Bromodomain and extra-terminal domain inhibition modulates the expression of pathologically relevant microRNAs in diffuse large B-cell lymphoma

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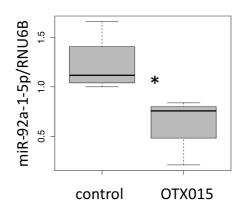
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Supplementary Data



Supplementary Figure 1

OTX015 modulates miRNA expression in an *in vivo* DLBCL model. miR-92a-1-5p expression was assessed in SU-DHL-2 xenografts from NOD-SCID mice previously treated with vehicle (control) or OTX015 for three days ¹. The thick black line in each boxplot denotes the median expression. Expression of RNU6B was used for normalisation.

Supplementary Tables Legends

Supplementary Table S1. Canonical pathways and hallmarks associated with OTX015-modulated miRNAs.

Supplementary Table S2. miRNAs with at least one BRD4-binding event within their regulatory regions and miRNAs with decreased BRD4-binding after exposure to JQ1, as determined via data mining of the ChIP-Seq datasets obtained in the ABC-DLBCL cell line HBL-1 (SRP043524) ² and in the GCB-DLBCL cell line OCI-LY-1 (SRP022129) ³.

Supplementary Table S3. Names and histological derivation of cell lines used for baseline miRNA profiling. Cell lines have already been used and described in previous studies ^{4, 5}.

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