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POSTER ABSTRACTS

637.MYELODYSPLASTIC SYNDROMES - CLINICAL AND EPIDEMIOLOGICAL

Molecular Measurable Residual Disease (MRD) Clearance (<1%) Is Associated with Improved Clinical Outcomes in Patients with Higher-Risk Myelodysplastic Neoplasms (HR-MDS): An Exploratory Analysis of Stimulus-MDS1 in Patients Receiving Sabatolimab or Placebo + Hypomethylating Agent (HMA)

Amer M. Zeidan, MBBS, MHS¹, Pierre Fenaux, MD PhD², Xia Han, MSc³, David A. James, MSc³, Kamel Malek, MDMPH⁴, Pedro Marques Ramos, PhD⁴, Yasushi Miyazaki, MDPhD⁵, Uwe Platzbecker, MD⁶

Background: Sabatolimab (MBG453) is a novel immunotherapy targeting TIM-3, an immuno-myeloid regulator expressed on both immune and leukemic stem cells. In the randomized, double-blind, placebo-controlled, Phase (Ph) II STIMULUS-MDS1 study (NCT03946670) in patients (pts) with HR-MDS, although improvements in complete remission (CR) and progression-free survival (PFS) were not statistically significant, sabatolimab + HMA was associated with longer duration of response (DoR), and the data suggest a delayed-onset benefit in the sabatolimab arm, as well as a potential treatment effect in pts with lower disease burden (Zeidan AM, et al. ASH 2022). The clinical significance of MRD in HR-MDS is not well established. We report an exploratory analysis of mutational clearance and MRD by next-generation sequencing (NGS) to evaluate the association with clinical outcomes.

Methods: Treatment-naive pts aged >18 years with intermediate (+ >5% bone marrow [BM] blasts), high or very high risk MDS by Revised International Prognostic Scoring System were randomized to sabatolimab or placebo added to azacitidine/decitabine (Zeidan AM, et al. ASH 2022). Peripheral blood (PBMC) and/or BM mononuclear cell (BMMC) samples were collected and NGS performed on genomic DNA extracted from baseline (BL) and on-treatment samples using a 38-gene NGSerror-corrected panel with sensitivity of 2% and 0.2% variant allelic frequency (VAF), respectively. Concordance and correlation between PBMC and BMMC were assessed. The prognostic value of clonal clearance at different VAF cut-offs (0.2%, 0.5% or 1%) was assessed. MRD-x status was defined as any mutation call above VAF cut-off x, excluding mutations in DNMT3A, TET2 and ASXL1 (DTA). Association between MRD status and clinical outcomes was analyzed regardless of treatment to increase sample and used best overall response, 6-month landmark analysis, and a time-dependent Cox model to rule out immortal bias.

Results: NGS was run on 332 BMMC samples from 112 pts and 439 PBMC samples from 123 pts (127 pts randomized). 112 pts had >1 qualifying BL mutation (NGS cohort) and 106 pts (sabatolimab + HMA, N=56; placebo + HMA, N=50) had >1 at follow-up (MRD cohort). BL characteristics were balanced between NGS/MRD cohorts and between treatments in the MRD cohort.

Overall agreement in mutation calling at variant level between BMMC and PBMC was >97% (8312/8566 total mutation calls), with >88% positive (817/926) and >98% negative agreement (7495/7640). Across reported mutated variants (N=817), there was good correlation (pearson cor=0.88) in VAF between BMMC and PBMC. When considering all paired samples, concordance of best MRD status using PBMC or BMMC results at the pt level (N=95) was: MRD-0.2, 97%; MRD-0.5, 89%; MRD-1, 90%. Given good PBMC and BMMC agreement, results were combined (higher VAF used if both) for outcomes analysis. In the MRD cohort, 11% (N=12), 18% (N=19) and 27% (N=29) of pts achieved MRD-0.2, MRD-0.5 and MRD-1 negativity, respectively. Most pts with MRD-0.2, -0.5 and -1 clearance had documented response of CR/marrow CR [mCR] (>80%) or CR/partial remission [PR]/hematologic improvement [HI] (>75%). Best overall response of MRD-1 negativity achieved anytime on treatment was associated with improved PFS and overall survival (OS) in the overall population and for responding pts (CR/mCR or CR/PR/HI). Similar results were obtained in a landmark analysis at 6 months (Fig 1). This was supported by a

¹ Yale School of Medicine and Yale Cancer Center, Yale University, New Haven, CT

²Department of Hematology, Université de Paris, Saint-Louis Hospital, Paris, France

³Novartis Pharmaceuticals Corporation, East Hanover, NJ

⁴Novartis Pharma AG, Basel, Switzerland

⁵Nagasaki University, Nagasaki, Japan

⁶Department of Hematology, University Hospital of Leipzig, Leipzig, Germany

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time-dependent Cox-model indicating that MRD-x negativity was associated with a lower hazard ratio for PFS and OS vs MRD-x positivity (Fig 2).

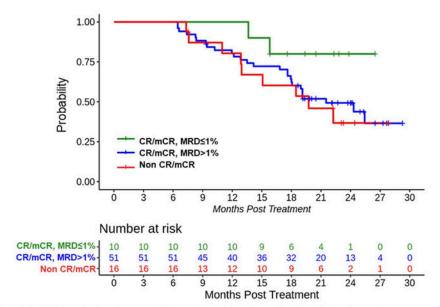
A consistently higher proportion of pts in the sabatolimab (N=56) vs placebo (N=50) arms had on-treatment mutation clearance at different cut-offs: MRD-0.2, 16.1% vs 6.0%; MRD-0.5, 25.0% vs 10.0%; MRD-1, 35.7% vs 18.0%, respectively.

Conclusions: We present the first results from a prospective, controlled, randomized study demonstrating the potential prognostic value of MRD for PFS and OS in HR-MDS. Our results demonstrate high NGS concordance between PBMC and BMMC on-treatment samples, indicating that PBMC could represent an alternative for clonal monitoring. Interestingly, more pts treated with sabatolimab + HMA reached MRD negativity while in remission, potentially explaining the longer DoR in pts receiving sabatolimab + HMA vs placebo + HMA. These findings will be further explored in the ongoing Ph III STIMULUS-MDS2 study (NCT04266301).

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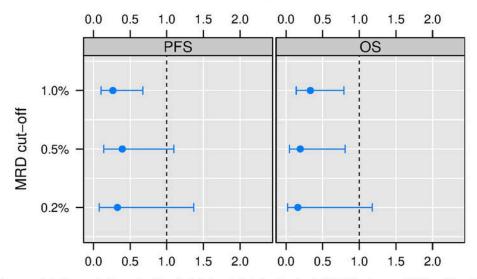
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Figure 1. Overall Survival by CR/mCR and MRD 1% Cut-Off Status at 6 Months



Kaplan-Meier plot of OS in patients who were MRD-positive at baseline with CR/mCR (best overall response of either CR or mCR as per investigator assessment according to modified IWG-MDS criteria) and MRD ≤1% at 6 months, CR/mCR and MRD >1% at 6 months, or without CR/mCR. MRD was calculated using highest VAF with combined BM and PB samples. BM, bone marrow; CR, complete remission; mCR, marrow CR; IWG, International Working Group; MDS, myelodysplastic syndromes; MRD, measurable residual disease; OS, overall survival; PB, peripheral blood; VAF, variant allele frequency.

Figure 2. Hazard Ratios of Clinical Endpoints by MRD Status at Different Cut-Offs



Time-varying covariate Cox model hazard ratios of clinical endpoints in all patients (N=112) who were MRD-positive at baseline at different MRD cut-offs. MRD was calculated at each visit using highest VAF with combined BM and PB samples. Bars represent 95% confidence intervals. BM, bone marrow; MRD, measurable residual disease; OS, overall survival; PB, peripheral blood; PFS, progression-free survival; VAF, variant allele frequency.

Figure 1

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