Daratumumab displays *in vitro* and *in vivo* anti-tumor activity in models of B-cell non-Hodgkin lymphoma and improves responses to standard chemo-immunotherapy regimens

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**ABSTRACT**

CD38 is expressed in several types of non-Hodgkin lymphoma (NHL) and constitutes a promising target for antibody-based therapy. Daratumumab (Darzalex) is a first-in-class anti-CD38 antibody approved for the treatment of relapsed/refractory (R/R) multiple myeloma (MM). It has also demonstrated clinical activity in Waldenström macroglobulinaemia and amyloidosis. Here, we have evaluated the activity and mechanism of action of daratumumab in preclinical *in vitro* and *in vivo* models of mantle cell lymphoma (MCL), follicular lymphoma (FL) and diffuse large B-cell lymphoma (DLBCL), as monotherapy or in combination with standard chemo-immunotherapy. *In vitro*, daratumumab engages Fc-mediated cytotoxicity by antibody-dependent cell cytotoxicity and antibody-dependent cell phagocytosis in all lymphoma subtypes. In the presence of human serum, complement-dependent cell cytotoxicity was marginally engaged. We demonstrated by Selective Plane Illumination Microscopy that daratumumab fully penetrated a three-dimensional (3D) lymphoma organoid and decreased organoid volume. *In vivo*, daratumumab completely prevents tumor outgrowth in models of MCL and FL, and shows comparable activity to rituximab in a disseminated *in vivo* model of blastic MCL. Moreover, daratumumab improves overall survival (OS) in a mouse model of transformed CD20<sup>+</sup> FL, where rituximab showed limited activity. Daratumumab potentiates the antitumor activity of CHOP and R-CHOP in MCL and FL xenografts. Furthermore, in a patient-derived DLBCL xenograft model, daratumumab anti-tumor activity was comparable to R-CHOP and the addition of daratumumab to either CHOP or R-CHOP led to full tumor regression. In summary, daratumumab constitutes a novel therapeutic opportunity in certain scenarios and these results warrant further clinical development.
Introduction

B-cell non-Hodgkin’s lymphoma (B-NHL) constitutes 4-5% of all hematologic neoplasia with increasing incidence in western countries. DLBCL and FL represent the most frequent aggressive and indolent NHL, accounting for approximately 55% and 20% of all lymphomas, respectively. Moreover, roughly one third of FL patients develop a histologic transformation (tFL) to DLBCL leading to a dismal prognosis. Both entities are currently treated with chemo-immunotherapy including a rituximab backbone. FL responses are usually high, although recurrence occurs in the majority of the cases. In DLBCL, currently classified into germinal center type (GCB) or activated B-cell type (ABC), treatment is not guided by subtype, and responses to chemo-immunotherapy are normally higher in the GCB subtype. Nevertheless, a portion of DLBCL (20%) do not respond to this regimen. Several second-generation anti-CD20 antibodies, such as the Food and Drug Administration-approved obinutuzumab have been clinically tested to overcome these limitations. However, an alternative evolving therapeutic approach is to target a different antigen. In this regard, both FL and DLBCL originate in the germinal center (GC) and consequently express high levels of CD38, making this molecule an attractive therapeutic target.

MCL is a rare NHL (6% of all NHL) with an aggressive evolution and clinically challenging. Its frontline therapy, although heterogeneous, typically consists of rituximab-based chemo-immunotherapy followed by autologous stem cell transplantation and/or rituximab maintenance. Even with intensive therapy, MCL patients ultimately relapse. Novel targeted therapies currently approved for R/R MCL include the mTOR inhibitor temsirolimus, the immunomodulatory agent lenalidomide, the proteasome inhibitor bortezomib, also approved in front-line, and the BTK inhibitor ibrutinib that achieves the highest response rates. However, MCL patients failing ibrutinib treatment have very limited therapeutic options. In this situation, where virtually all MCL cases express some level of CD38, this antigen represents a potential alternative target to be explored. Moreover, CD38 is associated with nodal disease and poorer survival, and high CD38 expression correlates with poor in vitro response to bortezomib. Thus, targeting CD38 could hold promise as a strategy for MCL, also in bortezomib-resistant tumors.

CD38 is present at high levels in bone marrow (BM) precursor cells and it is downregulated in resting normal B cells. The molecule is re-expressed at high density once naïve B lymphocytes are activated, and peaks when B cells enter the GC. Terminally differentiated plasma cells and their pathological counterparts express the highest surface density among human cells, while it is completely absent in memory B cells. CD38 behaves simultaneously as an enzyme and as a receptor. The extracellular domain of CD38 contains an enzymatic site that can generate cyclic ADP ribose (cADPR) and ADPR from nicotine adenine dinucleotide (NAD⁺). This control of adenosine synthesis by CD38 may be important in the context of the characteristic immunosuppressive tumor microenvironment.

Daratumumab (Darzalex) is a first-in-class, human IgG1κ monoclonal antibody (mAb) that targets the CD38 epitope. It was approved by the Food and Drug Administration in 2015 as a monotherapy for patients with MM, who have received at least three prior therapies. Currently, daratumumab has been approved in combination with dexamethasone plus either lenalidomide or bortezomib, or pomalidomide for the treatment of relapsed MM patients. Daratumumab has a broad-spectrum killing activity in MM engaging complement-dependent cytotoxicity (CDC), antibody-dependent cellular cytotoxicity (ADCC), antibody-dependent cellular phagocytosis (ADCP), and apoptosis. Moreover, daratumumab modulates the enzymatic activity of CD38 and induces an immunomodulatory role in MM by depleting CD38-immune suppressive cells contributing to its antitumor activity. In chronic lymphocytic leukemia (CLL), we have demonstrated that daratumumab induces cytotoxic activity in vitro via ADCC and ADCP in primary CLL cells and cell lines. In vivo, daratumumab significantly prolongs OS of animals in systemic CLL murine models. Daratumumab also affects tumor-microenvironment interactions by blocking CLL homing and dissemination to secondary lymphoid organs in vitro and in vivo.

In the present study, we aimed to investigate in vitro and in vivo activity of daratumumab on MCL, FL, and DLBCL cells as monotherapy and in combination with standard therapies.

Methods

Therapeutic drugs

Daratumumab (Darzalex, anti-CD38 mAb, IgG1) and the isotype control mAb (CNTO 3930, IgG1) were provided by Janssen. Rituximab (Mabthera, anti-CD20 mAb, IgG1) and the chemotherapy regimen CHOP (cyclophosphamide, doxorubicin, vincristine, and prednisone) were obtained from the department of pharmacy of the Hospital Clínic of Barcelona.

Subcutaneous pre-emptive mouse models

SCID mice (Janvier Laboratories) were subcutaneously (sc) injected with 10x10⁶ RL-luc cells in the FL model or 10x10⁶ REC-1 cells in the MCL model, respectively, following a protocol approved by the Animal Testing Ethic committee of the University of Barcelona and Generalitat de Catalunya (Protocol # 9971). Mice were randomly assigned into cohorts of six mice per group and received one intraperitoneal (ip) injection of 10 mg/kg of daratumumab or isotype control every other week, starting the day of cell inoculation.

Patient-derived DLBCL xenograft model

ST1361 is a DLBCL patient-derived xenograft model developed using a DLBCL tumor originated in a metastatic site from a 58-year-old chemotherapy-naïve Hispanic male. ST1361 was GCB subtype transformed from an Epstein-Barr virus negative FL tumor.

The DLBCL tumor was homogeneously chipped into fragments of similar size and injected sc into SCID mice. Treatment was initiated when mean tumor volume was approximately 150-250 mm³. Daratumumab 20 mg/kg was administered weekly for three weeks alone or in combination with CHOP (20 mg/kg cyclophosphamide, 1.25 mg/kg doxorubicin, 0.2 mg/kg vincristine [intravenously on Day 0], and 0.15 mg/kg prednisone [Days 0-4]; once a day (QD) for five days) or R-CHOP (10 mg/kg rituximab [intraperitoneally on Day 0]+ CHOP as before).

Statistical analysis

Unpaired and paired t-tests were used to assess statistical differences between two groups using GraphPad Prism software 4.0. For Kaplan-Meier survival curves, SPSS19 software was used.
### Table 1. B-cell non-Hodgkin’s lymphoma cell lines characterization and daratumumab activity.

| Cell line     | NHL subtype | CD38 sABC | %CD38 | MFR C D38 | %CD20 | MFR C D20 | %CD46 | %CD55 | %CD59 | %ADCC | %ADCP | %CDC |
|---------------|-------------|-----------|-------|-----------|-------|-----------|-------|-------|-------|-------|-------|------|
| Jeko          | MCL         | 16253     | 84    | 22        | 100   | 46        | 92    | 98    | 61    | 58    | 41    | 7    |
| REC-1         | MCL         | 106234    | 100   | 155       | 100   | 77        | 99    | 99    | 88    | 60    | 46    | 18   |
| HBL2          | MCL         | 25317     | 100   | 92        | 100   | 99        | 98    | 96    | 98    | 28    | 21    | 0    |
| Mino          | MCL         | 127458    | 100   | 51        | 100   | 42        | 97    | 90    | 96    | 32    | 35    | 0    |
| UPN1          | MCL         | 54089     | 99    | 97        | 97    | 8         | 98    | 99    | 63    | 36    | 73    | 0    |
| Z138          | MCL         | 186239    | 100   | 120       | 100   | 28        | 100   | 75    | 69    | 53    | 89    | 0    |
| DOHH2         | FL          | 22369     | 87    | 20        | 99    | 31        | 100   | 76    | 98    | 25    | 38    | 0    |
| SC-1          | FL          | 71726     | 100   | 40        | 32    | 4         | 95    | 96    | 83    | 41    | 81    | 1    |
| RL            | FL          | 132684    | 96    | 76        | 93    | 49        | 100   | 94    | 97    | 52    | 45    | 6    |
| WSU-FSCCL     | FL          | 89707     | 97    | 354       | 8     | 2         | 82    | 81    | 94    | 67    | 92    | 7    |
| Toledo        | DLBCL       | 125339    | 99    | 186       | 89    | 13        | 94    | 99    | 97    | 34    | 50    | 14   |
| WSU-DLCL2     | DLBCL       | 14926     | 94    | 46        | 99    | 137       | 95    | 95    | 98    | 45    | 15    | 1    |
| SU-DHL-6      | DLBCL       | 174033    | 92    | 383       | 98    | 172       | 97    | 44    | 31    | 64    | 56    | 5    |
| SU-DHL-4      | DLBCL       | 96623     | 96    | 108       | 100   | 45        | 93    | 95    | 61    | 34    | 8     |
| Daudi         | BL          | 415667    | 100   | 276       | 99    | 30        | 88    | 17    | 9     | 63    | 79    | 96   |

1Percentage of positive cells for CD38, CD20, CD46, CD55 and CD59 determined by flow cytometry, referred to isotype control; 2 sABC: number of surface antibodies bound per cell evaluated by Quantibrite CD38-PE; 3M FIR: mean fluorescence ratio referred to isotype control; 4 Percentage of antibody-dependent cellular cytotoxicity (ADCC) and antibody-dependent cellular phagocytosis (ADCP) induction at 1 µg/mL daratumumab. Percentage of CDC induction at 10 µg/mL daratumumab MCL mantle cell lymphoma, FL: follicular lymphoma; DLBCL: diffuse large B-cell lymphoma; NHL: non-Hodgkin lymphoma; BL: Burkitt lymphoma.

Additional methodologic details are described in the Online Supplementary Materials and Methods.

### Results

**Daratumumab induces cell killing in the presence of external effectors in B-NHL**

We first assessed by calcein-AM release assay, the ability of daratumumab to induce ADCC, ADCP and CDC in MCL, FL and DLBCL cell lines. Using peripheral blood mononuclear cells (PBMC) from healthy donors as external effectors we demonstrated that daratumumab engages ADCC in a panel of B-NHL cell lines (Table 1). Daratumumab induced a significant dose-dependent cell lysis reaching its maximum antitumor activity at 1 µg/mL on MCL cell lines (mean ± standard deviation (SD)=45±14%) (Figure 1A), at 0.1 µg/mL on FL cell lines (mean ± SD=48±18%) (Figure 1B), and at 0.1 µg/mL on DLBCL cell lines (mean ± SD=49±11%) (Figure 1C). The degree of ADCC induction did not correlate with the number of CD38 molecules per NHL cell specific antibody binding capacity (sABC) (r²=0.1988; Online Supplementary Figure S1A and Table 1). A phagocytosis assay using mouse macrophages as effector cells was set up to determine daratumumab-mediated ADCP. Daratumumab induced significant ADCC at 1 µg/mL in MCL cell lines (mean ± SD=54±24; Figure 1D), FL cell lines (mean ± SD=64±26%; Figure 1E) and in DLBCL cell lines (mean ± SD=39±18%: Figure 1F). The degree of ADCP induction did not correlate with the number of CD38 sABC on NHL cell lines (r²=0.2270; Online Supplementary Figure S1B and Table 1).

Interestingly, the daratumumab-opsonized cells that were exposed to macrophages and not phagocytosed, suffered a lost in CD38 surface expression, irrespective of the degree of ADCP induction (Online Supplementary Figure S2), in a process that may resemble trogocytosis as previously described in MM studies.31,32

We next evaluated the CDC activity and observed that, similar to that reported for CLL cells,33 daratumumab induced low to marginal cell death in the presence of normal human serum (NHS 10%) (Table 1). Very similar results were obtained in these CDC experiments when 50% NHS was applied. The high expression of the complement regulatory proteins (CRP) CD46, CD55 and CD59, and an insufficient number of CD38 molecules per cell on NHL cells may explain the low CDC activity (Table 1).

Collectively, daratumumab effectively kills CD38⁺ NHL cells through engagement of ADCC and ADCP but not CDC, independently of the expression levels of CD38.

**Daratumumab enters and distributes homogeneously within a 3D lymphoma model**

To model the compact aggregates of lymphoma cells growing in the patients’ lymph nodes, 3D spheroids were generated by the hanging drop method,34 using RL-GFP cells previously reported as an useful tool for antibody therapy studies.34 We first sought to determine the degree to which daratumumab can penetrate these lymphoma aggregates depending on dose (1 and 10 µg/mL mAb) and time of exposure (4, 24 and 48 hours). First, we measured the percentage of mAb diffusion in the spheroid, representing the total amount of the mAb in the spheroid. The 3D reconstruction images obtained by SPIM (Figure 2A) revealed a maximum diffusion of daratumumab at 1 µg/mL after 48 hours of treatment (Figure 2B). Next, maximum depth of daratumumab in the spheroids was determined under the same conditions and represented as percentage of mAb penetration in the spheroid. As observed for the diffusion, a maximum penetration of daratumumab at 1 µg/mL after 48 hours of treatment was achieved.

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**References:**

1. Percentage of positive cells for CD38, CD20, CD46, CD55 and CD59 determined by flow cytometry, referred to isotype control; 2 sABC: number of surface antibodies bound per cell evaluated by Quantibrite CD38-PE; 3M FIR: mean fluorescence ratio referred to isotype control; 4 Percentage of antibody-dependent cellular cytotoxicity (ADCC) and antibody-dependent cellular phagocytosis (ADCP) induction at 1 µg/mL daratumumab. Percentage of CDC induction at 10 µg/mL daratumumab MCL mantle cell lymphoma, FL: follicular lymphoma; DLBCL: diffuse large B-cell lymphoma; NHL: non-Hodgkin lymphoma; BL: Burkitt lymphoma.
Moreover, a significant dose-dependent reduction of the spheroid volume was observed after daratumumab treatment compared to isotype control (mean ± SD=15±6%, 20±7%, respectively at 1 and 10 µg/mL) (Figure 2D) showing a direct effect of daratumumab on tumor growth in 3D. This observation was validated in additional lymphoma cell lines. Daratumumab moderately but significantly (P<0.0001) reduced the spheroid volume in all the cell lines analyzed, albeit in a different degree (Online Supplementary Figure S3A-B), that was accompanied in most of cases by a concomitant decrease in the cell number. However, this observation did not apply to HBL-2 and WSU-DLCL2 cell lines, where daratumumab may just generate a tighter spheroid (Online Supplementary Figure S3C).

These data demonstrate that daratumumab efficiently penetrates a lymphoma 3D structure and moderately reduces the growth of spheroids in the absence of external effectors.

Daratumumab prevents tumor outgrowth in MCL and FL xenografts

As a first test of daratumumab activity in vivo in NHL models, we inoculated SCID mice sc with two cell lines that we had previously set up: RL cells to model tFL,35 and REC-1 cells to model MCL.36 SCID mice have active natural killer (NK) cells and macrophages that can act as effector cells to engage daratumumab activity, as we have shown using in vitro models. Mice were treated with daratumumab or an isotype control (20 mg/kg) starting on the day of tumor injection. Subsequently, animals received antibody injections every other week (10 mg/kg) for a total of three doses and were sacrificed at day 30 (REC-1) (Figure 3A) or day 34 (RL) (Figure 3B). Daratumumab prevented tumor formation in all mice injected with REC-1, and in 5 of 6 mice injected with RL cells. In this latter model, a 20 mm3 RL tumor was found in one daratumumab-treated mouse at the endpoint (day 34). As the RL cells inoculated expressed the luciferase gene, bioluminescence analysis at different time points further supported these results (Figure 3B). All animals receiving isotype control antibody formed tumors of at least 100 mm3 within a mean of 18 days (REC-1) and 21 days (RL) and were bigger than 1,000 mm3 at the end of the experiment.

Daratumumab prolongs OS and reduces tumor infiltration in systemic xenograft models of MCL and FL

Next, we generated a model for blastic MCL by inoculating Z-138 cells intravenously into SCID mice, and one week later mice were treated weekly with either daratumumab (D), rituximab (R) or isotype (I) control (20/10/10/10 mg/kg). Seven of ten isotype-treated mice had to be sacrificed between days 57 - 89 due to systemic signs of disease (Figure 4). Daratumumab significantly improved OS (I vs. D: ***P<0.001), with 90% of the mice treated with daratumumab surviving for over 107 days, (Figure 4A) compared to just 10 % alive in the controls. OS for the daratumumab group was slightly superior but did not reach statistical significance (D vs. R: ns, P=0.2907), compared to that observed in the rituximab-treated mice.
(I vs. R: **P<0.01), where 70% of the mice were alive at day 107 (Figure 4). Mean survival was 73 days for the control group, 97 for rituximab and 103 for daratumumab groups. On autopsy, the disease had spread to the brain, BM, spleen and lungs (Online Supplementary Figure S4A). Daratumumab treatment reduced the disease burden in the brain, BM, and spleen by 93%, 63%, and 48%, respectively (Online Supplementary Figure S4B). Rituximab-treated mice showed a similar reduction in tumor load to that achieved by daratumumab in the brain (73%) and BM (69%) compared to that observed with the isotype control, while no effect was observed in spleen infiltration. Daratumumab treatment reduced the disease burden in the brain, BM, and spleen by 93%, 63%, and 48%, respectively (Online Supplementary Figure S4B). Rituximab-treated mice showed a similar reduction in tumor load to that achieved by daratumumab in the brain (73%) and BM (69%) compared to that observed with the isotype control, while no effect was observed in spleen infiltration.

In the FL systemic model, CD20™ WSU-FSCCL cells (Table 1) were intravenously inoculated in SCID mice and one week later mice were treated weekly with daratumumab, rituximab or isotype control (20/10/10/10 mg/kg). All control-treated and rituximab-treated mice rapidly succumbed to the disease and died or had to be euthanized due to severe disease signs by day 40 (control group) and day 62 (rituximab group), respectively (Figure 4). In contrast, in the daratumumab-treated group, OS was significantly prolonged (I vs. D: ***P<0.001; I vs. R: **P<0.01; D vs. R *P<0.05). Mean survival was 35 days for the control group, 41 for rituximab and 60 for daratumumab. By the end of the experiment (day 90), 30% of mice treated with daratumumab were cured (Figure 4). On autopsy, WSU-FSCCL cells showed systemic dissemination of disease in the brain, BM and spleen (identified as huCD45+/CD19+CD10+; Online Supplementary Figure S4C). Daratumumab robustly decreased dissemination to the brain (90%, P<0.01) compared to the control group, while rituximab was not effective (Online Supplementary Figure S4D). Homing of WSU-FSCCL cells to the spleen and BM was less prominent. There were no significant differences in the reduction of dissemination to the spleen and BM by daratumumab compared to the rituximab-treated group (Online Supplementary Figure S4C).

Altogether, these data confirm that daratumumab shows comparable to superior activity with respect to rituximab in systemic models of MCL and FL.

**Daratumumab synergizes with R-CHOP to reduce tumor burden in pre-clinical models of MCL and FL.**

The synergy between daratumumab and rituximab and/or chemotherapy regimens was evaluated in MCL (REC-1) and tFL (RL) xenograft heterotopic models. SCID mice were inoculated subcutaneously with NHL cells and when tumors became palpable, mice were randomly assigned into the following groups: i) Isotype ii) daratumumab iii) CHOP iv) D-CHOP v) R-CHOP and vi) R-D-CHOP. Tumor volume was assessed twice a week and tumor weights at sacrifice. In the MCL model (Figure 5A), daratumumab induced significant anti-tumor activity.
compared to the isotype control (42%, *P*< 0.01). CHOP alone marginally reduced tumor growth (*P*=0.182) and no difference was observed between daratumumab and D-CHOP groups (*P*=0.451). Importantly, adding rituximab to D-CHOP treatment (R-D-CHOP) induced a significant reduction in tumor volume compared to the isotype control (86%, *P*<0.001) and also compared to R-CHOP (73%, *P*<0.001) and D-CHOP (51%, *P* = 0.05) alone. Similarly, in the tFL model (Figure 5C-D), daratumumab treatment inhibited tumor growth compared to isotype control (36%, *P*<0.001). On the other hand, the combination D-CHOP inhibited tumor growth significantly better than daratumumab as a single agent (57%, *P*<0.001), resembling that achieved by R-CHOP (67%, *P*<0.001). In addition, the combination regimen of R-D-CHOP had a remarkable effect in impeding tumor growth (84% inhibition, *P*<0.001), which was superior than that achieved by each combination regimen separately (*P*<0.001). The results of the RL-luc model were further validated by bioluminescence and quantification of the signal captured at the endpoint (Online Supplementary Figure S5A-B).

Moreover, tissue sections of selected mice of each group were analyzed by immunohistochemistry for the proliferation marker pH3 and the angiogenesis marker CD31, further confirming these results (Online Supplementary Figure S5C).

In conclusion, these data demonstrate the value of the combination R-D-CHOP as a new therapeutic strategy in MCL and tFL superior to the standard-of-care (R-CHOP).

**Daratumumab induces tumor growth inhibition and prolongs survival in a patient-derived DLBCL xenograft model in combination with R-CHOP**

To further evaluate the activity of daratumumab alone or in combination with the standard-of-care therapy we used a clinically more relevant patient-derived DLBCL xenograft model (ST1361) with detectable CD38 expression by immunohistochemistry (Online Supplementary Figure S6). Identical to the MCL and tFL models, all individual treatments were capable of reducing tumor volume over time (*P*<0.001, Figure 6A); however, daratumumab alone was strikingly efficient with an anti-tumor activity...
comparable to R-CHOP ($P<0.001$, Figure 6A). Furthermore, the addition of daratumumab to either CHOP or R-CHOP led to full tumor regression ($P<0.001$, Figure 6A). Importantly, tumor volume negatively correlated very well with the OS of xenograft-bearing mice, regardless the treatment group (Figure 6B). Thus, mice treated with either daratumumab alone or R-CHOP displayed identical OS and, in line with the full tumor regression achieved, daratumumab combined with either CHOP or R-CHOP resulted in 100% survival at the end of the experiment ($P<0.05$ after Bonferroni multiplicity correction, Figure 6A). Remarkably, full tumor regression and 100% survival at day 60 were observed even when daratumumab treatment was ceased as early as 14 days (total of three doses) after tumor cell inoculation. Collectively, these results indicate that tumor cells were completely eliminated within a short time and that no tumor escape occurred in this DLBCL patient-derived model.

**Discussion**

In this study we have demonstrated that daratumumab engages Fc-mediated cell killing of NHL cells by ADCC and ADCP. Variation in the extent of ADCC and ADCP could not be explained by the differences in CD38 expression levels. Similar observations were also reported for daratumumab-induced ADCC and ADCP in CLL cells. The capacity of daratumumab to induce ADCC and phagocytosis may to some extent be related to other factors, such as the expression of different types of FcR, KIRs and Natural Cytotoxic Receptors (NCR) on NK cells. Moreover, the expression of the so-called ‘don’t eat me’ signals, such as CD47, on tumor cells plays an important role in regulating phagocytosis. In fact, CD47 expression has been found to be increased on MCL, FL and DLBCL cells compared to normal cells, conferring a worse clinical prognosis. Moreover, in agreement with previous results in MM, the daratumumab-opsonized cells that were exposed to macrophages and not phagocytosed, suffered a loss in CD38 surface expression. This event may reduce the ability of daratumumab to kill lymphoma cells via CDC and ADCC, compromising the therapeutic efficacy of daratumumab. However, MM studies demonstrated that this phenomenon occurs early in the treatment irrespective of their clinical responses. Therefore, CD38 reduction via trogocytosis should not necessarily be considered as an escape mechanism from daratumumab treatment. On the contrary, trogocytosis may be beneficial and represent a novel mechanism of action of daratumumab, as there is also a transfer of tumor cell membrane fragments containing important signaling molecules such as CD56, CD49d and CD138 in this process. The decreased expression of these adhesion molecules in tumor cells may
compromise their interaction with the tumor microenvironment as we reported in a previous work.\textsuperscript{3}\textsuperscript{4}

Daratumumab did not induce CDC in FL, or DLBCL or MCL cell lines. Flow cytometry analysis demonstrated that MCL and FL cells show medium to high expression of CD38 while DLBCL cells tend to express higher levels of CD38. However, the number of molecules per cell in these NHL cell lines was lower than that found on the CDC-sensitive Daudi Burkitt lymphoma cell line, suggesting a threshold for CD38-targeted CDC lysis. In addition, this low induction of CDC was also associated with high expression of the CRP CD55 and CD59 that were lower in Daudi cells. These observations are in line with our previous data obtained for CLL cell lines and primary cells.\textsuperscript{30}

Overall, the baseline expression of CD38/CD55/CD59 appears to be associated with the response to daratumumab-induced CDC in vitro in FL, DLBCL, and MCL cells. These results mirrored those obtain with the anti-CD20 rituximab in CLL cells.\textsuperscript{39}

We have shown for the first time that daratumumab, at clinically achievable doses, effectively penetrates a lymphoma organoid in vitro model. A moderate but significant reduction of the sphere volume was observed with daratumumab in 3D lymphoma models, even though no significant effect was observed in a 2D model (data not shown). These results agree with daratumumab efficacy against subcutaneous tumors in vivo (Figure 5 and Figure 6).

In vivo results support the ability of daratumumab to prevent the outgrowth of MCL and FL cells, when administered in a prophylactic setting, prior to tumor development. These results open a window of opportunity for daratumumab as an alternative to rituximab maintenance therapy in these models, in the context of a complete response after the induction therapy.\textsuperscript{26,41} Moreover, we have demonstrated that daratumumab shows single agent activity both in systemic and subcutaneous models of NHL, when administered after disease onset. More importantly, the data from the FL model using WSU-FSCCL cells point to a possible role for daratumumab as a therapeutic alternative to rituximab in NHL with reduced CD20 expression. Previous studies have shown a substantial and rapid reduction of CD20 in lymphoma patients treated with rituximab, which has been linked to the development of acquired resistance.\textsuperscript{40,46} Thus, daratumumab treatment may be considered in these scenarios where anti-CD20 resistance has developed due to antigen shaving.\textsuperscript{44} In addition, our results indicate that daratumumab significantly improves long-term survival when used as a single agent, indicating that it may be also an alternative to rituximab in NHL CD20\textsuperscript{high}, evidenced by the comparable activity of both antibodies in the blastic MCL model using Z-138 cells. We have also observed a remarkable effect of daratumumab in the tumor cell dissemination to the brain in FL and MCL systemic models. This may be related to the capacity of daratumumab to cross the hematooencephalic barrier. In fact, daratumumab has shown efficacy in CNS plasmocytoma\textsuperscript{45} and extramedullary myeloma.\textsuperscript{46}

Finally, we have examined the combination of daratumumab with the standard-of-care therapy R-CHOP. Our data suggest that daratumumab significantly potentiates R-CHOP activity, inducing higher rates of tumor regression in MCL and FL. Moreover, we analyzed the efficacy of daratumumab in a DLBCL patient derived mouse xenograft, generated from a transformed FL, and usually less responsive to chemotherapy, as shown in our results where indeed the CHOP regimen displayed limited activity. In this model, daratumumab in combination only with CHOP, in the absence of rituximab leads to a complete abrogation of tumor growth. This strong synergy may be explained by the induction of stress-related cytokines by chemotherapeutic agents that can effectively target cancer cells for removal by the innate immune system through macrophage infiltration and phagocytic activity,\textsuperscript{47} potentially augmenting the described ability of daratumumab to engage ADCP.\textsuperscript{23}

In the phase II clinical trial CARINA, LYM2001 in R/R NHL, daratumumab monotherapy did not yield the expected results in heavily pretreated FL and DLBCL patients, while no results could be concluded in MCL because of insufficient patient recruitment. However, we have identified certain NHL scenarios were daratumumab shows comparable antitumor activity as rituximab (blastic

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**Figure 6. Daratumumab combined with R-CHOP in a patient-derived diffuse large B-cell lymphoma xenograft model.** Fragments from a patient-derived diffuse large B-cell lymphoma (DLBCL) were injected subcutaneously into SCID mice (n=8-10) and staged to approximately 150-250 mm\(^3\) mean tumor volume to measure (A) tumor growth and (B) overall survival (OS). 20 mg/kg daratumumab was administered weekly alone or in combination with CHOP (20 mg/kg cyclophosphamide, 1.25 mg/kg doxorubicin, 0.2 mg/kg vincristine, and 0.15 mg/kg prednisone) or R-CHOP (10 mg/kg rituximab+ CHOP) for a total of three weeks.
MCL) and other where exhibits superiority (transformed CD20dim FL). Moreover, daratumumab potentiates the antitumor activity of CHOP and R-CHOP in all three models.

In addition, as demonstrated in MM, it is likely that daratumumab may modulate the activity and frequency of CD38 expressing immune suppressive cell populations present in NHL, an effect not exerted by rituximab, and therefore may offer a superior overall antitumor effect. These may be of special relevance in FL and DLBCL where infiltration of diverse T and myeloid subpopulations is specially prominent. The immune-compromised mouse models used in this study preclude the analysis of this immune-modulating activity of daratumumab.

In conclusion, our findings warrant further clinical development of daratumumab in NHL providing a strong rationale for examining its clinical efficacy in different scenarios, including maintenance therapy after induction therapy, cases with anti-CD20 resistance, FL histologic transformation, and as frontline in combination with the standard of care (CHOP/R-CHOP).

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