

# New IMS/IMWG Consensus Genomic Staging Risk Criteria by Next-Generation Sequencing: Analysis of Daratumumab Benefit in Both High- and Standard-Risk Patients in the PERSEUS/EMN017 Study

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## Key Takeaway

These data further support D-VRd induction/consolidation followed by D-R maintenance as standard of care for transplant-eligible patients with NDMM regardless of risk status

## Conclusions

D-VRd improved PFS versus VRd for both standard- and high-risk patients, as defined by PP-FISH or IMS/IMWG CGS criteria

In both standard- and high-risk patients according to IMS/IMWG CGS criteria, D-VRd increased the rates of MRD negativity ( $\geq$ CR;  $10^{-5}$  and  $10^{-6}$ ) and sustained MRD negativity lasting  $\geq$ 12 months ( $\geq$ CR;  $10^{-5}$  and  $10^{-6}$ ) and improved PFS compared with VRd

The UMA-NGS panel successfully stratified cytogenetic risk according to the new IMS/IMWG CGS criteria in the PERSEUS/EMN017 study, improving risk stratification

### Acknowledgments

We thank the patients who participated in this study, the staff members at the study sites, the data and safety monitoring committee, the staff members involved in data collection and analyses, the Rotterdam Myeloma Group from the Department of Hematology at Erasmus MC, and groups at IRCCS AOU di Bologna and University of Milan for UMA-NGS panel development. PERSEUS was sponsored by EMN in collaboration with Johnson & Johnson. The current analysis was a collaborative project between EMN and Johnson & Johnson. Medical writing support was provided by Carlee M Cunningham, PhD, of Eloquent Scientific Solutions, part of Envision Spark, an Envision Medical Communications agency, a part of Envision Pharma Group, and was funded by Johnson & Johnson. © 2026 American Society of Clinical Oncology, Inc. Reused with permission. This abstract was accepted and previously presented at the 2026 ASCO Annual Meeting. All rights reserved.

### Disclosure

MAD serves as a consultant for Amgen, BeiGene, Bristol Myers Squibb, Celgene Corporation, GSK, Janssen, Menarini Silicon Biosystems, Regeneron Pharmaceuticals, Sanofi, and Takeda.

### Introduction

- The International Myeloma Society/International Myeloma Working Group (IMS/IMWG) recently proposed Consensus Genomic Staging (CGS) risk criteria for newly diagnosed multiple myeloma (NDMM), including genetic alterations not identifiable by fluorescence in situ hybridization (FISH), by utilizing next-generation sequencing (NGS; eg, *TP53* mutations)<sup>1</sup>
- This expanded post hoc analysis of the PERSEUS/EMN017 study investigated clinical outcomes in patients based on the presence of high-risk disease, as defined by the new IMS/IMWG CGS risk criteria utilizing the recently developed NGS-based Unique Molecular Assay (UMA) target panel<sup>2</sup>

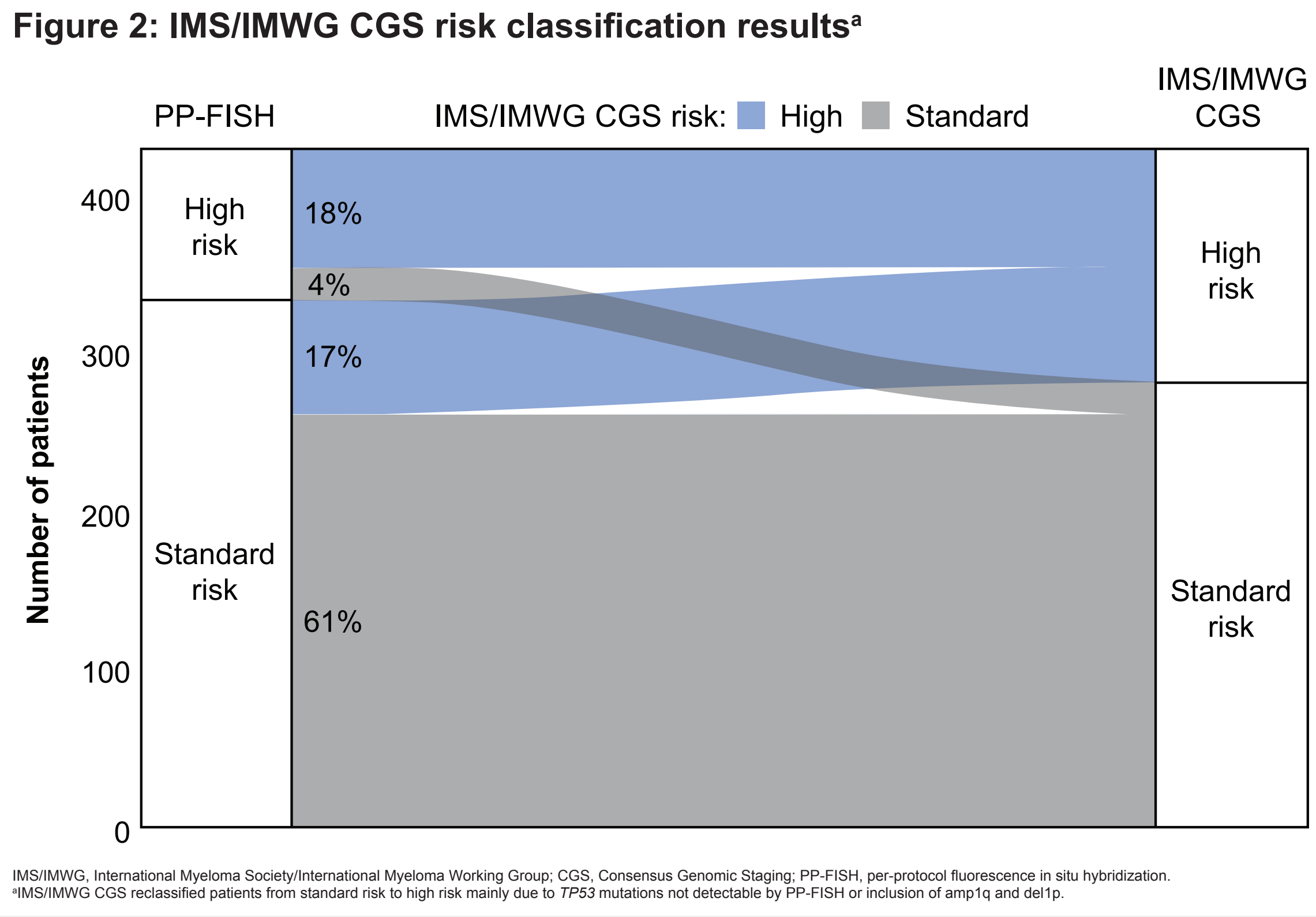
| Risk subgroup                       | Definition                                                                                                                                                                                                                                                                                                                                                                                                |
|-------------------------------------|-----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| PP-FISH high risk                   | Defined per protocol as $\geq$ 1 del17p, t(4;14), and/or t(14;16)                                                                                                                                                                                                                                                                                                                                         |
| IMS/IMWG CGS high risk <sup>1</sup> | Defined as the presence of $\geq$ 1 of the following: <ul style="list-style-type: none"><li>• del17p<sup>a</sup> and/or <i>TP53</i> mutation<sup>b</sup></li><li>• t(4;14), t(14;16), and/or t(14;20) with additional amp1q or biallelic del1p</li><li>• Monoallelic del1p with additional amp1q, or biallelic del1p<sup>c</sup></li><li>• High <math>\beta</math>2M with normal renal function</li></ul> |

PP-FISH, per-protocol fluorescence in situ hybridization; IMS/IMWG, International Myeloma Society/International Myeloma Working Group; CGS, Consensus Genomic Staging;  $\beta$ 2M,  $\beta$ 2 microglobulin; CCF, cancer cell fraction; NGS, next-generation sequencing.  
<sup>a</sup>CCF  $\geq$ 20%, by analyses conducted on CD138<sup>+</sup>/purified cells.  
<sup>b</sup>Assessed using an NGS-based method.

### Results

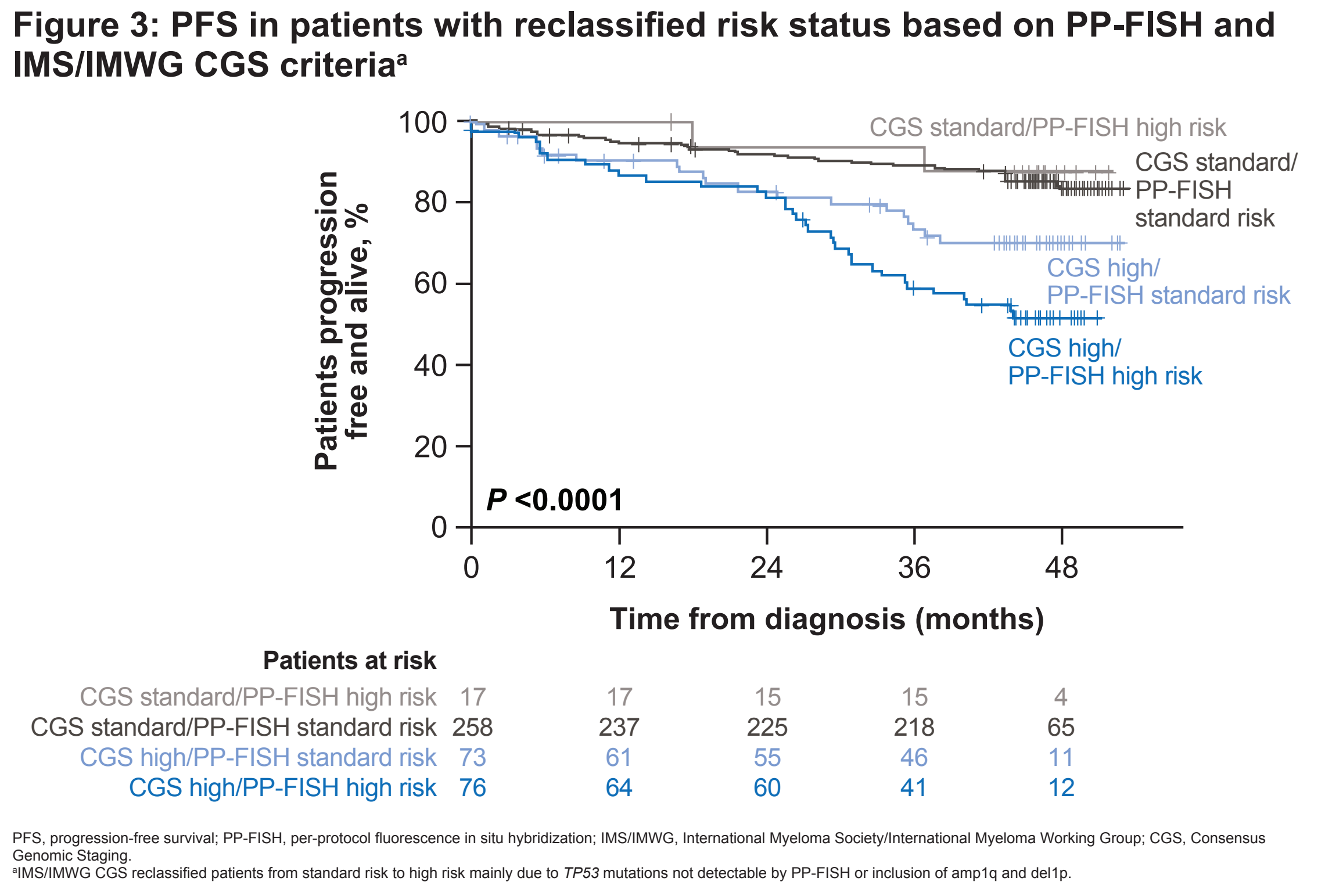
#### Risk classification

- The UMA-NGS panel successfully stratified risk according to IMS/IMWG CGS criteria, identifying  $\geq$ 12% more high-risk patients in the PERSEUS/EMN017 study versus those identified using per-protocol FISH (PP-FISH; **Figure 1**)
- The most common genetic lesions were del17p/*TP53* (14%), IgH translocation with amp1q/del1p (12%), and del1p with amp1q (9%); 2% of patients had biallelic focal del1p
- IMS/IMWG CGS reclassified 17% of patients from standard to high risk, with only 4% reclassified from high to standard risk with improved risk classification (**Figure 2**)



#### IMS/IMWG CGS: PFS in reclassified patients

- In patients reclassified from PP-FISH high risk to IMS/IMWG CGS standard risk, PFS outcomes did not differ from those observed in patients classified as standard risk by both methods (**Figure 3**)

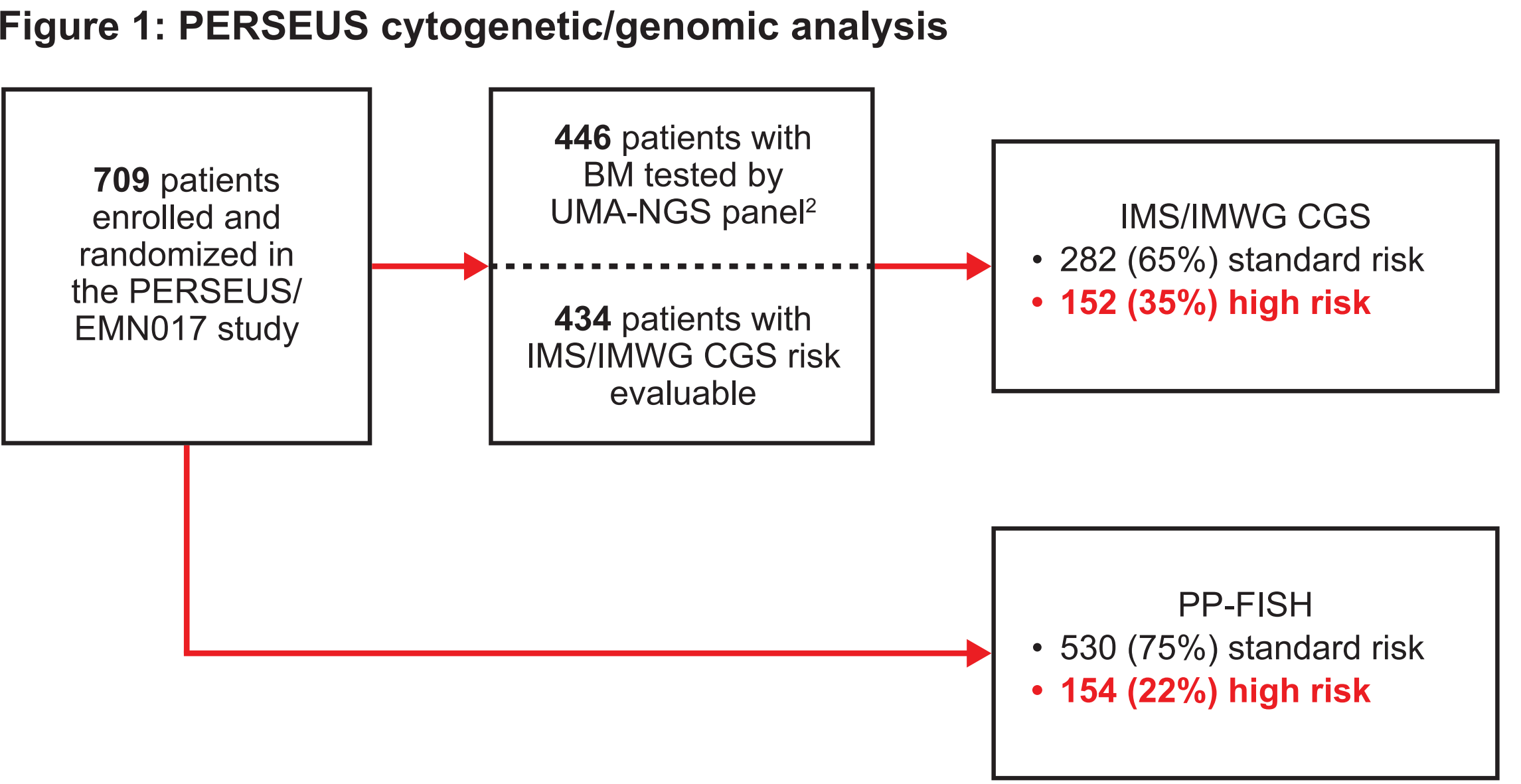


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### Methods

- In the phase 3 PERSEUS/EMN017 study (ClinicalTrials.gov Identifier: NCT03710603), 709 transplant-eligible patients with NDMM were randomized 1:1 to daratumumab plus bortezomib, lenalidomide, and dexamethasone (D-VRd) induction/consolidation followed by daratumumab and lenalidomide (D-R) maintenance or VRd induction/consolidation followed by lenalidomide maintenance<sup>3</sup>
- The primary endpoint was progression-free survival (PFS)
- Minimal residual disease (MRD)–negativity rate, defined as the proportion of patients who achieved both MRD negativity and complete response or better ( $\geq$ CR) in the intent-to-treat population, was a key secondary endpoint. Bone marrow MRD was assessed using NGS (clonoSEQ<sup>®</sup>)
- Sustained MRD negativity was defined as MRD negativity with  $\geq$ CR  $\geq$ 12 months apart and without MRD positivity in between
- For cytogenetic/genomic analyses, the UMA-NGS target panel was used on DNA extracted from CD138<sup>+</sup> bone marrow cells (**Figure 1**)



#### MRD negativity and PFS

- D-VRd was associated with increased overall (**Figure 4A**) and sustained (**Figure 4B**) MRD-negativity rates compared with VRd in both IMS/IMWG CGS standard- and high-risk patients at sensitivity thresholds of  $10^{-5}$  and  $10^{-6}$
- D-VRd led to improved PFS in both IMS/IMWG CGS standard- and high-risk patients (**Figure 5**)
- Notably, achieving  $\geq$ 12-month sustained MRD negativity removes the negative prognostic impact of an IMS/IMWG CGS high-risk classification at diagnosis (**Figure 6**)
- D-VRd improved PFS versus VRd for both standard- and high-risk patients classified by PP-FISH or IMS/IMWG CGS criteria (**Figure 7**)

