



## Correction to: Risk-adapted therapy and biological heterogeneity in pineoblastoma: integrated clinico-pathological analysis from the prospective, multi-center SJMB03 and SJYC07 trials

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The original version of this article unfortunately contained a typesetting error in Fig 3c. The corrected Fig. 3 is given in the following page.

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The original article can be found online at <https://doi.org/10.1007/s00401-019-02106-9>.

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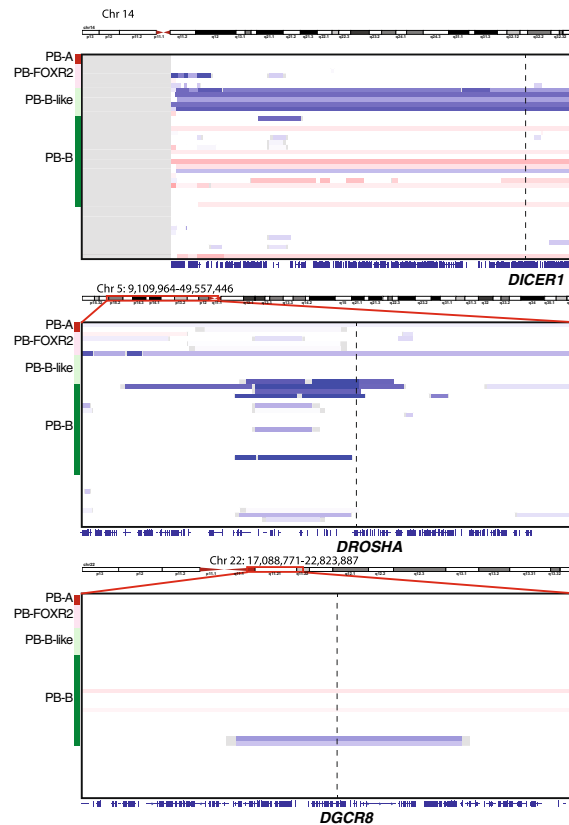
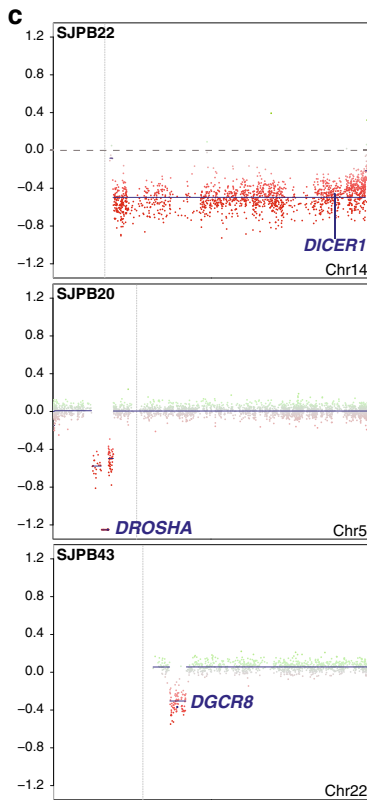
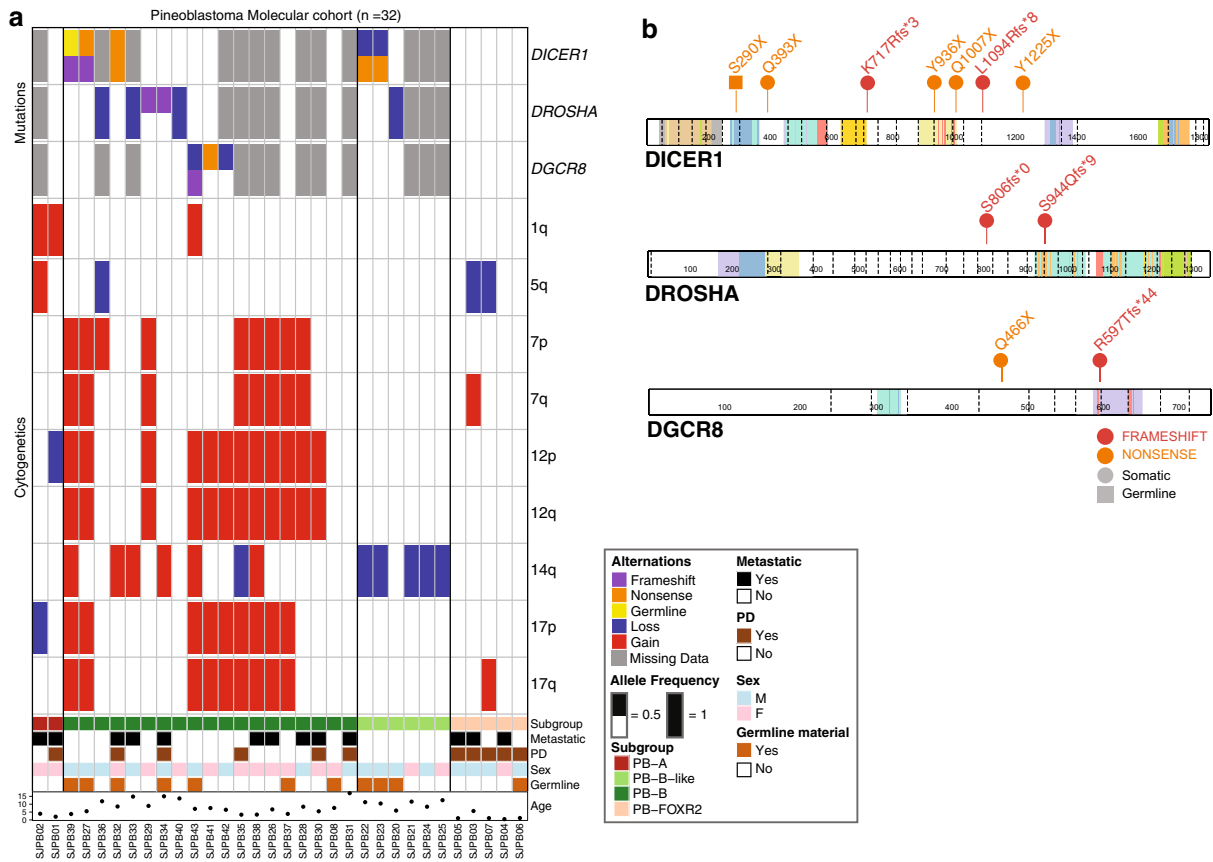
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**Fig. 3** Genomic alterations in molecularly defined pineoblastomas. **a** Oncoprint depicting loss-of-function mutations and/or allelic deletions in miRNA pathway genes *DICER1*, *DROSHA*, and *DGCR8* in addition to subgroup-specific chromosomal arm-level copy-number alterations, patient demographics, and outcome. **b** Lollipop plots depicting deleterious somatic and germline mutations in miRNA pathway genes *DICER1*, *DROSHA*, and *DGCR8*. chr, chromosome; fs, frameshift; X, stop codon. **c** Three exemplary copy-number plots highlighting 14q loss involving the *DICER1* locus, focal 5p loss involving the *DROSHA* locus, and focal 22q loss involving the *DGCR8* locus. Corresponding chromosomal regions of all samples from the molecular cohort, showcasing the restriction of these cytogenetic events to PB-B and PB-B-like subgroups. *Chr* chromosome, *F* female, *M* male, *PD* progressive disease

The original article has been corrected.