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Targeting tumors with nanobodies for cancer imaging and therapy 2

Sabrina Oliveira^{a,b}, Raimond Heukers^a, Jirawas Sornkom^a, Q1 Robbert J. Kok^c, Paul M.P. van Bergen en Henegouwen^{a,*} 4

^a Division of Cell Biology, Department of Biology, Faculty of Science, Utrecht University, Padualaan 8, 3584 CH Utrecht, The Netherlands

- ^b Department of Pathology, University Medical Center Utrecht, Heidelberglaan 100, 3584 CX Utrecht, The Netherlands 6
- ^c Department of Pharmaceutics, Utrecht Institute for Pharmaceutical Sciences, Faculty of Science, Utrecht University, Universiteitsweg 99, 3584 CG Utrecht, The Netherlands

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ABSTRACT

The use of monoclonal antibodies has revolutionized both cancer therapy and cancer imaging. Antibodies have 25 been used to directly inhibit tumor cell proliferation or to target drugs to tumors. Also in molecular imaging, 26 monoclonal antibodies have found their way to the clinic. Nevertheless, distribution within tumors is hampered 27 by their size, leading to insufficient efficacy of cancer treatment and irregular imaging. An attractive alternative 28 for monoclonal antibodies are nanobodies or VHHs. These are the variable domain of heavy-chain antibodies 29 from animals from the Camelidae family that were first discovered in 1993. Stimulated by the ease of nanobody 30 Q3 selection, production, and low immunogenicity potential, a number of nanobodies specific to different disease- 31 related targets have been developed. For cancer therapy, nanobodies have been employed as antagonistic 32 drugs, and more recently, as targeting moieties of effector-domaINS and of drug delivery systems. In parallel, 33 nanobodies have also been employed for molecular imaging with modalities such as nuclear and optical imaging. 34 In this review, we discuss recent developments in the application of nanobodies as targeting moieties in cancer 35 therapy and cancer imaging. With such a wide range of successful applications, nanobodies have become much 36 more than simple antagonists. 37

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Corresponding author, Tel.: + 31 30 253 3349. E-mail address: p.vanbergen@uu.nl (P.M.P. van Bergen en Henegouwen).

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1. Introduction

The use of monoclonal antibodies (mAbs) for cancer therapy has been 65 established extensively for over 15 years, with a number of impressive 66

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successes both for hematological malignancies and solid tumors treat-67 68 ments [1]. So far, there are 23 mAbs approved by the US Food and Drug Administration (FDA) on the market. Among these, six products are 69 70 specific for cancer, namely, rituximab (anti-CD20), trastuzumab (directed to HER2), bevacizumab (anti-VEGF), alemtuzumab (anti-CD52), 71 cetuximab, panitumumab, and matuzumab (all targeted to EGFR) 72 [2,3]. These mAbs interfere with the functioning of their target proteins 73 74in cancer, either by binding to transmembrane receptors or - in the 75case of bevacizumab - to the soluble ligand, thereby inhibiting tumor 76cell proliferation or tumor angiogenesis. As they all possess an intact 77 fragment crystallizable domain, i.e. Fc domain, they can interact with human complement or effector cells of the immune system, which 78 also contributes to their therapeutic effect. mAbs have also found their 79 80 way to the clinic for molecular imaging. In this case, mAbs are used to target radioactive or fluorescent tracers to the tumor, for either PET/ 81 SPECT or optical imaging, respectively [4-6]. Lastly, mAbs are used in 82 a variety of targeted nanomedicines, aiming at tumor cell directed de-83 livery of a cytotoxic payload [7]. It is however fair to state that the appli-84 cation of mAbs in both cancer therapy and imaging needs further 85 improvements. mAbs have a molecular weight of ~150 kDa and dimen-86 sions of 14.2 nm \times 8.5 nm \times 3.8 nm [8], which together with the 87 'binding site barrier' [9] limit mAb distribution and penetration into 05 89 the tumor. mAbs typically have several days of half-life in the bloodstream, which for molecular imaging results in high background levels. 90 Moreover, an important concern of mAbs application is their potential 91to induce immunogenic responses. To avoid unwanted immune re-92sponses in patients, mAbs are either completely humanized or pro-93 94duced as a chimeric protein. Altogether, these aspects have urged 95pharmaceutical companies and scientists to find new antibody formats 96 that provide the same binding specificity of mAbs, but with some of the 97 desired improvements.

As many of the mentioned drawbacks of mAbs are related to 98 99 their size, large efforts have been made towards the development of smaller antibody formats [10,11]. Naturally derived or synthetic 100 antigen-binding fragment (Fab; ~50 kDa), variable fragment (Fv; 101 ~15 kDa) and single-chain variable fragment (scFv; ~30 kDa) 102 were vastly tested and engineered to overcome the restrictions of 103 the full-length mAbs (Fig. 1) [10,12]. Nevertheless, their average 104 activities are still suboptimal due to lower affinities and limited 105stability, which is especially the case of scFv [13]. Apart from 106 those mentioned above, the minibody – an engineered antibody 107 108 fragment made by genetically fusing scFv binding domain to human CH3 - was introduced as another candidate for cancer immu-109 notherapy [14]. Furthermore, synthetic molecules or scaffold pro-110 teins, such as affibodies and DARPins (designed ankyrin repeat 111 proteins) have been developed, with important successes [15–17]. 112 113 However, no report has addressed their potential to induce immunological responses and their added value, compared to the other 114 platforms, still needs to be determined. 115

By serendipity, a special type of antibody was discovered in 116 animals from the Camelidae family by Hamers-Casterman and co-117 118 workers in 1993 [18]. These so-called heavy-chain antibodies 119 (HcAbs, ~95 kDa) are fully functional and, despite the absence of light chain and of the first constant domain (CH1), they bind their 120antigens with similar affinities to those of conventional antibodies 121[19]. Apart from Camelidae, some primitive fish species were also 122123found to produce different types of HcAbs, such as nurse shark and ratfish [20,21]. Interestingly, the variable domain alone of HcAbs 124(i.e. VHH) was proven to have sufficient antigen binding properties 125 and, as such, can be considered as the smallest naturally derived 126antigen-binding fragment with the approximate molecular weight 127 of 15 kDa [22,23]. The term 'nanobodies' was employed with respect 128 to their size in nanometer range by the Belgian company Ablynx®, 129and particularly refers to the VHH from camelid species [23-25]. 130Another term used for nanobodies is the single domain antibody 131 132 (sdAb) [26].



Fig. 1. Antibodies and their fragments. Schematic representation and corresponding molecular weight of (left) a monoclonal antibody, mAb, and its fragments, i.e., Fab', Fv, scFv; and of (right) a heavy chain only antibody, HcAb, together with its antigenbinding fragment, i.e. nanobody or VHH.

2. Nanobodies: Structure and characteristics

In 1994, the first detailed sequence of nanobody encoding genes was 134 published by Muyldermans and co-workers, providing more molecular 135 insights regarding their interaction and binding interface [22]. The 136 nanobody sequences were shown to have a high degree of identity 137 with the human type 3 VH domain (VH3), which most likely accounts 138 for the low immunological potential of nanobodies, as demonstrated 139 in mice [27]. In addition, humanization of nanobodies has been 140 performed before these were translated into the clinic (Ablynx) 141 [25,28,29], further minimizing their immunological potential. In this 142 line, Vincke and colleagues have presented the humanization of 143 dromedary-derived nanobodies resulting in a universal humanized 144 nanobody scaffold [30]. A number of distinctive amino acid substitu- 145 tions are specifically found in framework 2. In conventional antibodies, 146 this region serves as a part of the hydrophobic VL interface and, conse- 147 quently, substitutions that have occurred in HcAbs are thought to be the 148 main reason for the high hydrophilicity, stability and higher solubility of 149 VHHs as compared to conventional VH domains, including scFvs. Anoth- 150 er interesting difference between VHH and human VH domain is the 151 length of CDRs, which contributes to an increase of the antigen- 152 interacting surface [25,31]. A longer CDR3 in nanobodies allows it to 153 form a fingerlike structure able to extend into cavities on target pro- 154 teins, which causes nanobodies to bind to unique epitopes [32,33]. In 155 contrast, the binding interfaces of Fabs' and other mAbs' derived frag- 156 ments are more flat and less flexible, limiting the interactions of mAbs 157 and antibody fragments solely to the surface of antigens [34]. Recently, 158 we have determined the crystal structure of an anti-EGFR nanobody 159 (7D12) in complex with the EGFR ectodomain [33]. This nanobody 160 binds directly to domain III thereby sterically blocking EGF-binding. In- 161 terestingly, the 7D12 paratope that is binding to EGFR consists of CDR1 162 and 3, and the CDR2 makes no contact with EGFR (Fig. 2). Moreover, 163 nanobodies have a high refolding capacity even after being exposed to 164

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