HER2/NEU GENE EXPRESSION PRODUCTS EVALUATED WITH SUPERPARAMAGNETIC, GENETICALLY ENGINEERED ANTIBODIES

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ABSTRACT

HER2/neu is an oncogene amplified and overexpressed in ovary and breast cancer cells. In heterodimers, HER2/neu receptors stimulate signal transduction pathways leading to increased cell proliferation. The level of its expression is associated with cancer malignancy. Therefore, HER2/neu is an important indicator of cancer malignancy, as well as, a target for antibody guided cancer therapies. The main objective of this work was to develop molecular probes, which would report levels of HER2/neu gene expression and anatomical distribution of its products in vivo. Magnetic resonance imaging (MRI) is particularly suitable for this endeavor, as it offers not only the best spatial resolution from all in vivo imaging modalities currently available, but also the topographic reference for location of these probes within anatomy of the human body. Superparamagnetic single chain variable fragment (scFv) antibodies targeting HER2/neu were genetically engineered. They warranted high labeling specificity and affinity revealed with EDXSI, as well as, induced significant changes in relaxivity detected with NMR. This study demonstrated a proof of concept for using superparamagnetic scFv in diagnostic evaluation of levels of gene expression products with NMR and MRI for planning receptor targeted therapy.

Keywords

HER2/neu, ovarian cancer, breast cancer, signal transduction, genetically engineered antibodies

INTRODUCTION

Her2/neu is an oncogene amplified and overexpressed in ovarian and breast cancer cells (Di Fiore et al 1988, Berger et al 1988, Guerin et al 1988, van de Vijver et al 1988, Slamon et al 1989, Nielsen et al 2007). The level of its expression is associated with cancer malignancy (Berchuck et al 1990, King et al 1992, Zagouri et al 2007, Robert & Favret 2007). The ovarian or breast cancer cells may have approximately 3x10^6 expressed from multiple copies of the gene, while healthy cells in these organs may have approximately 2x10^4 HER2/neu receptors
on their surfaces. This leads to great increase in stimulations of signal transduction pathways, thus accelerated cell cycles and increased cell proliferation (King et al 1988, Lahusen et al 2007). Her2/neu positive cancers are the most invasive and have the worst prognosis. Therefore, levels of gene expression products and their distribution determined with monoclonal antibodies are of great diagnostic and prognostic value (Harris et al 1989). Furthermore, they are currently the primary target for antibody-guided, receptor-targeted therapies (Hudziak et al 1989, Jorgensen et al 2007, Park et al 2007, Allen et al 2007).

Ex vivo surgical biopsies are the primary material used currently for diagnostic analysis in immuno-histopathology laboratories (Shin et al 2007, Tischkowitz et al 2007, Tuma 2007, Carney et al 2007). Many techniques dealing with the evaluation of gene expression and its products include real time qualitative PCR, DNA microarray, differential display, blotting, serial analysis of gene expression (SAGE), etc. Each technique relies upon testing ex vivo of a small tissue or cell sample from a particular anatomical location at the time of biopsy only. However, the cancer gene expression profiles change rapidly, so are the levels and distributions of gene expression products (Fink-Retter et al 2007, Moon et al 2007). Diagnosis and prognosis would be far more accurate, if they would be based upon the images of the entire cancer and projections of its kinetics upon the whole patient’s pathophysiology.

In vivo molecular imaging of these antibodies would greatly facilitate such a diagnosis as well as reduce the patient’s trauma. This could be done with antibody guided contrast in vivo in magnetic resonance imaging (MRI). MRI offers not only the best spatial resolution from all in vivo imaging modalities currently available, but also the topographic reference for location of these probes within anatomy of the human body. The antibody guided probes, could provide the information concerned not only with antigenicity per se, but also report quantitative differences in levels of expression, as well presence of mutations, within architecture of the whole patient’s body at once.

The main objective of this work was to develop antibody guided molecular probes suitable for studying functions and locations of the HER2/neu gene expression products in vivo with MRI.

For developing of new probes for in vivo MRI, it is worth to consider that registered contrast differences between various tissue compartments are generated by local differences in relaxivities of water protons between those compartments. These are translated into varying brightness of the image details on the MRI scanner’s screen. Therefore, it is not as much the strength of the resonance signal itself, but rather the relative differences in signal intensity between various structures and/or in the signal to noise ratios that are the most essential properties in successful visualization of the analyzed features. Gadolinium (Gd) or Europium (Eu) atoms affect water proton relaxivity in their very immediate vicinity. $10^{-5}$ M of Gd is considered to be the threshold for inducing such a change in relaxivity of water, that it will be detected in NMR. If chelated into antibodies, these atoms indirectly report the presence of molecules that were targeted by antibodies. Attempts to realize that idea were conveyed by randomly attaching reporters: Gd chelates, dendrimers, or Fe nanoparticles to monoclonal IgG antibodies, thereby introducing paramagnetic properties (Curtet et al 1985, Mendonca et al 1986,
Linger et al 1986, Weissleder 1991, Unger et al 1999, Kobayashi et al 2003). Two main factors contributed why these attempts did not succeed. Random incorporation of reporters into IgG molecules leads to compromised specificity of antibodies up to their denaturation, thus low specific binding signal and high background due to non-specific binding. Significant increase in size of antibodies due to incorporation of reporters and change of their properties led to steric hindrance and repulsion forces. I have promoted an entirely different approach to improving labeling effectiveness by genetic engineering heterospecific, poly-functional molecules. They were engineered to contain multiple highly specific, separate domains assigned to their functions: scFv guided targeting domain, metal atoms chelating domains, signaling sequences, etc. Upon incorporation of Gd or Eu these molecules were gaining superparamagnetic properties without affecting their targeting functions.

The main hypothesis was, that proportional increase in the number of HER2/neu receptors per cell would result in the proportional increase in Gd atoms anchored via scFv to this cell, and that would result in a proportional increase in relaxivity of the surrounding water leading to the proportional increase in the signal strength recorded with NMR of ex vivo samples or MRI in vivo.

METHODS AND MATERIALS

Cell cultures

The cell lines TOV-112D CRL-11731 and CRL-11732 OV-90 were derived from primary malignant adenocarcinomas of the ovary at grade 3, stage IIIC. They were cultured in a 1:1 mixture of MCDB 105 medium and Medium 199, 85%; donor bovine serum 15% (ATCC). The cells were tumorigenic in nude mice. Cultured in soft agar they formed colonies and spheroids. The cells tested positive for HER2/neu and p53 mutation. The cell line NIH OVCAR-3 HTB-161 was derived from the cells in ascites of a patient with malignant adenocarcinoma of the ovary. The cell line is was grown in RPMI-1640 Medium (ATCC) supplemented with 0.01 mg/ml bovine insulin and donor bovine serum to a final concentration of 20%. The epithelial cells are positive for estrogen and progesterone receptor. They form tumors in nude mice. They form colonies and spheroids grown in soft agars. The cell line CRL-2340 HCC2157 was derived from the ductal carcinoma of the mammary gland tumor classified as TNM stage IIIA, grade 2, with lymph node metastasis. The cells are grown in a 1:1 mixture of Ham’s F12 medium with 2.5 mM L-glutamine and Dulbecco’s Modified Eagle’s Medium adjusted to contain 1.2 g/L sodium bicarbonate with additional supplements (ATCC).

The cell line MCF7 HTB-22. The cells are positive for estrogen receptor and express WNT7B oncogene. The medium to culture this cell line is Eagle’s Minimum Essential Medium (ATCC) with added following components: 0.01 mg/ml bovine insulin; donor bovine serum to a final concentration of 10%. The cell line 184A1 CRL-8798 was originally established from normal mammary tissue and transformed with benzopyrene. The line appears to be immortal,
but is not malignant. The line grows in Mammary Epithelial Growth Medium (MEGM) (Clonetics) supplemented with 0.005 mg/ml transferrin and 1 ng/ml cholera toxin. The normal, adherent fibroblast cell line Detroit 573 CCL-117 was derived from skin. It is grown in Minimum essential medium (Eagle) in Earle’s BSS with non-essential amino acids (ATCC), sodium pyruvate (1 mM) and lactalbumin hydrolysate (0.1%), 90%; fetal bovine serum, 10%. The cells were grown into spheroids within synthetic extracellular matrix.

Superparamagnetic single chain variable fragment (scFv) antibodies

Plasmid constructs were described in the details (Malecki et al. 2002). Coding sequences for variable fragment antibodies (scFvs) targeting HER2/neu selected from the surface displayed libraries were cloned in pM vectors designed with CMV immediate early promoter, SV40 poly(A) termination, hexahistidine, pentaglutamate, and selection neomycin-resistance coding sequences. Constructs for these bi-functional antibodies were electroporated into human myelomas (Malecki 1995; Malecki et al. 2002). Expressions of these constructs resulted in secretion of hetero-specific, poly-functional, mono-valent antibodies. Chelating sites were saturated with metal ions: Gd, Eu. Purification from non-bound metal was performed on affinity columns. The antibodies were produced in modified roller bottles.

Freezing and freeze-substitution of cell spheroids

The details of the cryoimmobilization by freezing were described previously and are only briefly presented here. The cells injected into the chambers were rapidly frozen in nitrogen slurry down to down to -196°C. The frozen samples were into methanol precooled to -90°C in the freezer (ThermoNoran). Temperatures were maintained at -90°C, -35°C, and 0°C for 48 hour. Infiltration with Lowicryl preceded polymerization with UV at-35°C and ultramicrotomy. Alternatively, critical point drying was followed by fast atom beam sputter coating.

Immunolabeling

Cell spheroids grown in culture were spun down at 300xg. The cells were resuspended in donor serum to which superparamagnetic antibodies were added. Upon completion of labeling the cells were rinsed with PBS. They were studied with NMR or processed by freezing in preparation for LSCM or EDXSI. Alternatively, cell lysates electrotransferred onto PVDF membranes were immunolabeled with antibodies with or without chelated Gd or Eu atoms.

Determination of metal atoms incorporated into chelating sites

The number of atoms chelated into the metal binding domains of the scFv was determined by using a titration method based on a competition between GdCl3 and radioactive, carrier-free GdCl3. Several aliquots of antibody (25 ul)
were incubated for 30 min with 100 ul on GdCl3 at various concentrations. The same aliquots were incubated for 30 min with 100 uCi of 153 GdCl3. Free Gd ions were complexed with 10 ul of 0.1 M DTPA. An aliquot of each solution was chromatographed on a silica ITLC (Gelman) support, using 0.1 M sodium citrate, pH5, as eluent. Alternatively, the chelated sites were saturated with Gd. Subsequently, these samples were purified on the gels as outlined above. Finally, they were analyzed with electron energy loss spectral imaging to determine total C to Gd ratio or in other words, the number of Gd atoms per scFv molecule. Alternatively, the scFvs were altered through carboxyl terminal derivatization with 125 I and their chelated sites saturated with 153 Gd. Subsequently, these samples were purified on the gels as outlined above. They were analyzed on a multi-channel analyzer which can display live full-energy spectrum 125 I at energy of 35 keV and 153 Gd at energy of 99 keV and was able to distinguish these two isotopes (Packard Cobra Gamma Counter).

Native electrophoresis

2% agarose gel was poured using a 10 mM Tris, 31 mM NaCl buffer of varying pH, that did not contain any denaturing agents. The samples in their native state were loaded after mixing with glycerol to add density without denaturating the proteins. The gel was run in the same buffer used for pouring the agarose at 60 mAmps until the desired separation was reached. The gel was then stained for 30 minutes in Sypro Tangerine Gel Stain (Invitrogen) diluted in the running buffer before imaging using a FluorImager (Molecular Dynamics).

SDS-PAGE

Electrophoresis was run on 12% polyacrylamide gel. 0.75 thick combs with the 2mm lanes were loaded with standard, cell culture lysates. The samples, after mixing with SDS and DTT containing sample buffers (Sigma) were loaded into the wells. The gels were run using a Tris/Glycine/SDS/DTT running buffers. After the run, the gels were stained with colloidal silver or Sypro Tangerine for imaging using a FluorImager (Molecular Dynamics).

Electrotransfer

After electrophoresis, the samples were immediately onto PVDF. The immunoblot was performed with the Mini Trans-Blot Cell (Bio-Rad) within CAPS: 10 mM 3-[Cyclohexylamino]-1-propanesulfonic acid (CAPS), Tris/glycine transfer buffer 25 mM Tris base, 192 mM glycine, pH 8.3. Prior the transfer the cooling units were stored with deionized water at -20 C. Immediately after electrophoresis the gel, membrane, filter papers and fiber pads were soaked in transfer buffer for 5-10 min. The pre-cooled transfer units were filled with cooled transfer buffer and electrotransfer proceeded at 350 mA.
Laser scanning confocal microscopy

The three-dimensional stacks of the cells labeled with scFv against HER2/neu were imaged with the laser scanning confocal system - Odyssey on the inverted microscope – Olympus. Excitation wavelengths were used: 337, 488, 543, and 588nm. Images were acquired with Kernel filtration and deconvolution of the data was followed by 3D or album display for analysis.

Nuclear magnetic resonance

The wide-bore nuclear magnetic resonance spectrometer operated at 9T (Brucker) with a mouse-cage resonator was used to evaluate relative relaxivity of the samples based upon T1 measurements. T1 spin lattice relaxation time calculated using inversion recovery pulse sequence were measured using inversion recovery imaging with Tl= 50-4000 ms in 100 ms increments. T1 was also calculated from T1-weighted fluid-attenuated inversion recovery (T1-FLAIR) sequence (Tr/Te/Flip = 2210/9.6/90), as well as standard T1-weighted imaging sequences (Tr/Te/Flip = 400/6/90).

Energy dispersive x-ray spectral imaging

Supramolecular of the scFv against HER2/neu was performed with Scanning Electron Microscope with Energy Dispersive X-Ray Spectral Imaging System (EDXSI) - Hitachi 3400. Complete elemental spectra were acquired for every pixel of the scans to create the elemental databases. From them, after selecting an element specific energy window, the map of this element atoms' distribution was calculated with ZAF correction (NIST). As the antibodies were tagged with atoms of Gd, Eu - exogenous elements incorporated into their structure, so was the location of antibodies determined based upon the elemental maps (Malecki 1995, Malecki et al 2002).

RESULTS

The major problem with designing new contrast agents for molecular imaging was lack of methods providing information concerned with their cell surface distribution and subcellular trafficking at the supramolecular level. This situation changed since the introduction of the very sensitive methods of their detection in situ with EELSI and EDXSI (Malecki 1995, Malecki et al 200). There, genetically engineered antibodies tagged with atoms of selected exogenous elements were localized within three-dimensional architecture of cells and cell organelles to determine molecular mechanisms governing their bio-distribution and bio-compatibility. In this study, TOV-112D CRL-11731, OV-90 CRL-11732, CRL-2340 HCC2157, NIH OVCAR-3 HTB-161, MCF7 HTB-22, 184A1 CRL-8798, Detroit 573 CCL-117 cell spheroids were cultured and labeled with anti- HER2/neu superparamagnetic scFv antibodies. In cultured cells were labeled with antibodies chelating Gd or Eu atoms. They were rapidly frozen. Frozen
Cells were freeze-substituted with no metal incorporation, infiltrated, and embedded. Distribution of antibodies, harboring metal atoms, in ultrathin sections or cell whole mounts were examined with elemental mapping systems. The antibodies chelating Gd atoms were anchored to the cell surface receptors. Therefore, they were visualized by mapping Gd (Figure 1). That could be only possible due to acquisition of the full spectrum for every pixel of the scan to create the elemental data base. Thereafter, an energy window selected for Gd allowed for extracting element distribution within the entire image to create element distribution map. This elemental map based antibody distribution was projected onto the cell surface ultrastructure to determine localization of superparamagnetic antibodies at the molecular level.

High specificity of superparamagnetic antibodies was also confirmed on Western blots from cell lysates. Exquisitely specific bands were present with antibodies chelating Gd or Eu atoms. Intentionally the space below and above the bands are not cut to show absence of any non-specific binding, but only specific bands are present. Chelation did not change the specificity of scFv antibodies.
site single bands were clear indications of high specificity of the engineered antibodies (Figure 2). All the combinations resulted in the same labeling patterns. Importantly, the blots demonstrated that no other proteins in the entire cell lysate were labeled with our Gd chelated GE antibodies. The antibodies retained specificity towards targeted receptors, even after Gd coordination. Moreover, the background was entirely label free.

The ultimate test for attaining the project objective was the effect, which superparamagnetic antibodies anchored to the receptors on cell surfaces might have on local relaxivity.

Table 1 shows data from a representative experiment. Refined measurements were conducted on wide-bore Bruker (Table 1). Importantly, we observed significant increase in water relaxivity $r$. That resulted in change in relaxivity proportional to the number of Gd chelated scFv antibodies attached. Relaxivity of water protons was about 200 mM $^{-1}$ s$^{-1}$ at 9.4T. The high relaxivity have to result in MRI contrast changes at antibody concentrations as little as 0.1 uM, which is sufficient for imaging of receptors in vivo. It was demonstrated that the antibodies with Gd are capable of labeling cells in vitro. In our studies in cell culture, we have observed a significant contrast-to-noise ratio (CNR) enhancement due to superparamagnetic antibodies. Therefore, these scFv-based receptor targeting contrast agents created a clinically relevant change in relaxivity detectable in NMR (Table 1).

To summarize, in this initial study, striking differences were noticed in the signal strength generated between unlabeled ECM, fibroblasts, ovarian and breast cancer cells after labeling with the scFv antibody chelating Gd – antibody guided contrast.

**DISCUSSION**

This work provides the proof of concept for using superparamagnetic antibodies in detecting differences in levels of gene expression products in cells in vitro and in vivo. In practice, labeling of cell receptors with scFv antibodies resulted

| Table 1. Differences in T1 relaxation times, between unlabeled physiological fluids and tissues versus GE paramagnetic antibodies labeled cells. |
|---|---|
| Water | 3.210 +/- 0.031 s |
| Serum | 2.273 +/- 0.024 s |
| Detroit fibroblasts culture | 1.598 +/- 0.015 s |
| Ovarian cancer TOV-112D CRL-11731 | 1.303 +/- 0.011 s |
| Ovarian cancer TOV-112D CRL-11731 + anti HER2/neu scFv$_{Gd}$ | 393.626 +/- 0.028 ms |
| Breast cancer CRL-2340 HCC2157 | 1.219 +/- 0.013 s |
| Breast cancer CRL-2340 HCC2157 + anti HER2/neu scFv$_{Gd}$ | 428.327 +/- 0.039 ms |

Measurements of T1 relaxation times change induced by GE paramagnetic antibodies in [s] by inversion recovery with 400MHz at 9.4T on 28mm wide-bore Bruker.
in a dramatic shortening of T1. It was proportional to the number of Gd or Eu atoms harbored by scFv antibodies and anchored to the cell surface receptors. The significant difference between the number of the receptors on surfaces of cancer and normal cells translated into the significant difference in the signal intensity between these cells. This work opens new avenues for in vivo studies involving antibody guided contrast.

Success of this work can be primarily attributed to the high specificity, affinity, and small size of the engineered scFv. Their high specificity resulted not only in heavy labeling of the HER2 neu receptors, but also in reduced non-specific labeling of other cells. Therefore, the signal to noise ratio was remarkably high. The high affinity of these antibodies was shifting the dynamic on/off balance; thus enhancing conditions for T1 acquisition. Finally, the small size of these antibodies helped in their penetration into the depth of the cell spheroid cultures, as well as, in their packing onto the receptors. That increase in packing or labeling density was also seen on the images from Phosphorimager, LSCM, and EDXSI. The labeling density was much higher with scFv, than it was with Fab or IgG. In this study, it translated into the significant concentration of Gd or Eu atoms on surfaces of the cells. Higher number of antibodies, each harboring Gd or Eu atoms, resulted in significant changes of the relaxivity reflected in shortening of T1 and strengthening of the generated signal. This will be perceived as the bright spots on the screen of MRI scanner.

Specific signal to background noise ratio is the main factor to discriminate, the structure labeled with the element tagged recombinant antibody guided contrast agent from the unlabeled structures surrounding it. Therefore, the primary objective of this effort was to bioengineer antibodies in such a way that they would generate label-free background i.e., no non-specific labeling. As described earlier and applied here, it has been accomplished by selecting clones using short receptor domain sequence libraries, purification prior to and after derivatization, evaluation of antibody affinity on native electrophoresis and blue blots, and validation of the data with EDXSI. This complex approach resulted in very specific localization of the superparamagnetic antibodies on targeted HER2/neu receptors.

Improved packing of Gd atoms into chelating domains may enhance local relaxivity of water. Here, it has been accomplished by engineering metal binding domains into the scaffold of GE antibodies. Contrary to all of the other methods of antibody derivatization based upon random incorporation of chelating agents, which are changing properties of these antibodies, in this work the highly specific domains are specific integral parts of superparamagnetic antibodies, but completely separate from antigen binding domains. Therefore, they retain their bio-kinetic properties and antibody binding properties after incorporation of Gd or Eu into their scaffolds. Further, affinity purification, which follows derivatization, secures elimination of all molecules, which might have altered properties.

To summarize, we demonstrated a proof of concept for using antibody guided contrast agents for evaluating gene expression products. They warrant pursuing studies involving superparamagnetic antibodies in vitro and in vivo. The conclusions outlined above should serve as guides for their streamlining into in vivo molecular imaging endeavors.
Above all, the main challenge, before engineering superparamagnetic antibodies for in vivo MRI, is to secure thermodynamic stability of Gd or Eu within chelating pockets. Harboring these metals within antibodies has to be extremely stable, so that release and/or transmetallation do not induce toxic effects.

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LITERATURE CITED


