



INTERIM RESULTS FROM THE CARO STUDY AND THE MYRIAM REGISTRY COMPARED WITH DATA FROM ACADEMIC INSTITUTION

IMPLEMENTATION OF THE REVISED MYELOMA COMORBIDITY INDEX (R-MCI) IN PATIENTS WITH MULTIPLE MYELOMA IN REAL WORLD

BACKGROUND

Risk-adapted treatment strategies to further improve outcome in multiple myeloma (MM) are crucial, especially for elderly patients. The R-MCI is a validated prognostic risk tool for MM patients developed at the University Clinic Freiburg (UKF)¹⁻⁵ based on 5 risk factors determined from multivariate analysis, being widely available in the clinics: impaired lung and kidney function, Karnofsky Performance Status, frailty, age and, if available, cytogenetics (Figure 1). Based on these risk factors and an easy-to-use webpage (www.myelomacomorbidityindex.org), patients can be classified into fit (R-MCI: 0-3),

intermediate-fit (R-MCI: 4-6) or frail (R-MCI: 7-9) (Figure 2) with substantially different prognosis, PFS, OS, hematologic and non-hematologic side effects (SAEs) and treatment endurance. Whereas the R-MCI has been successfully performed and validated in a small multicenter validation in academic cohorts (other referral and university DSMM study group centers) in Germany⁶, the external validation of the R-MCI in real world is missing, i.e., data on the feasibility of the R-MCI from secondary and primary sites. This was therefore performed in this analysis and compared to UKF data.

CONCLUSION

In real world, the R-MCI was adapted for lung function and frailty assessment to make it feasible for routine use. With these minor adjustments, the real world R-MCI risk group distribution was comparable with the R-MCI risk groups in the academic settings.

Next steps will be to correlate the R-MCI risk groups with treatment decision and outcome, plus learn which and why differences in primary, secondary and tertiary centers in terms of the R-MCI may exist. In the future, real world data on R-MCI may play a central part in trial design and risk-adapted treatment approaches in the elderly and/or impaired younger MM patients. Taken together, the real world data demonstrate that the R-MCI is of value also in real world settings and, thus, may be widely applicable in daily practice, e.g. for tailoring myeloma therapies.

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Figure 1

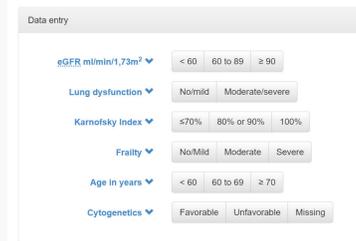


Figure 1. Factors of the revised myeloma comorbidity index (R-MCI). The first five of the 6 factors are mandatory and should be available. Cytogenetics can be available. (https://www.myelomacomorbidityindex.org)

Figure 2

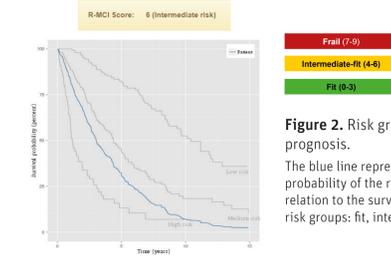


Figure 2. Risk groups of the R-MCI and prognosis. The blue line represents the survival probability of the respective patient in relation to the survival probability of the 3 risk groups: fit, intermediate-fit and frail.

METHODS

The prospective, multicenter, non-interventional study CARO evaluates the combination of carfilzomib with a) lenalidomide/dexamethasone, b) dexamethasone or c) dexamethasone/daratumumab in patients with at least one prior therapy, and the registry MYRIAM collects longitudinal data on routine clinical care of primary and secondary myeloma treatment in hospitals and private practices in Germany (Sponsor: iOMEDICO).

Both studies collect parameters to calculate the R-MCI and, therefore can serve as external validation cohorts for the R-MCI. We here compared the frequency of R-MCI risk-groups from CARO and MYRIAM with historical UKF data (>1,000 patients), thus from a tertiary referral center. Descriptive statistics were used for the analysis.

Academic Setting

Categorization of lung function

Hilfe/Spezifikation Lungenfunktionsstörung

Mild
FEV1 <80%

Moderat
FEV1 <50% oder Diffusionskapazität <51%

Schwer
FEV1 <50%

Einfluss auf Score (Maximalpunktzahl: 9)

Kein / Mild: 0
Moderat / Schwer: 1

Frailty Assessment

Hilfe/Spezifikation Karnofsky Index

Bitte nutzen Sie die Einteilung des Karnofsky Index diese Tabelle.

Einfluss auf Score (Maximalpunktzahl: 9)

KI 100%: 0
KI 80% oder 90%: 2
KI <70%: 3



Real World Adjustment

Categorization of lung function

If FEV1 and/or diffusion capacity are not available, lung dysfunction is categorized as:

| Lung dysfunction | Lung disease | | | |
|------------------|--------------|---|---|--|
| | A | B | C | D |
| None | None | Known lung disease without symptoms (i.e. no dyspnea) | Known lung disease and dyspnea on effort OR Known lung disease without symptoms (i.e. no dyspnea) but under medical therapy | Known lung disease and dyspnea at rest |
| Smoking status | non-smoker | >0 - <20 py | ≥20 - <40 py | ≥40py |
| | 0 (none) | 0 (mild) | 1 (moderate) | 1 (severe) |
| | 0 (mild) | 0 (mild) | 1 (moderate) | 1 (severe) |
| | 1 (moderate) | 1 (moderate) | 1 (moderate) | 1 (severe) |
| | 1 (severe) | 1 (severe) | 1 (severe) | 1 (severe) |

Smoking status is only required for the evaluation of lung dysfunction of patients with the lung disease categories A, B.

Frailty Assessment

| Frailty assessment done by treating physician | Frailty Score |
|---|---------------|
| Mild | 0 |
| Moderate | 1 |
| Severe | 1 |

No frailty assessment by treating physician available, therefore approximation by Karnofsky Performance Status (KPS)

| KPS | Frailty Score |
|---------|----------------|
| ≥80% | 0 (no or mild) |
| 50- 70% | Not evaluable |
| ≤ 40% | 1 (severe) |

Table 1

| Parameters | MYRIAM n=1,236 | CARO n=312 | UKF n=1,080 |
|--|-------------------|---------------|----------------|
| Age at inclusion (years), median (range) | 72 (29-97) | 74 (35-90) | 72 (31-99) |
| Sex, n (%) | | | |
| Female | 556 (45) | 141 (45) | 455 (42) |
| Male | 680 (55) | 171 (55) | 625 (58) |
| Type of MM, n (%) | | | |
| IgG | 713 (58) | 173 (56) | 604 (56) |
| IgA | 227 (18) | 69 (22) | 204 (19) |
| κ- / λ- light chain only | 263 (21) | 63 (20) | 202 (19) |
| Other | 33 (3) | 7 (2) | 70 (6) |
| Cytogenetics*, n (%) | | | |
| Favorable | 406 (33) | 191 (61) | 430 (40) |
| Unfavorable | 308 (25) | 52 (17) | 316 (29) |
| Missing | 522 (42) | 69 (22) | 334 (31) |

*Cytogenetics: favorable = Hyperdiploidy, t(11;14), normal Karyotype, del(13q14); unfavorable = t(4;14), t(14;16), t(14;20), del(17p), Hypodiploidy, c-myc, Chromosome1-Aberrations. (www.myelomacomorbidityindex.org / Sonneveld et al. Blood 2016)

Table 1: Patient characteristics

Patient characteristics were comparable throughout all 3 cohorts for age, sex and myeloma paraprotein subtypes. Smaller, albeit expected differences were determinable in cytogenetics.

Table 2

| | Academic Setting | Real world Setting |
|---------------|---|--|
| Lung Function | Pulmonary function test (FEV1) | Pre-existing lung disease +/- Medication +/- Smoking Status |
| Frailty | Timed up and go Instrumental activities of daily living (IADL) etc. | Simple estimation by treating physician: <ul style="list-style-type: none"> • KPS • Mobility assessment (walking into consultation room) • Performance regarding daily tasks • Overall fitness grade (1= very good – 6 = unsatisfactory) |

Table 2: Lung function and frailty assessment in real world

Lung and frailty assessment was adapted for real world. With this adaption, it was possible to determine the R-MCI of real world patients more easily.

RESULTS

In CARO, 334 patients have been enrolled from 65 sites across Germany. Thereof, at database cut for this interim analysis, data of 312 (93%) patients were available for R-MCI assessment. In MYRIAM, at database cut for this interim analysis, 1,917 patients had been enrolled from 150 sites in Germany and for 1,236 (65%) patients the R-MCI could as yet be obtained (Table 1).

In order to implement the R-MCI in the real world setting, we did adapt two assessments to make its usage even more feasible: Firstly, the lung function was more easily defined as moderately or severely impaired and secondly, the frailty assessment according to Fried⁷ as exemplified for both in Table 2. These two minor adjustments made it possible to use the R-MCI in real world settings even more swiftly.

Notably, the R-MCI results, as assessed from the webpage and exemplified in Fig. 1 & 2, revealed similar results in all 3 cohorts. In patients with evaluable R-MCI, the risk distribution in MYRIAM, CARO and UKF was similar with 31%, 31% and 27% of fit, 57%, 53% and 55% of intermediate-fit and 11%, 16% and 18% of frail patients, respectively (Figure 3).

Figure 3

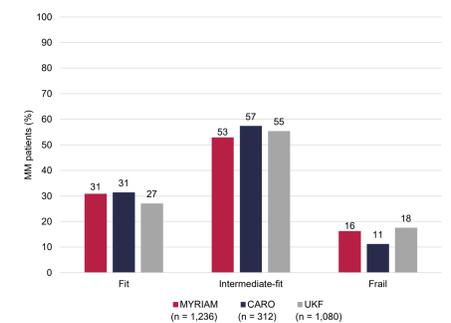


Figure 3. Distribution of R-MCI risk groups. Distribution of R-MCI risk groups was similar throughout the 3 cohorts.

