

Supplemental Data Table 5. Detected mutation frequency in samples

| Sample | Median VAF, % (IQR) | | <i>P</i> |
|------------------|------------------------------------|-------------------------------------|----------|
| | Complete remission (BM blasts <5%) | Residual disease (BM blasts >5%) | |
| Peripheral blood | 0.000 (0.000–0.000), variant N=35 | 2.155 (0.000–4.310), variant N=2 | 0.401 |
| BM | 0.000 (0.000–0.159), variant N=141 | 41.524 (0.000–47.086), variant N=22 | <0.001 |
| Overall | 0.000 (0.000–0.127) | 39.731 (0.000–46.096) | <0.001 |

Abbreviations: BM, bone marrow; VAF, variant allele frequency; IQR, interquartile range.