

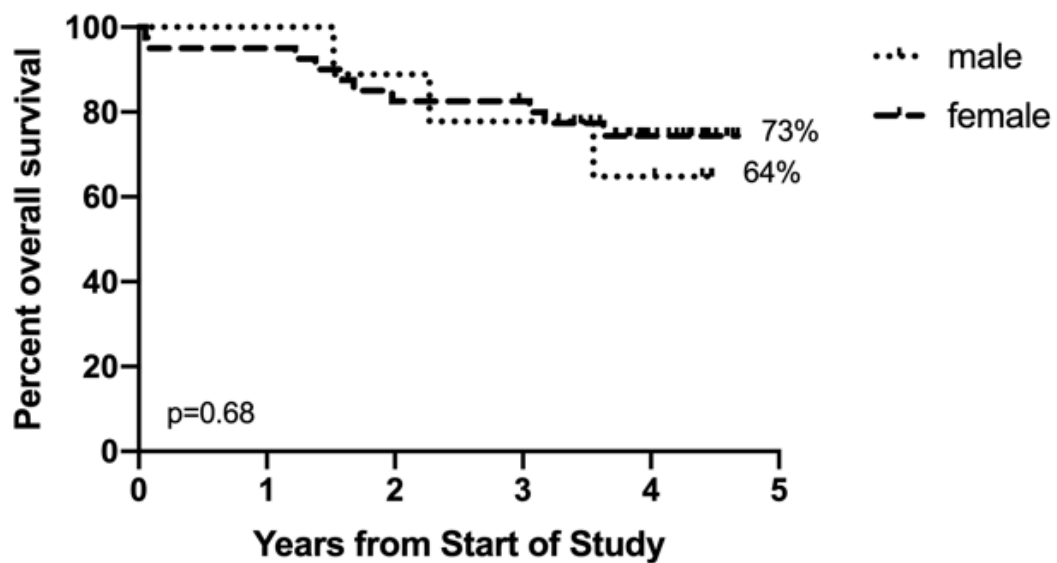
1 **Supplemental Material for Manuscript “Genome-wide DNA methylation analysis**
2 **pre- and post-lenalidomide treatment in patients with myelodysplastic syndrome**
3 **with isolated deletion (5q): results from a prospective multicenter study of the**
4 **German MDS study group”, Annals of Hematology**

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33 **Supplemental Figure 1:** Overall survival of the 51 patients depending on gender
34 estimated using the Kaplan-Meier method.



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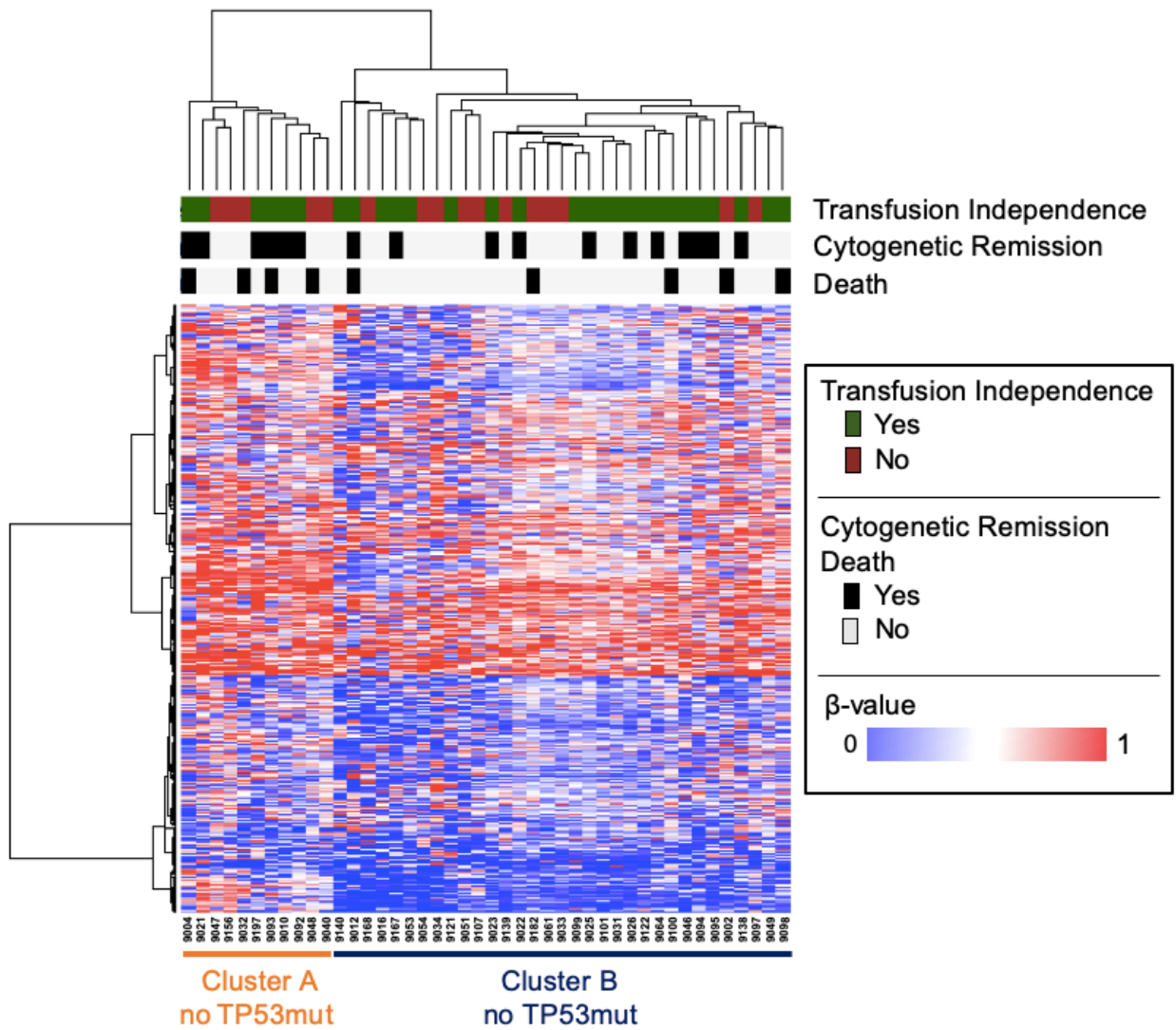
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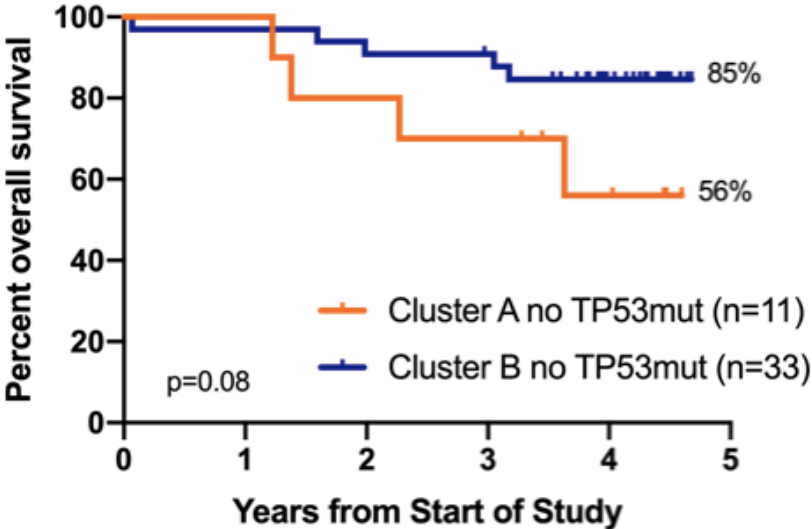
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51 **Supplemental Figure 2:** Unsupervised clustering of 44 patient bone marrow samples
 52 prior to lenalidomide treatment. Patients with mutations in TP53 were excluded from
 53 analysis. After exclusion of TP53mut patients, each of the TP53wt patients still clusters
 54 in the same way as before when analyzing the whole cohort. Patients are displayed on
 55 the X axis and the 1534 most variable CpG sites are displayed on the Y axis.
 56 Information on transfusion independence, cytogenetic remission and death is shown at
 57 the top of the figure.



59 **Supplemental Figure 3.** Overall survival of the 44 patients without TP53 mutations was
60 estimated using the Kaplan-Meier method. Patients are differentiated by Cluster A and
61 B (compare to Supplemental Figure 2).



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