

ERRATUM

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Erratum to: Extensive next-generation sequencing analysis in chronic lymphocytic leukemia at diagnosis: clinical and biological correlations

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Erratum

*n.b. The error described below was mistakenly carried forward by the production team handling this article, and thus was **not** the fault of the authors.*

The original version of this article [1] had a duplication of Fig. 1 in place of where Fig. 2 should have been, resulting in two displays of Fig. 1 and the absence of Fig. 2.

The article has now been updated to remove the duplicate of Fig. 1 and to insert the correct Fig. 2 into its appropriate place.

The two figures in question can be seen below.

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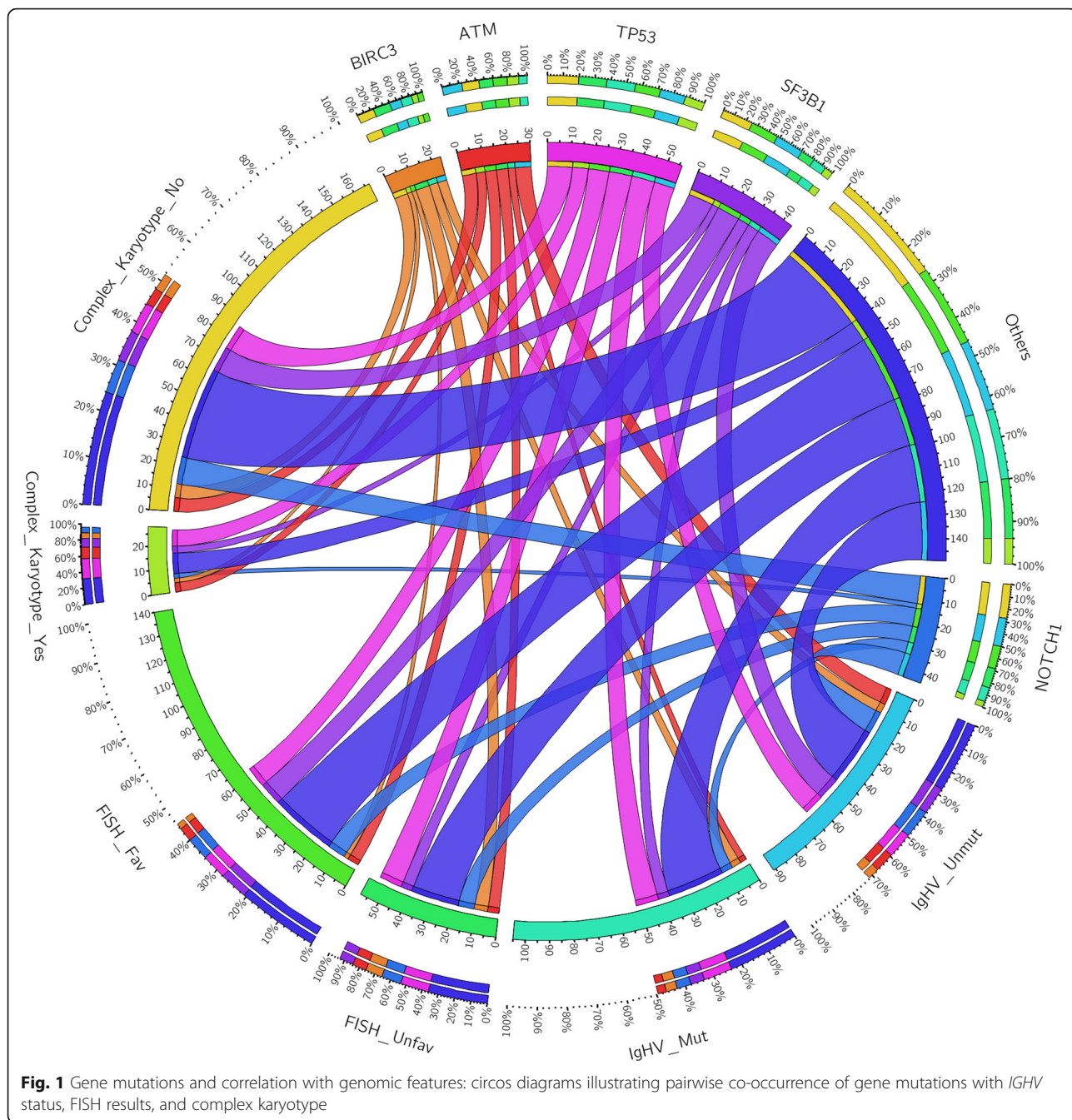
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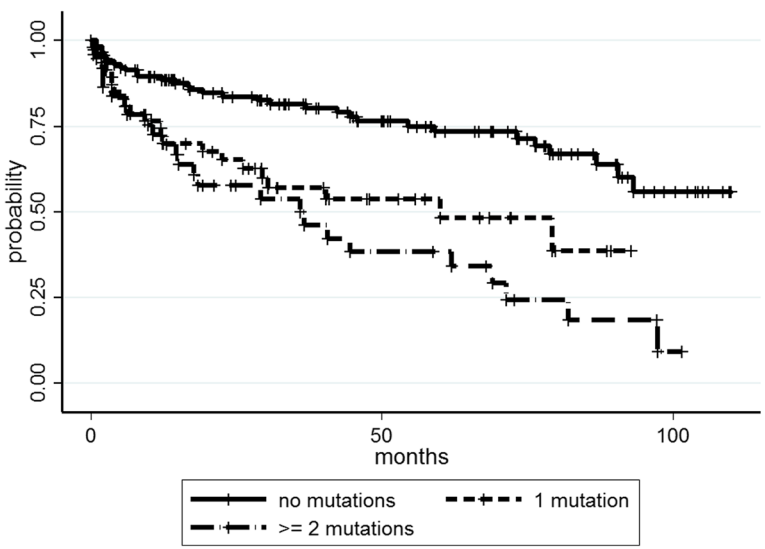


Fig. 2 TTFT according to number of mutations by NGS analysis ($p < 0.001$)