

## Development of a Solid-Phase Assay for Analysis of Matrix Metalloproteinase Activity

Janelle L. Lauer-Fields,<sup>a</sup> Hideaki Nagase,<sup>b</sup> and Gregg B. Fields<sup>a</sup>

<sup>a</sup>Department of Chemistry and Biochemistry, Florida Atlantic University, Boca Raton, FL;

<sup>b</sup>Kennedy Institute of Rheumatology Division, Imperial College London, Hammersmith, London

Proteases play fundamentally important roles in normal physiology and disease pathology. Methods for detection of active proteolysis may greatly aid in the diagnosis of disease progression, and suggest modes of therapeutic intervention. Most assays for proteolytic potential are limited by a lack of specificity and/or quantification. We have developed a solid-phase activity assay for members of the matrix metalloproteinase (MMP) family that is specific and can be used to quantify active enzyme concentration. The assay has two principal components: a capture antibody that immobilizes the MMP without perturbing the enzyme active site, and a fluorescence resonance energy transfer substrate for monitoring proteolysis at low enzyme concentrations. The assay was standardized for MMP-1, MMP-3, MMP-13, and MMP-14. The efficiency of the assay was found to be critically dependent upon the quality of the antibodies, the use of substrates exhibiting high specific activities for the enzymes, and enzyme samples that are fresh. The assay was applied to studies of constitutive and induced MMP activity in human melanoma cells. Analysis of several melanoma cell lines, and comparison with prior studies, correlated higher constitutive MMP-13

ADDRESS CORRESPONDENCE AND REPRINT REQUESTS TO: Gregg B. Fields, Department of Chemistry and Biochemistry, Florida Atlantic University, 777 Glades Road, Boca Raton, FL 33431–0991 (phone: 561–297–2093; fax: 561–297–2759; email: fieldsg@fau.edu).

activity with higher levels of the cell surface receptor CD44. Ligands to two different melanoma cell surface receptors (the  $\alpha_2\beta_1$  integrin or CD44) were found to induce different proteolytic profiles, suggesting that the extracellular matrix can modulate melanoma invasion. Overall, the solid-phase MMP activity assay was found to be valuable for analysis of protease activity in cellular environments. The solid-phase assay is suitably flexible to allow studies of virtually any proteolytic enzyme for which appropriate substrates and antibodies are available.

**KEY WORDS:** matrix metalloproteinase, fluorogenic substrate, melanoma, solid-phase assay, ELISA

The metzincin subclan of enzymes—which includes the matrix metalloproteinase (MMP), ADAM (a disintegrin and metalloproteinase), and ADAMTS (a disintegrin and metalloproteinase with thrombospondin motifs) families—has been associated with many normal physiological functions.<sup>1</sup> Pathologies such as arthritis and tumor metastasis also feature prominent roles for the metzincins. Most MMP, ADAM, and ADAMTS members of the metzincin subclan are expressed in inactive form as zymogens. Once activated, their ability to process substrate is regulated by protease inhibitors that may be general ( $\alpha_2$ -macroglobulin) or specific (tissue inhibitor of metalloproteinase; TIMP). The vast majority of studies on MMP, ADAM, and ADAMTS regulation have focused on modulation of protein production. However, given the numerous pathways by which these enzymes are regulated, it is important to extend such studies to include the measurement of active enzyme. Ultimately, the ability to quantify enzymatic activity is necessary to discern which MMP, ADAM, and ADAMTS family members are associated with a disease process.

Numerous approaches have emerged by which activity of these proteases can be evaluated. Use of substrate-based assays often involves radiolabeling, dyes, or fluorescent tags.<sup>2–5</sup> Alternatively, polyacrylamide gel electrophoresis can be used in combination with densitometry or substrate zymography to analyze activity,<sup>6–10</sup> while enzyme-linked immunosorbent

assay (ELISA) and Western blotting can be utilized to differentiate between active enzyme and zymogen forms.<sup>8,11</sup> Substrate hydrolysis can also be monitored by using antibodies to neoepitopes exposed only after cleavage<sup>12–15</sup> or release of hydrolysis products from co-polymerized polyacrylamide beads.<sup>15,16</sup> A continuous assay method, such as one that utilizes an increase in fluorescence upon hydrolysis, allows for rapid and convenient kinetic evaluation of proteolytic activity. Quenched fluorescent substrates that rely on fluorescence resonance energy transfer (FRET)/intramolecular fluorescence energy transfer (IFET) have been constructed by incorporation of a fluorophore and a quencher on the same peptide chain. Fluorogenic substrates have several advantages over other methods, in that they (a) can be monitored continuously and at reasonably low concentration ranges, (b) can be used to determine individual kinetic parameters, and (c) are readily accommodated by high-throughput analytical techniques.

Several different fluorogenic substrates have been described for MMPs and ADAMs, and the relative merits of these substrates have been discussed previously.<sup>17–20</sup> In several cases, fluorogenic substrates that are selective for specific MMPs have been described.<sup>21–23</sup> However, it is unrealistic to expect that singly selective substrates can be designed for each member of the MMP family. Selectivity can be obtained with antibodies, and thus the present studies have examined the use of antibodies in combination with fluorogenic substrates to create assays selective for MMP activity.

## MATERIALS AND METHODS

### Peptides

The synthesis and characterization of the general fluorogenic triple-helical peptide (THP) substrates fTHP-4, fTHP-5, and fTHP-7, and the MMP-3-specific fluorogenic substrate NFF-3 have been described previously.<sup>20,21,24,25</sup> The synthesis and characterization of the triple-helical ligands for melanoma  $\alpha_2\beta_1$  integrin [ $C_{10}$ -( $\alpha 1$ (IV)382–393)-NH<sub>2</sub> THP,  $C_{16}$ -( $\alpha 1$ (IV)382–393)-NH<sub>2</sub> THP] and CD44 [ $C_{16}$ -( $\alpha 1$ (IV)1263–1277)-NH<sub>2</sub> THP] have also been described.<sup>22,26–28</sup>

### Matrix Metalloproteinases

ProMMP-1 and proMMP-3 were expressed in *Escherichia coli* and folded from the inclusion bodies as described previously.<sup>29</sup> ProMMP-1 was activated by reacting with 1 mM 4-aminophenylmercuric acetate

(APMA) and an equimolar amount of MMP-3 at 37°C for 6 h. After activation, MMP-3 was completely removed from MMP-1 by affinity chromatography using an anti-MMP-3 IgG Affi-Gel 10 column. ProMMP-3 was activated by reacting with 5  $\mu$ g/mL chymotrypsin at 37°C for 2 h. Chymotrypsin was inactivated with 2 mM diisopropylfluorophosphate. ProMMP-13 was a generous gift from Dr. P.G. Mitchell, Pfizer, Inc. (Groton, CT). ProMMP-13 was activated with 1 mM APMA. The amounts of active MMP-1, MMP-2, MMP-3, and MMP-13 were determined by titration with recombinant TIMP-1<sup>30</sup> over a concentration range of 0.1 to 3  $\mu$ g/mL. Recombinant MMP-14 with the linker and C-terminal hemopexin-like domains deleted [residues 279–523; designated MMP-14( $\Delta_{279-523}$ )] was purchased from Chemicon International (Temecula, CA). MMP-14( $\Delta_{279-523}$ ) was expressed in the active form with Tyr112 at the N-terminus. MMP-14( $\Delta_{279-523}$ )—which, in contrast to MMP-14, does not undergo rapid autoproteolysis—was used in the present studies due to the relatively small differences in MMP-14( $\Delta_{279-523}$ ) and MMP-14 triple-helical peptidase activities noted previously.<sup>31</sup> The amount of active MMP-14 was determined by comparison of activity measurements between several different MMP-14 preparations that had been directly titrated with TIMP-2.<sup>23,31</sup>

### Cell Culture

The human melanoma cell lines M14P, M14#5, and M14#11 were obtained as a generous gift from Dr. Barbara M. Mueller (La Jolla Institute for Molecular Medicine, La Jolla, CA). Tissue culture reagents were obtained from Fisher Scientific (Atlanta, GA) unless otherwise stated. All immunologicals were supplied by Chemicon International (Temecula, CA). M14 cells were routinely cultured with RPMI 1640 and supplemented with 10% fetal bovine serum (BioWhittaker, Walkersville, MD) and the following antibiotics: 0.1 mg/mL gentamicin sulfate, 50 units/mL penicillin, and 0.05 mg/mL streptomycin sulfate. Cells were cultured for up to 8 passages and then replaced by frozen stocks to minimize phenotypic drift. Human aortic endothelial cells (Clonetics/Biowhittaker/Cambrex, Walkersville, MD) were cultured according to the supplier's recommendations (cultured in EGM-2 and discarded at passage 9). All cells were maintained at 37°C in a humidified incubator containing 5% CO<sub>2</sub>.

### Induction of Melanoma Cells

Pro-Bind 96-well assay plates were conditioned at room temperature overnight prior to initiation of the

induction experiment with 10  $\mu\text{M}$   $\text{C}_{10}$ - $(\alpha 1(\text{IV})382\text{--}393)\text{-NH}_2$  THP,  $\text{C}_{16}$ - $(\alpha 1(\text{IV})382\text{--}393)\text{-NH}_2$  THP, or  $\text{C}_{16}$ - $(\alpha 1(\text{IV})1263\text{--}1277)\text{-NH}_2$  THP. Previous studies had determined that these were optimal ligand concentrations for cell adhesion.<sup>22,27,28,32,33</sup> The plates were then blocked by adding 2 mg/mL bovine serum albumin (BSA) in phosphate-buffered saline (PBS), pH 7.4, and incubated overnight at room temperature. M14P cultures used for induction experiments were typically 60–80% confluent before release from growth flasks with PBS containing 5 mM ethylenediaminetetraacetic acid (EDTA) (pH 7.3). Subsequent to release, cells were washed with adhesion media [RPMI-1640 containing 20 mM N-(2-hydroxyethyl)piperazine-N'-(2-ethanesulfonic acid) (HEPES)] and seeded at approximately 7500 cells/well. Melanoma cells were allowed to adhere to ligands for 60 min at 37°C. Three washes of adhesion media were used to remove nonadherent cells. Aliquots of the melanoma-conditioned media were harvested at regular intervals ( $T = 0, 6, 12, 18,$  and 24 h) for later determination of metalloproteinase levels. Samples were stored at  $-20^\circ\text{C}$  until analyses could be performed. Conditioned media was isolated by withdrawal of the media from growing cells and centrifuging at  $1000 \times g$  (to remove any floating cells).

### Matrix Metalloproteinase Assay

MMP capture antibodies were from Chemicon and were as follows: MMP-1, monoclonal antibody (mAb) MAB1346; MMP-3, polyclonal antibody (pAb) AB810; MMP-13, mAb MAB3321; and MMP-14, pAb AB815. The general assay protocol was as follows: A 96-well plate was incubated with the appropriate MMP Ab for at least 18 h at 4°C. Nonspecific binding sites were blocked by incubating with PBS containing 0.05% Tween 20 and 2 mg/mL BSA for at least 4 h at 4°C. Either MMP standards or unknown samples were added to each well and the plate gently shaken for at least 18 h at 4°C. Unbound enzyme was removed by washing wells 3 times with enzyme assay buffer (50 mM Tricine, 50 mM NaCl, 10 mM  $\text{CaCl}_2$ , 0.05% brij-35). The appropriate fluorogenic substrate was added to each well, and the plate incubated at 37°C in a humidified atmosphere for 0–18 h. Fluorescence readings were taken at appropriate intervals. More specific details for optimized assay conditions are given in the Results and Discussion section. For calculation purposes, the following MMP molecular weight values were used: MMP-1, 41.0 kDa; MMP-3, 43.0 kDa; MMP-13, 42.0 kDa; and MMP-14( $\Delta_{279\text{--}523}$ ), 20.5 kDa<sup>34,35</sup>. Assays were performed in at least duplicate.

## RESULTS AND DISCUSSION

### Standardization of Solid-Phase Assay

The solid-phase assay was standardized using four MMP family members: MMP-1, MMP-3, MMP-13, and MMP-14. MMP-1, MMP-13, and MMP-14 assays were developed using general triple-helical substrates fTHP-4, fTHP-5, or fTHP-7 (Fig. 1). The MMP-3 assay was developed using the MMP-3 specific substrate NFF-3 (Fig. 1). The general principle of the assay is given in Figure 2. For a traditional solid-phase indirect sandwich ELISA,<sup>36</sup> three antibodies are needed: (a) one to immobilize (trap/capture) the MMP, typically a pAb; (b) one to detect the MMP, typically an mAb; and (c) a secondary antibody to detect the mAb, which, in turn, is linked to an enzyme. Colorimetric detection is achieved by conversion of exogenous substrate. Alternatively, a fluorescently labeled secondary antibody can be used. ELISA is typically used to measure protein amounts, but does not discriminate for enzyme activity. For our solid-phase assay, only the immobilization antibody is needed. However, this antibody must bind the MMP selectively and efficiently without significantly inhibiting catalytic activity. MMPs can be divided into four domains: propeptide, catalytic, linker, and hemopexin-like (Fig. 3). The appropriate MMP capture antibodies were thus chosen based on prior application in ELISA and recognizing regions other than the MMP active site within the catalytic domain. After examining a variety of antibodies, the ones specifically selected were as follows: MMP-1 clone 3A9.3, a non-inhibitory mAb raised against human MMP-1; an MMP-3 pAb raised against the human MMP-3 linker region; MMP-13 clone 181–15A12, a non-inhibitory mAb raised against human MMP-13; and an MMP-14 pAb raised against the human MMP-14 linker region.

MMP hydrolysis of the fluorogenic substrates used herein had previously been shown to follow Michaelis-Menton kinetics.<sup>21,37</sup>  $K_M$  values for MMP-1 hydrolysis of fTHP-4, fTHP-5, and fTHP-7, MMP-3 hydrolysis of NFF-3, MMP-13 hydrolysis of fTHP-4, and MMP-14( $\Delta_{279\text{--}523}$ ) hydrolysis of fTHP-4 have previously been determined and fall in the range of 11 to 48  $\mu\text{M}$ .<sup>20,21,23–25</sup> Assays were performed using an excess of substrate compared with enzyme but with substrate concentrations typically below enzyme  $K_M$  values, approximating first-order kinetic conditions. Under these conditions, one can directly correlate enzyme activity to enzyme concentration by comparison with known concentrations of active enzyme.

Initial studies were performed with MMP-1 and fTHP-7. First, the appropriate capture mAb concentration was determined by comparing 0.125, 0.25, 0.5, 1.0, 2.0, and 5.0  $\mu\text{g/mL}$  of MMP-1 mAb over a range of

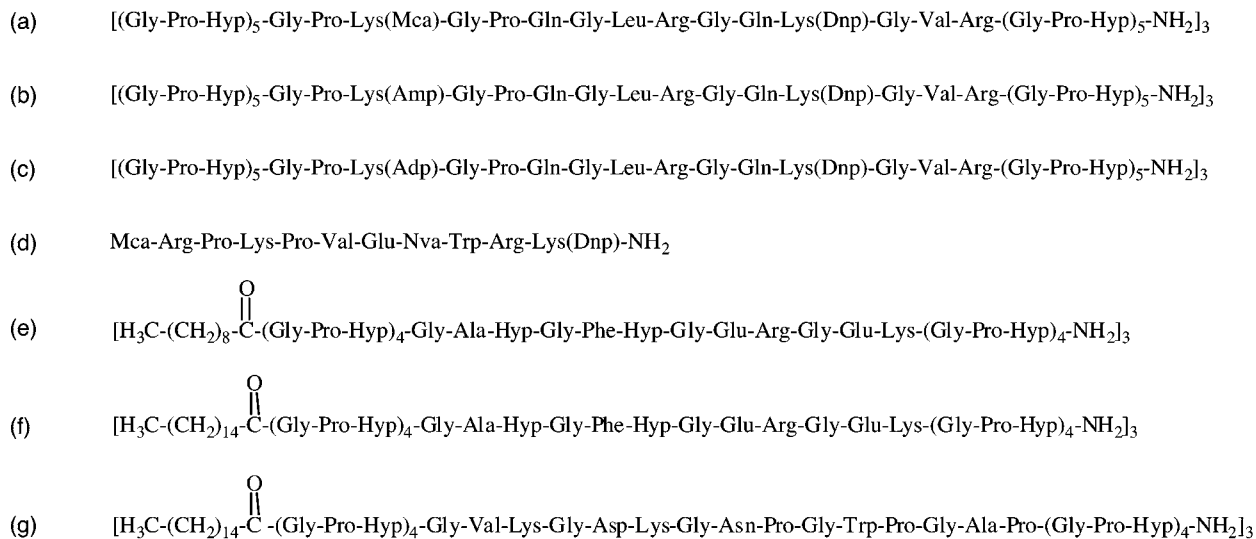


FIGURE 1

Sequences of substrates and ligands. The four substrates used in this study are (a) fTHP-4, (b) fTHP-5, (c) fTHP-7, and (d) NFF-3. The three peptide-amphiphile ligands used for induction of melanoma cell MMPs are (e) C<sub>10</sub>-(α I (IV)382–393)-NH<sub>2</sub> THP ligand for the α<sub>2</sub>β<sub>1</sub> integrin, (f) C<sub>16</sub>-(α I (IV)382–393)-NH<sub>2</sub> THP ligand for the α<sub>2</sub>β<sub>1</sub> integrin, and (g) C<sub>16</sub>-(α I (IV)1263–1277)-NH<sub>2</sub> THP ligand for CD44. Sequence numbering for the ligands corresponds to the human α I (IV) gene-derived sequence.<sup>57</sup>

MMP-1 concentrations. The capture mAb concentrations were chosen based on literature precedents for MMP ELISA.<sup>38,39</sup> The 2.0 μg/mL and 5.0 μg/mL mAb concentrations were effective, but the highest concentration of mAb captured the higher concentrations of MMP-1 slightly more efficiently (Fig. 4). If the anticipated enzyme production is below approximately 50 ng/mL, 2.5 μg/mL capture mAb is likely sufficient for analysis. The limit of detection was then evaluated using substrate fTHP-5, in which the fluorophore (Amp) has a higher quantum yield than the fluorophore of fTHP-7 (Adp).<sup>20</sup> The sensitivity of the assay allowed for detection of 1 × 10<sup>-15</sup> moles of active MMP-1 in a 200 μL volume (5.0 pM) (data not shown). The specificity of the assay was examined using an irrelevant mAb (one against the β<sub>1</sub> integrin subunit) or mouse IgG. MMP-1 activity was not observed over background when the anti-β<sub>1</sub> mAb was used as the capture antibody (Fig. 5). Subsequent use of a mouse IgG instead of the anti-β<sub>1</sub> mAb as a negative control produced a similar result (data not shown). The cross-reactivity of the MMP-1 capture mAb was examined by comparing its ability to recognize MMP-1 with its ability to recognize MMP-13. MMP-13 activity was not observed over background, demonstrating that the mAb was not cross-reactive and did specifically immobilize active MMP-1 (Fig. 6). Thus, the solid-phase assay was found to specifically quantify MMP-1 activity with good sensitivity over an MMP-1 range of 1.0 fmol to 1.0 pmol.

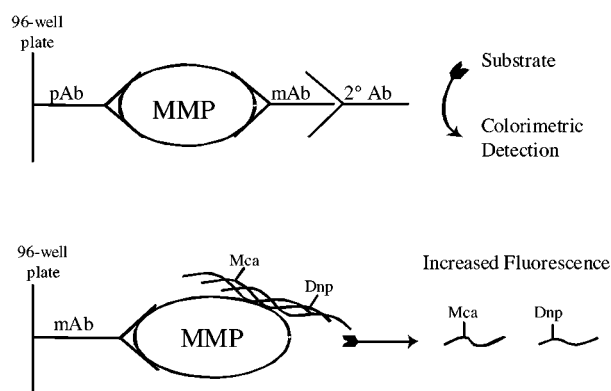


FIGURE 2

Schematic diagrams for (top) conventional indirect sandwich ELISA and (bottom) solid-phase MMP activity assay. For both assays, the MMP is captured by an antibody on the solid phase. The indirect sandwich ELISA then requires a detection antibody and an enzyme-labeled antibody. The sandwich ELISA can also be performed directly, in which case the detection antibody is also enzyme-labeled. In the solid-phase MMP activity assay, the fluorogenic substrate serves as the detection system. Dnp, 2,4-dinitrophenyl; Mca, (7-methoxycoumarin-4-yl)acetyl.

Based on the MMP-1 results, standard conditions were developed for the solid-phase MMP activity assay. The 96-well plate was incubated with the 100 μL of 5 μg/mL of the appropriate MMP capture Ab in PBS for

Collagenase (MMP-1, MMP-13)  
Stromelysin (MMP-3)

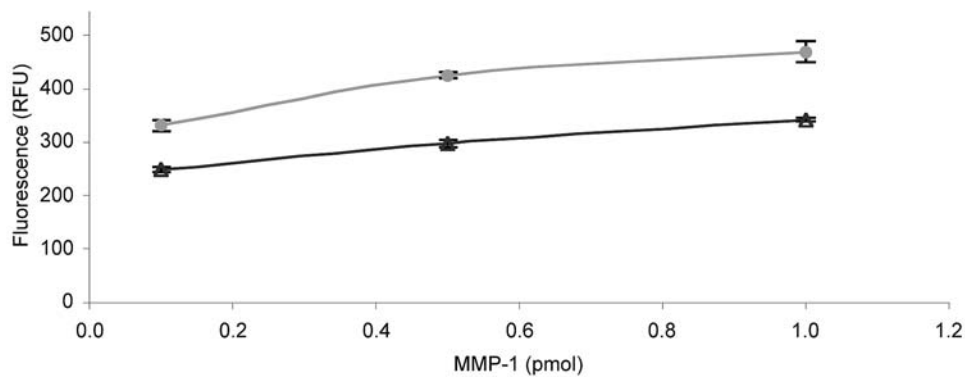


Membrane-bound (MMP-14)



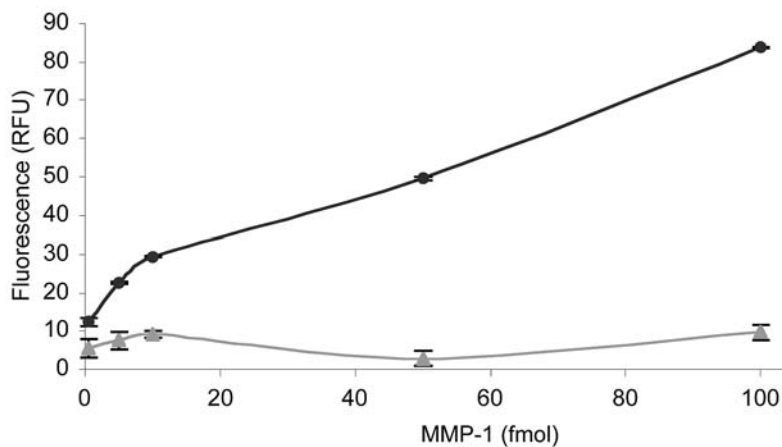
**FIGURE 3**

Domain structures of MMP-1, MMP-3, MMP-13, and MMP-14.



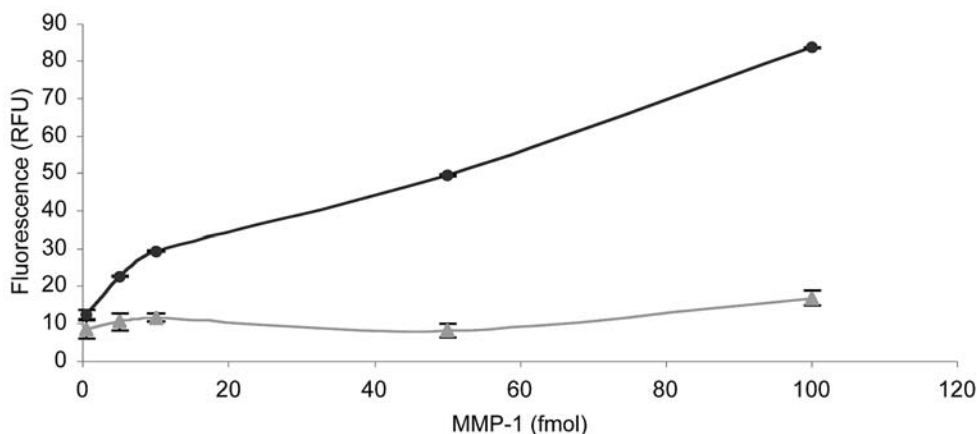
**FIGURE 4**

Efficiency of capture antibody for solid-phase MMP activity assay. The change in fluorescence upon MMP-1 hydrolysis of fTHP-7 was compared after 18 h using 2.5 µg/mL (triangles) or 5 µg/mL (circles) MMP-1 capture mAb. The amounts of MMP-1 added ranged from 0.1–1.0 pmol. Methods were as described in Materials and Methods.



**FIGURE 5**

Specificity of the solid-phase MMP activity assay based on the nature of the antibody. The change in fluorescence upon MMP-1 hydrolysis of fTHP-7 was compared after 18 h using 2.5 µg/mL MMP-1 mAb (circles) or 2.5 µg/mL of an anti-β1 integrin subunit mAb (triangles). The amounts of MMP-1 added ranged from 0.5 to 100 fmol. Methods were as described in Results and Discussion.

**FIGURE 6**

Specificity of the solid-phase MMP activity assay based on the nature of the MMP. The change in fluorescence upon MMP-1 (circles) or MMP-13 (triangles) hydrolysis of fTHP-7 was compared after 18 h using 2.5  $\mu\text{g/mL}$  MMP-1 mAb. The amounts of MMP-1 added ranged from 0.5 to 100 fmoles. Methods were as described in Results and Discussion.

at least 18 h at 4°C with mixing. Nonspecific binding sites were blocked by incubating with 100  $\mu\text{L}$  PBS containing 0.05% Tween 20 and 2 mg/mL BSA for at least 4 h at 4°C with mixing, and the plate washed 3 times with enzyme assay buffer. One-hundred-fifty microliters of either MMP standards or unknown samples were added to each well, followed by 50  $\mu\text{L}$  PBS containing 0.2% Tween 20 and 8 mg/mL BSA (to create final concentrations of 0.05% Tween 20 and 2 mg/mL BSA), and the plate was mixed for at least 18 h at 4°C. All liquid was removed and 200  $\mu\text{L}$  enzyme assay buffer or 2 mM APMA (where applicable) was added to each well. This was followed by a 2 h incubation at 37°C. The wells were washed 3 times with enzyme assay buffer and 200  $\mu\text{L}$  of the appropriate fluorogenic substrate (5–10  $\mu\text{M}$  of fTHP-4, fTHP-5, or fTHP-7; 2.5–5  $\mu\text{M}$  of NFF-3) was added to each well. The plate was incubated at 37°C in a humidified atmosphere for 0 to 18 h. Fluorescence readings ( $\lambda_{\text{excitation}} = 324 \text{ nm}$  and  $\lambda_{\text{emission}} = 393 \text{ nm}$  for NFF-3, fTHP-4, and fTHP-5;  $\lambda_{\text{excitation}} = 348 \text{ nm}$  and  $\lambda_{\text{emission}} = 436 \text{ nm}$  for fTHP-7) were taken at appropriate intervals and a standard curve was created by plotting the increase in fluorescence versus concentration of active enzyme. If enzyme activity is anticipated to be high, data collection is best optimized by monitoring activity over 1 to 4 h. The samples analyzed herein (see later discussion) have relatively low levels of active enzyme. Thus, fluorescence was monitored over an extended period of time (18–24 h).

The efficiency of enzyme capture by the MMP antibodies was evaluated by comparing the solid-phase assay with our previously described solution assay.<sup>20–22,24</sup> It is assumed that the change in fluores-

cence obtained from the solution assay represents 100% of the active enzyme, where active enzyme concentration is determined initially by TIMP-1 titration (see Materials and Methods). Change in fluorescence was thus compared directly between the solution and solid-phase assays. The capture efficiency was 63, 37, and 40% for MMP-1, MMP-3, and MMP-13, respectively, over an MMP concentration range of 10 to 20 ng/mL (data not shown). Below this range, capture efficiency approached quantitative values. The capture efficiency for MMP-14 could not be accurately evaluated due to limitations in commercially available MMP-14 reagents. The MMP-14 standard curve (see below) was generated in the presence of the MMP-14 capture pAb. The modulation of activity due to pAb binding was thus taken into consideration, while capture efficiency was not.

Standard curves were subsequently obtained for MMP-1, MMP-3, MMP-13, and MMP-14 ( $\Delta_{279-523}$ ) (Fig. 7). In all cases, reasonable changes in fluorescence were observed with increasing MMP concentrations. The curves were not linear, as capture efficiencies change at higher MMP concentrations (see above). Therefore, standard curves should be optimized based on the levels of active enzymes in a given set of samples. If the sample activity is unknown, a standard curve covering a wide range of known concentrations should be constructed and outlying concentrations can be adjusted upon completion of the experiment. The most sensitive assay observed herein was for MMP-3, where active enzyme could be readily detected at less than 1 ng/mL (less than 23 pM). The high  $k_{\text{cat}}/K_M$  value for MMP-3 hydrolysis of NFF-3<sup>21</sup> resulted in substantial changes in

fluorescence at even very low MMP-3 concentrations. The next most sensitive assay was the one for MMP-1, where active enzyme could be readily detected at 1–2 ng/mL (23–47 pM), followed by conservative limits of 3–5 ng/mL for MMP-13 (71–119 pM) and MMP-14( $\Delta_{279-523}$ ) (146–244 pM).

These results were somewhat surprising, considering that both MMP-13 and MMP-14( $\Delta_{279-523}$ ) hydrolyze fTHPs more efficiently than MMP-1.<sup>23,24,31</sup> This suggests that during an extended capture and assay process, MMP-13 and MMP-14 lose some of their activity. Loss of activity was probably due to limitations in antibody recognition, or circumstances where the capture antibody obscures the MMP active site and/or destabilizes enzyme tertiary structure. Alternatively, activity loss could result from MMP autoproteolysis and/or dissociation of structural metals.<sup>34,35,40,41</sup> Since MMP-14 capture efficiency could not be calculated, the apparently low MMP-14 sensitivity may reflect inefficient capture. With standard curves generated for MMP-1, MMP-3, MMP-13, and MMP-14, the solid-phase MMP assay was utilized to examine constitutive and inducible MMP activity in melanoma cells.

#### Constitutive Production of Soluble, Active MMP-3, MMP-13, and MMP-14 by Melanoma Cells

MMP-1, -2, -3, -8, and -13 are expressed in numerous human melanoma cell lines, but not melanocytes, with MMP-2 and MMP-8 expression at higher constitutive levels than other MMPs.<sup>42,43</sup> Conversely, MMP-14 is expressed in both melanocytes and melanoma.<sup>42</sup> MMP-14 activity has been correlated with the invasion of numerous tumor types, including melanoma.<sup>44–46</sup> MMP-14 activity may be membrane-bound or soluble, where nonautocatalytic shedding of MMP-14 is responsible for generation of soluble activity.<sup>35</sup> The solid-phase assay was utilized to examine constitutive MMP-1, MMP-3, MMP-13, and soluble MMP-14 (sMMP-14) activity in human melanoma cells as well as in human aortic endothelial cells, which serve as a normal control cell line. MMP activity was quantified for M14P, M14#5, and M14#11 melanoma and aortic endothelial cells by assaying conditioned media. The melanoma cells differ in their surface expression of certain receptors. M14#5 cells have higher levels of CD44 than M14P, while M14#11 cells have lower levels of CD44 than M14P.<sup>27</sup> In contrast, the levels of the  $\beta_1$  integrin subunit are similar for M14P, M14#5, and M14#11.<sup>27</sup> Finally, M14#5 expresses CD44 but does not express MPG/MCSP—another cell surface proteoglycan receptor.<sup>47</sup> It is of interest to learn if receptor distribution has a role in active MMP production.

All three melanoma cell lines produced equivalent, low amounts of active MMP-1 and MMP-3, and levels were overall slightly higher than for endothelial cells (Table 1). M14P melanoma cells had the lowest levels of active MMP-13 and sMMP-14, followed by M14#11 and then M14#5 (Table 1). The accuracy of enzyme quantification was confirmed by spiking melanoma cell conditioned media with known concentrations of MMPs (data not shown). It is interesting to note that the cell line with the highest levels of CD44 (M14#5) had the highest levels of MMP-13 and sMMP-14 activity, since CD44 and MMP-14 are known to associate at the tumor cell migration front<sup>46,48</sup> and MMP-14 can serve as an activator of proMMP-13.<sup>49</sup> It will be interesting to learn from future studies if increased surface-bound MMP-14 activity also correlates with CD44 and/or MMP-13 levels. Endothelial cells exhibited constitutive MMP-13 activity that was lower than the M14#5 melanoma cell line, further suggesting that the level of this enzyme in M14#5 melanoma cells is functionally significant.

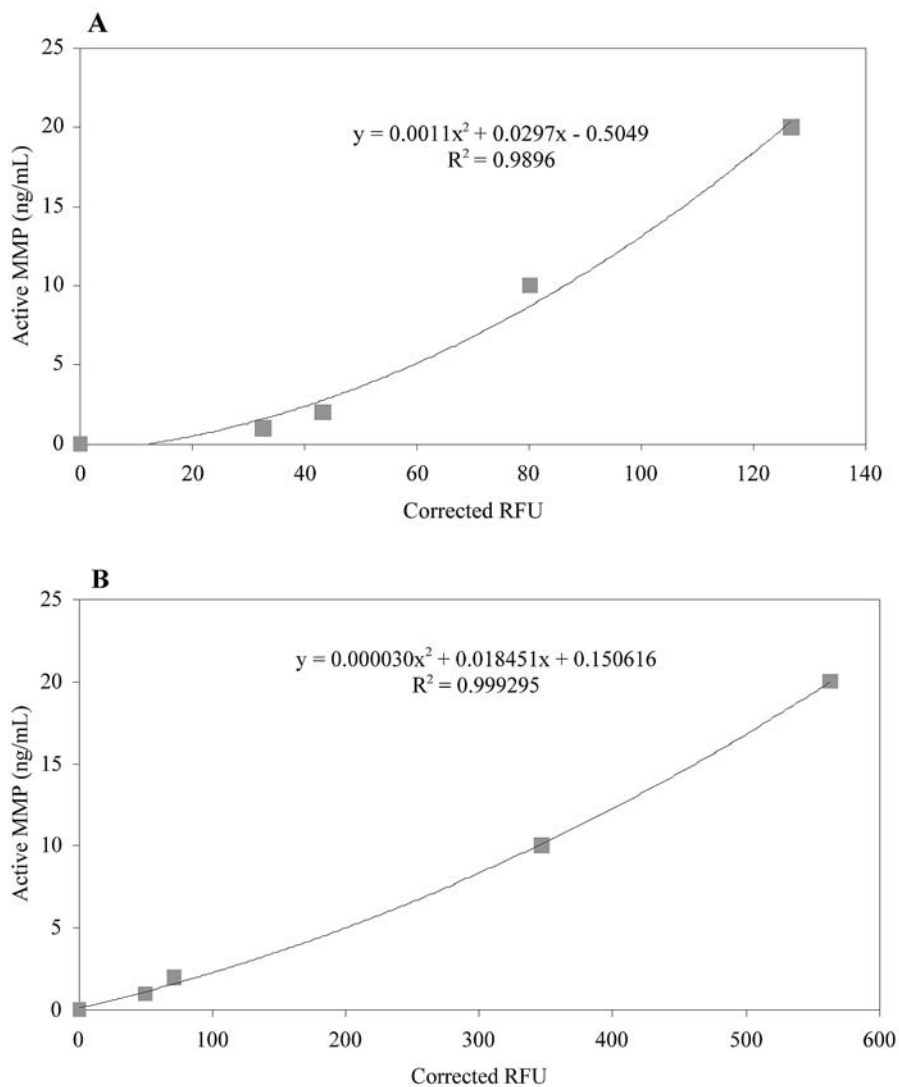
#### Induction of Active MMP-1 by Engagement of Melanoma Cell Receptors

Engagement of the  $\alpha_2\beta_1$  integrin and CD44 by extracellular matrix ligands results in intracellular signaling.<sup>50,51</sup> However, the precise nature of this signaling within melanoma cells has not been documented. The solid-phase assay was further utilized to examine induction of proteolysis upon melanoma cell  $\alpha_2\beta_1$  integrin and CD44 binding to “peptide-amphiphiles.” Peptide-amphiphiles consist of peptide sequences covalently linked to hydrocarbon chains, and are designed to house discrete binding sites (such as those found in various extracellular matrix proteins). These small biomolecules assume stable “mini-protein” structures ( $\alpha$ -helical or triple-helical) at physiological temperatures, allowing for the study of cellular behaviors in response to discrete binding sites that maintain the three-dimensional structure of the parent protein. For the present studies, melanoma cells were bound to triple-helical peptide-amphiphile ligands that incorporated either an  $\alpha_2\beta_1$  integrin or a CD44 binding site from type IV collagen.<sup>22,27,28,33</sup>

Induction of active MMP-1 was quantified by the solid-phase assay using fTHP-4. Active enzyme induction was compared for three ligands: C<sub>10</sub>-( $\alpha$ 1(IV)382–393)-NH<sub>2</sub> THP, C<sub>16</sub>-( $\alpha$ 1(IV)382–393)-NH<sub>2</sub> THP, and C<sub>16</sub>-( $\alpha$ 1(IV)1263–1277)-NH<sub>2</sub> THP. The first two ligands are specific for the  $\alpha_2\beta_1$  integrin, while the third ligand binds CD44. The solid-phase assay showed more activity induced by the  $\alpha_2\beta_1$  integrin than by CD44 (4.0–7.0 versus 2.5 ng/mL) (Fig. 8).

**FIGURE 7**

Standard curves for solid-phase MMP activity assay. Active MMP concentration was correlated to a change in substrate (fTHP-4 or fTHP-5) fluorescence using (A) MMP-1, (B) MMP-3, (C) MMP-13, and (D) MMP-14( $\Delta_{279-523}$ ). The amounts of MMP-1, MMP-3, and MMP-13 added ranged from 1.0 to 20 ng/mL, while a range of 0.5 to 10 ng/mL of MMP-14( $\Delta_{279-523}$ ) was used. Methods were as described in Results and Discussion. Corrected RFU refers to the relative fluorescence units minus background (no enzyme).



This result may be related to increased MMP-3, seen in response to the  $\alpha_2\beta_1$  integrin ligand.<sup>28</sup> MMP-3 is an activator of proMMP-1.<sup>34</sup> Prior studies had shown that binding of the  $\alpha_2\beta_1$  integrin to collagen results in increased production of MMP-1<sup>50,52</sup>.

Between the two  $\alpha_2\beta_1$  integrin ligands, the one with the C<sub>10</sub> modification induced higher levels of MMP-1 activity than did the C<sub>16</sub> modified ligand (7.0 versus 4.0 ng/mL) (Fig. 8). Differing MMP-1 induction based on the length and subsequent clustering of the  $\alpha_2\beta_1$  integrin ligand is reminiscent of prior studies showing different melanoma cell responses (adhesion, spreading, signaling) based on the length or density and thus accessibility of ligand.<sup>53-56</sup> Melanoma responses to ligand are clearly dependent upon how much of the ligand is accessible for cell binding.

**TABLE I**

Analysis of Constitutive MMP Activity in Melanoma and Endothelial Cell Media

Cell line	MMP-1 (ng/mL)	MMP-3 (ng/mL)	MMP-13 (ng/mL)	sMMP-14 (ng/mL)
M14P melanoma	1.37	1.47	1.71	0.89
M14#5 melanoma	0.89	1.69	27.1	2.19
M14#11 melanoma	0.72	1.49	14.3	1.73
Aortic endothelial	0.79	0.59	7.09	2.17



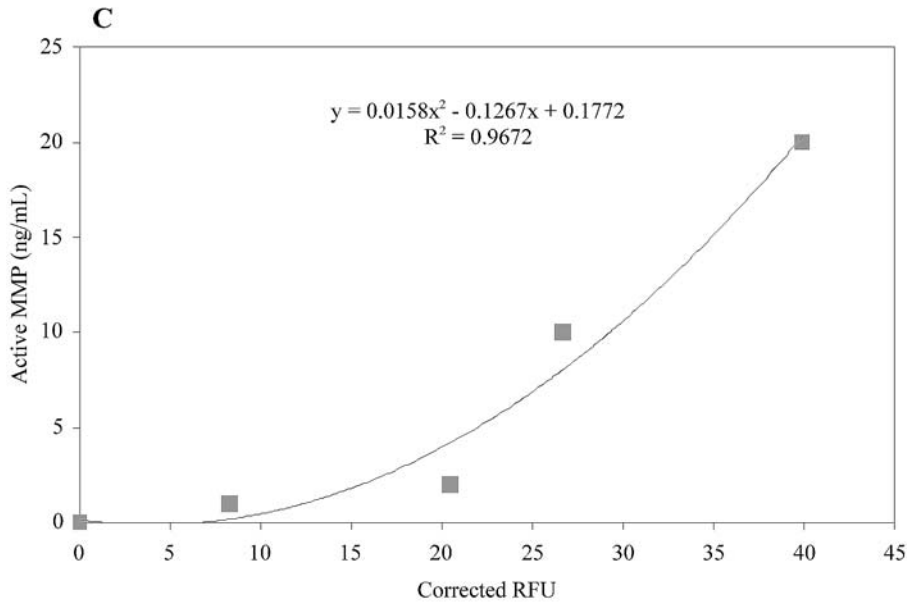


FIGURE 7 (CONT.)

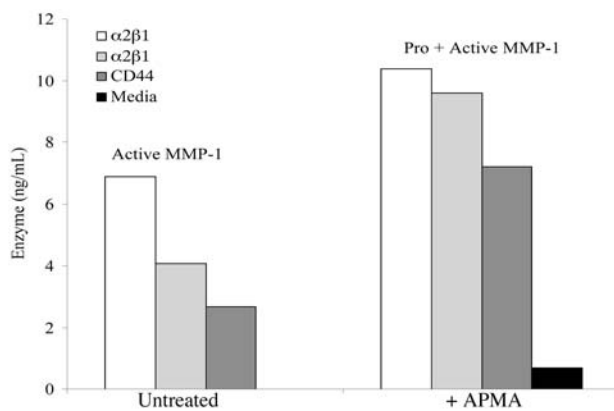
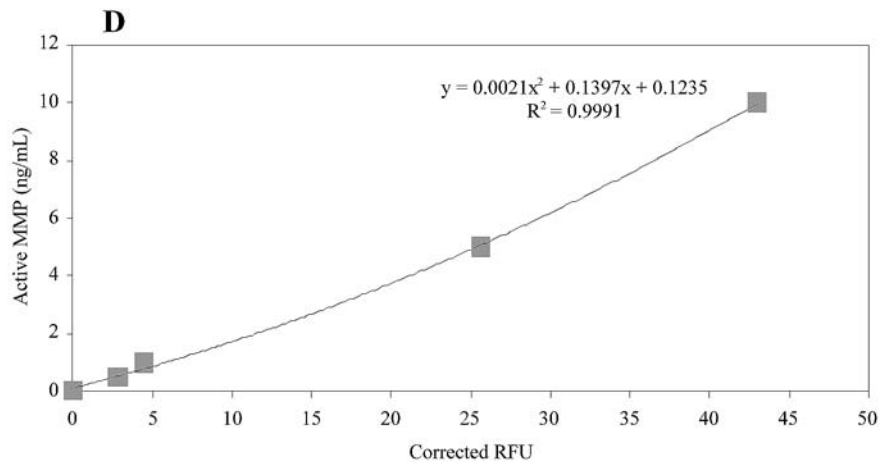


FIGURE 8

MMP-1 activity levels in conditioned media following melanoma cell adhesion to the  $\alpha_2\beta_1$  integrin specific  $C_{10}$ -( $\alpha 1$ (IV)382–393)-NH<sub>2</sub> THP (white bar) and  $C_{16}$ -( $\alpha 1$ (IV)382–393)-NH<sub>2</sub> THP (light grey bar) and the CD44/CSPG specific  $C_{16}$ -( $\alpha 1$ (IV)1263–1277)-NH<sub>2</sub> THP (dark grey bar). Cell supernatants were added to 96 well plates containing 2.5  $\mu$ g/mL MMP-1 mAb. Samples were either untreated (left panels) or activated with 2 mM APMA (right panels). Three washes of enzyme assay buffer were used to remove unbound proteins. The fTHP-4 substrate was added, and hydrolysis was monitored at 18 h. The black bar at the far right was media alone. Methods are as described in Materials and Methods. Results are the mean of triplicate assays.

Treatment of samples with an activator of proMMPs (APMA) resulted in a further increase in MMP-1 activity on the order of 3.5 to 5.5 ng/mL (Fig. 8). Thus, engagement of either the  $\alpha_2\beta_1$  integrin or CD44 induces production of both MMP-1 and proMMP-1.

## CONCLUSIONS

The solid-phase MMP activity assay has numerous advantages over prior methods for analyzing MMP activity, including combining (a) high sensitivity, (b) enzyme selectivity, and (c) high-throughput analysis. The efficiency of the assay was critically dependent upon the quality of the antibodies and enzyme standards, the use of a substrate with high specific activity, and enzyme samples that were fresh (immediately stored at  $-20^\circ\text{C}$  and analyzed within one month). The assay did require the use of pure enzyme as a control to ensure proper quantification, although comparisons between samples can readily be made without a standard if the overall goal is to assess relative activities. Antibodies to MMP linker regions, while avoiding the enzyme active site, may be problematic if MMP autolysis occurs within the linker but the MMP retains substantial activity (i.e., MMP-3).

The solid-phase MMP activity assay was utilized to demonstrate differences in constitutive and inducible levels of active MMP production by melanoma cells. Of particular interest was the low constitutive level of MMP-1 and MMP-3 activity by all cell types while one melanoma cell line had significant MMP-13 constitutive activity. These initial studies warrant further investigation of the interactive role of CD44, MMP-13, and MMP-14 in melanoma progression and comparison between surface-bound and soluble active MMP-14 levels. Future analysis of surface-bound MMP-14 will be especially valuable, as current methods cannot quantify MMP-14 levels in cellular environments. Analysis of surface-bound MMP-14 will probably require cell lysis and inhibition of nonmetalloproteases to minimize processing of capture antibodies.

Differences in MMP-1 induction by engagement of the  $\alpha_2\beta_1$  integrin compared with CD44 suggested that extracellular matrix components such as type IV collagen can modulate melanoma invasion. The solid-phase MMP activity assay appears to be useful for studies on microenvironmental regulation of cellular proteolytic potential.

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The solid-phase MMP activity assay was initially presented at Proteases, Extracellular Matrix and Cancer: American Association for Cancer Research Special Conference in Cancer Research, Hilton Head Island, SC, October 9–13, 2002. Solid-phase MMP activity assay kits are currently commercially available for MMP-1, MMP-8, and MMP-13 (Fluorokine E kits, R&D Systems, Minneapolis, MN).

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