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Targeted next generation sequencing reveals high mutation frequency of CREBBP, BCL2 and KMT2D in high-grade B-cell lymphoma with MYC and BCL2 and /or BCL6 rearrangements

Solène M Evrard, Sarah Péricart, David Grand, Nadia Amara, Frédéric Escuidé, Julia Gilhodes, Pierre Bories, Alexandra Traverse-Glehen, Romain Dubois, Pierre Brousset, Marie Parrens, and Camille Laurent

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Contributions: S.M.E. and F.E. investigated molecular mutations; A.T.G, R.D, M.P. and C.L provided samples; S.M.E. , D.G. and C.L. conceived and designed the experiments; S.M.E, S.P., D.G., N.A., F.E., P.Bo. and C.L. analysed the data; S.M.E and J.G. performed statistical analysis; S.M.E., P.Br. and C.L contributed to the writing and the critical review of the manuscript.