

Figure 1. Identified fusion transcripts. (A) Histogram of identified transcripts, assigned to fusion groups. On top of each bars the percentage of all cases. *PICALM-MLLT10* (4%), *SET-NUP214* (3.4%) and *NUP214-ABL1* (3.4%) are the most recurrent. (B) Circos plot representing the identified fusion transcripts. (C) Schematic representation of *MLLT10* fused polypeptide chains with relevant domains. OM-LZ and nuclear addressing signals are always conserved. (D) Histograms of incidence of cases by age classes and fusion groups (absolute count on y axis). ETS fusion group is constituted with only pediatric cases. Abbreviations: ENTH, epsin N-terminal homology; NES, nuclear export domain; PZP, pregnancy zone protein.

