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3 Article

4 **Freeze-dried kit formulation and physicochemical assessment of a**
5 **daratumumab-based radiopharmaceutical**

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ABSTRACT

33 Daratumumab is a fully human anti-CD38 monoclonal antibody with strong potential as a targeting
34 vector for therapeutic radionuclides. This study aimed to develop a freeze-dried daratumumab
35 immunoconjugate kit by selecting a suitable chelator (DOTA-NHS, *p*-SCN-Bn-DOTA, or *p*-SCN-Bn-
36 1B4M-DTPA) for ¹⁷⁷Lu-labeling, optimizing the freeze-drying formulation, and evaluating the
37 physicochemical properties and purity profiles. Conjugation performed in carbonate buffer at elevated
38 temperature enhanced chelator incorporation, and supported selection of daratumumab-*p*-SCN-Bn-
39 DOTA as the most suitable candidate, achieving radiolabeling yield up to 99.8 % without additional
40 purification. Among the evaluated freeze-dried formulations, a saline-based, buffer-free sucrose-
41 mannitol formulation containing polysorbate 20 (S.F5) provided the most favorable characteristics,
42 including minimal residual moisture and the highest monomer purity with non-detectable HMWS
43 species under the applied SE-HPLC conditions. ATR-FTIR and Raman spectroscopy confirmed
44 preservation of the antibody structural integrity after conjugation and freeze-drying. In an *in vitro* study
45 using human serum, [¹⁷⁷Lu]Lu-daratumumab-*p*-SCN-Bn-DOTA was shown to maintain higher
46 radiochemical purity over 168 h compared to [¹⁷⁷Lu]Lu-daratumumab-*p*-SCN-Bn-1B4M-DTPA,
47 indicating greater stability. These results support the feasibility of a ready-to-use freeze-dried
48 daratumumab-*p*-SCN-Bn-DOTA kit for ¹⁷⁷Lu-labeling.

49 *Keywords:* daratumumab, immunoconjugate, ¹⁷⁷Lu-radiopharmaceutical, ready-to-use kit formulation,
50 freeze-drying, physicochemical characterization

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INTRODUCTION

56 Recent advances in cancer therapy have demonstrated the central role of monoclonal
57 antibodies (mAbs), both as stand-alone therapeutics and as a key component of innovative
58 combination strategies (1). Beyond their immunological functions, mAbs can serve as selective
59 carriers for therapeutic radionuclides, an approach known as radioimmunoconjugate (RIC)
60 therapy. This strategy allows targeted delivery of radionuclide-conjugated antibodies to tumor
61 cells, while minimizing radiation exposure to surrounding healthy tissues (2). The clinical value

62 of this concept has been previously established and confirmed by ⁹⁰Y-ibritumomab tiuxetan
63 and ¹³¹I-tositumomab (3, 4).

64 Multiple myeloma is a malignant clonal plasma cell disorder characterized by a
65 biologically and clinically heterogeneous spectrum ranging from asymptomatic (smoldering)
66 to symptomatic disease, defined by the presence of myeloma-defining events, including CRAB
67 features and SLiM biomarkers. Despite advances in classification and risk stratification,
68 multiple myeloma remains challenging to treat due to its biological heterogeneity, the
69 development of resistance to standard therapies, and the cumulative toxicity of conventional
70 chemotherapy (5-7).

71 Daratumumab, a fully human IgG1κ monoclonal antibody (Darzalex, approved for *i.v.*
72 administration by the EMA in 2016; Darzalex Faspro, approved for *s.c.* administration by the
73 FDA in 2020), and isatuximab, a humanized IgG1 monoclonal antibody (Sarclisa, approved for
74 *i.v.* administration by the EMA in 2020), target CD38, a transmembrane ectoenzyme highly
75 expressed on multiple myeloma cells (8-11). Their antitumor activity is mediated through
76 multiple immune-dependent mechanisms, including antibody-dependent cellular cytotoxicity
77 (ADCC), complement-dependent cytotoxicity (CDC), antibody-dependent cellular
78 phagocytosis (ADCP), and direct induction of apoptosis. Both agents exhibit a comparable,
79 generally manageable safety profile characterized by infusion-related reactions, infections, and
80 hematologic toxicities (12-16).

81 Beyond its use as a native antibody, daratumumab has also been investigated as a vector
82 for targeted radionuclide therapy. Preclinical studies with β⁻-emitting ¹⁷⁷Lu-DOTA-
83 daratumumab, α-emitting ²²⁵Ac-daratumumab, and ²¹²Pb-daratumumab have demonstrated
84 enhanced antitumor effects in disseminated multiple myeloma models (17-19). For diagnostic
85 application, conjugates with positron-emitting radionuclides, such as ⁸⁹Zr-DFO-daratumumab
86 and ⁶⁴Cu-DOTA-daratumumab, have enabled specific immunoPET imaging of CD38-positive
87 malignancies (20, 21). Furthermore, the integration of diagnostic and therapeutic radionuclides
88 within a single vector molecule constitutes the basis of the theranostic approach, a key concept
89 in modern precision medicine, that embodies the principle of “see what you treat and treat what
90 you see” (22). Paired ⁸⁹Zr/¹⁷⁷Lu-daratumumab has demonstrated the potential of a theranostic
91 approach that integrates immunoPET imaging with targeted radioimmunotherapy in preclinical
92 studies (23).

93 In this context, a freeze-dried kit formulation offers a practical strategy for standardized
94 preparation of antibody-based immunoconjugates, enabling reproducible radiolabeling at the
95 point of use. Overall, these findings provide a strong rationale for the present study, which aims
96 to develop a daratumumab-based immunoconjugate kit suitable for ^{177}Lu labeling. These
97 chelators were selected to represent both macrocyclic (DOTA) and acyclic (DTPA-based)
98 systems commonly used for radiometal labeling of antibodies, enabling comparison of
99 radiolabeling efficiency, complex stability, and impact on antibody integrity. The
100 physicochemical properties and structural integrity of the immunoconjugate were evaluated
101 using chromatographic, spectroscopic, and electrophoretic techniques, as well as radiolabeling
102 efficiency.

103

104

EXPERIMENTAL

105 *Conjugation of daratumumab with BFCs*

106 The commercially available antibody daratumumab (400 mg/20 mL solution for *i.v.*
107 injection, Janssen Biologics B.V., The Netherlands) was purified by repeated ultrafiltration (30
108 min, 5000 rpm) on a 30 kDa cut-off filter (Amicon[®] Ultra, Merck KGaA, Germany). Buffer
109 exchange was performed with 0.1 mol L⁻¹ phosphate buffer (pH 8.0). Three commercially
110 available bifunctional chelators (BFCs), *p*-SCN-Bn-DOTA (2-(4-isothiocyanatobenzyl)-
111 1,4,7,10-tetraazacyclododecane-1,4,7,10-tetraacetic acid), DOTA-NHS ester (1,4,7,10-
112 tetraazacyclododecane-1,4,7,10-tetraacetic acid mono-*N*-hydroxysuccinimide ester) and *p*-
113 SCN-Bn-1B4M-DTPA (2-(4-isothiocyanatobenzyl)-6-methyl-diethylene-triamine pentaacetic
114 acid), were evaluated to identify the most suitable chelator for subsequent ^{177}Lu -labeling.

115 Conjugation reactions were performed using 20-, 30-, and 50-fold molar excess of each
116 BFC. According to the first protocol, the reaction mixtures were gently shaken and incubated
117 at 4 °C for 16 h (24). In a second protocol, the phosphate buffer was replaced with 0.1 mol L⁻¹
118 carbonate buffer at pH 8.5, and the samples were incubated at 37 °C for 1.5 h (18). After
119 completion of the reaction, the resulting immunoconjugates were purified by repeated
120 ultrafiltration (30 min, 5000 rpm) using 0.15 mol L⁻¹ ammonium acetate (pH 7.0) to remove
121 the unbound chelator. The final immunoconjugate concentration was determined
122 spectrophotometrically at 280 nm (UV-1600PC, VWR, USA).

123 *Freeze-drying formulations (F1-F5)*

124 Two batches, each comprising five formulations, were prepared and adjusted to the
125 immunoconjugate concentration of 1 mg mL^{-1} . In both batches, formulations F1 and F2 differed
126 only in buffer strength, and both contained 1 % (*m/V*) mannitol as a bulking agent. All
127 remaining formulations were buffer-free. Formulations F3 and F4 contained mannitol and
128 sucrose at different ratios, whereas formulation F5 additionally included polysorbate 20 as a
129 surfactant. Ultra-pure water produced in-house (TKA MicroPure-ST system, Germany) was
130 used as the main solvent for the first batch (Batch W.F1-5). In the second batch (Batch S.F1-
131 5), the identical formulations were prepared using 0.9 % (*m/V*) saline solution (Alkaloid A.D.,
132 North Macedonia) as the main solvent. The detailed composition of all formulations, expressed
133 as molar ratios, is provided in Table I.

134 *Determination of chelator-to-antibody ratio (CAR)*

135 The average number of chelator molecules conjugated per antibody (chelator-to-antibody
136 ratio, CAR) was determined by matrix-assisted laser desorption/ionization time-of-flight
137 (MALDI-TOF) mass spectrometry using an Axima Performance (Shimadzu, Japan).

138 The pure freeze-dried antibody and various preparations of bifunctional chelating agents
139 were suspended in 1 mL of ultrapure H_2O and washed 3 times using Amicon[®] 30 kDa (Merck)
140 to remove salts. The pure samples were reduced to 20 μL , and 80 μL of resuspension solution
141 (30 % ACN/70 % 0.1 mol L^{-1} TFA) was added. An aliquot of 1 μL of the final sample was
142 applied to the sample target and immediately covered by 1 μL of matrix (20 mg mL^{-1} sinapinic
143 acid in 50 % ACN/50 % 0.1 mol L^{-1} TFA). Operational conditions for the MALDI-TOF
144 instrument were set as follows: mode of operation - linear, polarity - positive, max laser rep
145 rate 1, laser power 85, pulsed extraction: 150000, acquisition mass range 100-300000 Da. Peak
146 processing parameters were as follows: peak width 80 chans, chosen to accommodate the
147 relatively broad peak shapes characteristic of linear mode measurements, smoothing method -
148 average, peak detection method - gradient centroid.

149 *Freeze-drying protocol*

150 Freeze-drying was performed using a FreeZone 6 liter freeze dryer equipped with a
151 stoppering tray (Labconco, USA). Type I glass vials containing 1 mL of each formulation were
152 partially stoppered with 13-mm butyl rubber stoppers and loaded on precooled shelves at $4 \text{ }^\circ\text{C}$.
153 The shelf-temperature, chamber pressure, hold times for each segment, and a graphical
154 representation are summarized in Table II and Fig. S1.

155 *Freeze-dried immunoconjugates sample analysis*

156 Physicochemical and structural characterization was performed to evaluate the quality of
157 the freeze-dried immunoconjugates and to assess the impact of the formulations and freeze-
158 drying process on the final product.

159 *Product appearance and reconstitution time*

160 The freeze-dried formulations were photographed in a black photo box, and their
161 macroscopic appearance was visually assessed. Reconstitution time was determined by
162 dissolving the products in 1 mL of 0.9 % (*m/V*) saline.

163 *Residual moisture*

164 Residual moisture, as a key quality attribute of the freeze-dried product, was determined
165 by Karl Fischer volumetric titration using a Compact V20S titrator (Mettler Toledo, USA).
166 Samples of ~20 mg were prepared under controlled-humidity conditions at 25 °C. Sample
167 handling time was minimized, and vials were opened immediately before analysis to reduce
168 moisture uptake.

169 *Purity*

170 The purity and aggregation profile were analyzed by size-exclusion high-performance
171 liquid chromatography (SE-HPLC) using a Waters Alliance e-2695 system coupled with a
172 UV/Vis Waters 2489 detector (Waters Corporation, USA). Isocratic elution was performed on
173 an XBridge Premier Protein SEC column (250 Å, 2.5 µm, 7.8 i.d. x 300 mm) at 25 °C, for 25
174 minutes. The mobile phase consisted of 50 mmol L⁻¹ phosphate buffer (pH 6.8), 150 mmol L⁻¹
175 NaCl, 50 mmol L⁻¹ KCl, with 10 % (*V/V*) ACN, at a flow rate of 0.5 mL min⁻¹. A volume of 10
176 µL of reconstituted sample was injected after filtration through a low protein-binding PVDF
177 membrane filter. Data were processed in Empower 3 software.

178 *Structure integrity – ATR-FTIR and Raman*

179 To assess secondary structural changes, Fourier-transform infrared (FTIR) spectra of
180 freeze-dried samples were recorded using a Cary 630 equipped with a diamond attenuated total
181 reflection (ATR) accessory (Agilent Technologies, USA) in the 2000–400 cm⁻¹ spectral range.
182 Spectral resolution was set to 4 cm⁻¹ with 64 accumulated scans. MicroLabPC software was
183 used for data acquisition and initial spectral processing. The placebo sample was recorded
184 under the same conditions.

185 In parallel, Raman spectroscopy was used to further verify the higher structural integrity
186 of the immunoconjugates, following a previously published protocol (25).

187 *Electrophoresis*

188 Sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE) was performed
189 using the MultiGel Long system (Biometra, USA) under reducing and non-reducing conditions.
190 Freeze-dried immunoconjugate and daratumumab (control) were reconstituted in 0.9 % saline
191 and mixed with loading buffer. For reducing conditions, β -mercaptoethanol was added to the
192 samples, and the samples were heated at 95 °C for 5 minutes. Non-reducing samples were
193 incubated at 30 °C without β -mercaptoethanol. A prestained protein ladder (10–180 kDa,
194 Thermo Fisher Scientific, USA) served as the molecular mass standard.

195 Discontinuous gels were prepared as 4 % stacking gel and either a 12 % (reducing) or a
196 6 % (non-reducing) resolving gel, using an acrylamide/bis-acrylamide solution (29:1). Proteins
197 were visualized by staining with Coomassie Brilliant Blue R-250 followed by destaining until
198 a clear background was obtained.

199 *Radiolabeling and in vitro stability*

200 Radiolabeling was performed after reconstitution of the freeze-dried samples in various
201 solvents: a solution of ascorbic acid (0.1 g mL⁻¹, *m/V*), 0.1 mol L⁻¹ sodium acetate and saline.
202 The pH was adjusted to 4.5, and the immunoconjugates were labeled by adding approximately
203 11-20 MBq n.c.a. ¹⁷⁷LuCl₃ in aqueous 0.04 mol L⁻¹ HCl (EndolucinBeta 40 GBq mL⁻¹, ITM,
204 Germany n.c.a), at 37 °C with gentle shaking, for a period of 60 minutes.

205 The labeling efficiency was estimated by the ratio of total-to-labeled antibody
206 radioactivity with [¹⁷⁷Lu]Lu-daratumumab-*p*-SCN-Bn-DOTA as a representative sample, by
207 instant thin-layer chromatography (iTLC) using silica gel (SG) strips (Agilent Technologies)
208 and 0.1 mol L⁻¹ sodium acetate buffer (pH 4.5) as the mobile phase. Aliquots (10 μ L) were
209 spotted, and the chromatograms were scanned with a Scan-RAM radio-TLC scanner (LabLogic
210 Group, UK) and analyzed using Laura 6 software. The same iTLC procedure was used to
211 determine the radiochemical yield of the labeled product.

212 Subsequently, the *in vitro* stability of [¹⁷⁷Lu]Lu-daratumumab-*p*-SCN-Bn-DOTA and
213 [¹⁷⁷Lu]Lu-daratumumab-*p*-SCN-Bn-1B4M-DTPA were assessed in human serum at room
214 temperature. Samples were examined every 24 h up to 168 h, and radiochemical purity was
215 determined as previously described.

216

217 RESULTS AND DISCUSSION

218 To achieve efficient radiolabeling, the average CAR was first determined as a key
219 predictor of conjugation efficiency (26). Accordingly, the CAR number was quantified by
220 MALDI-TOF and compared across BFCs, molar excesses and conjugation conditions, and was
221 related to the ¹⁷⁷Lu radiolabeling yield of the daratumumab-immunoconjugates. Based on a
222 previously published approach for daratumumab, the initial conjugation was performed using
223 a 10-fold molar excess of DOTA-NHS in phosphate buffer (pH 8.0) (21). However, neither the
224 conjugation reaction nor the subsequent radiolabeling provided satisfactory results. Therefore,
225 additional conjugations were performed using macrocyclic *p*-SCN-Bn-DOTA, DOTA-NHS,
226 and an acyclic *p*-SCN-Bn-1B4M-DTPA, with 20-, 30- and 50-fold molar excesses for each of
227 the three BFCs, and extended to different conditions, using a carbonate buffer (pH 8.5).

228 The results presented in Table III and Fig. 1 clearly demonstrate that a carbonate buffer
229 and elevated temperature (37 °C) with gentle shaking, compared with a phosphate buffer,
230 improved conjugation, resulting in consistently higher CAR values. Under these conditions,
231 daratumumab-DOTA-NHS showed low CAR overall, with the highest value at a 50-fold excess
232 (reaching CAR value of 0.46 in carbonate buffer pH 8.5), indicating that lower ratios may not
233 give efficient labeling. Considering the published findings that optimal CAR range is
234 commonly within 1-3 (26-28), daratumumab-*p*-SCN-Bn-DOTA conjugates prepared at 20-
235 and 30-fold molar excess were selected for formulation and further experiments. Although
236 daratumumab-*p*-SCN-Bn-1B4M-DTPA showed a similarly low CAR, the 20- and 30-fold
237 excesses were selected for further evaluation, whereas the 50-fold excess was excluded because
238 its CAR was comparable to that of the 30-fold excess. Furthermore, a related immunoconjugate
239 (daratumumab-*p*-SCN-Bn-CHX-DTPA) has been reported using only a 5- to 10-fold excess,
240 supporting that a high excess is not desired (23).

241 Overall, MALDI-TOF comparison of native daratumumab and its corresponding
242 immunoconjugates showed CAR values ranging from 0 to 3.77, and only immunoconjugates
243 prepared under selected molar ratios were taken forward for characterization.

244 The impact of formulation composition and applied freeze-drying protocol on residual
245 moisture was first evaluated. As shown in Table IV, the percentage of residual moisture varies
246 with formulation composition, notably between the water- and saline-based F2-5 formulations.
247 The water-based batch consistently showed higher moisture levels (5.10-6.06 %), while the
248 saline-based batch exhibited lower moisture levels (1.90-5.01 %). This trend suggests that the
249 presence of saline facilitates freezing and drying, thereby more efficiently removing residual
250 water, as indicated by Matejtschuk *et al.* (29).

251 Formulation F1 showed residual moisture of ~5 % in both batches, which is in line with
252 a buffer–mannitol system where 1 % mannitol predominantly acts as a crystalline bulking
253 agent. To minimize the risk of pH shifts, formulation F2 consists of reduced buffer
254 concentration (10 mmol L⁻¹) while retaining 1 % mannitol (30). Moisture remained unchanged
255 in W.F2 but decreased to 2.73 % in S.F2, confirming that the presence of NaCl favors drying
256 and results in lower residual moisture. To further protect the antibody integrity and reduce
257 moisture, formulations F3-F5 were prepared without buffer salts (31). In these buffer-free
258 formulations, sucrose was added at different ratios to modulate mannitol crystallization, and in
259 F5, polysorbate 20 was added as a surfactant (32). These optimizations resulted in significantly
260 lower moisture in S.F5, reaching a desirable level below 3 % (33).

261 Complete reconstitution was defined as the time required to obtain a clear solution,
262 without visible particles, upon gentle swirling. Under these criteria, all formulations
263 reconstituted rapidly (≤ 30 s) in saline, confirming the literature findings for low protein
264 concentrations and formulations containing a crystalline bulking agent (34). The initial
265 macroscopic appearance of the freeze-dried formulations resembles a compact cake, which
266 readily collapses into a powder-like appearance after minimal agitation. This indicates limited
267 cake robustness under the applied freeze-drying protocol and formulation composition, despite
268 achieving the desired dryness. Nevertheless, the integrity of the formulations was confirmed
269 by subsequent analytical characterization, indicating that the observed collapse did not
270 adversely affect product quality.

271 As a preliminary experiment, radiolabeling with ¹⁷⁷Lu was performed on all freeze-dried,
272 water- and saline-based daratumumab immunoconjugates. Among the tested conjugates, the
273 30-fold daratumumab-*p*-SCN-Bn-DOTA showed the highest radiolabeling yield (99.8 %),
274 with no observable difference between the batches. Based on these results, the subsequent
275 evaluation of formulation-related effects on the antibody purity was limited to the 30-fold
276 daratumumab-*p*-SCN-Bn-DOTA conjugate and assessed by SE-HPLC.

277 SE-HPLC results of the reconstituted formulations showed batch-dependent differences
278 in purity between water- and saline-based preparations. Quantification was based on peak-area
279 normalization, and the results were expressed as relative percentages of the main peak, low-
280 molecular-mass species (LMWS, fragments), and high-molecular-mass species (HMWS,
281 aggregates). The purity results are summarized in Table V, and corresponding overlay
282 chromatograms are presented in Fig. 2. Taken together, HMWS formation was identified as the
283 primary selection criterion, whereas LMWS levels remained consistently low and stable, and

284 monomer content remained above 95 % (35). Aggregates were detected in the first two
285 formulations in both batches, with HMWS levels ranging from 4.80–5.90 % for F1 and 6.67–
286 8.55 % for F2. In contrast, formulations F3 and F5 exhibited the highest purity profile, with
287 monomer contents above 98.00 % in both batches, non-detectable HMWS, and minimal
288 LMWS. The immunoconjugate in F4 remained predominantly as a monomer in water, with no
289 detectable HMWS, but showed a slight HMWS formation in saline. These purity profiles
290 highlight F3 and F5 as the most promising candidates for subsequent experiments and for
291 optimizing radiolabeling efficiency.

292 Based on the results for formulation composition, residual moisture and purity, the saline-
293 based formulation containing mannitol:sucrose (2:1) and polysorbate 20 (S.F5) was selected as
294 the promising formulation for further evaluation.

295 As mentioned above, based on CAR, five candidates: 50-fold daratumumab–DOTA–
296 NHS, 20- and 30-fold daratumumab–*p*-SCN-Bn-DOTA, and 20- and 30-fold daratumumab–*p*-
297 SCN-Bn-1B4M-DTPA, were examined. For radiolabeling optimization, 0.1 mol L⁻¹ sodium
298 acetate (pH 4.5) was selected as the reaction solvent for subsequent radiolabeling experiments
299 based on overall performance. The results from the 50-fold daratumumab–DOTA–NHS
300 showed a low radiolabeling yield (<30 %). The 20-fold [¹⁷⁷Lu]Lu-daratumumab–*p*-SCN-Bn-
301 DOTA yielded 86.0 %, whereas the 30-fold reached 99.8 %. The radiolabeling yield for both
302 daratumumab–*p*-SCN-Bn-1B4M-DTPA immunoconjugates exceeded 98.0 %, and the lower
303 chelator excess was preferred to minimize chemical modification of the antibody, while also
304 reducing excess reagent use and associated chemical waste, in line with green chemistry
305 considerations (Fig. 3).

306 The radiolabeling yield for both compounds, 30-fold [¹⁷⁷Lu]Lu-daratumumab–*p*-SCN-
307 Bn-DOTA and 20-fold [¹⁷⁷Lu]Lu-daratumumab–*p*-SCN-Bn-1B4M-DTPA, exceeded 98.0 %.
308 Because of the high radiochemical purity and minimal presence of free ¹⁷⁷Lu³⁺, no additional
309 purification was required, and both were used in subsequent *in vitro* stability evaluation.

310 The *in vitro* stability of the two radioimmunoconjugates, [¹⁷⁷Lu]Lu-daratumumab–*p*-
311 SCN-Bn-DOTA and [¹⁷⁷Lu]Lu-daratumumab–*p*-SCN-Bn-1B4M-DTPA, was assessed by
312 measuring the percentage of radioactivity bound to the antibody relative to the initial value at
313 the start of the incubation. More than 95.93 % of the ¹⁷⁷Lu remained attached to daratumumab-
314 DOTA-SCN within 168 hours. In contrast, [¹⁷⁷Lu]Lu-daratumumab–*p*-SCN-Bn-1B4M-DTPA

315 showed significantly lower stability, with a notable decrease already after 48 hours, and only
316 51.91 % of the activity remaining bound to the antibody after one physical half-life of ^{177}Lu .

317 The labeling efficiency was evaluated by incubation of the daratumumab-*p*-SCN-Bn-
318 DOTA conjugate with $^{177}\text{LuCl}_3$ for a period of 5–60 min at 37 °C with gentle shaking. The
319 developed iTLC chromatograms showed clear separation between the radiolabeled
320 immunoconjugate retained at the origin ($R_f = 0$) and [^{177}Lu]Lu-*p*-SCN-Bn-DOTA migrating
321 toward the solvent front ($R_f = 1$), and free [^{177}Lu]Lu $^{3+}$ ($R_f = 0.3-0.5$).

322 Time-dependent incubation experiments showed an increase in labeling efficiency over
323 time, reaching a plateau at 50–60 min, indicating effective coordination of the metal under the
324 applied conditions (Fig. 4).

325 To verify that the antibody structure remains intact after conjugation and freeze-drying,
326 ATR-FTIR spectra were evaluated, focusing on the Amide I (1700–1600 cm^{-1}) and Amide II
327 (1580–1510 cm^{-1}) regions. Since excipient composition and residual moisture can significantly
328 affect the mid-IR region, particularly where the water bending vibration overlaps with the
329 Amide I band, placebo spectra were acquired as a control and used for spectral subtraction to
330 improve interpretation of the protein bands. To further enhance band resolution, the second
331 Savitzky–Golay derivative spectra were processed (36). The analysis of the Amide I band
332 showed that the spectra of the purified freeze-dried daratumumab and the daratumumab-*p*-
333 SCN-Bn-DOTA immunoconjugate were almost identical between 1700 and 1600 cm^{-1} (Figs.
334 S2, S3). In the second derivative spectra (Fig. 5), the dominant component remained centered
335 around 1640 cm^{-1} and no new assignments were identified, suggesting that the β -sheet-rich
336 secondary structure of the IgG was preserved after both conjugation and freeze-drying.
337 Importantly, no additional components in the 1610–1620 cm^{-1} region, commonly associated
338 with aggregation-related intermolecular β -sheet formation, were observed.

339 Raman spectra were evaluated in the Amide I (1600–1700 cm^{-1}) and Amide III (1200–
340 1350 cm^{-1}) regions to confirm the structural integrity. Importantly, neither relevant peak shifts
341 nor new bands were observed in the spectra of the purified, freeze-dried daratumumab and the
342 daratumumab-*p*-SCN-Bn-DOTA immunoconjugate, which are sensitive to changes in
343 secondary structure (Fig. 6).

344 Overall, the findings from both ATR-FTIR and Raman spectroscopy indicate that
345 conjugation of DOTA-SCN and freeze-drying did not result in detectable structural changes or
346 antibody aggregation.

347 To further confirm antibody integrity and distinguish fragmentation from aggregation,
348 SDS-PAGE was performed under both reducing and non-reducing conditions. Protein bands
349 were evaluated to assess the integrity of the intact antibody (~150 kDa) and its heavy (~50 kDa)
350 and light (~25 kDa) chains. Under reducing conditions, all samples showed the expected heavy-
351 and light-chain bands, with no additional LMWS. Under non-reducing conditions, a dominant
352 band at ~150 kDa corresponding to intact IgG was observed, with no evident high-molecular-
353 weight bands. Finally, the electrophoretic profiles support preserved structural integrity of the
354 immunoconjugates, consistent with the SE-HPLC and vibrational spectroscopic findings (Fig.
355 S4).

356

357

CONCLUSIONS

358

359 Conjugation conditions for daratumumab were optimized in carbonate buffer for both
360 BFCs: *p*-SCN-Bn-DOTA and *p*-SCN-Bn-1B4M-DTPA. Daratumumab-*p*-SCN-Bn-DOTA was
361 selected for further physicochemical characterization based on its ¹⁷⁷Lu radiolabeling yield and
362 improved *in vitro* stability compared with the *p*-SCN-Bn-1B4M-DTPA analogue. The quality
363 assessment of freeze-drying formulations showed that excipient and solvent selection strongly
364 influence product characteristics. A saline-based, buffer-free sucrose–mannitol formulation
365 containing polysorbate 20 (S.F5) showed the most favorable characteristics with low residual
moisture, rapid reconstitution, and non-detectable HMW species.

366

367 From a pharmaceutical perspective, these findings support the development of a ready-
368 to-use radiopharmaceutical kit and provide the basis for extending the approach to other
369 diagnostic and therapeutic radionuclides. In addition, the developed formulation contributes to
370 the translational advancement of theranostic strategies. Future work will focus on long-term
stability evaluation, supported by predictive modeling approaches, and *in vivo* validation.

371

372 *Acronyms.* – ADCC – antibody-dependent cellular cytotoxicity, ADCP – antibody-dependent
373 cellular phagocytosis, BFC – bifunctional chelator, CAR – chelator-to-antibody ratio, CDC –
374 complement-dependent cytotoxicity, FTIR – Fourier transform infrared spectroscopy, HMWS – high
375 molecular weight species, iTLC – instant thin-layer chromatography, LMWS – low molecular weight
376 species, mAb – monoclonal antibody, PBS – phosphate-buffered saline, RCP – radiochemical purity,
377 RIC – radioimmunoconjugate, SDS-PAGE – sodium dodecyl sulfate–polyacrylamide gel
378 electrophoresis, SE-HPLC – size exclusion high-performance liquid chromatography

379 *Authors contributions.* – Conceptualization: PA, MAL, EJI; methodology: PA, SVĐ; EJI;
380 investigation: PA, MAL, KD, MA, DK, PM, ISS; writing – original draft: PA, MAL; writing - review
381 and editing: PA, MAL, KD, MA, DK, PM, IM, AD, SVĐ, EJI; project administration: EJI; supervision:
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Table I. Composition of freeze-drying formulations

Formulation 1	0.1 mol L ⁻¹ PBS, pH=8.0; 1 % (m/V) mannitol
Formulation 2	0.01 mol L ⁻¹ PBS, pH=8.0; 1 % (m/V) mannitol
Formulation 3	Sucrose:mAb = 1:450; mannitol:sucrose = 2:1
Formulation 4	Sucrose:mAb = 1:450; mannitol:sucrose = 4:1
Formulation 5	Sucrose:mAb = 1:450; mannitol:sucrose = 2:1; 0.02 % (V/V) polysorbate 20

Batch W: ultra-pure water as solvent

Batch S: 0.9 % NaCl as solvent

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Table II. Freeze-drying protocol parameters (temperature, ramp rate, and hold time) for each segment

SEGMENT 1	Freezing: -40 °C
	Ramping 1 °C min ⁻¹
	Hold 3 h
SEGMENT 2	Annealing: -10 °C
	Ramping 0.5 °C min ⁻¹
	Hold 2 h
SEGMENT 3	Freezing: -40 °C
	Ramping 1 °C min ⁻¹
	Hold 1.5 h
SEGMENT 4	Primary drying: -10 °C
	Ramping 0.15 °C min ⁻¹
	Hold 18 h
SEGMENT 5	Secondary drying: +25 °C
	Ramping 0.15 °C min ⁻¹
	Hold 8 h
Total ca 50 hours	

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Table III. Average CAR values of daratumumab immunoconjugates

Immunoconjugate	Conjugation in phosphate buffer (pH 8.5)	Conjugation in carbonate buffer (pH 8.5)
Daratumumab-DOTA-NHS 1:20	0.00	0.18
Daratumumab-DOTA-NHS 1:30	0.00	0.10
Daratumumab-DOTA-NHS 1:50	0.20	0.46
Daratumumab-DOTA-SCN 1:20	1.27	2.31
Daratumumab-DOTA-SCN 1:30	1.75	2.66
Daratumumab-DOTA-SCN 1:50	2.76	3.77
Daratumumab-1B4M DTPA 1:20	0.40	0.36
Daratumumab-1B4M DTPA 1:30	0.65	0.76
Daratumumab-1B4M DTPA 1:50	0.78	0.85

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Table IV. Residual moisture in freeze-dried formulations

Formulation	Residual moisture (%)	
	Batch W	Batch S
F1	5.54	5.01
F2	5.60	2.73
F3	6.06	2.17
F4	6.05	2.83
F5	5.10	1.90

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Table V. Purity profile of water-based (W) and saline-based (S) batches

Formulation	Main monomer (%)		High molecular mass species (%)		Low molecular mass species (%)	
	W	S	W	S	W	S
	1	93.86	94.98	5.90	4.80	0.24
2	93.01	91.23	6.67	8.55	0.23	0.22
3	99.76	99.75	n.d.*	n.d.	0.24	0.25
4	99.73	98.90	n.d.	0.85	0.27	0.25
5	99.76	99.75	n.d.	n.d.	0.24	0.25

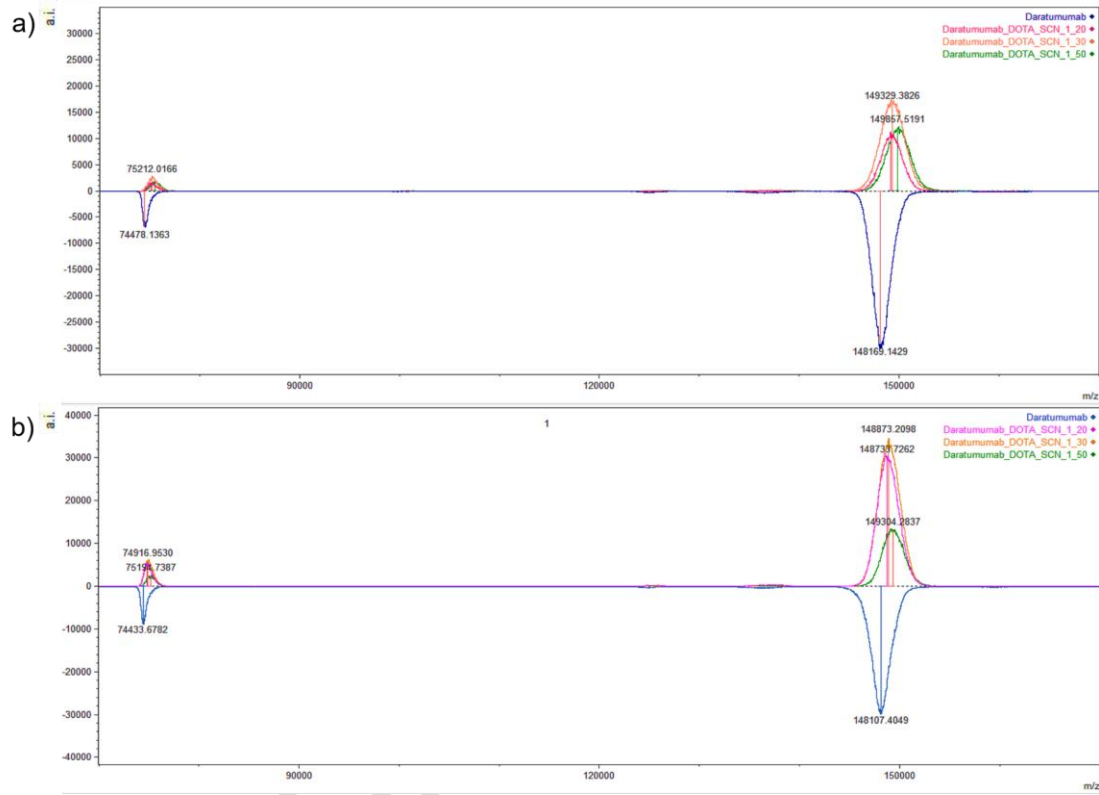
*n.d. – non-detectable (no peak observed in the predefined HMWS retention-time region under the applied SE-HPLC conditions)

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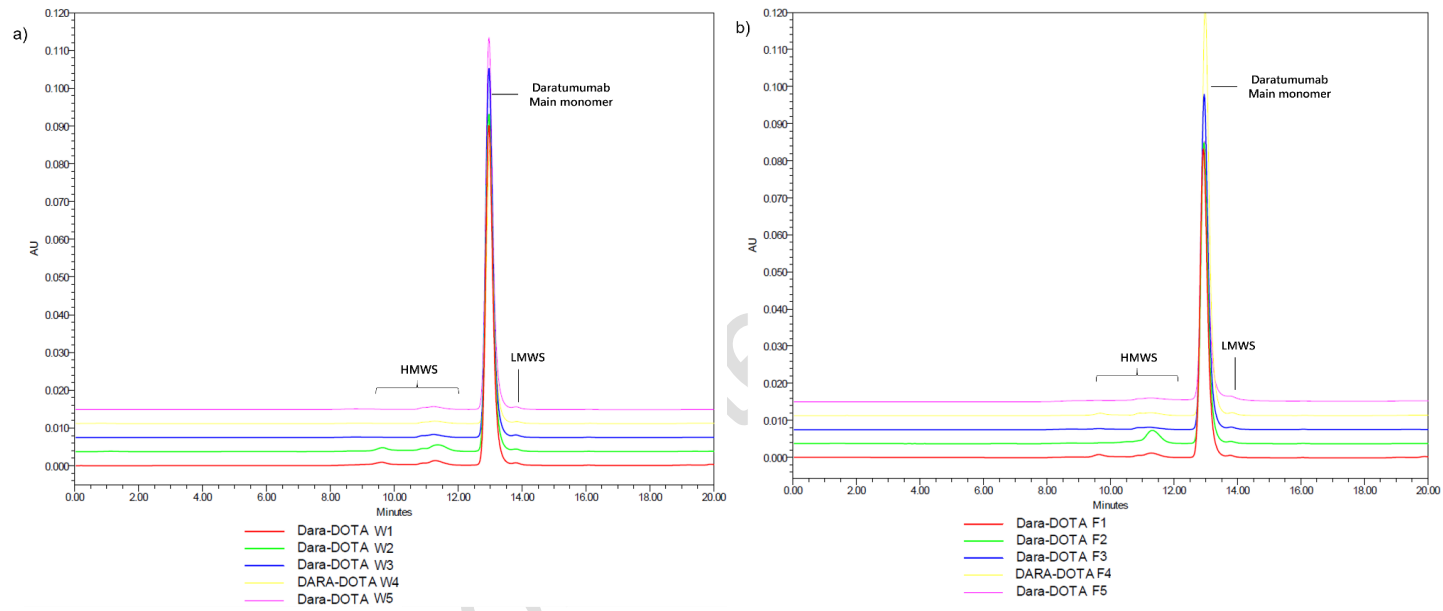
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557 Fig. 1. MALDI-TOF MS of daratumumab-*p*-SCN-Bn-DOTA conjugates in: a) phosphate buffer and b) carbonate buffer. Numbers shown refer to the center of
558 the peaks. Peaks with ~75 kDa are the doubly charged molecular ions displaying the half-MWs daratumumab-*p*-SCN-Bn-DOTA conjugates.

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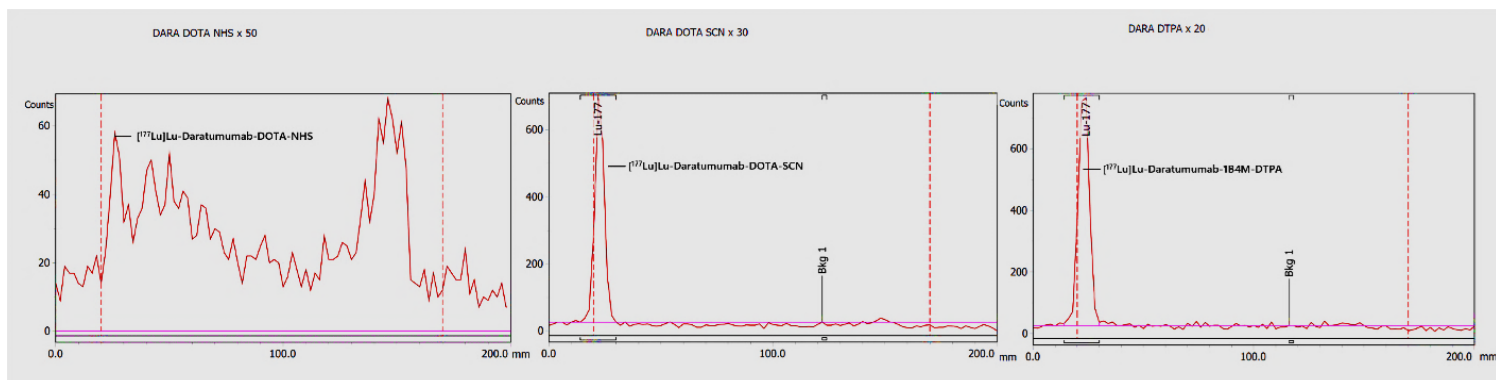


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562 Fig. 2. Overlay SE-HPLC chromatograms: a) W.F1-5 and b) S.F1-5 batches. The main monomer peak at $t_R \approx 13$ min, HMWS at $t_R \approx 9$ and 11 min, and LMWS
563 at $t_R \approx 14$ min.

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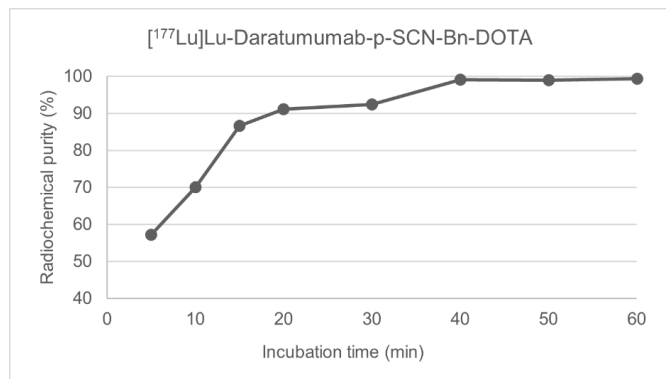
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567 Fig. 3. Representative radiochromatograms of: a) 50-fold $[^{177}\text{Lu}]\text{Lu}$ -daratumumab-DOTA-NHS, b) 30-fold $[^{177}\text{Lu}]\text{Lu}$ -daratumumab-*p*-SCN-Bn-DOTA and c)
 568 20-fold $[^{177}\text{Lu}]\text{Lu}$ -daratumumab-*p*-SCN-Bn-1B4M-DTPA.

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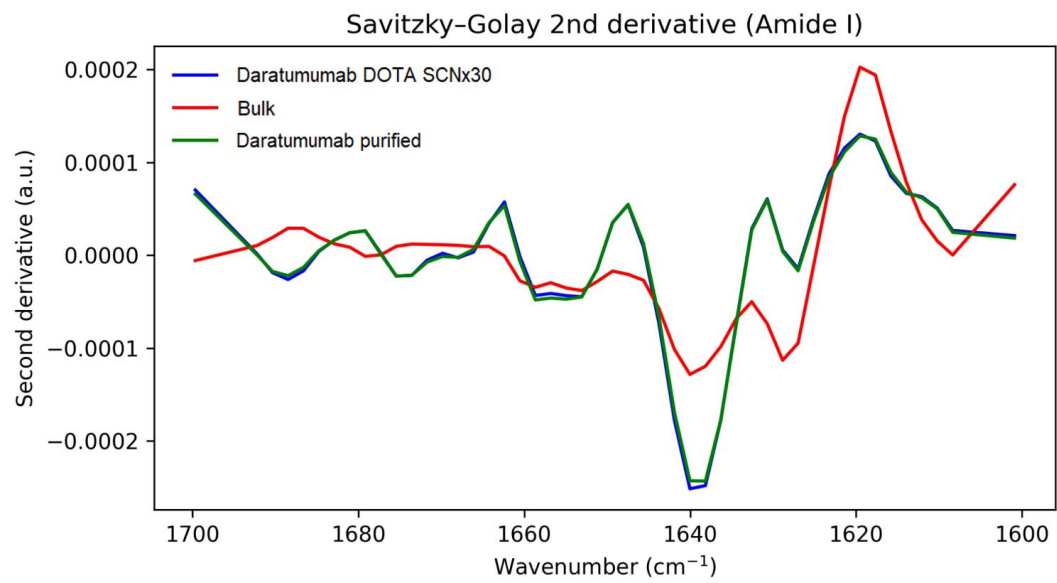
572 Fig. 4. ¹⁷⁷Lu-labelling efficiency of daratumumab-*p*-SCN-Bn-DOTA (representative measurements).

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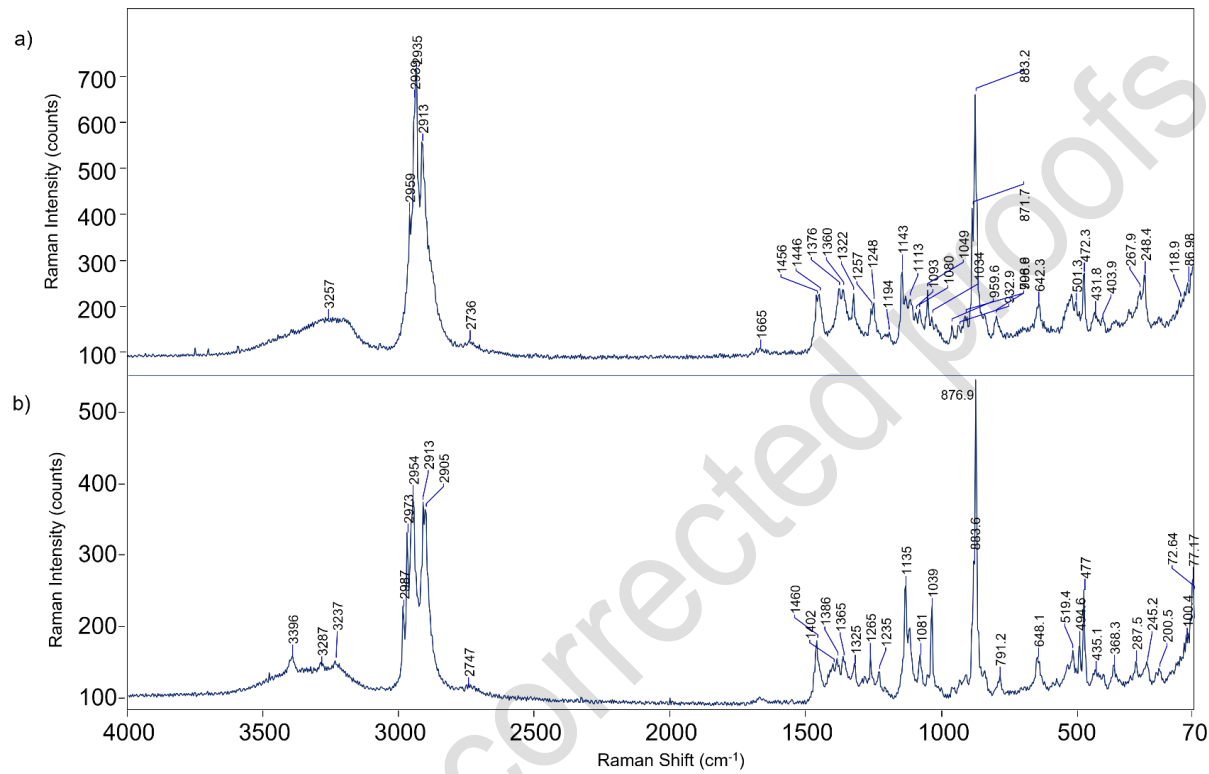
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576 Fig. 5. Second derivative spectra.

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579 Fig. 6. Representative Raman spectra of: a) purified daratumumab and b) daratumumab-*p*-SCN-Bn-DOTA immunoconjugate.

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