

# Reclassification of ASCERTAIN (ASTX727-02) Myelodysplastic Syndrome (MDS) Patients: Outcomes Including Clinical Response, Overall Survival (OS), and Leukemia Free Survival (LFS) based on IPSS-R and IPSS-M Scoring Systems

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## BACKGROUND

Oral decitabine/cedazuridine (ASTX727) is a fixed-dose combination of the hypomethylating agent (HMA) decitabine (DEC) and the cytidine deaminase (CDA) inhibitor cedazuridine. ASTX727 has been developed as monotherapy treatment for MDS/CMML and AML (NCT03306264).

- Oral DEC has poor oral bioavailability, but when co-administered with cytidine deaminase inhibitor cedazuridine (CED) has increased oral bioavailability
- Results of the Phase III ASCERTAIN (ASTX727-02) study demonstrated a fixed dose combination of DEC (35 mg) and CED (100 mg) given once daily X 5 days on a 28-day cycle is pharmacokinetic (PK) exposure equivalent to the standard intravenous (IV) DEC regimen of 20 mg/m<sup>2</sup> daily X 5 days on a 28-day cycle<sup>1</sup>
- To facilitate PK-bridging with existing historical efficacy data, ASCERTAIN study subjects were classified using the International Prognostic Scoring System (IPSS)
- IPSS considers three main prognostic indicators (marrow blasts, cytogenetics, and cytopenias) in determining risk level<sup>2</sup>
- Many of the relevant hematologic parameters were collected as part of ASCERTAIN allowing reclassification of many patients by more recently developed prognostic criteria (IPSS-R, IPSS-M)
- Here we present additional analyses which characterize the ASCERTAIN population as possible using IPSS-R and IPSS-M

Table 1. IPSS-R Scoring

IPSS-R defines five IPSS-R risk groups: very low, low, intermediate, high, and very high

- IPSS-R enhances the prognostic model with more detailed cytogenetic subgroups (five vs three risk groups), separate subgroups within the “marrow blasts <5%” group, and a depth of cytopenias measurement defined with cut-offs for hemoglobin counts, platelet counts, and neutrophil counts<sup>3</sup>

Prognostic category	Prognostic score value						
	0	0.5	1	1.5	2	3	4
Cytogenetics	Very good	Good	Intermediate	Poor	Very poor		
BM blasts, %	≤ 2	> 2 to < 5	5-10	> 10			
Hgb, g/dL	≥ 10	8 to < 10	< 8				
Platelets, × 10 <sup>9</sup> /L	≥ 100	50 to < 100	< 50				
ANC, × 10 <sup>9</sup> /L	≥ 0.8	< 0.8					
Cytogenetic group	Characteristics						
Very good	-Y, del(11q)						
Good	Normal, del(5q), del(2p), del(20q), del(5q) + 1 additional abnormality						
Intermediate	del(7q), +8, +9, i(17q), other abnormalities not in other groups						
Poor	-7, inv(3)(q3q), -7del(7q) + 1 additional abnormality, complex (3 abnormalities)						
Very poor	Complex (> 3 abnormalities)						

Table 2. IPSS-M Scoring

IPSS-M incorporates molecular data into its risk assessment

- IPSS-M considers the effects of molecular profiling on adverse patient outcome, in particular the impact of variations in the TP53<sup>mut</sup>, MLL<sup>del</sup>, and FLT3<sup>mut</sup> mutations<sup>4</sup>

Table 1. IPSS-M Risk Score Construction from an Adjusted Cox Multivariable Regression for Leukemia-Free Survival.<sup>4</sup>

Category and Variable	Adjusted Hazard Ratio (95% CI)	Model Weight
<b>Clinical</b>		
Bone marrow blasts — %	1.07 (1.05-1.09)	0.0704
min(Platelets,250) — ×10 <sup>9</sup> /l	0.998 (0.997-0.999)	-0.00222
Hemoglobin — g/dl	0.84 (0.81-0.88)	-0.171
<b>Cytogenetic</b>		
IPSS-R cytogenetic category	1.33 (1.21-1.47)	0.287
<b>Gene main effects (17 variables, 16 genes)<sup>4</sup></b>		
TP53 <sup>mut</sup>	3.27 (2.38-4.48)	1.18
MLL <sup>del</sup>	2.22 (1.49-3.32)	0.798
FLT3 <sup>mut</sup> /T3D	2.22 (1.11-4.45)	0.798
SF3B1 <sup>del</sup>	1.66 (1.03-2.66)	0.504
NPM1	1.54 (0.78-3.02)	0.430
RUNX1	1.53 (1.23-1.89)	0.423
NRAS	1.52 (1.05-2.20)	0.417
ETV6	1.48 (0.98-2.23)	0.391
IDH2	1.46 (1.05-2.02)	0.379
CBL	1.34 (0.99-1.82)	0.295
EZH2	1.31 (0.98-1.75)	0.270
UZAF1	1.28 (1.01-1.61)	0.247
SRSF2	1.27 (1.03-1.56)	0.239
DNMT3A	1.25 (1.02-1.53)	0.221
ASXL1	1.24 (1.02-1.51)	0.213
KRAS	1.22 (0.84-1.77)	0.202
SF3B1*	0.92 (0.74-1.16)	-0.0794
Gene residuals (1 variable, 15 genes; possible values of 0, 1, or 2) <sup>4</sup>		
min(Nres,2)	1.26 (1.12-1.42)	0.231

Figure 1. Risk Classifications By IPSS Version

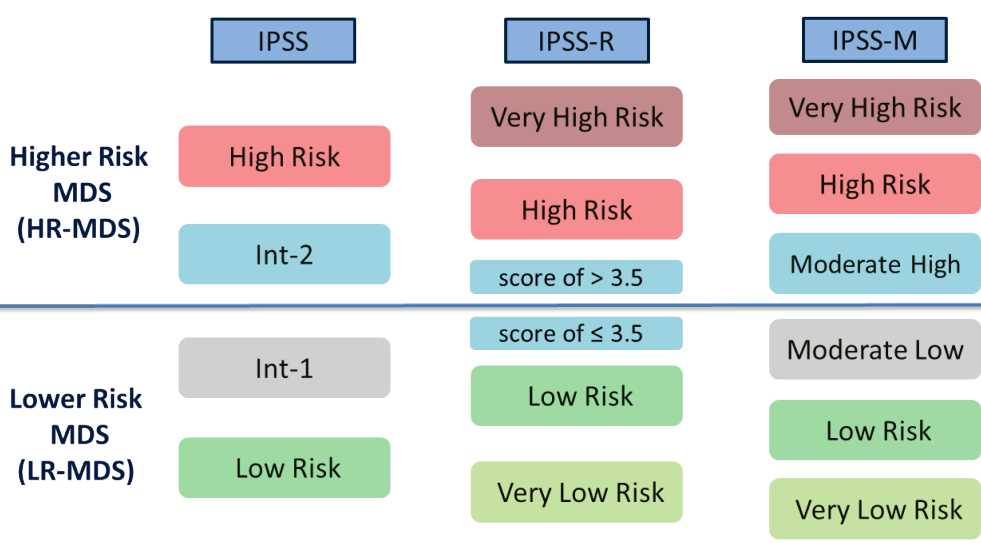
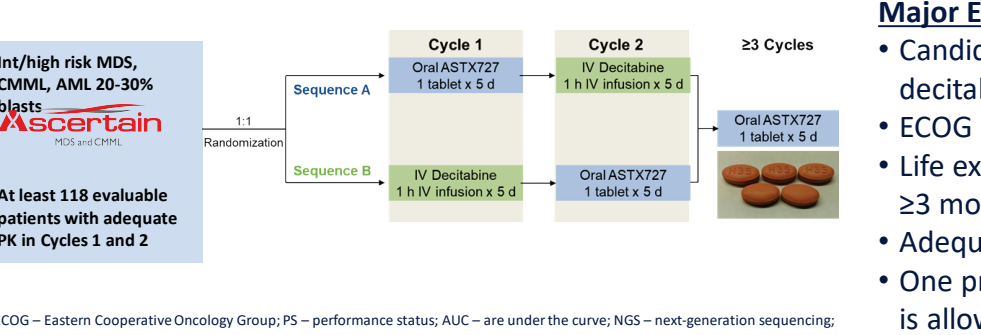


Figure 2. Phase 3 Study Design



**Major Entry Criteria:**

- Candidates for IV decitabine
- ECOG PS 0-1
- Life expectancy of ≥3 months
- Adequate organ function
- One prior cycle of HMA is allowed

**Previously Presented:**

- Demonstrated PK AUC equivalence to IV DEC at 20 mg/m<sup>2</sup>
- Oral safety profile consistent with IV, no marked GI toxicity<sup>1</sup>
- Clinical activity (CR rate 22%, mOS ~32 months)<sup>2</sup>

**Current Analyses:**

- 133 subjects enrolled and whole blood collected for NGS analyses
- NGS (179 genes) analyses available for 125 subjects
- Focus on genes from 2022 MDS NCCN Clinical Practice Guidelines in Oncology (NCCN Guidelines)<sup>9</sup> (30 genes)

## ASCERTAIN STUDY PATIENT CHARACTERISTICS

Table 3. Patient Characteristics

MDS, IPSS classification	Total Treated N=133 (%)	
	High	21 (16%)
Int-2	27 (20%)	
Int-1	64 (48%)	
Low	5 (4%)	

- Of the 117 MDS subjects treated, 104 and 105 had the necessary data for reclassification by IPSS-R and IPSS-M, respectively

MDS, IPSS-R classification	Total Treated N=117 (%)	
	Very High Risk	23 (20%)
High Risk	24 (21%)	
Intermediate	31 (26%)	
Low Risk	23 (20%)	
Very Low	3 (2%)	

- As a result of re-classification from IPSS to IPSS-R, 28 subjects upgraded to HR-MDS, 3 downgraded to LR-MDS

MDS, IPSS-M classification	Total Treated N=117 (%)	
	Very High	25 (21%)
High	24 (21%)	
Moderate High	18 (15.5%)	
Moderate Low	13 (11%)	
Low	21 (18%)	
Very Low	4 (3.5%)	

- As a result of re-classification from IPSS to IPSS-M, 29 subjects upgraded to HR-MDS, 5 downgraded to LR-MDS

## RECLASSIFICATION RESULTS

Figure 3. Reclassification Results: Sankey Diagram

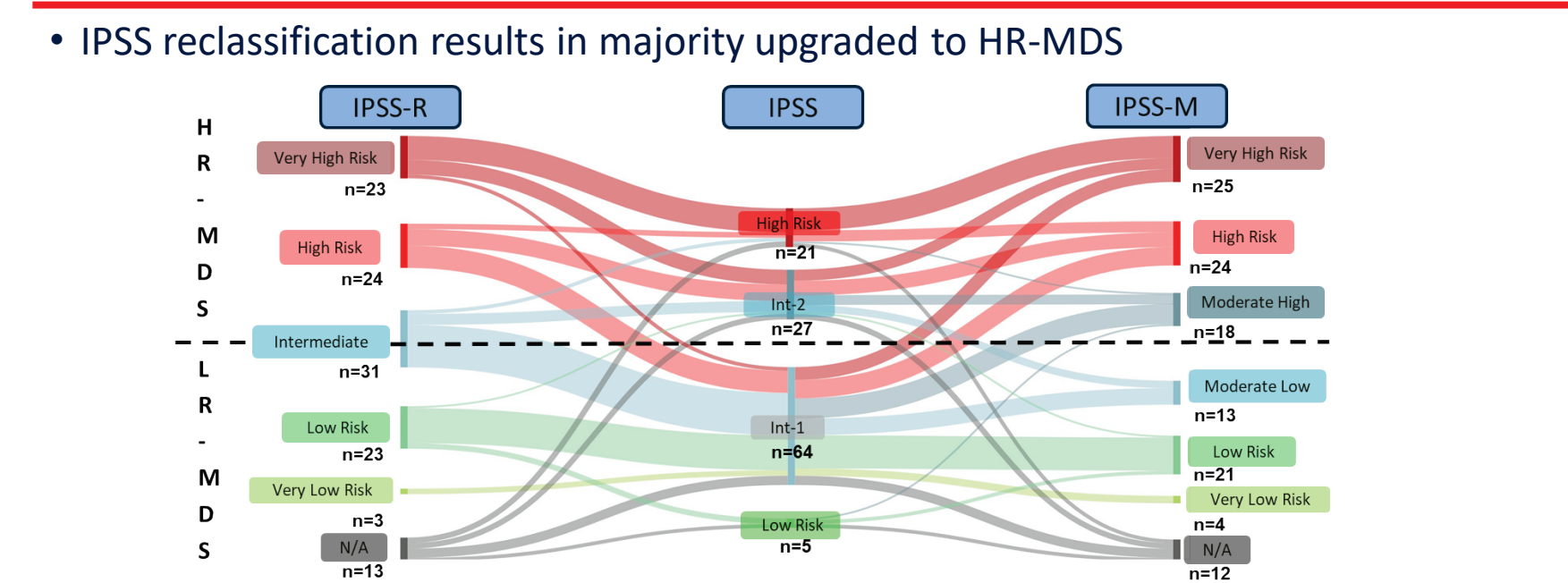


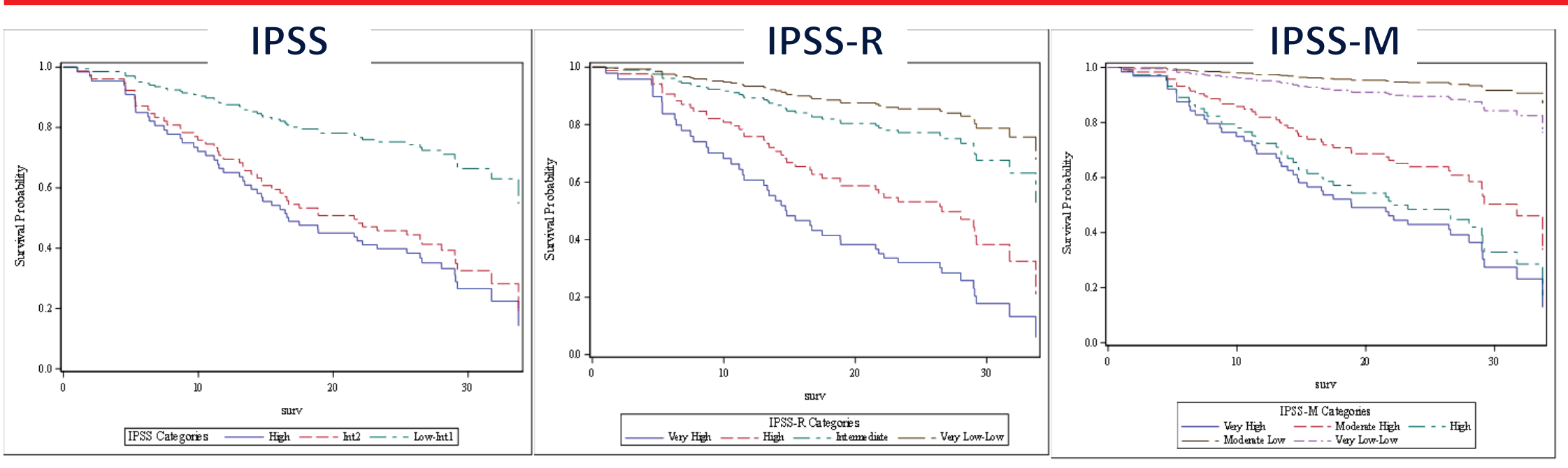
Table 4. Reclassification Results: OS and LFS

- Overall Survival (OS) and Leukemia Free Survival (LFS)

IPSS	Risk Category	OS Median Estimate (Months)	OS Median Estimate (Months)		LFS Median Estimate (Months)	
			IPSS-R	IPSS-M	IPSS-R	IPSS-M
IPSS (n=117)	HR-MDS	High	15.44	10.87		
	LR-MDS	Int-2	23.26	16.66		
IPSS-R (n=104)	HR-MDS	Very High	26.58	14.69	18.86	11.70
		High	28.02	18.09		
	LR-MDS	Intermediate	NE	33.74	NE	33.74
IPSS-M (n=105)	HR-MDS	Very High	13.44	9.95		
		High	21.78	17.20		
	LR-MDS	Mod High	29.21	29.21		

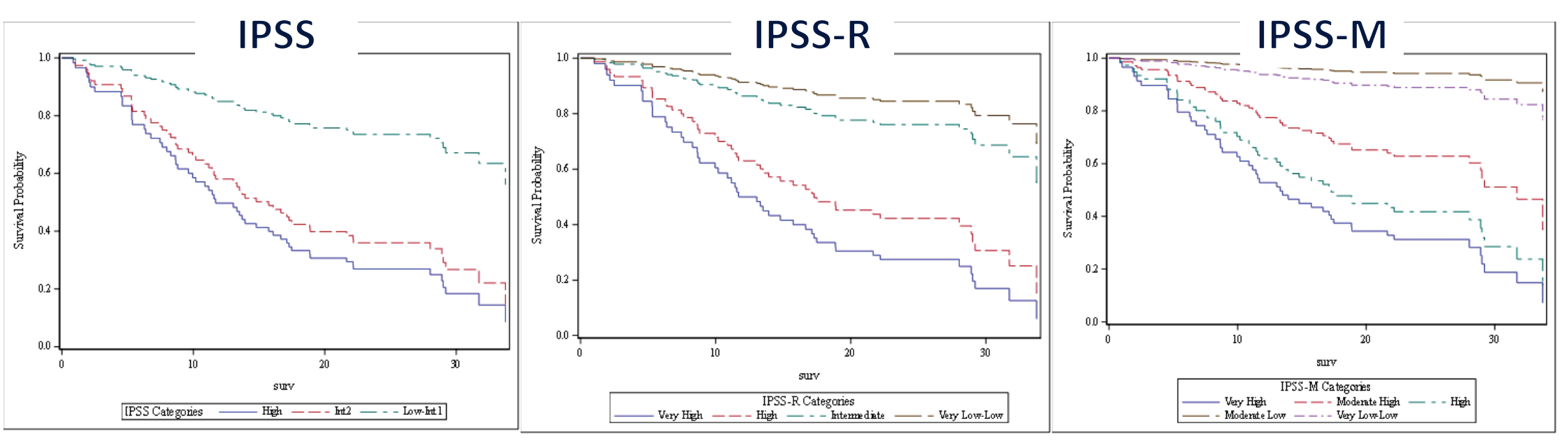
## RECLASSIFICATION RESULTS

Figure 4. Kaplan-Meier Plot: Overall Survival



- IPSS, IPSS-R, and IPSS-M showed a clear separation of the different risk categories for OS

Figure 5. Kaplan-Meier Plot: Leukemia Free Survival



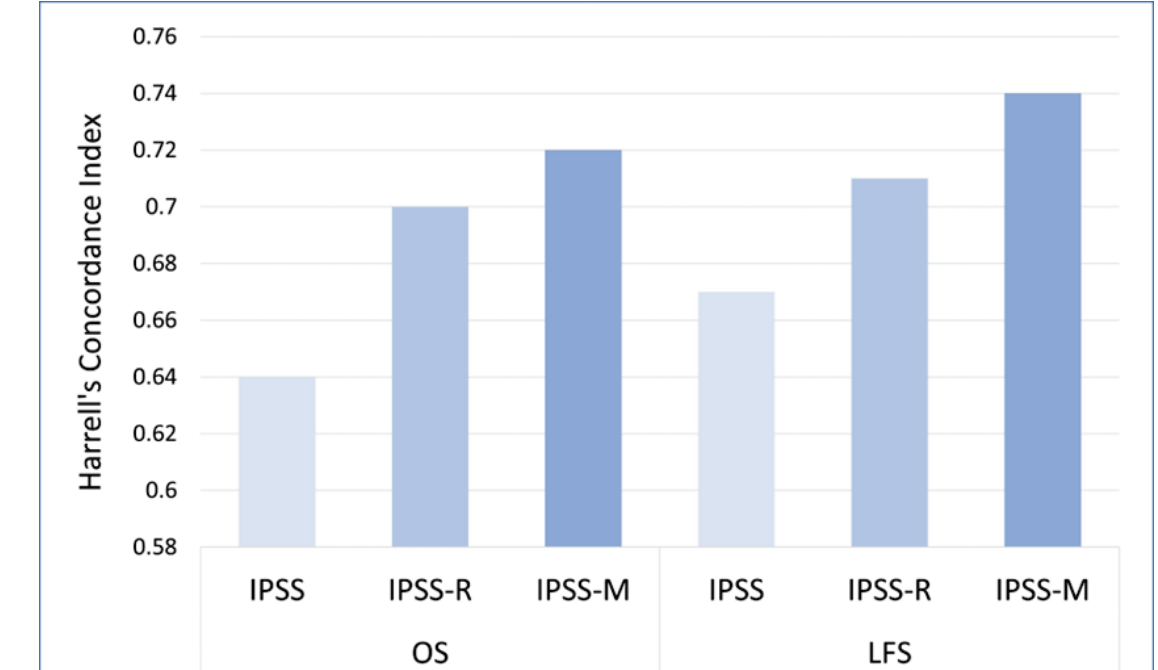
- IPSS, IPSS-R, and IPSS-M showed a clear separation of the different risk categories for LFS

Table 5. Clinical Response

IPSS	Risk Category	CR (%)	
		HR-MDS	High
IPSS-R (n=104)	HR-MDS	Very High	16/67 (23.9)
		High	Risk Level >3.5
IPSS-M (n=105)	HR-MDS	Intermediate	8/37 (21.6)
		Very Low/Low	Risk Level ≤3.5

- No significant difference in CR rates was observed from redefining HR and LR risk categories after migration from IPSS to IPSS-R and IPSS-M

Figure 6. c-index for OS and LFS



- Harrell's concordance index (c-index) for both OS and LFS improves with migration from IPSS to IPSS-R to IPSS-M
- IPSS-M has the highest c-index for both OS and LFS, indicating better ability to predict patient outcome compared to IPSS and IPSS-R

## CONCLUSIONS

### Impact of risk stratification by IPSS-R and IPSS-M on the IPSS-defined ASCERTAIN patient population

- Reclassification from IPSS to IPSS-R or IPSS-M upgraded multiple subjects from a LR to a HR category, describing the ASCERTAIN patient population as a *majority higher-risk population with worse prognosis* than previously assumed from IPSS
- The efficacy as measured by the CR rates *did not change* when the LR and HR categories were defined by the different risk stratification systems
- The c-index improves with migration from IPSS to IPSS-R to IPSS-M, suggests that the IPSS-M score is more effective in predicting patient outcomes compared to IPSS and IPSS-R

## REFERENCES

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