

The improved prognosis of *FLT3*-internal tandem duplication but not tyrosine kinase domain mutations in acute myeloid leukemia in the era of targeted therapy: a real-world study using large-scale electronic health record data

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Supplementary Table 1: Coefficient of post-midostaurin x *FLT3* ITD interaction term (the DiD term), which reflects the relative change in prognosis of *FLT3* ITD mutations, after adjusting for different variables. Each model below adjusts for the “Baseline model,” which is shown in Table 2. In addition to the Baseline model, most regressions here adjust for another variable, such as patient performance status.

<i>Variables adjusted for in model</i>	<i>HR of DiD term (95% CI)</i>	<i>P-value of DiD term</i>
Baseline model	0.39 (0.22 – 0.68)	0.001
Baseline model + venetoclax	0.39 (0.20 – 0.77)	0.007
Baseline model + performance status	0.34 (0.16 – 0.69)	0.003
Baseline model + log(WBC + 1)	0.40 (0.23 – 0.70)	0.001
Baseline model + FLT3 inhibitor	0.62 (0.30 – 1.3)	0.19