



Extended Data Figure 9 | Validation of ependymoma transcription factors. **a, b,** Gene expression of ‘high activity’ transcription factors (ranked <50) (**a**) and ‘low activity’ transcription factors (ranked >50) (**b**) in ependymoma ($n = 83$ independent samples) versus normal brain tissue ($n = 172$ independent samples). Box plots showing median value

(horizontal bar), interquartile range and dotted line representing the data range. Comparison between groups was assessed using a two-sided Wilcoxon rank-sum test. **c,** Constituent enhancer activity in the central nervous system (CNS) of developing zebrafish embryos derived from subgroup-specific super enhancers identified in ependymomas.