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Journal Name

ARTICLE

Cite this: DOI: 10.1039/x0xx00000x

Received ooth January 2015, Accepted ooth January 2015

DOI: 10.1039/x0xx00000x

www.rsc.org/

Structure-activity relationship study for design of highly active covalent peroxidase-mimicking DNAzyme

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We synthesized a series of conjugates of hemin and its aptamer EAD2, named covalent peroxidasemimicking DNAzymes (PMDNAzymes), varying the length, rigidity and 5'-/3'-position of the linker between the oligonucleotide and hemin. Systemic structure-activity relationship study of these PMDNAzymes showed that covalent PMDNAzyme with hemin bound to the 5'-end of EAD2 via T_{10} spacer (PMDNAzyme(T_{10})) demonstrated the highest activity in luminol oxidation assay. Its activity was significantly higher in comparison to the non-covalent complex of hemin and aptamer EAD2 (noncovalent PMDNAzyme). Comparison of the detection limit values for the PMDNAzyme(T_{10}) in the reactions of oxidation of luminol and ABTS, which were equal to 0.2 and 1.6 pM, respectively, showed that the chemiluminescent method of PMDNAzyme(T_{10}) detection is preferred over the colorimetric one. Similarity of the detection limit values for the PMDNAzyme(T_{10}) and horseradish peroxidase, whose activity was measured in an enhanced chemiluminescence reaction (0.25 pM), opens up very promising perspectives for the development of highly sensitive PMDNAzyme(T_{10})-based assays and devices.

1. Introduction

Nowadays, horseradish peroxidase (HRP, EC 1.11.1.7) is extensively used in analytical practice due to its high activity, stability and commercial availablility.¹⁻³ However, the high cost of the enzyme has stimulated a search of its alternatives. Some peroxidase mimetics such as hemin, metal-containing porphyrins and phthalocyanines, Fe^{III}-TAML activators and nanoparticles of different chemical nature have been reported to possess peroxidaselike activity.⁴⁻⁷ Among them, complexes of hemin and its Gquadruplex aptamers attract great interest, since their catalytic activity is significantly higher than that of hemin itself.^{8,9} Such complexes named peroxidase-mimicking **DNAzymes** (PMDNAzymes) have been used in construction of various analytical assays and biosensors.10-14

Nevertheless, the widespread application of non-covalent PMDNAzymes is restricted because of their relatively low sensitivity. Detection limit of non-covalent PMDNAzyme formed by hemin and its EAD2 aptamer, even in the chemiluminescent assay, was only 350 pM,¹⁵ whereas the detection limit of the native HRP was reported to be 0.25 pM.¹⁶ One of the main reasons of the low sensitivity of non-covalent PMDNAzyme determination is dissociation of the PMDNAzyme molecules to the initial components (DNA and hemin) at nano- and picomolar concentrations because of a poor affinity of hemin to its aptamers (K_d was ca. 10⁻⁷ M).^{15,17-19} However, this drawback could be overcome by a covalent PMDNAzyme.

The first covalent PMDNAzyme synthesis has been reported in 2010, when Thirstup and Baird prepared a conjugate of hemin and its aptamer PS2.M.²⁰ Hemin was bound to the 3'-amino group of 5'-biotin-dA₁₅-PS2.M-NH₂. A detection limit of this PMDNAzyme estimated with 3,3'-diaminobenzidine as a substrate was 10 nM,

whereas the detection limit of a noncovalent complex of hemin and PS2.M was 100 nM. The obtained covalent PMDNAzyme was applied in the immunohistochemical analysis of prostate specific antigen in prostate tissue as an alternative of HRP.²⁰

Later Nakavama et al. have synthesized covalent PMDNAzymes by covalent coupling of hemin to aptamers with parallel and antiparallel G-quadruplex topology.21 Hemin was attached to the 5'-ends of the aptamers or internally to the G-quadruplex loop. All obtained PMDNAzymes were active catalysts, their catalytic activity depending strongly on substrates used and experimental conditions. Although covalent PMDNAzymes demonstrate significantly lower detection limits compared to the non-covalent PMDNAzymes, our present knowledge about the effect of the covalent PMDNAzyme structure on their catalytic activity is extremely limited. The high peroxidase activity of PMDNAzymes is considered to result from the efficient end stacking of hemin with its aptamer having Gquadruplex structure.²¹ In all covalent PMDNAzymes mentioned above hemin was bound to amino modified aptamers without any spacer. This manner of the hemin coupling limits its mobility and can therefore lead to serious steric hindrances for 3D organization of catalytically active covalent PMDNAzymes. Such hindrances could be removed by insertion of spacers between hemin and its aptamers. In the present work we described first the systemic structure-activity relationship study of a covalent PMDNAzyme based on the EAD2 aptamer (CTGGG(AGGG)₃). This aptamer was chosen for the covalent PMDNAzyme construction, since its complex with hemin is one of the most active non-covalent PMDNAzymes.^{15,22,23} The length and rigidity of spacers between hemin and EAD2 as well as hemin position at 3'- or 5'-end of the aptamer were varied. This approach allowed producing covalent PMDNAzymes with extremely high catalytic activity. Among all the hemin-EAD2 conjugates

obtained, the PMDNAzyme with hemin bound at the 5'-end of

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Fig. 1 Scheme of the synthesis of covalent PMDNAzymes

EAD2 via dT₁₀ spacer (PMDNAzyme(T₁₀)) was the most active catalyst. The activity of PMDNAzyme(T₁₀) measured towards both luminol (chemiluminescent substrate) and 2,2'-azino-bis(3-ethylbenzothiazoline-6-sulphonic acid) (ABTS, colorimetric substrate) turned out to be significantly higher than that of the noncovalent hemin/EAD2 complex. The detection limit value for the PMDNAzyme(T₁₀) was similar to that one for the native HRP and significantly lower than that for the parent non-covalent PMDNAzyme.

2. Experimental

2.1. Chemicals

Luminol, hemin, NaCl, KCl, and tris(hydroxymethyl)aminomethane (Tris), dicyclohexylcarbodiimide (DCC), N-hydroxysuccinimide (NHS), ABTS and *N*,*N*-diisopropylethylamine (DIPEA) were obtained from Sigma (USA). Universal Controlled Pore Glass (CPG) (loading 40 μ mol/g), protected 2'-deoxynucleoside (A(Bz), G(iBu), C(Ac), T) phosphoramidites, C3 (1,3-propanediol) linker and 5'-amino-modifier C6 phosphoramidites were purchased from ChemGenes (USA). C7-amino CPG (loading 45 μ mol/g) was purchased from GlenResearch (USA).

Hydrogen peroxide (H₂O₂, 30%) was obtained from ChemMed (Russia). The concentration of H₂O₂ was estimated by measuring absorbance using $\varepsilon_{240} = 43.6$.

Amino modified oligonucleotides (Table 1) were synthesized by standard phosphoramidite method under conditions recommended by manufacturer on an ABI 3400 DNA synthesizer (Applied Biosystems, USA) with minor modifications: detrytilation was carried out with 6% dichloroacetic acid in 1,2-dichloroethane, coupling of all modified phosphoramidites was carried out using 4,5-dicyanoimidazole as an activator for 5 minutes. The oligonucleotides obtained were deprotected using ammonium hydroxide/methylamine solution for 20 min at 65°C.

The oligonucleotides electroeluted from gel by Elutrap electroelution system (Whatman) were additionally purified by HPLC on an ÄKTA Purifier (GE Healthcare, USA) equipped with a Jupiter C18 column (Phenomenex, Jupiter 5 μ m, 300 Å, 250 x 4.6 mm) and a

Table 1 Comparison of the chemiluminescent activity of covalent PMDNAzymes containing spacers of different lengths and hemin in different positions. Experimental conditions: 25 mM Tris buffer, pH 8.6, 200 mM NaCl, [luminol] = 5 μ M, [H₂O₂] = 2 mM, [PMDNAzyme] = 2 x 10⁻⁷ M.

N	covalent PMDNAzyme 5^{2} \searrow 2^{2}	Chemiluminescent activity	
	5 - 5	10 ⁶ RLU	%
1	Hemin-T ₁₅ -EAD2	7.9	71
2	Hemin-T ₁₀ -EAD2	11.1	100
3	Hemin-T ₅ -EAD2	10.0	90
4	Hemin-EAD2	6.7	60
5	EAD2-T ₁₅ -Hemin	7.5	68
6	EAD2-T ₁₀ -Hemin	8.6	77
7	EAD2-T ₅ -Hemin	5.5	50
8	EAD2-Hemin	5.2	47
9	Hemin-Prd ₁₀ -EAD2*	8.1	73
10	EAD2-Prd ₁₀ -Hemin*	2.2	20

*- Prd – 1,3-propandiol

UV-Vis detector. The oligonucleotides were eluted using 0.05 M ammonium acetate (pH 7) with an acetonitrile gradient (0 - 80%). The oligonucleotides were desalted by ethanol precipitation and redissolved in distilled water.

2.2. Synthesis of hemin-oligonucleotide conjugates

The NHS ester of hemin was prepared accordingly to²¹ (Fig. 1) by adding 6 equiv each of DCC, NHS and DIPEA in 100 μ L anhydrous dimethylformamide (DMF) to 3 μ mole hemin and the reaction was carried out overnight at room temperature. Then a 10-fold volume of isopropanol was added to the reaction mixture, and the black precipitate was collected by centrifugation for 10 min at 10 000 g and washed twice with 1 mL of isopropanol. The obtained activated ester of hemin was dissolved in 400 μ L of DMF and used for preparation of hemin-DNA conjugates instantly.

Hemin-DNA conjugates were prepared by adding the activated ester of hemin (600 nmoles) to amino-functionalized oligonucleotides (10 nmoles) in 200 µL of 50 mM NaHCO₃, pH 9.0 containing 50% DMF, and incubating at room temperature for 20 h (Fig. 1). The obtained conjugates were precipitated with 1 mL of 2% LiClO₄ solution in acetone at room temperature for 10 min. The precipitate was collected by centrifugation for 10 min at 10 000 g and redissolved in 80% aqueous formamide. The hemin-DNA conjugates were purified by electrophoresis in a 20% polyacrylamide/7 M urea gel. The main band corresponded to a hemin-DNA conjugate was cut out, eluted from gel using aqueous 2 M LiClO₄ and then precipitated adding 1 mL of acetone. Finally the precipitate obtained by centrifugation (10 min, 10 000 g) was re-dissolved in distilled water. The band corresponded to the initial oligonucleotide was also cut and treated as described above. The DNA amount eluted from both bands was determined by UV spectroscopy, and on the base of these data the conjugation efficiency was estimated as 80-90%. The low yield (< 5%) of a conjugate with a molar ratio of hemin/EAD2=1:2 (see Fig. S1 as example) is explained by the use of a 60-fold molar excess of hemin in the reaction.

The absorbance spectra of the hemin-DNA conjugates were recorded on Carry 50 Bio (Varian). The extinction coefficients of DNA at 260 nm were calculated as a sum of the corresponding nucleoside extinction coefficients. The extinction coefficients of hemin at 260 nm and 405 nm were taken from the paper published by Travascio et al.⁸ Using these extinction coefficients, the absorbance spectra of the covalent derivatives were consistent with a 1:1 DNA-hemin complex. The concentration of the covalent PMDNAzymes prepared was determined measuring absorbance at 405 nm. The overall yields of the covalent PMDNAzyme preparations varied between 15 and 20%.

2.3. Characterization of hemin-oligonucleotide conjugates

2.3.1. MS analysis. ESI MS spectra were recorded on q-TOF Maxis Impact (Bruker Daltonics, USA) equipped with a 1260 (Agilent) HPLC system using acetonitrile gradient (2 - 80%) in 10 mM diisopropylamine/15 mM hexafluoroisopropanol as described previously.²⁴ The data obtained are presented in Table 1S.

2.3.2. Circular dichromism experiments (CD). CD experiments were performed on a JASCO J-815 spectropolarimeter at 25 °C in a 1 cm path length cuvette. The concentration of DNA was 2 μ M. Before the measurement, the sample was heated to 88 °C, and then cooled to room temperature for 60 min. CD measurements from 220 to 340 nm were taken. The data pitch was 1 nm and the scan speed was 50 nm/min. The response was 2 sec. The band width was 1 nm. Each spectrum was corrected by subtracting the CD of the buffer.

2.3.3. Determination of the catalytic activity of PMDNAzyme(T_{10}). Covalent PMDNAzyme solutions (50 nM) in 25 mM Tris buffer (pH 7.0 – 9.0) were heated at 88°C for 15 min and then cooled to room temperature for 60 min. Then KCl, NaCl and Triton X-100 were added to the DNA-hemin conjugates solutions up to concentrations of 20 mM, 200 mM and 0.1%, respectively, and incubated for 40 min at room temperature. The non-covalent PMDNAzyme was prepared by an interaction of hemin and aptamer EAD2 as described previously.¹⁵

Colorimetric determination of the covalent PMDNAzyme was carried out in wells of transparent polystyrene plates for enzyme immunoassay (Costar, USA). For this, covalent PMDNAzyme samples were mixed with aqueous solutions of ABTS and H_2O_2 . The final concentrations of ABTS and H_2O_2 in the wells were 4 mM and 12 mM, respectively, unless otherwise stated. Change of adsorption at 405 nm was measured for 20 min after the initiation of ABTS oxidation at room temperature on a reader (Zenyth 3100, Austria). The determination of the non-covalent PMDNAzyme was carried out under the same conditions.

Chemiluminescent determination of the covalent PMDNAzyme was carried out in wells of black polystyrene plates for enzyme immunoassay (Costar, USA). For this, covalent PMDNAzyme samples were mixed with aqueous solutions of luminol and H_2O_2 . The final concentrations of luminol and H_2O_2 in the wells were 5 μ M and 2 mM, respectively, unless otherwise stated. CL intensity was measured 1 min after the initiation of luminol oxidation at room temperature on a luminometer (Spectra Max L, USA). Chemiluminescent activity of the non-covalent PMDNAzyme was measured as described previously.¹⁵ The light intensity was expressed in relative luminescence units (RLU).

3. Results and discussion

3.1. Optimization of the covalent PMDNAzyme structure

Limited knowledge about the structure-activity relationships for covalent PMDNAzymes stimulated optimization of their structure. We synthesized two sets of covalent PMDNAzymes composed of hemin and its aptamer EAD2 covalently bound through spacers of different lengths and nature. The role of the spacer within covalent PMDNAzyme structure is to covalently attach hemin to its aptamer thus providing formation of an intramolecular complex of hemin and EAD2 with high peroxidase-like activity.

The conjugation was carried out by acylation of amino-modified oligonucleotides with hemin NHS ester (Fig. 1). In the first set of covalent PMDNAzymes hemin was attached to the 5'-end, whereas covalent PMDNAzymes from the second set contained hemin at the 3'-end of DNA (Table 1). All the covalent PMDNAzymes prepared were purified by electrophoresis in 20% polyacrylamide gel/7 M urea and their structure was confirmed by the electrophoresis mobility (Fig. S1), LC-MS (Table 1S, ESI) and absorbance spectra. The absorbance spectrum of one of the conjugates (Table 1, *item 2*) is presented in Fig. 1. The data obtained clearly indicated that a molar ratio of hemin and DNA in the synthesized conjugates was 1:1. Interestingly, the spectra of the covalent PMDNAzymes in the range of 350-370 nm have no shoulder that is characteristic of free hemin in aqueous solutions (Fig.2, curve 2). The existence of such shoulder for water solution of hemin was also reported earlier.8,25 This means that hemin within the covalent PMDNAzymes exists in monomeric form unlike free hemin. This is an important characteristic of any hemin/aptamer complex, because a transfer of hemin from dimeric to monomeric form increases the catalytic activity of the complex compared to free hemin.⁸

We measured catalytic activity of the EAD2/hemin conjugates prepared towards luminol since it usually gives an ultrasensivity in peroxidase-based assays.²⁶⁻²⁸ All the covalent PMDNAzymes possessed HRP-like activity (Table 1). Nevertheless, we observed a clear effect of the hemin position on the covalent PMDNAzymes catalytic properties. All covalent PMDNAzymes with hemin bound at the 5'-end had higher catalytic activity than their analogs bearing hemin at the 3'-end. This effect may arise from different conformational changes in the proper G-quadruplex structure induced with its 5'- or 3'-end modifications. Further our work will focus on the detailed study of this question. It should be also noted that our data correlate with results of Zhang et al. who showed that adding dT_n fragment to the 5'-end of hemin aptamer DZ (5'- $GGTA(G)_3C(G)_3TT(G)_3-3'$ improved the catalytic activity of a noncovalent complexes of such modified aptamers with hemin, whereas the addition of dT_n to the 3'-end decreased their activity.²⁹ In addition, the covalent PMDNAzyme activity was found to depend



Fig. 2 UV/vis spectra of PMDNAzyme(T_{10}) (1), amino-containing aptamer NH₂- T_{10} -EAD2 (2) and hemin alone (3) dissolved in distilled water.

on the spacer length. In both series of the conjugates (Table 1, *items 1-4 and 5-8*) the highest activity was observed for the covalent PMDNAzymes with T_{10} spacer (Table 1, *items 2 and 6*). It appears that this spacer is optimal to provide the intramolecular stacking interactions between hemin and G-quanduplex structure of EAD2 necessary for the organization of the catalytically active conformation of the PMDNAzyme.

To evaluate the effect of the spacer flexibility on the covalent PMDNAzyme activity, we replaced the spacer dT_{10} with (1,3-propanediol)₁₀ spacer (Prd₁₀), which had a similar length, but higher mobility. It turned out that increasing the flexibility of the spacer decreased the activity of the covalent PMDNAzyme, and this effect was observed for the compounds containing hemin at both 5'- and 3'-ends (Table 1, *items 1, 8*). Taking into account the abovementioned, the further work was focused on the study of catalytic properties of the most active covalent PMDNAzyme with hemin bound at 5'-end via dT_{10} spacer (PMDNAzyme(T_{10})).

3.2. Study of the synthesized covalent PMDNAzymes topology

The most active G-quadruplex aptamers of hemin in the presence of K^+ cations are known to have parallel or mixed-hybrid topology.^{21,30,31} We proceeded to investigate the topology of the synthesized covalent PMDNAzymes using circular dichroism (CD) spectroscopy and gel shift assay.

CD spectra of PMDNAzyme(T_{10}) and its parent aptamer NH₂- T_{10} -EAD2 are shown in Fig. 3. In the presence of K^+ cations CD spectrum of PMDNAzyme(T_{10}) had a positive signal at around 265 nm and a negative signal at 242 nm, which are characteristic of parallel G-quadruplexes.²¹ The similar spectrum was measured for the aptamer NH_2 -T₁₀-EAD2 itself. Therefore, these results demonstrated that EAD2 after its covalent binding with hemin remained the parallel G-quadruplex structure found earlier for the studied aptamer.^{21,32} Moreover, we showed that CD spectra of PMDNAzyme(T_{10}) in the presence and in the absence of K⁺ cations were practically identical, whereas CD spectra of NH₂-T₁₀-EAD2 changed (Fig. 3). The invariability of CD-spectrum of PMDNAzyme(T_{10}) in the presence and absence of K⁺ cations allowed us to suppose that its G-quadruplex structure is stabilized by Na⁺ ions and hemin together, hemin stabilization being due to its interactions with the adjacent G-quartet.



Fig. 3 CD spectra of (*1*,*3*) PMDNAzyme(T_{10}) and (*2*,*4*) NH₂- T_{10} -EAD2 aptamer in the presence (*1*,*2*) and absence (*3*,*4*) of 20 mM KCl. The samples were dissolved in 25 mM Tris buffer, pH 8.6, 200 mM NaCl. Oligonucleotides concentration was 2 μ M.

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Capacity of the synthesized covalent PMDNAzymes to maintain quadruplex topology in the absence of potassium was confirmed by gel shift assay experiments (Fig. 2S). The conjugate EAD2- T_{10} -hemin and its parent aptamer EAD2- T_{10} -NH₂ were used, both 5'-end ³²P-labelled to simplify the analysis. As seen in Fig. 2S, the migration rate of EAD2- T_{10} -hemin rate did not depend on the presence or absence of K⁺ cations, whereas the mobility of EAD2- T_{10} -NH₂ decreased in the absence of potassium because of destruction of its compact quadruplex structure.

Of note, the existence of a parallel G-quadruplex structure in the absence of potassium (but in presence of sodium) ions has been shown for a covalent PMDNAzyme prepared on the base of another hemin aptamer $(5'-(G)_3TA(G)_3C(G)_3TT(G)_3T-3')$.²¹

3.3. Optimization of conditions of PMDNAzyme(T₁₀) catalysis

It is well known that both native enzymes and their mimetics implement their catalytic potential only under defined conditions. In for order to optimize the experimental conditions PMDNAzyme(T₁₀), we varied concentrations of a substratehydrogen donor (luminol and ABTS used in chemiluminescent and colorimetric detection of the PMDNAzyme, respectively) and a substrate-oxidant (hydrogen peroxide) as well as pH value of the substrate solution (Figs. S3-S8). The most favorable conditions for PMDNAzyme(T_{10}) catalysis of luminol oxidation were found to be 25 mM Tris buffer, pH 8.6 containing 200 mM NaCl, 5 µM luminol and 2 mM H_2O_2 . The ABTS oxidation by PMDNAzyme(T_{10}) was the most efficient in 25 mM citrate-phosphate buffer, pH 6.0 containing 200 mM NaCl, 4 mM ABTS and 12 mM H₂O₂. It should be noted that contrary to the non-covalent PMDNAzyme, the activity of PMDNAzyme(T_{10}) did not depend on the presence of potassium cations in the reaction solution (data not shown), and this is in good accordance with the above-mentioned CD experiments. In addition, Triton X100 in the concentration up to 0.1% did not affect the PMDNAzyme(T_{10}) activity. This surfactant is commonly used upon preparation of non-covalent PMDNAzymes to transfer hemin from dimeric to monomeric state. Therefore, the absence of Triton X100 effect additionally points to the monomeric form of hemin within PMDNAzyme(T_{10}) (see above).

3.4. Analytical parameters of PMDNAzyme(T₁₀) detection

The dependence of the initial rate of ABTS oxidation on the PMDNAzyme(T_{10}) concentration studied under favorable conditions is presented in Fig. 4. The obtained results demonstrated clearly that hemin in the covalent PMDNAzyme is significantly more active than hemin within the non-covalent counterpart. The calibration curve for the PMDNAzyme(T_{10}) allowed calculating a detection limit value (3 σ) equal to 1.6 pM. Comparison of this value with the detection limit of the parent non-covalent PMDNAzyme (18 nM) showed that the sensitivity of the synthesized PMDNAzyme(T_{10}) determination was remarkably higher than that of its non-covalent analog (Fig. 4).

Although many researchers studying PMDNAzymes and applying them in analytical practice use the ABTS-based colorimetric method to estimate their catalytic activity,³³⁻³⁶ the chemiluminescent method based on luminol oxidation is well-known to be more sensitive. This fact is widespread used in the development of ultrasensitive analytical assays.²⁶⁻²⁸ Therefore, we studied the dependence of the chemiluminescence intensity produced upon luminol oxidation on the concentration of PMDNAzyme(T₁₀) and its parent non-covalent counterpart (hemin/EAD2 complex) and constructed the calibration curves for both studied HRP mimetics (Fig. 5). The detection limit value of PMDNAzyme(T₁₀) in the chemiluminescent method was 0.2 pM. Journal Name



Fig. 4 Dependence of the initial rate of catalytic oxidation of ABTS on the concentration of PMDNAzyme(T_{10}) (1) and the non-covalent PMDNAzyme (2). Experimental conditions: (1) 25 mM citrate-phosphate buffer, pH 6.0, 200 mM NaCl, 20 mM KCl and 0,1% Triton X-100, [ABTS] = 4 mM, [H₂O₂] = 12 mM.



Fig. 5 Dependence of the chemiluminescence intensity produced upon catalytic oxidation of luminol under favorable conditions on the concentration of PMDNAzyme(T_{10}) (1) and the non-covalent PMDNAzyme (2). Experimental conditions: (1) 25 mM Tris buffer, pH 8.6 with 200 mM NaCl, [luminol] = 5 μ M, [H₂O₂] = 2 mM; (2) 25 mM Tris buffer, pH 8.0 with 200 mM NaCl, 20 mM KCl, 0.05% Triton X-100, and 1% DMSO, [luminol] = 5 μ M, [H₂O₂] = 1.3 mM.

This value was remarkably lower than that of its non-covalent analog (350 pM). Moreover, the detection limit of the PMDNAzyme(T_{10}) estimated in the chemiluminescent method was one order lower than that of the same catalyst in the colorimetric method. These results clearly demonstrated that upon development of highly sensitive assays it is preferable to apply PMDNAzyme(T_{10}) instead of traditionally used non-covalent PMDNAzymes.

Conclusions

In this study we optimized the structure of a covalent PMDNAzyme, a conjugate of hemin with its aptamer EAD2. The influence of the hemin position at 3'- or 5'-end of the aptamer, as well as the effect of the length and nature of spacers between the oligonucleotide and hemin on the covalent PMDNAzyme catalytic activity towards luminol was studied in detail. Among all the conjugates studied, the covalent PMDNAzyme with hemin bound at the 5'-end of EAD2 via dT_{10} spacer possessed the highest activity. The chemiluminescent activity of the produced PMDNAzyme(T_{10}) was significantly higher in comparison to the non-covalent hemin/EAD2 complex. Taking into account that the detection limit for PMDNAzyme(T_{10}) is similar to that for HRP (0.25 pM), whose enzymatic activity was measured in an enhanced chemiluminescence reaction with the most active enhancers (3-(10'-phenothiazinyl)propane-1-sulfonate or 3-(10'-phenothiazinyl)propane-1-sulfonate or 3-(10'-phenothiazinyl)propane-1-sulfonate or 3-(10'-phenothiazinyl)propane-1-sulfonate or 3-(10'-phenothiazinyl)propinic acid with 4-morpholinopyridine) so far, ^{16,26} the PMDNAzyme(T_{10}) has very promising perspectives for its wide application to develop assays with improved sensitivity. Further our work will be focused on the development of covalent PMDNAzyme based analytical kits and assays.

Acknowledgements

The authors thank K. Sukhoverkov for help with CD measurements and the Russian Foundation for Basic Research (NK-13-04-91164/14) for financial support.

Supporting Information

Additional information as noted in the text (Table S1 and Figures S1-S6).

This material is available free of charge via the Internet at http://.

Notes and references

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Electronic Supplementary Information (ESI) available: [Additional information as noted in the text (Table S1 and Figures S1-S6)]. See DOI: 10.1039/b000000x/

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