AGENTS TARGETING CD138 AND USES THEREOF

CROSS REFERENCE TO RELATED APPLICATIONS

This application claims the benefit of U.S. provisional application 61/016,630, filed December 26, 2007, which is incorporated herein by reference in its entirety.

FIELD OF THE INVENTION

The present invention relates to improved targeting agents for the antigen CD138 as well as compositions comprising the targeting agent and methods employing them.

BACKGROUND

CD138, which acts as a receptor for the extracellular matrix, is overexpressed on multiple myeloma (MM) cells and has been shown to influence MM cell development and/or proliferation. CD138 is also expressed on cells of ovarian carcinoma, kidney carcinoma, gall bladder carcinoma, breast carcinoma, prostate cancer, lung cancer, colon carcinoma cells and cells of Hodgkin's and non-Hodgkin's lymphomas, chronic lymphocytic leukemia (CLL) to name just a few.

The publications and other materials, including patents, used herein to illustrate the invention and, in particular, to provide additional details respecting the practice are incorporated by reference. For convenience, the publications are referenced in the following text by author and date and/or are listed alphabetically by author in the appended bibliography.

Tassone et al. (2004) have reported excellent binding of the murine IgG1 antibody B-B4 to the CD138 antigen expressed on the surface of MM cells. Tassone also reported high cytotoxic activity of the immunoconjugate B-B4-DM1, which comprises the mytansinoid DM1 as an effector molecule, against multiple myeloma cells (see also US Patent Publ. 20070183971).

There remains a need for a targeting agent, in particular a targeting antibody based on B-B4 that is devoid of certain properties and/or functions associated with B-B4. Such a targeting antibody may comprise one or more antibody regions of a human antibody. There is, in particular a need for a chimerized antibody based on B-B4 that binds CD138 as effectively as B-B4 but can be administered to humans without significant side effects. There also remains a need for a targeting agent having binding affinity that exceeds the binding affinity of B-B4. There is also a need for such a B-B4 based targeting agent that shows one or more advantageous properties relative to its murine counterpart. Those properties include improved antigen binding, in particular of

CD138 expressing tumor cells and cells accessory thereto or more homogenous binding.

SUMMARY OF THE INVENTION

The present invention is directed at a method for homogenous binding to CD138 comprising:

providing an engineered targeting antibody, wherein said engineered targeting antibody comprising

an antigen binding region against CD138, wherein said antigen binding region is of a non-human antibody, and

a further antibody region, wherein at least part of said further antibody region is of a human antibody, and

administering said engineered targeting antibody to CD138 expressing cells,

wherein said engineered targeting antibody homogenously binds CD138 expressed on said CD138 expressing cells.

The present invention is also directed at an isolated polypeptide comprising an amino acid sequence of an immunoglobulin heavy chain or part thereof, wherein said immunoglobulin heavy chain or part thereof has at least 70%, at least 80%, at least 90%, at least 95% or at least 98% sequence identity with SEQ ID NO:1, wherein a targeting agent comprising said immunoglobulin heavy chain or part thereof targets CD138.

Said immunoglobulin heavy chain or part thereof may have at least 80%, at least 85%, at least 90%, at least 95% sequence identity with residues 31 to 35, residues 51 to 68 and residues 99 to 111 of SEQ ID NO:1 and said targeting agent may be an engineered targeting antibody.

A constant region of said immunoglobulin heavy chain or said part thereof may be an IgG4 isotype constant region.

Said targeting agent may be a mouse human chimeric antibody.

Said targeting agent or engineered targeting antibody may be a humanized antibody.

The isolated polypeptide may further comprise an amino acid sequence of an immunoglobulin light chain or part thereof, wherein said immunoglobulin light chain or part thereof may have at least 70%, at least 80%, at least 90%, at least 95% or least 98% sequence identity with SEQ ID NO:2.

The isolated polypeptide may further comprise an amino acid sequence of an immunoglobulin light chain or part thereof, wherein said immunoglobulin light chain or part thereof has at least 75%, at least 85%, at least 95% or at least 97% sequence identity with residues 24 to 34, residues 50 to 56 and residues 89 to 97 of SEQ ID NO:2.

Said immunoglobulin heavy chain may be identical to the sequence of SEQ ID NO:1.

Said immunoglobulin light chain may be identical to the sequence of SEQ ID NO:2.

The present invention is also directed at an engineered targeting antibody which recognizes CD138 comprising
an antigen binding region against CD138, wherein said antigen binding region is of a non-human antibody, and
a further antibody region, wherein at least part of said further antibody region is of a non-human antibody, wherein said engineered targeting antibody
(a) binds CD138 with a binding affinity that exceed the binding affinity of said non-human antibody; and/or
(b) provides for homogenous binding to CD138 of CD138 expressing cells.

Said further antibody region may be at least one constant region comprising a heavy chain constant region or a part thereof that is of a human antibody, and wherein said engineered antibody is of an IgG4 isotype.

Said engineered targeting antibody may be a chimeric antibody and said non-human antibody may be B-B4.

Said engineered targeting antibody may be a humanized antibody and said non-human antibody may be B-B4.

Said heavy chain may have at least 70%, at least 80%, at least 90%, at least 95% or at least 98% sequence identity with SEQ ID NO:1.

Said engineered targeting antibody may comprise at least one light chain, wherein said light chain has at least 70%, at least 80%, at least 90%, at least 95% or at least 98% sequence identity with SEQ ID NO:2.

Said heavy chain may have at least 80%, at least 85%, at least 90%, at least 95% or 100% sequence identity with residues 31 to 35, residues 51 to 68 and/or residues 99 to 111 of SEQ ID NO:1. Said heavy chain may have at least 75%, at least 85%, at least 95%, at least 97% or 100% sequence identity with residues 24 to 34, residues 50 to 56 and/or residues 89 to 97 of SEQ ID NO:2.

The further antibody region may comprise:
(a) amino acid residues 123 to 448 of SEQ ID NO: 1, and/or
(b) amino acid residues 108 to 214 of SEQ ID NO: 2, respectively and mutations thereof that maintain or lower the antibody-dependent cytotoxicity and/or complement-dependent cytotoxicity of the engineered targeting antibody mutations thereof that and/or stabilize the engineered targeting antibody.

Said further antibody region may be a constant heavy region of a human antibody.

Said engineered targeting antibody may bind CD138 with a targeting variation of less than 150%, 140%, 130%, 120%, 110%, 100%, 90%, 80%, 70%, 60% or 50%.

Said heavy chain may have at least 70%, at least 80%, at least 90%, at least 95% or at least 98% sequence identity with SEQ ID NO:1.

Said engineered targeting antibody may comprise at least one light chain, wherein said light chain has at least 70%, at least 80%, at least 90%, at least 95% or at least 98% sequence identity with SEQ ID NO:2.

Said heavy chain may have at least 80%, at least 85%, at least 90%, at least 95% sequence identity with residues 31 to 35, residues 51 to 68 and residues 99 to 111 of SEQ ID NO:1.

Said heavy chain may have at least 75%, at least 85%, at least 95% or at least 97% sequence identity with residues 24 to 34, residues 50 to 56 and residues 89 to 97 of SEQ ID NO:2.

The present invention is also directed at a pharmaceutical composition comprising or consisting essentially of the engineered targeting antibody and a pharmaceutically acceptable carrier.

A hybridoma which produces the engineered targeting is also part of the present invention.

The present invention also includes an antibody based assay comprising the engineered targeting antibody.

The present invention provides the engineered targeting antibody described herein for use in medicine. In particular, the engineered targeting antibody comprises:

an antigen binding region against CD138, wherein said antigen binding region is of a non-human antibody, and

a further antibody region, wherein at least part of said further antibody region is of a human antibody.

More particularly, the engineered targeting antibody is for use in a treatment targeting tumor cells.

The present invention also provides the use of the engineered targeting antibody described herein for the manufacture of a medicament for targeting tumor cells. In particular, the engineered targeting antibody comprises:

an antigen binding region against CD138, wherein said antigen binding region is of a non-human antibody, and

a further antibody region, wherein at least part of said further antibody region is of a human antibody.

More particularly, in these medical uses of the present invention the engineered targeting antibody is to be administered to an individual with CD138 expressing cells. Further, the engineered targeting antibody is capable of homogenously binding CD138 expressed on said CD138 expressing cells.

BRIEF DESCRIPTION OF THE FIGURES

FIG. 1 provides a schematic representation of nBT062 having effector molecules attached.

FIG. 2 is a chemical representation of BT062.

FIG. 3 shows the conversion of ansamitocin P-3 to maytansinol (stereochemistry is omitted for simplicity).

FIG. 4 shows a representative synthesis scheme of DM4.

FIG. 5 is a schematic representation of an antibody conjugation (nBT062 to DM4).

FIG. 6 shows an analysis of the binding of nBT062-SPDB-DM4, nBT062-SPP-DM1, nBT062-SMCC-DM1 and nBT062 antibody to OPM-2 cells. Different concentrations of nBT062 and conjugates were given to the cells and mean fluorescence was measured by FACS analysis.

FIG. 7(A)-(D) depict *in vitro* cytotoxicity of nBT062-DMx conjugates towards MOLP-8 ($CD138^+$) and BJAB ($CD138^-$) cells . The cells were cultured in flat bottom plates and incubated with the indicated concentrations of immunoconjugates for 5 days. WST reagent was added for further 3 hours to asses cell viability. In (D) cytotoxic activity of nBT062-SPDB-DM4 was analyzed in the presence or absence of blocking antibody (1 µM nBT062).

FIG. 8 shows tumor volumes for individual mice treated with (A) PBS, (B) nBT062 antibody, (C) free DM4 or (D) non-targeting conjugate huC242-DM4 over time (days) post-inoculation with MOLP-8 tumor cells.

FIG. 9 shows tumor volumes for individual mice treated with (A) PBS, (B) nBT062-SPDB-DM4, (C) B-B4-SPP-DM1 or (D) nBT062-SPP-DM1 over time (days) post-inoculation with MOLP-8 tumor cells.

FIG. 10 depicts mean tumor volume (+/- SD) of MOLP-8 human multiple myeloma xenografts in CB.17 SCID mice over time (days) post-inoculation.

FIGS. 11A and B show the anti-tumor activity of nBT062-DMx against $CD138^+$ MOLP-8 tumor cells in a bulky MOLP-8 tumor model in SCID mice. Tumor volume is given as mean (+/- SD) for each group.

DETAILED DESCRIPTION OF VARIOUS AND PREFERRED EMBODIMENTS

The present invention relates to targeting agents, in particular CD138 targeting antibodies, more in particular engineered CD138 targeting antibodies. Immunoconjugates comprising said targeting agents allow the delivery of the effector molecule(s) to target sites and the site specific release of effector(s) molecule in, at or near target cells, tissues and organs. The effector molecules may be activated by cleavage/dissociation from the targeting agent portion of the immunoconjugate at the target site.

The antibodies according to the present invention and/or immunoconjugates comprising the same may be administered to a subject in need of therapeutic treatment or to cells isolated from such a subject in need of therapeutic treatment. The effector molecule or molecules may be released from the immunoconjugate by cleavage/dissociation in, at or close to the target cell, tissue or organ.

As one example, the antibody nBT062 is employed in an chromographic assay. Formalin fixed, paraffin embedded tissue of a patient are provided. The antibody nBT062 is added as a primary antibody and surface expressed CD138 of the tissue binds the antibody. A detecting antibody is added to bind nBT062. In a final step, this binding of the detecting antibody, which comprises a chromogen is determined. The antibody nBT062 is used identy human plasmocytes among hemopoietic cells and thus allows diagnosis of a variety of hematological malignancies. The method also allows one to follow the progress of certain carcinomas. A reduction of unspecific detection due to a reduced cross reactivty with Fc receptors is observed when nBT062 is employed as opposed to its murine counterpart.

As a second example, the nBT062 antibody and an immunoconjugate that comprises the antibody nBT062 and at least one highly cytotoxic drug or an immunotoxin as an effector molecule are provided and is administered to a patient with cancer. In this example, an effective amount of nBT062 shields CD138 expressing non-

tumor cells from a therapeutically effective amount of the immunoconjugate that is later administered intravenously to a patient so that it concentrates in the cancerous cells. The effector molecule or molecules are released from the antibody target by an external means to induce cell death or continuous cell cycle arrest in the cancer cells.

CD138 or syndecan-1 (also described as SYND1; SYNDECAN; SDC; SCD1; CD138 ANTIGEN, SwissProt accession number: P18827 human) is a membrane glycoprotein that was originally described to be present on cells of epithelial origin, and subsequently found on hematopoietic cells (Sanderson, 1989). CD138 has a long extracellular domain that binds to soluble molecules (e.g., the growth factors EGF, FGF, HGF) and to insoluble molecules (e.g., to the extracellular matrix components collagen and fibronectin) through heparan sulfate chains (Langford, 1998; Yang, 2007) and acts as a receptor for the extracellular matrix. CD138 also mediates cell to cell adhesion through heparin-binding molecules expressed by adherent cells. It has been shown that CD138 has a role as a co-receptor for growth factors of myeloma cells (Bisping, 2006). Studies of plasma cell differentiation showed that CD138 must also be considered as a differentiation antigen (Bataille, 2006).

In malignant hematopoiesis, CD138 is highly expressed on the majority of MM cells, ovarian carcinoma, kidney carcinoma, gall bladder carcinoma, breast carcinoma, prostate cancer, lung cancer, colon carcinoma cells and cells of Hodgkin's and non-Hodgkin's lymphomas, chronic lymphocytic leukemia (CLL) (Horvathova, 1995), acute lymphoblastic leukemia (ALL), acute myeloblastic leukemia (AML) (Seftalioglu, 2003 (a); Seftalioglu, 2003 (b)), solid tissue sarcomas, colon carcinomas as well as other hematologic malignancies and solid tumors that express CD138 (Carbone et al., 1999; Sebestyen et al.,1999; Han et al., 2004; Charnaux et al., 2004; O'Connell et al.,2004; Orosz and Kopper, 2001).

Other cancers that have been shown to be positive for CD138 expression are many ovarian adenocarcinomas, transitional cell bladder carcinomas, kidney clear cell carcinomas, squamous cell lung carcinomas; breast carcinomas and uterine cancers (see, for example, Davies et al., 2004; Barbareschi et al., 2003; Mennerich et al., 2004; Anttonen et al., 2001; Wijdenes, 2002).

In the normal human hematopoietic compartment, CD138 expression is restricted to plasma cells (Wijdenes, 1996; Chilosi, 1999) and CD138 is not expressed on peripheral blood lymphocytes, monocytes, granulocytes, and red blood cells. In particular, $CD34^+$ stem and progenitor cells do not express CD138 and anti-CD138 mAbs do not affect the number of colony forming units in hematopoietic stem cell cultures (Wijdenes, 1996). In non-hematopoietic compartments, CD138 is mainly expressed on simple and stratified epithelia within the lung, liver, skin, kidney and gut. Only a weak staining was seen on endothelial cells (Bernfield, 1992; Vooijs, 1996). It has been reported that CD138 exists in polymorphic forms in human lymphoma cells (Gattei, 1999).

Monoclonal antibodies B-B4, BC/B-B4, B-B2, DL-101, 1 D4, MI15, 1.BB.210, 2Q1484, 5F7, 104-9, 281-2 in particular B-B4 have been reported to be specific to CD138. Of those B-B4, 1D4 and MI15 recognized both the intact molecule and the core protein of CD138 and were shown to recognize either the same or closely related epitopes (Gattei, 1999). Previous studies reported that B-B4 did not recognize soluble CD138, but only CD138 in membrane bound form (Wijdenes, 2002).

B-B4, a murine IgG1 mAb, binds to a linear epitope between residues 90-95 of the core protein on human syndecan-1 (CD138) (Wijdenes, 1996; Dore, 1998). Consistent with the expression pattern of CD138, B-B4 was shown to strongly react with plasma cell line RPMI8226, but not to react with endothelial cells. Also consistent with the expression pattern of CD138, B-B4 also reacted with epithelial cells lines A431 (keratinocyte derived) and HepG2 (hepatocyte derived). An immunotoxin B-B4-saporin was also highly toxic towards the plasma cell line RPMI8226, in fact considerably more toxic than free saporin. However, from the two epithelial cell lines tested, B-B4-saporin showed only toxicity towards cell line A431, although in a clonogenic assay B-B4 saporin showed no inhibitory effect on the outgrowth of A431 cells (Vooijs, 1996). Other researchers reported lack of specificity of MM-associated antigens against tumors (Couturier, 1999).

An antibody "consisting essentially of" certain components means in the context of the present invention that the antibody consists of the specified components and any

additional materials or components that do not materially affect the basic characteristics of the antibody.

The present invention uses the term "tumor cell" to include cancer cells as well as pre-cancerous cells which may or may not form part of a solid tumor.

A "targeting agent" according to the present invention is able to associate with a molecule expressed by a target cell and includes peptides and non-peptides. In particular, targeting agents according to the present invention include targeting antibodies and non-immunoglobulin targeting molecules, which may be based on non-immunoglobulin proteins, including, but not limited to, AFFILIN® molecules, ANTICALINS® and AFFIBODIES®. Non-immunoglobulin targeting molecules also include non-peptidic targeting molecules such as targeting DNA and RNA oligonucleotides (aptamers), but also physiological ligands, in particular ligands of the antigen in question, such as CD138.

A "targeting antibody" according to the present invention is or is based on a natural antibody or is produced synthetically or by genetic engineering and binds to an antigen on a cell or cells (target cell(s)) of interest. A targeting antibody according to the present invention includes a monoclonal antibody, a polyclonal antibody, a multispecific antibody (for example, a bispecific antibody), or an antibody fragment. The targeting antibody may be engineered to, for example, improve its affinity to the target cells (Ross, 2003) or diminish its immunogenicity. The targeting antibody may be attached to a liposomal formulation including effector molecules (Carter, 2003). An antibody fragment comprises a portion of an intact antibody, preferably the antigen binding or variable region of the intact antibody. Examples of antibody fragments according to the present invention include Fab, Fab', $F(ab')_2$, and Fv fragments, but also diabodies; domain antibodies (dAb) (Ward, 1989; United States Patent 6,005,079); linear antibodies; single-chain antibody molecules; and multispecific antibodies formed from antibody fragments. In a single chain variable fragment antibody (scFv) the heavy and light chains (VH and VL) can be linked by a short amino acid linker having, for example, the sequence $(glycine_4serine)_n$, which has sufficient flexibility to allow the two domains to assemble a functional antigen binding pocket. Addition of various signal sequences may allow for more precise targeting of the targeting antibody. Addition of the light chain constant region (CL) may allow dimerization via disulphide bonds, giving increased stability and

avidity. Variable regions for constructing the scFv can, if a mAb against a target of interest is available, be obtained by RT-PCR which clones out the variable regions from mRNA extracted from the parent hybridoma. Alternatively, the scFv can be generated de novo by phage display technology (Smith, 2001). As used herein, the term "functional fragment", when used in reference to a targeting antibody, is intended to refer to a portion of the targeting antibody which is capable of specifically binding an antigen that is specifically bound by the antibody reference is made to. A bispecific antibody according to the present invention may, for example, have at least one arm that is reactive against a target tissue and one arm that is reactive against a linker moiety (United States Patent Publication 20020006379). A bispecific antibody according to the present invention may also bind to more than one antigen on a target cell (Carter, 2003). An antibody according to the present invention may be modified by, for example, introducing cystein residues to introduce thiol groups (Olafsen, 2004).

In accordance with the present invention, the targeting antibody may be derived from any source and may be, but is not limited to, a camel antibody, a murine antibody, a chimeric human/mouse antibody or a chimeric human/monkey antibody, in particular, a chimeric human/mouse antibody such as nBT062.

Humanized antibodies are antibodies that contain sequences derived from a human-antibody and from a non-human antibody and are also within the scope of the present invention. Suitable methods for humanizing antibodies include CDR-grafting (complementarity determining region grafting) (EP 0 239 400; WO 91/09967; United States Patents 5,530,101; and 5,585,089), veneering or resurfacing (EP 0 592 106; EP 0 519 596; Padlan, 199; Studnicka et al., 1994; Roguska et al., 1994), chain shuffling (United States Patent 5,565,332) and DeImmunosation™ (Biovation, LTD). In CDR-grafting, the mouse complementarity-determining regions (CDRs) from, for example, mAb B-B4 are grafted into human variable frameworks, which are then joined to human constant regions, to create a human B-B4 antibody (hB-B4). Several antibodies humanized by CDR-grafting are now in clinical use, including MYLOTARG (Sievers et al., 2001) and HECEPTIN (Pegram et al, 1998).

The resurfacing technology uses a combination of molecular modeling, statistical analysis and mutagenesis to alter the non-CDR surfaces of antibody variable regions to resemble the surfaces of known antibodies of the target host. Strategies and methods for the resurfacing of antibodies, and other methods for reducing immunogenicity of antibodies within a different host, are disclosed, for example, in United States Patent

5,639,641. Human antibodies can be made by a variety of methods known in the art including phage display methods. See also United States Patents 4,444,887, 4,716,111, 5,545,806, and 5,814,318; and international patent application publications WO 98/46645, WO 98/50433, WO 98/24893, WO 98/16654, WO 96/34096, WO 96/33735, and WO 91/10741.

Targeting antibodies that have undergone any non-natural modification such as chimeric human/mouse antibodies or a chimeric human/monkey antibodies, humanized antibodies or antibodies that were engineered to, for example, improve their affinity to the target cells or diminish their immunogenicity but also antibody fragments, in particular functional fragments of such targeting antibodies that have undergone any non-natural modification, diabodies; domain antibodies; linear antibodies; single-chain antibody molecules; and multispecific antibodies are referred to herein as **engineered targeting antibodies**.

Chimerized antibodies, maintain the antibody binding region (ABR or Fab region) of the non-human antibody, e.g., the murine antibody they are based on, while any constant regions may be provided for by, e.g., a human antibody. Generally, chimerization and/or the exchange of constant regions of an antibody will not affect the affinity of an antibody because the regions of the antibody which contribute to antigen binding are not affected by this exchange. In a preferred embodiment of the present invention, the engineered, in particular chimerized, antibody of the present invention, may have a higher binding affinity (as expressed by K_D values) than the respective non-human antibody it is based on. In particular, the nBT062 antibody and antibodies based thereon may have higher antibody affinity than the murine B-B4. In another preferred embodiment of the present invention, immunoconjugates comprising those engineered/chimerized antibodies also display this higher antibody affinity. These immunconjugates may also display in certain embodiments other advantageous properties, such as a higher reduction of tumor load than their B-B4 containing counterparts. In a preferred embodiment, the engineered, in particular chimerized targeting antibodies display binding affinities that are characterized by dissociation constants K_D (nM) of less than 1.6, less than 1.5 or about or less than 1.4, while their murine counterparts are characterized by dissociation constants K_D (nM) of about or more than 1.6. Immunoconjugates comprising targeting agents such as targeting antibodies may be characterized by dissociation constants of K_D (nM) of less than 2.6,

less than 2.5, less than 2.4, less than 2.3, less than 2.2, less than 2.1, less than 2.0, less than or about 1.9 are preferred, while immunoconjugates comprising the murine counterpart antibodies may be characterized by dissociation constants K_D (nM) of about or more than 2.6 (compare Table 3, Materials and Methods).

Fully human antibodies may also be used. Those antibodies can be selected by the phage display approach, where CD138 or an antigenic determinant thereof is used to selectively bind phage expressing, for example, B-B4 variable regions (see, Krebs, 2001). This approach is advantageously coupled with an affinity maturation technique to improve the affinity of the antibody. All antibodies referred to herein are isolated antibodies.

In one embodiment, the targeting antibody is, in its unconjugated form, moderately or poorly internalized. Moderate internalization constitutes about 30% to about 75% internalization of antibody, poor internalization constitutes about 0.01% to up to about 30% internalization after 3 hours incubation at 37°C. In another preferred embodiment the targeting antibody binds to CD138, for example, antibodies B-B4, BC/B-B4, B-B2, DL-101, 1 D4, MI15, 1.BB.210, 2Q1484, 5F7, 104-9, 281-2 in particular B-B4. Hybridoma cells, which were generated by hybridizing SP02/0 myeloma cells with spleen cells of Balb/c mice have been deposited with the DSMZ-Deutsche Sammlung von Mikroorganismen und Zellkulturen GmbH, Mascheroder Weg 1, D-38124 Braunschweig on December 11, 2007. The identification number of these B-B4 expressing hybridoma cells is DSM ACC2874. In another embodiment, the targeting antibody does not substantially bind non-cell-surface expressed CD138. When, in the context of the present invention, the name of a specific antibody is combined with the term "targeting antibody" such as "nBT062 targeting antibody," this means that this targeting antibody has the binding specificity of the antibody nBT062. If a targeting antibody is said to be "based on" a specified antibody, this means that this targeting antibody has the binding specificity of this antibody, but might take any form consistent with the above description of a targeting antibody. When, in the context of the present invention, the name of a specific antigen is combined with the term "targeting antibody" such as "CD138 targeting antibody," this means that this targeting antibody has the binding specificity for CD138. If, in the context of the present invention, for example, a targeting antibody is said to do something "selectively" such as "selectively targeting cell-

surface expressed CD138" or, to be "selective" for something, this means that there is a significant selectivity (i.e. a higher affinity towards CD138-positive cells compared with CD138-negative cells) for, in the case of the example provided, cell-surface expressed CD138, compared to any other antigens. Adverse side effects in a given environment are substantially reduced or even avoided due to this selectivity.

"Non-immunoglobulin targeting molecules" according to the present invention include targeting molecules derived from non-immunoglobulin proteins as well as non-peptidic targeting molecules. Small non-immunoglobulin proteins which are included in this definition are designed to have specific affinities towards, in particular surface expressed CD138. These small non-immunoglobulin proteins include scaffold based engineered molecules such as Affilin® molecules that have a relatively low molecular weight such as between 10 kDa and 20 kDa. Appropriate scaffolds include, for example, gamma crystalline. Those molecules have, in their natural state, no specific binding activity towards the target molecules. By engineering the protein surfaces through locally defined randomization of solvent exposed amino acids, completely new binding sites are created. Former non-binding proteins are thereby transformed into specific binding proteins. Such molecules can be specifically designed to bind a target, such as CD138, and allow for specific delivery of one or more effector molecules (see, scil Proteins GmbH at www.scilproteins.com, 2004). Another kind of non-immunoglobulin targeting molecules are derived from lipocalins, and include, for example ANTICALINS®, which resemble in structure somewhat immunoglobulins. However, lipocalins are composed of a single polypeptide chain with 160 to 180 amino acid residues. The binding pocket of lipocalins can be reshaped to recognize a molecule of interest with high affinity and specificity (see, for example, Beste et al., 1999). Artificial bacterial receptors such as those marketed under the trademark Affibody® (Affibody AB) are also within the scope of the present invention. These artificial bacterial receptor molecules are small, simple proteins and may be composed of a three-helix bundle based on the scaffold of one of the IgG-binding domains of Protein A (Staphylococcus aureus). These molecules have binding properties similar to many immunoglobulins, but are substantially smaller, having a molecular weight often not exceeding 10kDa and are also comparatively stable. Suitable artificial bacterial receptor molecules are, for example, described in United States Patents 5,831,012; 6,534,628 and 6,740,734.

An "effector molecule" according to the present invention is a molecule or a derivative, or an analogue thereof that is attached to a targeting agent, in particular a targeting antibody and/or an engineered targeting antibody, and that exerts a desired effect, for example, apoptosis, or another type of cell death, or a continuous cell cycle arrest on the target cell or cells. Effector molecules according to the present invention include molecules that can exert desired effects in a target cell and include, but are not limited to, toxins, drugs, in particular low molecular weight cytotoxic drugs, radionuclides, biological response modifiers, pore-forming agents, ribonucleases, proteins of apoptotic signaling cascades with apoptosis-inducing activities, cytotoxic enzymes, prodrug activating enzymes, antisense oligonucleotides, antibodies or cytokines as well as functional derivatives or analogues/fragments thereof. Toxins may include bacterial toxins, such as, but not limited to, Diphtheria toxin or Exotoxin A, plant toxins, such as but not limited to, Ricin. Proteins of apoptotic signaling cascades with apoptosis-inducing activities, include, but are not limited to, Granzyme B, Granzyme A, Caspase-3, Caspase-7, Caspase-8, Caspase-9, truncated Bid (tBid), Bax and Bak.

In a preferred embodiment, the effector increases internal effector delivery of the immunoconjugate, in particular when the natural form of the antibody on which the targeting antibody of the immunoconjugate is based is poorly internalizable. In another preferred embodiment the effector is, in its native form, non-selective. In certain embodiments the effector has high non-selective toxicity, including systemic toxicity, when in its native form. The "native form" of an effector molecule of the present invention is an effector molecule before being attached to the targeting agent to form an immunoconjugate. In another preferred embodiment, the non-selective toxicity of the effector molecule is substantially eliminated upon conjugation to the targeting agent. In another preferred embodiment, the effector molecule causes, upon reaching the target cell, death or continuous cell cycle arrest in the target cell. A drug-effector molecule according to the present invention includes, but is not limited to, a drug including, for example, small highly cytotoxic drugs that act as inhibitors of tubulin polymerization such as maytansinoids, dolastatins, auristatin and crytophycin; DNA alkylating agents like CC-1065 analogues or derivatives (United States Patents 5,475,092; 5,585,499; 6,716,821) and duocarmycin; enediyne antibiotics such as calicheamicin and esperamicin; and potent taxoid (taxane) drugs (Payne, 2003). Maytansinoids and calicheamicins are particularly preferred. An effector maytansinoid includes maytansinoids of any origin, including, but not limited to synthetic maytansinol and maytansinol analogue and

derivative. Doxorubicin, daunomycin, methotrexate, vinblastine, neocarzinostatin, macromycin, trenimon and α-amanitin are some other effector molecules within the scope of the present invention. Also within the scope of the present invention are antisense DNA molecules as effector molecules. When the name of, for example, a specific drug or class of drugs is combined herein with the term "effector" or "effector molecule," reference is made to an effector of an immunoconjugate according to the present invention that is based on the specified drug or class of drugs.

Maytansine is a natural product originally derived from the Ethiopian shrub Maytenus serrata (Remillard, 1975; United States Patent 3,896,111). This drug inhibits tubulin polymerization, resulting in mitotic block and cell death (Remillard, 1975; Bhattacharyya, 1977; Kupchan, 1978). The cytotoxicity of maytansine is 200-1000-fold higher than that of anti-cancer drugs in clinical use that affect tubulin polymerization, such as Vinca alkaloids or taxol. However, clinical trials of maytansine indicated that it lacked a therapeutic window due to its high systemic toxicity. Maytansine and maytansinoids are highly cytotoxic but their clinical use in cancer therapy has been greatly limited by their severe systemic side-effects primarily attributed to their poor selectivity for tumors. Clinical trials with maytansine showed serious adverse effects on the central nervous system and gastrointestinal system.

Maytansinoids have also been isolated from other plants including seed tissue of Trewia nudiflora (United States Patent 4,418,064)

Certain microbes also produce maytansinoids, such as maytansinol and C-3 maytansinol esters (United States Patent 4,151,042).

The present invention is directed to maytansinoids of any origin, including synthetic maytansinol and maytansinol analogues which are disclosed, for example, in United States Patents 4,137,230; 4,248,870; 4,256,746; 4,260,608; 4,265,814; 4,294,757; 4,307,016; 4,308,268; 4,308,269; 4,309,428; 4,313,946; 4,315,929; 4,317,821; 4,322,348; 4,331,598; 4,361,650; 4,362,663; 4,364,866; 4,371,533; 4,424,219 and 4,151,042.

In a preferred embodiment, the maytansinoid is a thiol-containing maytansinoid and is more preferably produced according to the processes disclosed in United States Patent 6,333,410 to Chari et al or in Chari et al.(Chari, 1992).

DM-1 (N^2-deacetyl-N^2-(3-mercapto-1-oxopropyl)-maytansine) is a preferred effector molecule in the context of the present invention. DM1 is 3- to 10-fold more cytotoxic than maytansine, and has been converted into a pro-drug by linking it via

disulfide bond(s) to a monoclonal antibody directed towards a tumor-associated antigen. Certain of these conjugates (sometimes called "tumor activated prodrugs" (TAPs)) are not cytotoxic in the blood compartment, since they are activated upon associating with a target cells and internalized, thereby releasing the drug (Blättler, 2001). Several antibody-DM1 conjugates have been developed (Payne, 2003), and been evaluated in clinical trials. For example, huC242-DM1 treatment in colorectal cancer patients was well tolerated, did not induce any detectable immune response, and had a long circulation time (Tolcher, 2003).

Other particularly preferred maytansinoids comprise a side chain that contains a sterically hindered thiol bond such as, but not limited to, maytansinoids $N^{2'}$-deacetyl- $N^{2'}$-(4-mercapto-1-oxopentyl)-maytansine, also referred to as "DM3," and $N^{2'}$-deacetyl- $N^{2'}$-(4-methyl-4-mercapto-1-oxopentyl)-maytansine, also referred to as "DM4." The synthesis of DM4 is shown in **FIGS. 3** and **4** and is described elsewhere herein. DM4 differs from DM1 and DM3 in that it bears methyl groups at its αC. This results in a sterical hindrance when DM4 is attached via a linker in particular, but not limited to, a linker comprising a disulfide bond, to a targeting agent such as nBT062. A wide variety of maytansinoids bearing a sterically hindered thiol group (possessing one or two substituents, in particular alkyls substituents, such as the methyl substituents of DM4) are disclosed U.S. Patent Publication 2004/0235840, published Nov. 25, 2004, which is incorporated herein in its entirety by reference. As reported by Goldmahker et al. in U.S. Patent Publciation 2006/0233814, such a hindrance induces alkylation (e.g., methylation) of the free drug, once the drug is released at its target. The alkylation may increase the stability of the drug allowing for the so-called bystander effect. However, as the person skilled in the art will appreciate, other effector molecules comprising substitutents such as alkyl groups at positions that result in a sterical hindrance when the effector is attached to a targeting agent via a linker are part of the present invention. Preferably this hindrance induces a chemical modification such as alkylation of the free drug to increase its overall stability, which allows the drug to not only induce cell death or continuous cell cycle arrest in CD138 expressing tumor cells but, optionally, also to affect auxillary cells that, e.g., support or protect the tumor from drugs, in particular cells of the tumor stroma and the tumor vasculature and which generally do not express CD138 to diminish or lose their supporting or protecting function.

DNA alkylating agents are also particularly preferred as effector molecules and include, but are not limited to, CC-1065 analogues or derivatives. CC-1065 is a potent antitumor-antibiotic isolated from cultures of Streptomyces zelensis and has been shown to be exceptionally cytotoxic in vitro (United States Patent 4,169,888). Within the scope of the present invention are, for example the CC-1065 analogues or derivatives described in United States Patents 5,475,092, 5,585,499 and 5,739,350. As the person skilled in the art will readily appreciate, modified CC-1065 analogues or derivatives as described in United States Patent 5,846,545 and prodrugs of CC-1065 analogues or derivatives as described, for example, in United States Patent 6,756,397 are also within the scope of the present invention. In certain embodiments of the invention, CC-1065 analogues or derivatives may, for example, be synthesized as described in United States Patent 6,534,660.

Another group of compounds that make preferred effector molecules are taxanes, especially highly potent ones and those that contain thiol or disulfide groups. Taxanes are mitotic spindle poisons that inhibit the depolymerization of tubulin, resulting in an increase in the rate of microtubule assembly and cell death. Taxanes that are within the scope of the present invention are, for example, disclosed in United States Patents 6,436,931; 6,340,701; 6,706,708 and United States Patent Publications 20040087649; 20040024049 and 20030004210. Other taxanes are disclosed, for example, in United States Patent 6,002,023, United States Patent 5,998,656, United States Patent 5,892,063, United States Patent 5,763,477, United States Patent 5,705,508, United States Patent 5,703,247 and United States Patent 5,367,086. As the person skilled in the art will appreciate, PEGylated taxanes such as the ones described in United States Patent 6,596,757 are also within the scope of the present invention.

Calicheamicin effector molecules according to the present invention include gamma 1I, N-acetyl calicheamicin and other derivatives of calicheamicin. Calicheamicin binds in a sequence-specific manner to the minor groove of DNA, undergoes rearrangement and exposes free radicals, leading to breakage of double-stranded DNA, resulting in cell apoptosis and death. One example of a calicheamicin effector molecule that can be used in the context of the present invention is described in United States Patent 5,053,394.

An immunoconjugate according to the present invention comprises at least one targeting agent, in particular targeting antibody, such as an engineered targeting

antibody, and one effector molecule. The immunoconjugate might comprise further molecules for example for stabilization. For immunoconjugates, the term "conjugate" is generally used to define the operative association of the targeting agent with one or more effector molecules and is not intended to refer solely to any type of operative association, and is particularly not limited to chemical "conjugation". So long as the targeting agent is able to bind to the target site and the attached effector functions sufficiently as intended, particularly when delivered to the target site, any mode of attachment will be suitable. The conjugation methods according to the present invention include, but are not limited to, direct attachment of the effector molecule to the targeting antibody, with or without prior modification of the effector molecule and/or the targeting antibody or attachment via linkers. Linkers can be categorized functionally into, for example, acid labile, photolabile linkers, enzyme cleavable linkers, such as linkers that can be cleaved by peptidases. Cleavable linkers are, in many embodiments of the invention preferred. Such cleavable linkers can be cleaved under conditions present in the cellular environment, in particular, an intracellular environment and that have no detrimental effect on the drug released upon cleavage. Low pHs such as pH of 4 to 5 as they exist in certain intracellular departments will cleave acid labile linkers, while photolabile linkers can be cleaved by, e.g., infrared light. However, linkers that are cleaved by/under physiological conditions present in the majority of cells are preferred and are referred to herein as **physiologically cleavable linkers**. Accordingly, disulfide linkers are being preferred in many embodiments of the invention. These linkers are cleavable through disulfide exchange, which can occur under physiological conditions. Preferred heterobifunctional disulfide linkers include, but are not limited to, N-succinimidyl 3-(2-pyridyldithio)propionate (SPDP) (see, e.g., Carlsson et al.(1978)), N-succinimidyl 4-(2-pyridyldithio)butanoate (SPDB) (see, e.g., U.S. Pat. No. 4,563,304), N-succinimidyl 4-(2-pyridyldithio)pentanoate (SPP) (see, e.g., CAS Registry number 341498-08-6), N-succinimidyl 4-(N-maleimidomethyl)cyclohexane-1-carboxylate (SMCC) (see, e.g., Yoshitake et al., (1979)), and N-succinimidyl 4-methyl-4-[2-(5-nitro-pyridyl)-dithio]pentanoate (SMNP) (see, e.g., U.S. Pat. No. 4,563,304). The most preferred linker molecules for use in the inventive composition are SPP, SMCC, and SPDB.

Other suitable linkers may include "non-cleavable" bonds, such as, but not limited to Sulfosuccinimidyl maleimidomethyl cyclohexane carboxylate (SMCC), which is a heterobifunctional linker capable of linking compounds with SH-containing compounds. Bifunctional and heterobifunctional linker molecules, such as carbohydrate-directed

heterobifunctional linker molecules, such as S-(2-thiopyridyl)-L-cysteine hydrazide (TPCH), are also within the scope of the present invention (Vogel, 2004). The effector molecule, such as a maytansinoid, may be conjugated to the targeting antibody via a two reaction step process, including as a first step modification of the targeting antibody with a cross-linking reagent such as N-succinimidyl pyridyldithiopropionate (SPDP) to introduce dithiopyridyl groups into the targeting antibody. In a second step, a reactive maytansinoid having a thiol group, such as DM1, may be added to the modified antibody, resulting in the displacement of the thiopyridyl groups in the modified antibody, and the production of disulfide-linked cytotoxic maytansinoid/antibody conjugate (United States Patent 5,208,020). However, one-step conjugation processes such as the one disclosed in United States Patent Publication 20030055226 to Chari et al are also within the scope of the present invention. In one embodiment of the present invention multiple effector molecules of the same or different kind are attached to a targeting antibody.

CC-1065 analogues or derivatives may be conjugated to the targeting agent via for example PEG linking groups as described in United States Patent 6,716,821.

Calicheamicins may be conjugated to the targeting antibodies via linkers (United States Patent 5,877,296 and United States Patent 5,773,001) or according to the conjugation methods disclosed in United States Patent 5,712,374 and United States Patent 5,714,586. Another preferred method for preparing calicheamicin conjugates is disclosed in Unites States Patent Publication 20040082764. The immunoconjugates of the present invention may take the form of recombinant fusion proteins.

The term sequence identity refers to a measure of the identity of nucleotide sequences or amino acid sequences. In general, the sequences are aligned so that the highest order match is obtained. "Identity", per se, has recognized meaning in the art and can be calculated using published techniques. (See, e.g.: Computational Molecular Biology, Lesk, A. M., ed., Oxford University Press, New York, 1988; Biocomputing: Informatics and Genome Projects, Smith, D. W., ed., Academic Press, New York, 1993; Computer Analysis of Sequence Data, Part I, Griffin, A. M., and Griffin, H. G., eds., Humana Press, New Jersey, 1994; Sequence Analysis in Molecular Biology, von Heinje, G., Academic Press, 1987; and Sequence Analysis Primer, Gribskov, M. and Devereux, J., eds., M Stockton Press, New York, 1991). While there exist a number of methods to measure identity between two polynucleotide or polypeptide sequences, the term

"identity" is well known to skilled artisans (Carillo, H. & Lipton, D., SIAM J Applied Math 48:1073 (1988)).

Whether any particular nucleic acid molecule is at least 50%, 60%, 70%, 75%, 80%, 85%, 90%, 95%, 96%, 97%, 98% or 99% identical to, for instance, the nBT062 nucleic acid sequence, or a part thereof, can be determined conventionally using known computer programs such as DNAsis software (Hitachi Software, San Bruno, Calif.) for initial sequence alignment followed by ESEE version 3.0 DNA/protein sequence software (cabot@trog.mbb.sfu.ca) for multiple sequence alignments.

Whether the amino acid sequence is at least 50%, 60%, 70%, 75%, 80%, 85%, 90%, 95%, 96%, 97%, 98% or 99% identical to, for instance SEQ ID NO:1 or SEQ ID NO:2, or a part thereof, can be determined conventionally using known computer programs such the BESTFIT program (Wisconsin Sequence Analysis Package, Version 8 for Unix, Genetics Computer Group, University Research Park, 575 Science Drive, Madison, Wis. 53711). BESTFIT uses the local homology algorithm of Smith and Waterman, Advances in Applied Mathematics 2:482-489 (1981), to find the best segment of homology between two sequences.

When using DNAsis, ESEE, BESTFIT or any other sequence alignment program to determine whether a particular sequence is, for instance, 95% identical to a reference sequence according to the present invention, the parameters are set such that the percentage of identity is calculated over the full length of the reference nucleic acid or amino acid sequence and that gaps in homology of up to 5% of the total number of nucleotides in the reference sequence are allowed.

If, in the context of the present invention, reference is made to a certain sequence identity with a combination of residues of a particular sequence, this sequence identity relates to the sum of all the residues specified.

The basic antibody molecule is a bifunctional structure wherein the variable regions bind antigen while the remaining constant regions may elicit antigen independent responses. The major classes of antibodies, IgA, IgD, IgE, IgG and IgM, are determined by the constant regions. These classes may be further divided into

subclasses (isotypes). For example, the IgG class has four isotypes, namely, IgG1, IgG2, IgG3, and IgG4 which are determined by the constant regions. Of the various human antibody classes, only human IgG1, IgG2, IgG3 and IgM are known to effectively activate the complement system. While the constant regions do not form the antigen binding sites, the arrangement of the constant regions and hinge region may confer segmental flexibility on the molecule which allows it to bind with the antigen.

Different IgG isotypes can bind to Fc receptors on cells such as monocytes, B cells and NK cells, thereby activating the cells to release cytokines. Different isotypes may also activate complement, resulting in local or systemic inflammation. In particular, the different IgG isotypes may bind FcγR to different degrees. FcγRs are a group of surface glycoproteins belonging to the Ig superfamily and expressed mostly on leucocytes. The FcγR glycoproteins are divided into three classes designated FcγRI (CD64), FcγRII (CD32) and FcγRIII (CD16). While IgG1, IgG2 and IgG3 bind strongly to a variety of these classes of FcγR glycoproteins, IgG4 display much weaker binding. In particular, IgG4 is an intermediate binder of FcγRI, which results in relatively low or even no ADCC (antibody dependent cellular cytotoxicity), and does not bind to FcγRIIIA or FcγRIIA. IgG4 is also a weak binder of FcγRIIB, which is an inhibitory receptor. Furthermore, IgG4 mediates only weak or no complement fixation and weak or no complement dependent cytotoxicity (CDC). In the context of the present invention, IgG4 may be specifically employed to prevent Fc-mediated targeting of hepatic FcR as it displays no interaction with FcRγII on LSECs (liver sinusoidal endothelial cells), no or weak interaction with FcRγI-III on Kupffer cells (macrophages) and no interaction with FcRγIII on hepatic NK cells. Certain mutations that further reduce any CDC are also part of the present invention. For example IgG4 residues at positions 327, 330 and 331 were shown to reduce ADCC (antibody dependent cellular cytotoxicity) and CDC (Amour, 1999; Shields, 2001). One of more mutations that stabilize the antibody are also part of the present invention (also referred to herein as "stabilizing mutations"). Those mutations include in particular, leucine-to-glutamic acid mutations in the CH2 region of IgG4 and serine-to-proline exchanges in the IgG4 hinge core. These mutations decrease, in certain embodiments of the invention, the amount of half-molecules to less than 10%, less than 5% and preferably less than 2% or 1%. Moreover, the in vivo half life of so stabilized antibodies might be increased several days, including 1, 2, 3, 4 or more than 5 days (Schuurman, 1999).

Targeting agents, including targeting antibodies, in particular engineered targeting antibodies, disclosed herein may also be described or specified in terms of their **binding affinity** to antigen, in particular to CD138. Preferred binding affinities of targeting agents such as targeting antibodies, in particular engineerend targeting antibodies, are characterized by dissociation constants K_D (nM) of less than 1.6, less than 1.5 or about or less than 1.4. For immunoconjugates comprising said targeting agents such as targeting antibodies dissociation constants K_D (nM) of less than 1.6, less than 1.5 or less than 2.5, less than 2.4, less than 2.3, less than 2.2, less than 2.1, less than 2.0, less than or about 1.9 are preferred.

An **antigen binding region** (ABR) according to the present invention will vary based on the type of targeting antibody or engineered targeting antibody employed. In a naturally occurring antibody and in most chimeric and humanized antibodies, the antigen binding region is made up of a light chain and the first two domains of a heavy chain. However, in a heavy chain antibody devoid of light chains, the antigen binding region will be made up of, e.g., the first two domains of the heavy chain only, while in single chain antibodies (ScFv), which combine in a single polypeptide chain the light and heavy chain variable domains of an antibody molecule, the ABR is provided by only one polypeptide molecule. FAB fragments are usually obtained by papain digestion and have one light chain and part of a heavy chain and thus comprise an ABR with only one antigen combining site. On the other hand, diabodies are small antibody fragments with two antigen-binding regions. In the context of the present invention, however, an antigen binding region of an targeting antibody or engineered targeting antibody is any region that primarily determines the **binding specificity** of the targeting antibody or engineered targeting antibody.

If an ABR or another targeting antibody region is said to be "**of a certain antibody**", e.g., a human or non-human antibody, this means in the context of the present invention that the ABR is either identical to a corresponding naturally occurring ABR or is based thereon. An ABR is based on a naturally occurring ABR if it has the binding specificity of the naturally occurring ABR. However, such an ABR may comprise, e.g., point mutations, additions, deletions or posttranslational modification

such as glycosylation. Such an ABR may in particular have more than 70%, more than 80%, more than 90%, preferably more than 95%, more than 98% or more than 99% sequence identity with the sequence of the naturally occurring ABR.

Homogenous targeting of a targeting agent such as a targeting antibody, but in particular an immunoconjugate comprising the same, in the context of the present invention, is a measure of the variance associated with obtaining the desired result of said targeting with the targeting agent. In certain embodiments of the invention, the desired result is obtained by simple binding to the target. This is, for example, the case in embodiments in which a certain targeting agent provides a shield against subsequent binding. However, the homogeneity of a targeting agent can be readily assessed, e.g., via the efficacy of an immunoconjugate comprising said targeting agent. For example, the efficacy of said immunoconjugate against a tumor antigen such as CD138 that comprises an effector aimed at destroying tumor cells and/or arresting the growth of a tumor can be determined by the degree of growth suppression of a tumor comprising cells expressing the CD138 antigen. Such an immunoconjugate may display a high variance in its efficacy. It may, for example, arrest tumor growth sometimes with high efficacy, but other times with an efficacy that hardly exceeds the efficacy of the control. A low variance in the efficacy of an immunoconjugate, on the other hand, shows that the immunoconjugate and/or targeting agent, respectively, provide the desired result consistently. One way of quantifying the homogeneity of targeting is to calculate the **targeting variation**. In the context of tumor growth arrested by an immunoconjugate comprising a certain targeting agent, the targeting variation can be calculated by first determining the time for a tumor to reach a predetermined volume, e.g. 300mm^3. Preferably, the predetermined volume is chosen so that any tumor growth before and after reaching said predetermined volume is steadily increasing at about the same rate. After such time has been determined for a group of subjects the mean of these times (T_m) in the group of subjects (e.g., SCID mice or another suitable model displaying homogenous tumor growth) is calculated. T_m is then correlated to the observations made in the subject of the group showing the least efficacy in targeting and thus being associated with tumors that need the least time (T_f) to reach said predetermined volume, and, on the other hand, the subject in the group showing the highest efficacy in targeting and thus being associated with tumors that need the most time (T_s) to reach

said predetermined volume by calculating the targeting variation for the predetermined volume according to the following formula:

TARGETING VARIATION [%] =

Ts-Tf/Tm x 100

In a preferred embodiment, the targeting variation of the engineered targeting antibody of the present invention is less than 150%, less than 140%, less than 130%, less than 120%, less than 110%, less than 100%, less than 90%, less than 80%, less than 70%, less than 60%, or less than 50%, and in certain embodiments even less than 45%. Preferably, the targeting variation is between about 10% and about 150%, preferably between about 10% and about 100%, about 10% and about 80%, about 10% and about 70%, about 10% and about 60%, about 10% and about 50%.

The homogenity of targeting (also referred to herein as the homogenity of binding to a particular antigen) can be also quantified by other means such as determining the tumor growth delay. Also, as the person skilled in the art will readily understand tumor volume of a certain size is only one parameter on which basis targeting variation may be determined. Depending on the desired result, other parameters include time (for, e.g., measuring tumor growth delay) or % of binding may be employed. The person skilled in the art can readily determine such other parameters.

nBT062 (see also **FIG. 1**) is a murine human chimeric IgG4 mAb a chimerized version of B-B4. This chimerized version of B-B4 was created to reduce the HAMA (Human Anti-Mouse Antibody) response, while maintaining the functionality of the antibody binding region of the B-B4 for CD138. Surprisingly it was found that this chimeric antibody displays improved binding affinities relative to B-B4. Also surprisingly, the chimeric antibody has been associated with homogenous targeting, which reduces the variance in results obtained when using the antibody or immunoconjugate comprising the same. The protocol for producing nBT062 is specified below. Chinese hamster ovary cells expressing nBT062 have been deposited with the DSMZ-Deutsche Sammlung von Mikroorganismen und Zellkulturen GmbH, Mascheroder Weg 1, D-38124 Braunschweig on December 11, 2007. The identification number is DSM ACC2875. A CD138 specific chimeric antibody based on B-B4 is generically referred to herein as c-B-B4.

The amino acid sequence for both, the heavy and the light chains has been predicted from the translation of the nucleotide sequence for nBT062. The amino acid sequences predicted for the heavy chain and light chain are presented in **Table 1**. Predicted variable regions are bolded, predicted CDRs are underlined.

Table 1. Predicted Amino Acid Sequence for nBT062

- nBT062 heavy chain predicted sequence **(SEQ ID NO:1)**:

1	**QVQLQQSGSE LMMPGASVKI SCKATGYTFS NYWIEWVKQR PGHGLEWIGE**
51	**ILPGTGRTIY NEKFKGKATF TADISSNTVQ MQLSSLTSED SAVYYCARRD**
101	**YYGNFYYAMD YWGQGTSVTV SS**ASTKGPSV FPLAPCSRST SESTAALGCL
151	VKDYFPEPVT VSWNSGALTS GVHTFPAVLQ SSGLYSLSSV VTVPSSSLGT
201	KTYTCNVDHK PSNTKVDKRV ESKYGPPCPS CPAPEFLGGP SVFLFPPKPK
251	DTLMISRTPE VTCVVVDVSQ EDPEVQFNWY VDGVEVHNAK TKPREEQFNS
301	TYRVVSVLTV LHQDWLNGKE YKCKVSNKGL PSSIEKTISK AKGQPREPQV
351	YTLPPSQEEM TKNQVSLTCL VKGFYPSDIA VEWESNGQPE NNYKTTPPVL
401	DSDGSFFLYS RLTVDKSRWQ EGNVFSCSVM HEALHNHYTQKSLSLSLG(K)

The C-terminal lysine is prone to clipping and might be present due to incomplete clipping to a certain extent and is not part of SEQ ID NO: 1.

- nBT062 light chain predicted sequence **(SEQ ID NO:2)**:

1	**DIQMTQSTSS LSASLGDRVT ISCSASQGIN NYLNWYQQKP DGTVELLIYY**
51	**TSTLQSGVPS RFSGSGSGTD YSLTISNLEP EDIGTYYCQQ YSKLPRTFGG**
101	**GTKLEIK**RTV AAPSVFIFPP SDEQLKSGTA SVVCLLNNFY PREAKVQWKV
151	DNALQSGNSQ ESVTEQDSKD STYSLSSTLT LSKADYEKHK VYACEVTHQG
201	LSSPVTKSFN RGEC

Table 2. shows a comparision of the general CDR definitions of Krabat and Chothia and the predicted CDRs for BT062

Kabat CDR definition		**nBT062**
Light chain	CDR1: residues 24-34	CDR1: residues 24-34
	CDR2: residues 50-56	CDR2: residues 50-56
	CDR3: residues 89-97	CDR3: residues 89-97
Heavy chain	CDR1: residues 31-35	CDR1: residues 31-35
	CDR2: residues 50-56	CDR2: residues 51-68
	CDR3: residues 95-102	CDR3: residues 99-111

Chothia CDR definition		**nBT062**
Light chain	CDR1: residues 26-32	CDR1: residues 24-34
	CDR2: residues 50-52	CDR2: residues 50-56
	CDR3: residues 91-96	CDR3: residues 89-97
Heavy chain	CDR1: residues 26-32	CDR1: residues 31-35
	CDR2: residues 52-56	CDR2: residues 51-68
	CDR3: residues 96-101	CDR3: residues 99-111

BT062 is an immunoconjugate comprising the CD138 targeting chimeric antibody nBT062 that is attached via a linker, here SPDB, to the cytostatic maytansinoid derivative DM4. A chemical representation of BT062 is provided in **FIGS. 1** and **2.** Immunoconjugates comprising nBT062 and a maytansinoid effector molecule are often characterized in terms of their linker and maytansinoid effector, e.g., nBT062-SMCC-DM1, is an immunoconjugate comprising nBT062, SMCC (a "noncleavable" linker containing a thioester bond) and DM1 as an effector. More generically, an immunoconjugate containing nBT062 and an effector molecule may also be described as nBT062-linker-effector or just as nBT062-effector (nBT062N, wherein N is any effector described herein.

Reference is made herein to a unhindered counterpart **(UI: unhindered immunoconjugate)** of an immunoconjugate comprising an engineered targeting antibody against CD138 attached to an effector molecule via a **cleavable linker (CL)** and is described herein as UICL, which is contrasted to an immunoconjugate in which said effector molecule is sterically hindered, but contains a cleavable linker **(HICL)**. The UICL is an immunoconjugate equivalent to the HICL comprising an engineered targeting antibody in which the effector molecule is, however, not sterically hindered. Examples of

a pair of HICL/UICL are BT062 and nBT062-SPP-DM1. An unhindered counterpart of such a immunoconjugate comprising a **non-cleavable** linker **(UINCL)** refers to the equivalent immunoconjugate comprising an engineered targeting antibody in which the effector molecule is not sterically hindered and comprises a noncleavable linker. For BT062, nBT062-SMCC-DM1 would constitute an example of such an unhindered counterpart comprising an non-cleavable linker.

A growth of a tumor inhibiting activity (=tumor growth inhibiting activity) of an immunoconjugate is a relative measure. It describes the tumor growth inhibiting activity of a conjugate relative to the activity of the highest performing immunoconjugate whose activity is set as 100%. For example if the activity of the highest performing immunoconjugate, say, BT062, which causes a tumor growth delay (TGD) of 32 days, is set as 100%, the activity of, e.g., nBT062-DM1, which displays a tumor growth delay (TGD) of 18 days is calculated as follows:

Tumor Growth Inhibiting Activity=

$$100x\ (TGD_{nBT062\text{-}DM1}/TGD_{BT062}),$$

more generically:

Tumor Growth Inhibiting Activity=

$$100x\ (TGD_{Sample}/TGD_{Reference}).$$

Table 3 provides suitable examples from the results depicted in Fig. 11B:

	TGD* (days)	% Activity**
PBS	0	0
nBT062-SMCC-DM1	18	56
BT062	32	100
nBT062-SPP-DM1	13	40

Table 3: Tumor growth delay (TGD) and % Activity of nBT062-DMx against MOLP-8 tumor xenografts in SCID mice based on treatment groups receiving a 450 µg/kg dose.
(*) Tumor growth delay in days (TGD) as mean time in days for treatment group to reach a predetermined size (160 mm^3) minus the mean time for the control group to reach this predetermined size.
(**)Tumor Growth Inhibiting Activity =$100x(TGD_{Sample}/TGD_{BT062})$. The activity of BT062 is defined to be 100%.

In the example provided in Table 2, BT062 provides a growth of a tumor inhibiting activity that exceeds that of its unhindered counterpart (nBT062-SPP-DM1) by 60%, and a growth of a tumor inhibiting activity that exceeds that of its unhindered counterpart immunoconjugate comprising a non-cleavable linker (nBT062-SMCC-DM1) by 44%.

It was previously reported that a cleavable linker in an immunoconjugate provides for a so called bystander effect. Goldmahker et al. (U.S. Patent Publication 2006/0233814) also disclose that the bystander effect is particulary pronounced when the effector molecule is subject to further modification, in particular alkylation, upon cleavage from the targeting agent. Goldmahker et al. also showed that UICL displayed better TGD than the respective UINCL, a fact that was also attributed to what is referred to as the by-stander effect (see, e.g., Fig. 6 of U.S. Patent Publication 2006/0233814). Here, it was surprisingly found that an UICL in a high single dosage regime (250µg/kg) actually did not show any better results than the UINCL. In fact, the TGD in days that was observed in an UICL in such a regime was actually lower than that of the UINCL. This observation became more pronounced with an increase in dosage (450µg/kg). In sum, HICL outperformed UICL in single dose experiments to an unexpected degree. In addition, the UICL was unexpectedly outperformed by UINCL at higher dosages.

The targeting agents, in particular targeting antibodies, and/or immunoconjugates disclosed herein can be administered by any route, including intravenously, parenterally, orally, intramuscularly, intrathecally or as an aerosol. The mode of delivery will depend on the desired effect. A skilled artisan will readily know the best route of administration for a particular treatment in accordance with the present invention. The appropriate dosage will depend on the route of administration and the treatment indicated, and can readily be determined by a skilled artisan in view of current treatment protocols.

Pharmaceutical compositions containing an unconjugated targeting agent and the immunoconjugate of the present invention as active ingredients can be prepared according to conventional pharmaceutical compounding techniques. See, for example, Remington's Pharmaceutical Sciences, 17th Ed. (1985, Mack Publishing Co., Easton, Pa.). Typically, effective amounts of active ingredients will be admixed with a pharmaceutically acceptable carrier. The carrier may take a wide variety of forms depending on the form of preparation desired for administration, for example, intravenous, oral, parenteral, intrathecal, transdermal, or by aerosol.

For oral administration, the targeting agent and/or immunoconjugate can be formulated into solid or liquid preparations such as capsules, pills, tablets, lozenges, melts, powders, suspensions or emulsions. In preparing the compositions in oral dosage form, any of the usual pharmaceutical media may be employed, such as, for example, water, glycols, oils, alcohols, flavoring agents, preservatives, coloring agents, suspending agents, and the like in the case of oral liquid preparations (such as, for example, suspensions, elixirs and solutions); or carriers such as starches, sugars, diluents, granulating agents, lubricants, binders, disintegrating agents and the like in the case of oral solid preparations (such as, for example, powders, capsules and tablets). Because of their ease in administration, tablets and capsules represent the most advantageous oral dosage unit form, in which case solid pharmaceutical carriers are obviously employed. If desired, tablets may be sugar-coated or enteric-coated by standard techniques. The active agent must be stable to passage through the gastrointestinal tract. If necessary, suitable agents for stable passage can be used, and may include phospholipids or lecithin derivatives described in the literature, as well as liposomes, microparticles (including microspheres and macrospheres).

For parenteral administration, the targeting agent and/or the immunoconjugate may be dissolved in a pharmaceutical carrier and administered as either a solution or a suspension. Illustrative of suitable carriers are water, saline, phosphate buffer solution (PBS), dextrose solutions, fructose solutions, ethanol, or oils of animal, vegetative or synthetic origin. The carrier may also contain other ingredients, for example, preservatives, suspending agents, solubilizing agents, buffers and the like. When the unconjugated targeting agent and/or immunoconjugate are being administered intracerebroventricularly or intrathecally, they may also be dissolved in cerebrospinal fluid.

Dosages administered to a subject may be specified as amount, per surface area of the subject (which include humans as well as non-human animals). The dose may be administered to such a subject in amounts, preferably, but not exclusively from about 5 mg/m^2 to about 300 mg/m^2, including about 20mg/m^2, about 50mg/m^2, about 100mg/m^2, about 150mg/m^2, about 200mg/m^2 and about 250mg/m^2. The targeting agents/ immunoconjugates are suitably administered at one time or over a series of treatments. In a multiple dose regime these amounts may be administered once a day, once a week, once every two weeks, once every three weeks, once every four weeks, one every five weeks or once every six weeks. Loading doses with a single high dose or, alternatively,

lower doses that are administered shortly after one another followed by dosages timed at longer intervals constitute a preferred embodiment of the present invention. In a preferred embodiment, the timing of the dosages are adjusted for a subject so that enough time has passed prior to a second and/or any subsequent treatment so that the previous dose has been metabolized substantially, but the amount of immunoconjugate present in the subject's system still inhibits, delays and/or prevents the growth of a tumor. An exemplary "repeated single dose" regime comprises administering an initial dose of immunoconjugate of about 200mg/m^2 once every three weeks. Alternatively, a high initial dose may be followed by a biweekly maintenance dose of about 150µg/m^2. However, other dosage regimens may be useful. The progress of this therapy is easily monitored by known techniques and assays. Dosage may vary depending on whether they are administered for preventive or therapeutic purposes, the course of any previous therapy, the patient's clinical history and response to the targeting agent/immunoconjugate, and the discretion of the attending physician.

In accordance with the present invention, MM is treated as follows, with the use of nBT062 and BT062 as an example. This example is not intended to limit the present invention in any manner, and a skilled artisan could readily determine other immunoconjugate and nBT062 based systems that are within the scope of the present invention and other treatment regimes which could be utilized for the treatment of diseases such as MM.

Due to the selective expression of CD138 on patient MM cells on via the blood stream accessible cells, the specificity of nBT062 and the stability of BT062 in the bloodstream, BT062 remove the systemic toxicity of DM4 and provides an opportunity to target the delivery of the DM4-effector molecule(s). Administration of dosages of nBT062 is benefical to shield low expressing CD138 positive, non-tumor cells against BT062 binding of those cells and preferably destruction, while the immunoconjugates ~~of this invention~~ provide a means for the effective administration of the effector molecules to cell sites where the effector molecules can be released from the immunoconjugates. This targeted delivery and release provides a significant advance in the treatment of multiple myeloma, for which current chemotherapy methods sometimes provide incomplete remission.

The present invention is further described by reference to the following Examples, which are offered by way of illustration and are not intended to limit the invention in any manner. Standard techniques well known in the art or the techniques specifically described below are utilized.

Materials and Methods

Chimeric Antibody Construction (cB-B4: nBT062)

B-B4

Murine antibody B-B4 as previously characterized (Wijdenes *et al.*, Br J Haematol., 94 (1996), 318) was used in these experiments.

Cloning and expression of B-B4 and cB-B4 / nBT062

Standard recombinant DNA techniques were performed as described in detail in text books, for example in J. Sambrook; Molecular Cloning, A Laboratory Manual; 2nd Ed. (1989), Cold Spring Harbor Laboratory Press, USA, or as recommended by the manufacturer's instruction in the cases when kits were used. PCR-cloning and modification of the mouse variable regions have been conducted using standard PCR methodology. Primers indicated in the respective results section have been used.

Expression of cB-B4 / nBT062

Exponentially growing COS cells, cultured in DMEM supplemented with 10% FCS, 580 µg/ml L-glutamine, 50 Units/ml penicillin and 50 µg/ml streptomycin were harvested by trypsinisation and centrifugation and washed in PBS. Cells were resuspended in PBS to a final concentration of $1x10^7$ cells/ml. 700 µl of COS cell suspension was transferred to a Gene Pulser cuvette and mixed with heavy and kappa light chain expression vector DNA (10 µg each or 13 µg of Supervector). Cells were electroporated at 1900 V, 25 µF using a Bio-Rad Gene Pulser. Transformed cells were cultured in DMEM supplemented with 10% gamma-globulin free FBS, 580 µg/ml L-glutamine, 50 Units/ml penicillin and 50 µg/ml streptomycin for 72 h before antibody-containing cell culture supernatants were harvested.

Capture ELISA to measure expression levels of cB-B4 / nBT062

96 well plates were coated with 100 µl aliquots of 0.4 µg/ml goat anti-human IgG antibody diluted in PBS (4°C, overnight). Plates were washed three times with 200

µl/well washing buffer (PBS+0.1% Tween-20). Wells were blocked with 0.2% BSA, 0.02% Tween-20 in PBS, before addition of 200 µl cell culture supernatants containing the secreted antibody (incubation at 37°C for one hour). The wells were washed six times with washing buffer, before detection of bound antibody with goat anti-human kappa light chain peroxidase conjugate.

Purification of cB-B4 / nBT062 from cell culture supernatants

The cB-B4 antibody was purified from supernatants of transformed COS 7 cells using the Protein A ImmunoPure Plus kit (Pierce, Rockford, IL), according to the manufacturer's recommendation.

cB-B4 binding and competition assay

Analysis of binding activity of B-B4 and cB-B4 to CD138 was performed using the Diaclone (Besançon, France) sCD138 kit according to the manufacturer's recommendation, considering the changes described in the results section.

RNA preparation and cDNA synthesis

Hybridoma B-B4 cells were grown and processed using the Qiagen Midi kit (Hilden, Germany) to isolate RNA following the manufacturer's protocol. About 5 µg of B-B4 RNA was subjected to reverse transcription to produce B-B4 cDNA using the Amersham Biosciences (Piscataway, NJ) 1st strand synthesis kit following the manufacturer's protocol.

Cloning of B-B4 immunoglobulin cDNA

Immunoglobulin heavy chain (IgH) cDNA was amplified by PCR using the IgH primer MHV7 (5'-ATGGGCATCAAGATGGAGTCACAGACCCAGG-3') **[SEQ ID NO:3]** and the IgG1 constant region primer MHCG1 (5'-CAGTGGATAGACAGATGGGGG-3') **[SEQ ID NO:4]**. Similarly, immunoglobulin light chain (IgL) was amplified using the three different Igκ primers MKV2 (5'-ATGGAGACAGACACACTCCTGCTATGGGTG-3') **[SEQ ID NO:5]**, MKV4 (5'-ATGAGGGCCCCTGCTCAGTTTTTTGGCTTCTTG-3') **[SEQ ID NO:6]** and MKV9 (5'-ATGGTATCCACACCTCAGTTCCTTG-3') **[SEQ ID NO:7]**, each in combination with primer MKC (5'-ACTGGATGGTGGGAAGATGG-3') **[SEQ ID NO:8]**. All amplification products were directly ligated with the pCR2.1-TOPO vector using the

TOPO-TA cloning kit (Invitrogen, Carlsbad, CA) according to the manufacturer's instruction.

E. coli TOP10 bacteria (Invitrogen) transformed with the ligated pCR2.1 vector constructs were selected on LB-ampicillin-Xgal agar plates. Small scale cultures were inoculated with single white colonies, grown overnight and plasmids were isolated using the QIAprep Spin Miniprep kit according to the manufacturer's instruction.

cDNA sequence determination

Plasmids were sequenced using the BigDye Termination v3.0 Cycle Sequencing Ready Reaction Kit (ABI, Foster City, CA). Each selected plasmid was sequenced in both directions using the 1210 and 1233 primers cycled on a GeneAmp9600 PCR machine. The electrophoretic sequence analysis was done on an ABI capillary sequencer.

The complete cycle of RT-PCR, cloning and DNA sequence analysis was repeated to obtain three completely independent sets of sequence information for each immunoglobulin chain.

B-B4 Vκ DNA sequence

1st strand synthesis was performed in three independent reactions. The PCR products generated by using primers MKC and MKV2 (sequences given above) were ligated into pCR2.1-TOPO vectors according to the manufacturer's instruction. Clones from each independent set of RT-PCR reactions were sequenced in both directions. MKV2-primed product sequence was highly similar to sterile kappa transcripts originating from the myeloma fusion partner such as MOPC-21, SP2 and Ag8 (Carroll *et al.*, Mol Immunol., 25 (1988), 991; Cabilly *et al.*, Gene, 40 (1985); 157) and was therefore disregarded.
The PCR products using MKC with MKV4 and MKV9 primers were similar to each other and differed only at the wobble positions within the leader sequence primer.

B-B4 VH DNA sequence

1st strand synthesis was performed in three independent reactions and PCR products were cloned and sequenced from each 1st strand product. Five clones were sequenced from each 1st strand.

Construction of chimeric cB-B4 expression vectors

The construction of the chimeric expression vectors entails adding a suitable leader sequence to VH and Vκ, preceded by a *Bam*HI restriction site and a Kozak sequence. The Kozak consensus sequence is crucial for the efficient translation of a variable region sequence. It defines the correct AUG codon from which a ribosome can commence translation, and the single most critical base is the adenine (or less preferably, a guanine) at position –3, upstream of the AUG start. The leader sequence is selected as the most similar sequence in the Kabat database (Kabat *et al.*, NIH National Technical Information Service, 1991). These additions are encoded within the forward (For) primers (both having the sequence 5'-AGAGAAGCTT**GCCGCCACC**ATGATT-GCCTCTGCTCAGTTCCTTGGTCTCC-3' **[SEQ ID NO:9]**; restriction site is underlined; Kozak sequence is in bold type). Furthermore, the construction of the chimeric expression vectors entails introducing a 5' fragment of the human gamma1 constant region, up to a natural *Apa*I restriction site, contiguous with the 3' end of the J region of B-B4 and, for the light chain, adding a splice donor site and *Hind*III site. The splice donor sequence is important for the correct in-frame attachment of the variable region to its appropriate constant region, thus splicing out the V:C intron. The kappa intron + CK are encoded in the expression construct downstream of the B-B4 Vκ sequence. Similarly, the gamma-4 CH is encoded in the expression construct downstream of the B-B4 VH sequence.

The B-B4 VH and Vκ genes were first carefully analyzed to identify any unwanted splice donor sites, splice acceptor sites, Kozak sequences and for the presence of any extra sub-cloning restriction sites which would later interfere with the subcloning and/or expression of functional whole antibody. An unwanted *Hind*III site was found in the Vκ sequence which necessarily was removed by site-directed mutagenesis via PCR without changing the amino acid sequence. For this reactions, oligonucleotide primers BT03 (5'-CAACAGTATAGTAAGCTCCCTCGGACGTTCGGTGG-3') **[SEQ ID NO:10]** and BT04 (5'-CCACCGAACGTCCGAGGGAGCTTACTATACTGTTG-3') **[SEQ ID NO:11]** were used and mutagenesis was performed according to the Stratagene (La Jolla, CA) Quickchange Mutagenesis Kit protocol.

Kappa chain chimerization primers

The non-ambiguous B-B4 Vκ leader sequence, independent of the PCR primer sequence, was aligned with murine leader sequences in the Kabat database. The

nearest match for the B-B4 VH leader was VK-10 ARS-A (Sanz *et al.*, PNAS, 84 (1987), 1085). This leader sequence is predicted to be cut correctly by the SignalP algorithm (Nielsen *et al.*, Protein Eng, 10 (1997); 1). Primers CBB4Kfor (see above) and g2258 (5'-CGCGGGATCCACTCACGTTTGATTTCCAGCTTGGTGCCTCC-3' **[SEQ ID NO:12]**; Restriction site is underlined) were designed to generate a PCR product containing this complete leader, the B-B4 Vκ region, and *Hind*III and *Bam*HI terminal restriction sites, for cloning into the pKN100 expression vector. The forward primer, CBB4K introduces a *Hind*III restriction site, a Kozak translation initiation site and the VK-10 ARS-A leader sequence. The reverse primer g2258 introduces a splice donor site and a *Bam*HI restriction site. The resulting fragment was cloned into the *Hind*III/*Bam*HI restriction sites of pKN100.

Heavy chain chimerization primers

The non-ambiguous B-B4 VH leader sequence, independent of the PCR primer sequence, was aligned with murine leader sequences in the Kabat database. The nearest match for the B-B4 VK leader was VH17-1A (Sun *et al.*, PNAS, 84 (1987), 214). This leader sequence is predicted to be cut correctly by the SignalP algorithm. Primers cBB4Hfor (see above) and g22949 (5'-CGATGGGCCCTTGGTGGAGGCTGAGGA-GACGGTGACTGAGGTTCC-3' **[SEQ ID NO:13]**; Restriction site is underlined) were designed to generate a PCR product containing VH17-1A leader, the B-B4 VH region, and terminal *Hind*III and *Apa*I restriction sites, for cloning into the pG4D200 expression vector. The forward primer cBBHFor introduces a *Hind*III restriction site, a Kozak translation initiation site and the VH17-1A leader sequence. The reverse primer g22949 introduces the 5' end of the gamma4 C region and a natural *Apa*I restriction site. The resulting fragment was cloned into the *Hind*III/*Apa*I restriction sites of pG4D200, resulting in vector pG4D200cBB4.

Production of cBB4 antibody

One vial of COS 7 cells was thawed and grown in DMEM supplemented with 10% Fetal clone I serum with antibiotics. One week later, cells (0.7 ml at 10^7 cells/ml) were electroporated with pG4D200cBB4 plus pKN100cBB4 (10 µg DNA each) or no DNA. The cells were plated in 8 ml growth medium for 4 days. Electroporation was repeated seven times.

Detection of chimeric antibody

A sandwich ELISA was used to measure antibody concentrations in COS 7 supernatants. Transiently transformed COS 7 cells secreted about 6956 ng/ml antibody (data not shown).

Binding activity of cB-B4

To assay the binding activity of cB-B4 in COS 7 culture supernatants, the Diaclone sCD138 kit has been used, a solid phase sandwich ELISA. A monoclonal antibody specific for sCD138 has been coated onto the wells of the microtiter strips provided. During the first incubation, sCD138 and biotinylated B-B4 (bio-B-B4) antibody are simultaneously incubated together with a dilution series of unlabeled test antibody (B-B4 or cB-B4).

The concentrations of bio-B-B4 in this assay have been reduced in order to obtain competition with low concentrations of unlabeled antibody (concentration of cB-B4 in COS 7 cell culture supernatants were otherwise too low to obtain sufficient competition). Results from this assay reveal that both antibodies have the same specificity for CD138 (data not shown).

Purification of cB-B4

Chimeric B-B4 was purified from COS 7 cell supernatants using the Protein A ImmunoPure Plus kit (Pierce), according to the manufacturer's recommendation (data not shown).

K_D-determination: Comparison nBT062/ BB4

Purification of soluble CD138

Soluble CD138 antigen from U-266 cell culture supernatant was purified by FPLC using a 1 mL "HiTrap NHS-activated HP " column coupled with B-B4. Cell culture supernatant was loaded in PBS-Buffer pH 7.4 onto the column and later on CD138 antigen was eluted with 50 mM tri-ethylamine pH 11 in 2 mL fractions. Eluted CD138 was immediately neutralised with 375 µL 1 M Tris-HCl, pH 3 to prevent structural and/or functional damages.

Biotinylation of CD138

Sulfo-NHS-LC (Pierce) was used to label CD138. NHS-activated biotins react efficiently with primary amino groups like lysine residues in pH 7-9 buffers to form stable amide bonds.

For biotinylation of CD138, 50 µl of CD138 were desalted using protein desalting spin columns (Pierce). The biotinylation reagent (EZ-Link Sulfo NHS-LC-Biotin, Pierce) was dissolved in ice-cooled deionised H_2O to a final concentration of 0.5 mg/mL. Biotinylation reagent and capture reagent solution were mixed having a 12 times molar excess of biotinylation reagent compared to capture reagent (50 pmol CD138 to 600 pmol biotinylation reagent) and incubated 1 h at room temperature while shaking the vial gently. The unbound biotinylation reagent was removed using protein desalting columns.

Immobilization of bCD138

The sensorchip (SENSOR CHIP SA, BIACORE AB) used in the BIACORE assay is designed to bind biotinylated molecules for interaction analysis in BIACORE systems. The surface consists of a carboxymethylated dextran matrix pre-immobilized with streptavidin and ready for high-affinity capture of biotinylated ligands. Immobilization of bCD138 was performed on SENSOR CHIP SA using a flow rate of 10 µL/min by manual injection. The chip surface was conditioned with three consecutive 1-minute injections of 1 M NaCl in 50 mM NaOH. Then biotinylated CD138 was injected for 1 minute.

K_D-Determination of different antibodies using BIACORE

The software of BIACORE C uses pre-defined masks, so called "Wizards" for different experiments where only certain settings can be changed. As the BIACORE C was originally developed to measure concentrations, there is no wizard designed to carry out affinity measurements. However, with the adequate settings, the wizard for "non-specific binding" could be used to measure affinity rate constants and was therefore used for K_D-determination. With this wizard, two flow cells were measured and the dissociation phase was set to 90 s by performing the "Regeneration 1" with BIACORE running buffer. "Regeneration 2" which is equivalent to the real regeneration was performed with 10 mM Glycine-HCl pH 2.5. After this step, the ligand CD138 was in its binding competent state again. During the whole procedure HBS-EP was used as running and dilution buffer. To determine binding of the different antibodies (~150 kDa) to CD138, association and

dissociation was analysed at different concentrations (100, 50, 25 12.5, 6.25 and 3.13 nM). The dissociation equilibrium constants were determined by calculating the rate constants ka and kd. Afterwards, the K_D-values of the analytes were calculated by the quotient of kd and ka with the BIAevaluation software. The results are shown in **Table 4.**

Antibody	Affinity	
	K_D (nM)	mean K_D (nM)
nBT062	1.4 1.4 1.5	**1.4 +/- 0.06**
B-B4	1.7 1.7 1.6	**1.6 +/- 0.06**
nBT062-SPDB-DM4	1.9 1.9 1.9	**1.9 +/- 0.00**
B-B4-SPP-DM1	2.6 2.7 2.6	**2.6 +/- 0.06**

Table 4: Comparative analysis of K_D values of nBT062 and B-B4. Standard deviations are given for mean K_D values.

Discussion

Mean K_D values for each antibody were calculated from three independent experiments. The results show that in all measurements nBT062 exhibits slightly decreased K_D values compared to B-B4 (mean K_D values were 1.4 and 1.6 nM, respectively).

Preparation of Immunoconjugates

nBT062-DM1 and huC242-DM1

The thiol-containing maytansinoid DM1 was synthesized from the microbial fermentation product ansamitocin P-3, as previously described by Chari (Chari *et al.*, Cancer Res. 1 (1992), 127). Preparation of humanized C242 (huC242) (Roguska *et al.*, PNAS, 91 (1994), 969) has been previously described. Antibody-drug conjugates were prepared as previously described (Liu *et al.*, PNAS, 93 (1996), 8618). An average of 3.5 DM1 molecules was linked per antibody molecule.

nBT062-DM4

BT062 is an antibody-drug conjugate composed of the cytotoxic maytansinoid drug, DM4, linked via disulfide bonds through a linker to the nBT062 chimerized monoclonal antibody. Maytansinoids are anti-mitotics that inhibit tubulin polymerization and microtubule assembly (Remillard *et al.*, Science 189 (1977), 1002). Chemical and schematic representations of BT062 (nBT062-DM4) are shown in **FIGS. 1 and 2.**

Synthesis of DM4

DM4 is prepared from the well known derivative maytansinol (Kupchan *et al.*, J. Med. Chem., 21 (1978), 31). Maytansinol is prepared by reductive cleavage of the ester moiety of the microbial fermentation product, ansamitocin P3, with lithium trimethoxyaluminum hydride (see **FIG. 3**).

DM4 is synthesized by acylation of maytansinol with N-methyl-N-(4-methydithiopentanoyl)-L-alanine (DM4 side chain) in the presence of dicyclohexylcarbodiimide (DCC) and zinc chloride to give the disulfide-containing maytansinoid DM4-SMe. The DM4-SMe is reduced with dithiothreitol (DTT) to give the desired thiol-containing maytansinoid DM4 (see **FIG. 4** for the DM4 process flow diagram).

Immunoconjugate BT062

The procedure for the preparation of nBT062-DM4 is outlined in **FIG. 5**. The nBT062 antibody is modified with N-succinimidyl-4-(2-pyridyldithio) butyrate (SPDB linker) to introduce dithiopyridyl groups. DM4 is mixed with the modified antibody at a concentration in excess of the equivalents of dithiopyridyl groups. The BT062 conjugate forms by a disulfide exchange reaction between the thiol group of DM4 and the dithiopyridyl groups introduced into the antibody via the linker. Purification by chromatography and diafiltration removes the low molecular weight reactants (DM4) and reaction products (thiopyridine), as well as aggregates of conjugated antibody, to produce the bulk drug substance.

FACS analysis and WST cytotoxicity assays

FACS analysis

OPM-2 cells are plasma cell leukemia cell lines showing highly expressing CD138. OPM-2 cells were incubated with nBT062, nBT062-SPDB-DM4, nBT062-SPP-DM1 or

nBT062-SMCC-DM1 at different concentrations (indicated in **FIG. 6**). The cells were washed and CD138-bound antibody or conjugates were detected using a fluorescence-labeled secondary antibody in FACS analysis. The mean fluorescence measured in these experiments was plotted against the antibody concentration.

Cell viability assay

$CD138^+$ MOLP-8 cells were seeded in flat bottom plates at 3000 cells/well. $CD138^-$ BJAB control cells were seeded at 1000 cells/well. The cells were treated with nBT062-SPDB-DM4, nBT062-SPP-DM1 or nBT062-SMCC-DM1 at different concentrations (indicated in **FIG. 7**) for five days. WST reagent (water-soluble tetrazolium salt, ROCHE) was added in order to measure cell viability according to the manufacturer's instruction (ROCHE). The reagent was incubated for 7.5 h on MOLP-8 cells and for 2 h on BJAB cells. The fraction of surviving cells was calculated based on the optical densities measured in a microplate reader using standard procedures.

Discussion

Binding of nBT062-SPDB-DM4, nBT062-SPP-DM1, nBT062-SMCC-DM1 or nBT062 was analyzed by FACS. $CD138^+$ OPM-2 as target cells were incubated with nBT062 or immunoconjugates and cell-bound molecules were detected using a fluorescence-labeled secondary antibody. In **FIG. 6**, the mean fluorescences as measure for the amount of cell bound antibody is plotted against different antibody or conjugate concentrations. The results show, that nBT062-SPDB-DM4, nBT062-SPP-DM1 and nBT062-SMCC-DM1 show very similar binding characteristics. In addition, the results strongly suggest that the binding characteristics of the unconjugated antibody is not affected by the conjugated toxins.

In cell viability assays, the cytotoxic activity of the antibody against $CD138^+$ MOLP-8 target cells and against $CD138^-$ BJAB B-lymphoblastoma control cells were analyzed. Both cell lines were seeded in flat-bottom plates and incubated with increasing concentrations of the immunoconjugates. Unconjugated antibody was used as a control. The cytotoxic activity was analyzed five days after addition of the immunoconjugates by using WST reagent in order to measure cell viability. In **FIG. 7 (A)-(C)**, the fraction of surviving cells relative to control cells treated with vehicle control is plotted against increasing immunoconjugate concentrations. The results show that cytotoxic activity of nBT062-SPDB-DM4, nBT062-SPP-DM1 and nBT062-SMCC-DM1 against MOLP-8 cells

is very similar. As expected, CD138$^-$ BJAB control cells were not killed by the immunoconjugates, indicating that all immunoconjugates act via cell specific binding to CD138. In competition experiments, in which MOLP-8 cells were preincubated with a molar excess of unconjugated nBT062. Preincubation substantially blocked the cytotoxicity of nBT062-SPDB-DM4, providing further evidence that the immunoconjugates kill the cells via specific binding to CD138 onto the cell surface (**FIG. 7 (D)**).

Xenograft mouse experiments

To evaluate the importance of CD138 targeting on the anti-tumor activity of antibody-maytansinoid conjugates of a human chimeric version of the B-B4 antibody, nBT062, xenograft mouse experiments were performed. Two versions of nBT062-maytansinoid conjugates were prepared that may differ in the chemical stability of their disulfide linkages (nBT062-SPP-DM1 and nBT062-SPDB-DM4). The anti-tumor activity of these antibody-drug conjugates was compared to the activity of the B-B4-SPP-DM1 conjugate (comprising the murine parental antibody), as well as unconjugated free maytansinoid (DM4), native unmodified nBT062 antibody, and a non-targeting (irrelevant) IgG1-maytansinoid conjugate. The conjugates were evaluated in a CD138-positive xenograft model (MOLP-8) of human multiple myeloma in severe combined immunodeficient (SCID) mice.

In these mice, subcutaneous tumors were established (female CB.17 SCID mice) by inoculation with MOLP-8 cell suspensions. Treatment with a single bolus intravenous injection was conducted when tumor volumes reached an average 113 mm^3. Changes in tumor volume and body weight were monitored twice per week. Experiments were carried out over 68 days after tumor cell inoculation.

Xenograft mouse experiments A

Mice

Female CB.17 SCID mice, five weeks old, were obtained from Charles River Laboratories.

Human tumor cell lines

MOLP-8, a human multiple myeloma cell line, was supplied from ATCC. MOLP-8 cells, which express the CD138 antigen on their cell surface and develop xenograft

tumors in SCID mice, were maintained in RPMI-1640 medium supplemented with 4 mM L-glutamine (Biowhittaker, Walkersville, MD), 10% fetal bovine serum (Hyclone, Logan, Utah) and 1% streptomycin/penicillin, at 37°C in a humidified atmosphere that contained 5% CO_2.

PART I

Tumor growth in mice

Each mouse was inoculated with $1x10^7$ MOLP-8 cells subcutaneously into the area under the right shoulder. The total volume was 0.2 ml per mouse, in which the ratio of serum-free medium to matrigel (BD Bioscience, Bedford, MA) was 1/1 (v/v). Prior to treatment, the xenograft tumors were monitored daily and were allowed to become established. The tumor volume reached approximately 113 mm^3 about 11 days after tumor cell inoculation. Tumor take rate of CB.17 SCID mice was 100%.

Eleven days after tumor cell inoculation, 42 mice were selected based on tumor volumes and body weights. The tumor volume was in a range of 68.2 to 135.9 mm^3. The forty-two mice were randomly divided into seven groups (A-G) of six animals each based on tumor volume.

Each of six mice in Group A received 200 µl of PBS as vehicle control. Each mouse in group B received 13.8 mg/kg of nBT062 naked antibody. This dose is equivalent to the amount of nBT062 antibody component in 250 µg/kg of linked maytansinoid. The ratio of molecular weights of maytansinoids to nBT062 antibody in a conjugate molecule is approximate 1/55. Each mouse in Group C received 250 µg/kg of DM4. Each mouse in Group D received 250 µg/kg of huC242-DM4. Mice in groups E, F and G received 250 µg/kg of nBT062-SPDB-DM4, B-B4-SPP-DM1 and nBT062-SPP-DM1 each, respectively.

All agents were intravenously administered as a single bolus injection through a lateral tail vein with a 1 ml syringe fitted with a 27 gauge, ½ inch needle. Prior to administration, the stock solutions of nBT062 antibody, nBT062-SPDB-DM4 and nBT062-SPP-DM1 were diluted with sterile PBS to concentrations of 2 mg/ml, 28.1 µg/ml and 28.1 µg/ml, respectively, so that the injected volume for each mouse was between 120-220 µl.

PART II

In a second set of experiments, MOLP-8 cells ($1.5x10^7$ cells per mouse), suspended in a 50:50 mixture of serum free media and matrigel were injected subcutaneously in the area under the right shoulder in 100 µl. Tumor volumes reached about 80 mm^3 at day 11 and the mean of the controls was about 750 mm^3 at day 25, post cell inoculation. The tumor doubling time was estimated to be 4.58 days. Each mouse in the control group (n=6) received 0.2 ml of sterile PBS administered into the lateral tail vein (i.v.) in a bolus injection. All treatment doses were based on conjugated maytansinoid. Nine groups (n=6) were treated with a single intravenous injection of nBT062-SMCC-DM1, nBT062-SPDB-DM4, or nBT062-SPP-DM1, each at doses of 450, 250 and 100 µg/kg. An additional group (n=6) received 250 µg/kg nBT062-SMCC-DM1 in a repeated dosing (weekly for five weeks). Mice were randomized into eleven groups (n=6) by tumor volume using the LabCat Program. The tumor volumes ranged from 40.0 to 152.5 mm^3. The mice were dosed based on the individual body weight.

Tumor size was measured twice per week in three dimensions using the LabCat System (Tumor Measurement and Tracking, Innovative Programming Associated, Inc., Princeton, NJ). The tumor volume in mm^3 was calculated using the methodology described in Tomayko *et al.*, Cancer Chemother. Pharmacol, 24 (1989), 148:

Volume=Length x Width x Height x ½

Log_{10} cell kill was calculated with the formula described in Bissery *et al.*, Cancer Res., 51 (1991), 4845:

Log_{10} cell kill = (T-C) / T_d x 3.32

where (T-C) or tumor growth delay, is the median time in days required for the treatment group (T) and the control group (C) tumors, to reach a predetermined size (600 mm^3). T_d is the tumor doubling time, based on the median tumor volume in the control mice, and 3.32 is the number of cell doublings per log of cell growth.

Results

The tumor growth in individual mice is shown in **FIGS. 8** and **9**. The mean (+/- SD) tumor growth for each group is shown in **FIG. 10**.

As compared with tumor growth in the PBS-treated animals, treatment with nBT062 antibody, unconjugated free DM4 or the irrelevant non-targeting conjugate huC242-DM4 did not cause any significant inhibition of tumor growth.

All three CD138-targeting conjugates, nBT062-SPDB-DM4, B-B4-SPP-DM1 and nBT062-SPP-DM1, at a dose of 250 µg/kg caused marked delay in tumor growth. Based on the mean tumor volumes measured in the treatment groups, the DM4 conjugate nBT062-SPDB-DM4 was the most active one, while the nBT062-SPP-DM1 conjugate showed slightly increased activity as compared to its murine counterpart B-B4-SPP-DM1 (**FIG. 10**). The results obtained in individual mice show in addition that the anti-tumor activity obtained with B-B4-SPP-DM1 is more heterogeneously and therefore less predicable than that measure in mice treated with nBT062-SPP-DM1. In terms of homogeneity of anti tumor activity, the other conjugate that uses nBT062 as targeting antibody nBT062-SPDB-DM4 behaved similar to nBT062-SPP-DM1.

No body weight reduction was observed in any treatment group suggesting that the treatments were well tolerated.

Discussion

The results of the analysis of three CD138-targeting conjugates in experimental animals demonstrate the importance of targeted delivery for the anti-tumor activity. While the maytansinoid conjugates of the human chimeric nBT062 and the murine B-B4 antibodies show significant activity as measured by log cell kill, there was no significant impact on tumor growth from treatment with unconjugated DM4, unmodified native huBT062 antibody, or a non-targeting control conjugate (huC242-DM4).

The immunoconjugate prepared from the human chimeric antibody, nBT062-SPP-DM1, gave slightly higher anti-tumor activity then the conjugate prepared from its murine counterpart, B-B4-SPP-DM1. In addition, treatment with nBT062-SPP-DM1 and nBT062-SPDB-DM4 resulted in more homogenous responses in individual mice as compared to treatment with B-B4-SPP-DM1. The high binding variation of B-B4-SPP-DM1 explained that the measurement of the median tumor volume (+/- SD) of MOLP-8 human multiple myeloma xenografts in CB.17 SCID mice over time (days) post-inoculation actually provided for relatively better results for B-B4-SPP-DM1 than for nBT062-SPP-DM1 (data not shown).This feature of immunoconjugates using nBT062 as a targeting antibody seems to be beneficial especially for therapeutic use of the conjugates.

Lastly, the most potent of the maytansinoid conjugates, following single iv administration in the MOLP-8 xenograft models in SCID mice, was nBT062-SPDB-DM4.

Xenograft mouse experiments B

In this set of experiments, eighty-five mice were inoculated with MOLP-8 cells ($1.5x10^7$ cells/mouse) subcutaneously in the right shoulder. Tumor take rate was 100%. Sixty-six SCID mice bearing bulky MOLP-8 tumors with a mean tumor volume of about 80 mm^3 were randomized into eleven treatment groups (n=6). Mice were treated with a single dose of one of three conjugates (nBT062-SMCC-DM1, nBT062-SPDB-DM4 or nBT062-SPP-DM1). An additional group received five weekly doses of nBT062-SMCC-DM1 and a control group received a single dose of PBS. Mean tumor volumes are shown in **FIG. 11A**. A dose response was established for each conjugate. A median tumor volume of 750 mm^3 in the PBS-treated animals was reached on day 25. Tumor doubling time determined by the best-fit linear regression curve fit on a log-linear plot of control tumor growth was 4.58 days. Animals treated with nBT062-SPDB-DM4 at 450 µg/kg had the highest log cell kill (LCK=2.89), followed by animals treated with nBT062-SMCC-DM1 at 250 µg/kg weekly dosing (LCK=2.1; see **Table 5**). Comparison of the mean tumor growth curves for the treatment groups by repeated measures ANOVA performing Dunnett's Multiple Comparisopn Test showed a significant difference between the PBS control group and 450 µg/kg nBT062-SPDB-DM4 ($p<0.01$), 250 µg/kg nBT062-SPDB-DM4 ($p<0.05$) and five weekly doses of 250 µg/kg nBT062-SMCC-DM1 ($p<0.05$). No partial or complete tumor regression in any of the treatment groups occurred with the exception of one animal receiving 450 µg/kg nBT062-SPDB-DM4, which had partial regression of the tumor until day 85 post-inoculation.

Table 5. Log cell kill (LCK) values as measure for anti-tumor activity of different nBT062-DMx conjugates in different dosing schemes. Refer to the Materials and methods section for information on calculation of LCK values.

Test Material	Dose (µg/kg)	LCK	Dosing
PBS			single dose
nBT062-SMCC-DM1	450	0.85	single dose
nBT062-SMCC-DM1	250	0.53	single dose
nBT062-SMCC-DM1	100	0	single dose
nBT062-SPDB-DM4	450	2.89	single dose
nBT062-SPDB-DM4	250	1.05	sinle dose
nBT062-SPDB-DM4	100	0.39	single dose
nBT062-SPP-DM1	450	0.8	single dose
nBT062-SPP-DM1	250	0.39	single dose
nBT062-SPP-DM1	100	0.2	single dose
nBT062-SMCC-DM1	250	2.1	weekly for 5 weeks

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What is claimed is:

1. An engineered targeting antibody which recognizes CD138 comprising
an antigen binding region (ABR) against CD138, wherein said antigen binding region is of a non-human antibody, and
a further antibody region, comprising a human antibody region,
wherein the engineered targeting antibody:
(a) provides for homogenous binding to CD138 of CD138 expressing cells, and optionally,
(b) binds CD138 with a binding affinity that exceeds the binding affinity of said non-human antibody; and
wherein the antigen binding region comprises:
(a) a heavy chain variable region comprising amino acid residues 31 to 35 (CDR1), 51 to 68 (CDR2) and 99 to 111 (CDR3) of SEQ ID NO: 1, and
(b) a light chain variable region comprising amino acid residues 24 to 34 (CDR1), 50 to 56 (CDR2) and 89 to 97 (CDR3) of SEQ ID NO: 2.

2. The engineered targeting antibody of claim 1, wherein said further antibody region is at least one constant region comprising a heavy chain constant region or a part thereof that is of a human antibody, and wherein said engineered antibody is of an IgG4 isotype.

3. The engineered targeting antibody of claim 2, wherein said antibody binds CD138 with a binding affinity that exceeds the binding affinity of said non-human antibody.

4. The engineered targeting antibody of claim 2, wherein said antibody provides for homogenous binding to CD138 of CD138 expressing cells.

5. The engineered targeting antibody of claim 4, wherein said engineered targeting antibody binds CD138 with a targeting variation of less than 150%, 140%, 130%, 120%, 110%, 100%, 90%, 80%, 70%, 60% or 50%.

6. The engineered targeting antibody of any of claims 1 to 5, wherein said further antibody region comprises:
(a) amino acid residues 123 to 448 of SEQ ID NO: 1, and/or
(b) amino acid residues 108 to 214 of SEQ ID NO: 2, respectively
and mutations thereof that
(i) maintain or lower the antibody-dependent cytotoxicity and/or complement-dependent cytotoxicity of the engineered targeting antibody and/or

(ii) stabilize the engineered targeting antibody.

7. The engineered targeting antibody of claim 1 or 2, wherein said antigen binding region of said non-human antibody is of a mouse antibody.

8. The engineered targeting antibody of claim 1 or 2, wherein said engineered targeting antibody is a chimeric antibody and said non-human antibody is B-B4.

9. The engineered targeting antibody of claim 3, wherein said engineered targeting antibody comprises at least one heavy chain and one light chain and wherein
(a) said heavy chain has at least about 70% sequence identity with SEQ ID NO:1; and/or
(b) said light chain has at least about 70% sequence identity with SEQ ID NO:2.

10. The engineered targeting antibody of claim 6, wherein
(a) said heavy chain has at least about 90% sequence identity with SEQ ID NO:1; and
(b) said light chain has at least about 90% sequence identity with SEQ ID NO:2.

11. The engineered targeting antibody of claims 1 or 2, wherein the ABR consisting essentially of an ABR of a non-human antibody.

12. A pharmaceutical composition comprising the antibody of claims 1 or 2, and a pharmaceutically acceptable carrier.

13. A hybridoma which produces the antibody of claim 1 or 2.

14. An antibody based assay comprising the antibody of claim 1 or 2.

15. An isolated polypeptide comprising an amino acid sequence of an immunoglobulin heavy chain or part thereof, wherein said immunoglobulin heavy chain or part thereof has at least 70%, at least 80%, at least 90%, at least 95% or at least 98% sequence identity with SEQ ID NO:1, wherein a targeting agent comprising said immunoglobulin heavy chain or part thereof targets CD138.

16. The isolated polypeptide of claim 15, wherein a constant region of said immunoglobulin heavy chain or said part thereof is an IgG4 isotype constant region.

17. The isolated polypeptide of claim 15, wherein said targeting agent is a mouse human chimeric antibody.

18. An isolated polypeptide of claim 15 further comprising an amino acid sequence of an immunoglobulin light chain or part thereof, wherein said immunoglobulin light chain or part thereof has at least 70%, at least 80%, at least 90%, at least 95% or at least 98% sequence identity with SEQ ID NO:2.

19. An isolated polypeptide according to claim 15, wherein said immunoglobulin heavy chain is identical to the sequence of SEQ ID NO:1.

20. An isolated polynucleotide according to claim 18 or 19, wherein said immunoglobulin light chain is identical to the sequence of SEQ ID NO:2.

21. Use of an engineered targeting antibody according to any of claims 1 to 11 in the manufacture of a medicament for inhibiting tumor growth in an individual with CD138 expressing cells, wherein the engineered targeting antibody homogenously binds CD138 expressed on the CD138 expressing cells.

22. The use according to claim 21, wherein said engineered targeting antibody binds CD138 with a targeting variation of less than 150%, 140%, 130%, 120%, 110%, 100%, 90%, 80%, 70%, 60% or 50%.

SEQUENCE LISTING

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<130> 319936.WO/JND/CJS

<150> 61/016,630
<151> 2007-12-26

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Gly Glu Ile Leu Pro Gly Thr Gly Arg Thr Ile Tyr Asn Glu Lys Phe
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Lys Gly Lys Ala Thr Phe Thr Ala Asp Ile Ser Ser Asn Thr Val Gln
65                  70                  75                  80

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Ala Val Leu Gln Ser Ser Gly Leu Tyr Ser Leu Ser Ser Val Val Thr
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Val Pro Ser Ser Ser Leu Gly Thr Lys Thr Tyr Thr Cys Asn Val Asp
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His Lys Pro Ser Asn Thr Lys Val Asp Lys Arg Val Glu Ser Lys Tyr
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Arg Thr Pro Glu Val Thr Cys Val Val Val Asp Val Ser Gln Glu Asp
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Pro Glu Val Gln Phe Asn Trp Tyr Val Asp Gly Val Glu Val His Asn
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Val Ser Val Leu Thr Val Leu His Gln Asp Trp Leu Asn Gly Lys Glu
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Tyr Lys Cys Lys Val Ser Asn Lys Gly Leu Pro Ser Ser Ile Glu Lys
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Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro Gln Val Tyr Thr
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