Neuberg Oncopath, Mumbai, India; [7]Histopathology, SRL Diagnostics, Mumbai, India; [8]Institute of Pathology, University Hospital, Erlangen, Germany; [9]Institute of Pathology, Medical University of Vienna, Vienna, Austria; [10]Department of Pathology, Portuguese Institute of Oncology, Lisbon, Portugal; [11]Hematopathology Section, Hospital Clinic, IDIBAPS, University of Barcelona, Barcelona, Spain; [12]Division of Hematopathology, European Institute of Oncology (IEO) IRCCS, Milan, Italy and [13]Institute for General Pharmacology and Toxicology, Goethe University, Frankfurt, Germany

Correspondence: L. Lorenzi
luisa.lorenzi@unibs.it

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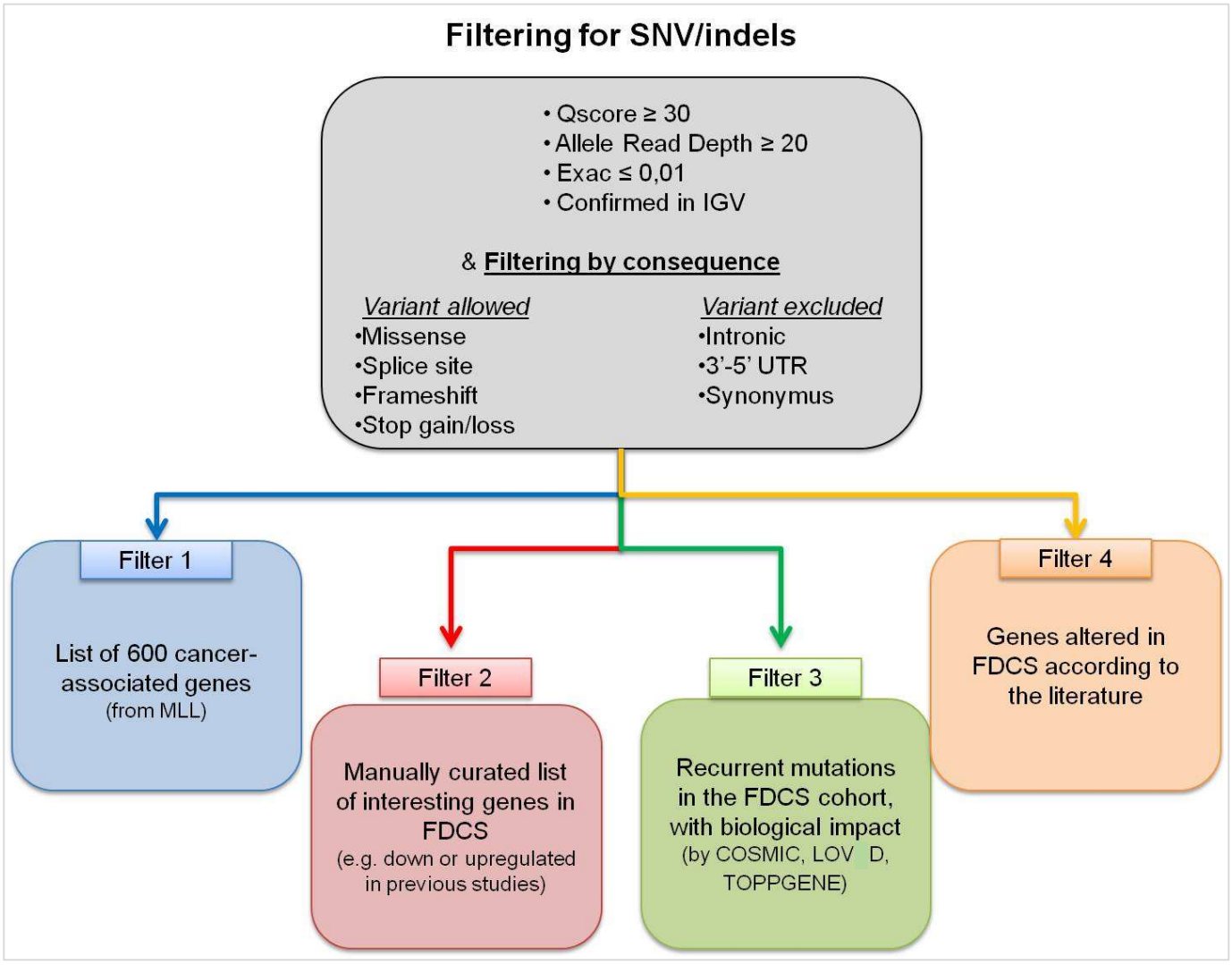
Supplementary Figure 1: Filters applied to prioritize data analysis and identify recurrent single nucleotide variations (SNV) in Follicular dendritic cell sarcoma identified by whole genome and whole exome sequencing.



Supplementary Table 1 (excel file): List of mutational variants detected by Whole Genome and While Exome sequencing in all the 13 samples of FDCS.

Supplementary Table 2 (excel file): List of Structural Variations (SV) and Copy number Variations (CNV) detected by Whole Genome Sequencing in case #13.