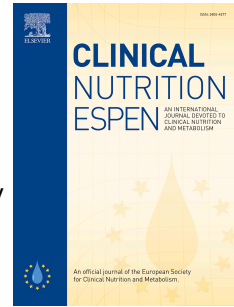


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Use of automated assessment for determining associations of low muscle mass and muscle loss with overall survival in patients with colorectal cancer – a validation study

Karel C. Smit, Jeroen W.G. Derksen, Sophie A. Kurk, Pim Moeskops, Miriam Koopman, Wouter B. Veldhuis, Anne M. May



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22 **Abstract**

23

24 **Background:** Low muscle mass and skeletal muscle mass (SMM) loss are associated with adverse
25 patient outcomes, but the time-consuming nature of manual SMM quantification prohibits
26 implementation of this metric in clinical practice. Therefore, we assessed the feasibility of automated
27 SMM quantification compared to manual quantification. We evaluated both diagnostic accuracy for
28 low muscle mass and associations of SMM (change) with survival in colorectal cancer (CRC) patients.

29

30 **Methods:** Computed tomography (CT) images from CRC patients enrolled in two clinical studies were
31 analyzed. We compared i) manual vs. automated segmentation of preselected slices at the third
32 lumbar [L3] vertebra ("semi-automated"), and ii) manual L3-slice-selection + manual segmentation
33 vs. automated L3-slice-selection + automated segmentation ("fully-automated"). Automated L3-
34 selection and automated segmentation was performed with Quantib Body Composition v0.2.1.
35 Bland-Altman analyses, within-subject coefficients of variation (WSCVs) and Intraclass Correlation
36 Coefficients (ICCs) were used to evaluate the agreement between manual and automatic
37 segmentation. Diagnostic accuracy for low muscle mass (defined by an established sarcopenia cut-
38 off) was calculated with manual assessment as the "gold standard". Using either manual or
39 automated assessment, Cox proportional hazard ratios (HRs) were used to study the association
40 between changes in SMM (>5% decrease yes/no) during first-line metastatic CRC treatment and
41 mortality adjusted for prognostic factors. SMM change was also assessed separately in weight-stable
42 (<5%, i.e. occult SMM loss) patients.

43

44 **Results:** In total, 1580 CT scans were analyzed, while a subset of 307 scans were analyzed in the fully-
45 automated comparison. Included patients (n=553) had a mean age of 63±9 years and 39% were
46 female. The semi-automated comparison revealed a bias of -2.41 cm², 95% limits of agreement [-9.02
47 to 4.20], a WSCV of 2.25%, and an ICC of 0.99 (95% confidence intervals (CI) 0.97 to 1.00). The fully-

48 automated comparison method revealed a bias of -0.08 cm^2 $[-10.91 \text{ to } 10.75]$, a WSCV of 2.85% and
49 an ICC of 0.98 (95% CI 0.98 to 0.99). Sensitivity and specificity for low muscle mass were 0.99 and
50 0.89 for the semi-automated comparison and 0.96 and 0.90 for the fully-automated comparison.

51 SMM decrease was associated with shorter survival in both manual and automated assessment
52 ($n=78/280$, HR 1.36 [95% CI 1.03 to 1.80] and $n=89/280$, HR 1.38 [95% CI 1.05 to 1.81]). Occult SMM
53 loss was associated with shorter survival in manual assessment, but not significantly in automated
54 assessment ($n=44/263$, HR 1.43 [95% CI 1.01 to 2.03] and $n=51/2639$, HR 1.23 [95% CI 0.87 to 1.74]).

55

56 **Conclusion:** Deep-learning based assessment of SMM at L3 shows reliable performance, enabling the
57 use of CT measures to guide clinical decision making. Implementation in clinical practice helps to
58 identify patients with low muscle mass or (occult) SMM loss who may benefit from lifestyle
59 interventions.

60

61 **Keywords:** colorectal cancer, body composition, muscle mass, survival analysis, deep-learning

62 Introduction

63 Cross-sectional skeletal muscle area (SMA), commonly measured through computed tomography
64 (CT) imaging at the level of the 3rd lumbar vertebra, serves as a reliable indicator of overall skeletal
65 muscle mass (SMM) (1,2). SMA assessment has garnered substantial attention due to its association
66 with various adverse clinical outcomes. Both low muscle mass and a decline in muscle mass over
67 time are associated with increased morbidity and mortality among diverse patient populations, but
68 especially among the elderly and cancer patients (3–9). While manual assessment of SMA currently
69 remains the gold standard, it is time-consuming and labor-intensive. This limits its feasibility in
70 routine clinical practice and large-scale studies. Several studies have already explored automated
71 segmentation techniques on a single axial CT slice, showcasing promising outcomes in accurately
72 delineating SMA (10). However, studies focusing on a fully-automated approach, combining
73 automated L3 localization with automated segmentation, are less prevalent, although also reporting
74 promising results (11–15). Importantly, research focusing on reproducibility and validity of these
75 methods through repeated measures within the same patient is sparse. This poses a limitation in the
76 translation of these techniques into clinical practice and longitudinal research. Additionally, in clinical
77 oncology practice there is a need for a diagnostic tool to identify sarcopenic and pre-cachectic
78 patients that – from the outside – seem to be weight stable and in good condition. In a retrospective
79 cohort of colorectal cancer (CRC) patients, it was found that incident sarcopenia occurs in 8.5% of
80 patients during a mean follow-up of 15 months and was associated with a higher risk of death (16).
81 Such patients easily remain unnoticed, while early interventions could prevent muscle depletion and
82 may improve clinical outcome.(16)

83 Therefore, the primary objective of this study was to evaluate and compare the accuracy of (fully-
84)automated SMA assessment at the third lumbar level against manual assessment in a clinically
85 relevant context. This included: i) analyzing the diagnostic accuracy of automated SMA assessment
86 compared to manual assessment for identifying low muscle mass and tracking muscle loss over time
87 using repeated measures within patients; and ii) comparing associations for changes in SMM and

- 88 overall survival (OS) between automated and manual assessment in patients with CRC, with
89 additional comparisons for weight stable and weight decreased patients separately.

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90 **Methods**

91 **Study population**

92 CT scans assessed in this study were from patients with CRC enrolled in two clinical studies. The first
93 is the Dutch Colorectal Cancer Group CAIRO3 study (17). This phase III study investigated the effect
94 of maintenance treatment with low dose capecitabine plus bevacizumab (CAP-B) versus observation
95 on overall survival in previously untreated metastatic CRC (mCRC) patients not progressing during
96 first-line systemic treatment with six cycles of treatment with capecitabine plus oxaliplatin and
97 bevacizumab (CAPOX-B). Patients were recruited from 64 Dutch hospitals between May 30, 2007 and
98 Oct 15, 2012. CT scans were available from 55/64 hospitals. CT scans were collected at four
99 timepoints: (i) before start of initial six cycles of CAPOX-B; (ii) at randomization between CAP-B and
100 observation; (iii) at first disease progression (PD1), before start of CAPOX-B or other reintroduction
101 treatment; (iv) at second disease progression (PD2), after progression on any systemic reintroduction
102 treatment. Depending on availability of the CT scans, up to four CT scans per patient were used.
103 The second clinical study is PLCRC PROTECT-Plus, which is a substudy of the Prospective Dutch
104 Colorectal Cancer (PLCRC) cohort (18,19). PLCRC PROTECT-Plus is an observational study investigating
105 the association between body composition and treatment toxicity among patients with stage II-III
106 colon cancer receiving adjuvant systemic therapy. Patients were recruited from three Dutch hospitals
107 from 15 Jun, 2017 onwards. CT scans were collected at three timepoints: (i) at first CRC diagnosis; (ii)
108 prior to the fourth adjuvant systemic therapy cycle; (iii) six months after surgery of primary tumor.
109 Up to three CT scans per patient were used, depending on availability of the CT scans. Survival data is
110 not yet available for PLCRC PROTECT-Plus.

111 A schematic overview of study characteristics and performed data analyses in both studies can be
112 found in **Supplement 1**. All patients of both studies provided informed consent. Protocols were in
113 accordance with the declaration of Helsinki.

114

115 **Manual segmentation**

116 All CT scans were manually segmented by trained researchers using the Slice-o-matic (version 5.0,
117 Tomovision) software package using pre-specified thresholds for Hounsfield units (HU; -29 to 150) to
118 identify and demarcate the muscle compartments. A single slice evaluation was performed at the
119 level of the third lumbar vertebra (L3). SMA was calculated in square centimeters by adding the
120 surface values of the psoas, abdominal and long spine muscle. SMA is highly correlated with total
121 SMM, which can be calculated based on SMA by linear transformation (20–22). Results from manual
122 assessment of skeletal muscle from the CAIRO3 studies have been previously published (5,23–27).

123

124 **Automated segmentation**

125 Automated segmentation was performed by the deep learning-based software Quantib Body
126 Composition (version 0.2.1, Quantib, Rotterdam, Netherlands) (28). The Quantib Body Composition
127 software is available online for testing through research.quantib.com. The automatic method
128 consists of two steps: first, the slice at the mid-L3 level is automatically selected from the full image
129 using a convolutional neural network. Second, 5 slices centered on the the selected mid-L3-level are
130 automatically segmented into visceral fat, subcutaneous fat, psoas muscle, abdominal muscle and
131 long spine muscle, using a second convolutional neural network. Before the analysis, the software
132 resamples CT scans to a standard slice thickness of 5 mm. Segmentation results are averaged over
133 the five segmented slices, i.e. over a standard 25 mm range centered on mid-L3.

134 The body composition tool was evaluated with two approaches:

- 135 1. The single L3 slices as selected during manual assessment were used as input , i.e., to
136 evaluate the segmentation independently (semi-automated comparison).
- 137 2. Manually assessed L3 slices were compared with subsequent automated L3 localization and
138 segmentation, (i.e., fully-automated comparison).

139 Differences between manual and automated L3 localization were calculated based on the middle of
140 the five selected L3 slices (mid-L3) in automated L3 localization. The fully-automated comparison was
141 performed with the subset of PLCRC PROTECT-Plus CT scans, for which full abdominal CT scans were

142 available. Results of fully-automated assessment of the psoas muscle at L3 compared to manual
143 segmentation have been published previously and yielded a Dice index of 0.93 (± 0.04 SD) (12). Four
144 random examples as well as the four biggest outliers based on absolute SMA differences will be
145 shown and possible causes for discrepancies between manual and automated assessment will be
146 discussed.

147 CT scans were excluded from comparison when noise-level or acquisition artifacts precluded manual
148 or automated segmentation, when manual segmentation was performed unilaterally due to the
149 abdominal being partly missing and/or abnormalities in the abdominal wall or when the original
150 DICOM source data could not be retrieved. Additionally, CT scans from PLCRC PROTECT-Plus were
151 excluded when the L3 distance from the most caudal slide could not be calculated based on the
152 available metadata.

153

154 **Statistical analyses**

155 SMA of manual and automated assessment was compared using Bland-Altman analyses with 95%
156 limits of agreement (LoA) for both the semi-automated and fully-automated comparison, with
157 adjustment for multiple observations per individual (29,30). Within-subject coefficients of variation
158 (WSCVs) were also calculated, considering the broad normal range of SMA values and the possibility
159 that within-subject variation is not uniform, but proportional to the magnitude of the measurement.
160 The root mean squared method was used for the calculation of WSCVs (31). Two-way mixed
161 Intraclass Correlation Coefficients (ICCs) were calculated as well. The same analyses were performed
162 to compare L3 slice location for manual and fully-automated assessment. For percentual SMA
163 changes between subsequent CT scans, WSCV will not be calculated, considering most patients will
164 have minor SMM changes. This results in possible large relative differences, despite small absolute
165 differences between manual and automated assessment, leading to large but uninformative WSCV
166 values.

167 Several stratified analyses were conducted to identify potential subgroups in which automated
168 assessment may be particularly (in)accurate. This was performed for sex (male/female), BMI (<18.5 /
169 18.5-25 / 25-30 / ≥ 30), and muscle mass (low muscle mass/normal muscle mass). Low muscle mass
170 was defined based on the skeletal muscle index (SMI) and BMI of patients, using an established cut-
171 off for sarcopenia, in accordance with prior research (5,32). SMI is calculated by dividing the SMA by
172 squared height in meters. Cut-offs for low muscle mass were a SMI <43 and a BMI <25 or a SMI <53
173 and a BMI ≥ 25 for males, and a SMI <41 for females. These thresholds were established based on a
174 large cohort of cancer patients by using optimal stratification to best separate patients based on
175 their survival time. Results of the two clinical studies were compared as well, since PLCRC PROTECT-
176 Plus is a more recent study and imaging quality might have improved thereby possibly influencing
177 accuracy. WSCV differences between subgroups were calculated with Welch's t-test for subgroups
178 with 2 categories, and with a one-way ANOVA with post-hoc Tukey-Kramer tests for subgroups with
179 >2 categories.

180 Diagnostic accuracy of possible clinical implementations was assessed for low muscle mass, and for
181 SMA change between subsequent CT scans. To improve generalizability, supplemental content will
182 be provided of the same using two alternative sarcopenia cut-offs which are used in prior research,
183 i.e. a SMI <55 for males and a SMI <39 for females (15,33), and a SMI <52.4 for males and a SMI <38.5
184 for females (14,34). SMA change was categorized as a >5% decrease (yes/no) in accordance with
185 prior research (16). Manual assessment was used as gold standard. For low muscle mass, subgroup
186 analyses were performed for sex, BMI, and study cohort. For SMA decrease, subgroup analyses were
187 performed for sex and weight change. Weight change was categorized as >5% decrease (yes/no), in
188 accordance with the cancer cachexia criteria by Fearon et al., and used in prior research (16,34).

189 Diagnostic accuracy terms (sensitivity, specificity, negative predictive value [NPV], and positive
190 predictive value [PPV]) were classified based on predefined thresholds. Values >0.7 were categorized
191 as moderate, >0.8 as good, and >0.9 as excellent.

192

193 *SMA – OS analysis (CAIRO3 study)*

194 For these analyses, we used CAIRO3 data, since survival data for PROTECT-Plus are not yet available.

195 To analyze associations between SMA and OS, Cox proportional hazards models were employed,

196 comparing manual and automated SMM assessments. For these analyses, we selected the first two

197 CT scans acquired before and after first-line treatment with six cycles of CAPOX-B. Median interval

198 between the two CT scans was 4.6 months (IQR 4.1 to 5.2). Survival time was calculated from the

199 time of randomization which can occur up to a few weeks after CT scan 2. It is worth noting that

200 inclusion in CAIRO3 required stable disease or better post-first-line treatment, meaning we

201 effectively evaluated associations with survival in mCRC patients who exhibited at least stable

202 disease after six cycles of CAPOX-B. Analyses were performed for SMA decrease (yes/no) and the

203 combination of weight decrease and SMA decrease (both stable/weight stable, SMA decrease >5%

204 (i.e. 'occult SMM loss')/weight decrease >5%, SMA stable/weight & SMA decrease >5%).

205 Furthermore, we conducted analyses for muscle mass (low muscle mass/normal muscle mass) for

206 both CT scans separately, and for changes between the two CT scans (normal muscle mass both CT

207 scans/new low muscle mass/persistent low muscle mass/remission of low muscle mass), in

208 accordance with prior research (16). Similar to our approach evaluating diagnostic accuracy for low

209 muscle mass, analyses will be repeated for two other commonly used sarcopenia cut-offs and

210 provided as supplemental content. All models were adjusted for age (continuously), sex

211 (male/female), WHO performance status (0/1), primary tumor site (colon/sigmoid/rectum), resection

212 primary tumor (yes/no), response to initial treatment (stable disease/partial or complete response),

213 lactate dehydrogenase (LDH) level at randomization (normal/elevated), metastasis pattern

214 (synchronous/metachronous), dose reductions during initial treatment (yes/no), and treatment arm

215 (CAP-B/observation).

216 All data were analyzed using R statistical software version 4.0.3 (35). All statistical tests were two-

217 sided with an alpha level of 0.05.

218 Results

219 Study population

220 In total, 553 patients (39% female) were included in the full study population including CAIRO3 and
221 PLCRC PROTECT-Plus patients. Baseline characteristics of both study populations can be found in
222 **table 1**. After exclusion of scans that were unsuitable for comparison, 1580 and 307 CT scans
223 remained for semi-automated and fully-automated comparison (**figure 1**). For the semi-automated
224 comparison, average SMA was $139.9 \pm 31.6 \text{ cm}^2$ for manual segmentation and $137 \pm 31.5 \text{ cm}^2$ for
225 automated segmentation, and $139.1 \pm 33.2 \text{ cm}^2$ and $139.1 \pm 31.8 \text{ cm}^2$ for the fully-automated
226 comparison (**table 2**).

227

228 Semi-automated comparison

229 Median difference between manual and automated assessment was -2.40 cm^2 [IQR: -3.80 to -1.10].
230 Bland-Altman analyses displayed a non-significant small negative bias of -2.41 cm^2 [95% LoA: -9.02 to
231 4.20] for the entire dataset ($n=1580$) with a WSCV of 2.25% and an ICC of 0.99 (95% CI 0.97 to 1.00).
232 (**figure 2, table 2**). Biases for subgroup analyses ranged from -1.81 cm^2 [95% LoA -8.90 to 5.27] to -
233 2.80 cm^2 [95% LoA -9.99 to 4.40] cm^2 . The degree of bias was not evidently correlated with SMA
234 magnitude. WSCV values in subgroup analyses ranged from 1.81% to 3.78% with a statistically
235 significant difference for PROTECT-Plus CT scans compared to CAIRO3 CT scans (1.81 vs 2.23%) and
236 for BMI 25-30 compared to ≤ 18.5 (1.91 vs 3.78%). ICC values ranged from 0.94 to 0.99, with the
237 lowest ICC for the $n=26$ BMI ≤ 18.5 subgroup. Four random examples and the four largest outliers
238 based on absolute SMA difference between manual and automated assessment can be found in
239 **figure 3**

240

241 Fully-automated comparison

242 *L3 slice selection*

243 Median difference between manual and automated mid-L3 slice selection was -0.50 mm [IQR -4.10
244 to 4.65]. Bland-Altman analyses showed a non-significant bias of 1.04 mm [95% LoA -22.17 to 24.25],
245 a WSCV of 3.18% and an ICC of 0.85 (95% CI 0.82 to 0.88). Based on the height of an average L3
246 corpus of 32 mm (36), 82.1% of the mid-L3 slices are within 16 mm below and 16 mm above the
247 manual level and are therefore expected to be on the same vertebra (**Figure 4**). No significant
248 differences between subgroups were observed. ICC values ranged from 0.39 to 0.92 with the lowest
249 ICC for the n=6 BMI \leq 18.5 subgroup.

250

251 *SMA assessment*

252 Median difference between manual and automated assessment was 0.27 cm² [IQR -2.01 to 2.89].
253 Bland-Altman analyses displayed a non-significant small negative bias of -0.08 cm² [95% LoA -10.91 to
254 10.75] for the entire dataset (n=307) with a WSCV of 2.85% and an ICC of 0.98 (95% CI 0.98 to 0.99).
255 (**figure 2, table 2**). Biases for subgroup analyses ranged from 1.26 cm² [95% LoA -10.63 to 13.15] to -
256 1.51 cm² [95% LoA -14.11 to 11.08]. In fully-automated SMA comparison, the degree of bias was also
257 not evidently correlated with SMA magnitude. WSCV values in subgroup analyses ranged from 2.58%
258 to 3.10%. No significant different differences between subgroups were observed. ICC values ranged
259 from 0.96 to 0.99, with the lowest ICC for both the n=157 male and n=150 female subgroups.

260

261 **Validation of SMA change between subsequent CT scans**

262 In the semi-automated comparison, SMA changes between subsequent CT scans could be calculated
263 for n=1027 changes. Median change between two subsequent CT scans was small with -0.62% (IQR -
264 5.08 to 3.57%) in manual assessment and -0.39% (IQR -5.48 to 4.37) in automated assessment, with a
265 median difference between manual and automated assessment of 0.18% (IQR -0.88 to 1.26). Bland-
266 Altman analyses displayed a non-significant bias 0.20% [95% LoA -6.13 to 6.53] for the entire dataset.
267 In the fully-automated comparison, SMA changes between subsequent CT scans could be calculated
268 for n=186 changes. Median change between two subsequent CT scans was small with 0.84% (IQR -

269 3.34 to 4.66) in manual assessment and 0.51% (IQR -3.07 to 4.36) in automated assessment, with a
270 median difference between manual and automated assessment of -0.11% (IQR -1.51 to 1.59). Bland-
271 Altman analyses displayed a non-significant bias 0.08% [95% LoA -6.48 to 6.64] for the entire dataset
272 **(figure 5, table 3).**

273

274 **Diagnostic accuracy for low muscle mass and SMA decrease between subsequent CT scans**

275 A comprehensive overview of the sensitivity, specificity, positive predictive value (PPV) and negative
276 predictive value (NPV) for both semi- and fully-automated comparisons for low muscle mass
277 detection can be found in **table 4**. This same overview can be found for two other commonly used
278 sarcopenia cut-offs in **supplement 2a and 2b**. In the semi-automated comparison, low muscle mass
279 occurred in 680/1262 (54%) of CT scans. Overall sensitivity and NPV were excellent (0.99 and 0.98),
280 with minor differences between subgroups. In the fully-automated comparison, low muscle mass
281 occurred in 105/187 (57%) of CT scans, with excellent sensitivity and NPV as well (0.96 and 0.95).
282 Compared to semi-automated assessment, greater differences across subgroups were seen, with
283 sensitivity and NPV being notably lower in males compared to females and for BMI >30 compared to
284 the other BMI subgroups.

285

286 The diagnostic accuracy for both semi- and fully-automated comparisons for SMA decrease can be
287 found in **table 5**. In the semi-automated analysis, SMA decrease occurred in 259/1027 (25%) of
288 subsequent CT scans. Overall sensitivity and NPV were excellent (0.90 and 0.97). Weight values were
289 available for n=732 changes between subsequent CT scans. SMA decrease occurred in 139/624 (22%)
290 patients with stable weight, and in 61/108 (56%) of patients with weight decrease. In the fully-
291 automated analysis, SMA decrease occurred in 33/186 (18%) of subsequent scans. Overall sensitivity
292 was moderate (0.73), with an excellent NPV (0.94). Weight values were available for n=68 changes
293 between subsequent CT scans. SMA decrease occurred in 20/61 (33%) of patients with stable weight,
294 and in 5/7 (71%) of patients with weight decrease. In general, diagnostic misclassification occurred

295 mainly for patients with a small SMA change in cm^2 , especially in the semi-automated comparison
296 **(Supplement 3)**.

297

298 **Survival analyses**

299 Median follow-up time in the CAIRO3 study population (n=432) was 20.3 months (IQR: 12.0 to 31.4),
300 during which a total number of 414 (96%) of patients died. CT scan and weight information was
301 accessible for 308 patients for the initial scan and for 341 for the subsequent scan. 280 patients had
302 scan data available for both scans, and 263 patients had both weight and CT scan data available at
303 both timepoints. Detailed patient characteristics can be found in **supplement 4**. Low muscle mass
304 both before and after completion of first-line CAPOX-B was not associated with shorter survival, nor
305 were muscle mass changes during first-line treatment in both manual and automated assessment
306 **(figure 6, supplement 5)**. A comparison of the multivariate analyses with two other commonly used
307 sarcopenia cut-offs can be found in **supplement 6**. SMA decrease during this time was associated
308 with shorter survival in both manual (adjusted HR [HR_{adj}] 1.36 [1.03 to 1.80]) and semi-automated
309 assessment (HR_{adj} 1.38 [1.05 to 1.81]). No significant associations with early death were observed
310 among patients with weight decrease and stable SMA compared to weight and SMA stable patients.
311 SMA decrease among weight stable patients compared to both weight and SMA stable patients was
312 associated with significantly shorter survival in manual assessment (HR_{adj} 1.42 [1.01 to 2.03]), but not
313 in automated assessment (HR_{adj} 1.23 [0.87 to 1.74]). A weight and SMA decrease compared to weight
314 and SMA stable patients was not associated with death in manual assessment (HR_{adj} 1.48 [0.98 to
315 2.22]), but with shorter survival in automated assessment (HR_{adj} 1.72 [1.15 to 2.58]).

316 Discussion

317 In this analysis, we found that deep-learning based assessment of skeletal muscle mass at the third
318 lumbar vertebra shows reliable performance. Furthermore, diagnostic accuracy was excellent for
319 ruling out low muscle mass and good for detecting SMM decrease above a pre-established threshold,
320 with semi-automated assessment performing notably better compared to fully-automated
321 assessment for the latter. Lastly, we found an association between (occult) SMM loss and shorter
322 survival during first-line treatment in patients with mCRC, with comparable hazard ratios between
323 manual and automated assessment of SMM.

324

325 Manual vs automatic SMA assessment

326 Our validation results are in line with an increasing number of studies showing reliable performance
327 of automated assessment of SMA (10). Generally we observed a small underestimation of skeletal
328 muscle compared to the current gold standard of manual segmentation, which has been reported by
329 others as well (37). Differences in performance among subgroups are small, with automated
330 assessment generally performing better in subgroups with higher SMA. Several explanations have
331 been given for this, including that small underestimations have less impact for patients with larger
332 muscle areas, and that in case of severe muscle wasting, the abdominal wall can be compromised,
333 leading to segmentation errors (12,37,38). Performance in the PROTECT-Plus cohort was slightly
334 better compared to the CAIRO3 cohort, which might be explained by improvements in imaging
335 modalities over time, since PROTECT-Plus is a more recent study. Detailed analyses comparing e.g.
336 different CT scan brands could not be performed, since this data was not available. Furthermore, we
337 found that automated L3 slice selection was accurate in the large majority of cases. Small changes in
338 L3 slice selection are not expected to lead to large differences, given that no substantial changes
339 occur in skeletal muscle area at the L3 level (39). This is indeed consistent with our observation that
340 biases, WSCV and ICC values were comparable between semi-automated and fully-automated
341 assessment in our analyses. Additionally, based on the visual inspection of outliers in our analyses, it

342 is important to exercise caution when applying automated assessment to patients with unusual intra-
343 abdominal anatomy. This is demonstrated by the four largest outliers in **Figure 3**. These outliers
344 include patients who exhibit unusual HU-densities in anatomical structures adjacent to the skeletal
345 muscle due to underlying pathological conditions, resulting in a less distinguishable abdominal wall.

346

347 **Using automated SMA assessment as a diagnostic tool**

348 To our knowledge, no other validation study has shown diagnostic accuracy of automated
349 assessment for changes of low muscle mass and skeletal muscle changes over time; i.e. analyzing
350 repeated assessments of the same patients, and limited research has been conducted on low muscle
351 mass (13–15). We found excellent diagnostic accuracy for low muscle mass in both semi-automated
352 and fully-automated assessment, with its main strength being its ability to identify true negative
353 cases. Our diagnostic results fit clinical implementation as a screening tool, where automated SMM
354 assessment could help identify patients who might benefit from exercise or dietary interventions,
355 and thereby also providing additional means to evaluate such interventions. Minor overdiagnosis is
356 not expected to lead to adverse consequences, since these interventions are recognized as safe and
357 feasible for patients with CRC, and potentially improve various health-related outcomes, benefitting
358 not only sarcopenic patients but CRC patients in general (40–42).

359 Total diagnostic accuracy for SMA decrease was excellent in both semi-automated and fully-
360 automated assessment as well, although sensitivity, specificity, PPV and NPV values showed larger
361 variety. In this context especially, it should be noted that using manual assessment as gold standard
362 has some limitations. Manual assessment of SMA is not perfectly accurate, and our reported
363 variability of a few percent between manual and automated assessment is in fact comparable with
364 interobserver variability of manual assessment performed by trained researchers (37,43). In other
365 words, a current limitation of both manual and automated assessment is an expected error rate of a
366 few percent per CT scan, resulting in increased diagnostic misclassification when assessing SMM
367 changes between subsequent CT scans. This is especially important for patients with low SMM,

368 illustrated by **supplement 2**, showing that most disagreement between automated and manual
369 assessment occurred with relatively small absolute SMM changes. Additionally, the generally lower
370 sensitivity values in fully-automated assessment compared to semi-automated assessment might
371 reflect minor differences in selection of the L3 slice compared to manual selection, resulting in a
372 subsequent small difference besides the inter-observer/intermodality segmentation error rate.
373 Automated assessment however, provides the opportunity to segment multiple CT slices, which
374 eventually might improve accuracy and enable future studies to focus on evaluating predictive value
375 of skeletal muscle volume, instead of SMA. Automated assessment also enables the possibility to
376 assess muscle mass without using a HU threshold, so that changes in intramuscular fat can also be
377 assessed, thereby more directly targeting muscle mass changes, which might prove to be an
378 alternative predictive marker (44).

379

380 **Survival analyses**

381 Whilst the primary focus for our survival analyses was to validate automated assessment of SMA, the
382 results of manual assessment of different SMM variables should be put into context first. An
383 increasing body of research shows associations between low muscle mass and shorter survival in
384 cancer patients (4,9). In our study however, and probably due to the study population, we did not
385 observe any associations between low muscle mass (change) and survival, which is in line with a
386 previous analysis with the CAIRO3 study population. CAIRO3 consists of mCRC patients with a
387 relatively good prognosis, because only patients with stable disease or better after 6 cycles of
388 CAPOX-B are included, and patients with disease progression and/or unacceptable toxicity during
389 first-line CAPOX-B were eligible for inclusion (5). Additionally, low muscle mass prevalence in our
390 study exceeded 50%, which is higher than other reported prevalences in CRC patients of around 40%,
391 further indicating our study population might be a specific CRC subgroup (9,45). Besides this, low
392 muscle mass prevalence differs widely between cancers while certain types of cancer do not show
393 significant associations, which might indicate that cancer type-specific cutoff values for low muscle

394 mass might be warranted (46). This hypothesis would also fit the differences we found on diagnostic
395 accuracy compared to Gu et al., who reported a lower sensitivity (0.718) compared to specificity
396 (0.876) in a group of females in China who underwent DXA examinations. After calculating an optimal
397 cutoff value based on their data instead of the pre-established cut-off by Prado et al., they reported a
398 higher AUC with higher sensitivity (0.875) compared to specificity (0.778) (15,33). The impact of
399 applying different sarcopenia cut-offs is also illustrated by the varying effect sizes between those cut-
400 offs as shown in **supplement 6**. Also, contrary to prior research (16), we did not find significant
401 associations with shorter survival for patients with newly diagnosed low muscle mass; although the
402 referenced study was conducted in stage I-III CRC patients and might thus not be directly
403 comparable.

404 In line with another study examining SMM decrease during first-line systemic therapy, we observed a
405 significant association with shorter survival for patients with SMM decrease (47). Results are also
406 similar compared to other CRC domains and/or other SMM cut-offs (48,49). Moreover, we found a
407 significant association with shorter survival for occult SMM loss, not for a decrease of weight only,
408 nor - unexpectedly - for a decrease of both, although numbers were small in both those groups.

409 Comparing manual assessment with automated assessment, we observed agreement in statistical
410 significance for most SMM variables. In general, automated assessment labels more patients as
411 having low muscle mass and/or decrease in SMM, thereby possibly overdiagnosing a subset of
412 patients. This might explain that most HRs of automated assessment are attenuated towards one
413 compared to manual assessment. The only exception here is the subgroup of patients who lost both
414 body weight and SMM, where automated assessment was significantly associated with shorter
415 survival, but manual assessment showed no significant association with death. This change was
416 based on 4/31 patients in automated assessment that were classified differently compared to
417 manual assessment (data not shown), indicating that subgroups categorized by weight/SMM lack
418 optimally stabilized hazard ratios in our study because of their small sample sizes.

419 Regardless, our results indicate that weight decrease might not always be indicative of SMM
420 decrease. Additionally, individuals experiencing weight loss alone appear to exhibit a lower risk
421 profile compared to those with SMM loss. This underscores the importance of evaluating SMM in
422 clinical practice—an aspect that is not commonly implemented, likely owing to time constraints, for
423 which automated assessment provides a viable solution. Future studies should concentrate on
424 establishing more robust criteria for SMM loss and further evaluate whether SMM decrease can be
425 stopped or even reversed.

426

427 **Strengths and limitations**

428 There are several strengths to this study. We analyzed a large sample of patients with longitudinal
429 data, providing the opportunity to validate performance to assess patterns of body composition; an
430 aspect that has received limited attention in prior research. Moreover, validation of fully-automated
431 assessment of SMA, which includes automated localization of L3, remains an area that has not been
432 thoroughly investigated in current literature. Furthermore, we provided a comprehensive overview
433 by both validating SMA assessment, and subsequently evaluating the performance of automated
434 SMM assessment in possible clinical contexts, while also focusing on low muscle mass and SMM
435 changes in weight stable patients, which has not been extensively evaluated.

436 Limitations are that we were only able to evaluate fully-automated SMM in a subset of our study,
437 due to missing full abdominal CT scan data, which resulted in small subsets for some stratified
438 analyses in the fully-automated comparison, especially for BMI subcategories. Furthermore, despite
439 having excellent agreement in most cases, automated assessment does not always produce optimal
440 segmentation, especially among patients with low BMI. We believe that the reported accuracy is
441 adequate for scientific research purposes, optionally supported by quality checks in selected
442 subgroups with poorer performance, thereby enabling and facilitating large scale studies. Based on
443 our results, we advise such quality checks for patients with low muscle mass and/or uncommon
444 intro-abdominal anatomy. These checks are especially important when assessing SMM change. For

445 application in clinical context, based on the current data, a quick visual inspection is advisable to
446 identify rare automated segmentation failures.

447

448 **Conclusion**

449 Deep-learning based assessment of skeletal muscle mass at the third lumbar vertebra shows reliable
450 performance, enabling the use of CT measures to guide clinical decision making. Our data suggests
451 implementation as a screening tool to help identify patients with low muscle mass or (occult) SMM
452 loss for whom lifestyle interventions might improve their outcomes.

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456

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462

463 Conflict of interest

464 MK reports institutional financial interests with Amgen, Bayer, BMS, Merck-Serono, Nordic Pharma,

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466 interests: an advisory role for ZON-MW, membership of the scientific board of the Dutch Cancer

467 Society (KWF), chairmanship of the Dutch Colorectal Cancer Group (DCCG), and principal investigator

468 (PI) of the Prospective Dutch Colorectal Cancer (PLCRC) cohort.

469 WV was a co-founder of Quantib-U.

470 PM was employed by Quantib-U.

471 All other authors declare no conflicts of interest.

472

473 Author contribution

474 All authors contributed to conceptualization, methodology and writing – review & editing of the

475 research. KS, JD, SK, PM, and WV contributed to data curation. KS performed formal analysis,

476 visualization and wrote the first draft under supervision of JD, MK, AM, and WV. All authors read and

477 approved the final version of the manuscript.

478

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