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Phenoxy radical reactivity of nucleic acids: practical implications for biotinylation

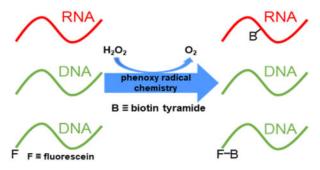
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Abstract

Recent advances in peroxidase-mediated biotin tyramide (BT) signal amplification technology have resulted in high-resolution and subcellular compartment-specific mapping of protein and RNA localization. Horseradish peroxidase (HRP) in the presence of H_2O_2 is known to activate phenolic compounds for phenoxy radical reaction with nucleic acids, where biotinylation by BT is a practical example. BT reactivity with RNA and DNA is not understood in detail. We report that BT phenoxy radicals react in a sequence-independent manner with guanosine bases in RNA. In contrast, DNA reactivity with BT is not detectable by our methods under the same conditions. Remarkably, we show that fluorescein conjugates of DNA rapidly and selectively react with BT phenoxy radicals, allowing convenient and practical biotinylation of DNA on fluorescein with retention of fluorescence.

Graphical Abstract



Practical applications of radical biotinylation: While recent advances in peroxidase-mediated biotinylation methods have resulted in impressive subcellular mapping, details of nucleic acid radical biotinylation remain undescribed. We assess nucleic acid reactivity with biotin tyramide radicals and describe novel reactivity with fluorescein, enabling DNA biotinylation methods for future application.

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conjugation; nucleic acids; peroxides; radicals; reaction mechanism

It has been shown that aryl radicals react at the C8 position of guanine and deoxyguanosine under certain conditions.^[1] Such reactions are of interest as examples of mutagenic DNA damage relevant to carcinogenesis. Aryl radical reactivity of RNA has not been studied in detail. The advent and rapid improvement of proximity-based labeling technologies such as biotin tyramide (BT) signal amplification and RNA localization mapping by the engineered ascorbate peroxidase APEX2 have highlighted the practical importance of RNA reactivity with phenoxy radicals.^[2] In another example, APEX2 polymerizes diaminobenzene in the presence of H_2O_2 and deposits it locally, providing focal osmium tetroxide binding for electron microscopy.^[3] However, the sequence specificity and relative reactivity of RNA and DNA with BT radicals have not been explored. We therefore studied BT phenoxy radical labelling of nucleic acids in vitro catalysed by horseradish peroxidase (HRP) in order to characterize these practical reactions in more detail.

We first aimed to confirm a previous report that RNA biotinylation by BT phenoxy radicals occurs at G₃ motifs.^[2f] LJM-6133 is an oligoribonucleotide 27-mer containing three separate G₃ motifs distributed within a mixed sequence (Table 1). We devised a streptavidin-dependent electrophoretic mobility shift assay for detection of biotinylation in vitro. A gel shift of RNA LJM-6133 was observed only when H₂O₂, BT, and HRP were incubated with the oligonucleotide for 90 min at 45 °C, followed by proteinase K digestion (Fig. 1A, lane 7; Fig. S1–S3). HRP contains heme, which is known to biotinylate certain G-quadruplex nucleic acids in the presence of H₂O₂ and BT^[4]. In order to demonstrate that observed biotinylation was HRP-dependent and would not occur with free heme, HRP was replaced by heme in an otherwise identical biotinylation reaction. Free heme was not capable of activating BT and biotinylating a target, demonstrating that 1) observed biotinylation is HRP-dependent and 2) the targets likely do not form stable G-quadruplexes that bind heme (Fig. S4). Mfold predictions^[5] did not identify significant secondary structures for these oligonucleotides (Fig. S5).

 G_3 motifs were then disrupted by substitution of a random nucleotide in the second position of each G_3 motif in RNA LJM-6135. This RNA was not detectably biotinylated under identical conditions (Fig. 1B, lane 7). To determine whether it is the number of guanosine residues, or their distribution, that confers BT reactivity, RNA LJM-6247 was synthesized containing the same number of guanosine residues as LJM-6133 but lacking G_3 motifs (Table 1). Strikingly, RNA LJM-6247 was biotinylated (Fig. 1C, lane 7; Fig. S6). These results demonstrate that RNA biotinylation does not depend solely on the presence of G_3 motifs, but instead on the total guanosine content of an RNA. We further demonstrated Gspecificity of RNA biotinylation by adding NTP competitor to biotinylation reactions, finding that only the addition of excess GTP reduced biotinylation of a target oligonucleotide (Fig. S7). In order to confirm that biotinylation of LJM-6247 was not influenced by the position of G relative to other RNA residues, we generated three

oligoribonucleotides with G residues in identical positions but with all other bases randomly mutated. These randomly-substituted targets were all similarly biotinylated (Fig. S8).

Finally, we confirmed by mass spectrometric analysis that a BT-guanosine adduct is formed at the C8 position of guanosine by BT phenoxy radical addition to LJM-6247 (Fig. S9). This structure is consistent with the literature for guanine and deoxyguanosine C8 reactivity with other phenoxy radical-containing compounds ^[1], but to our knowledge is the first direct evidence of such biotinylation of guanosine. Importantly, BT adducts with other bases were not identified (Fig. S10). Given the context-independent reactivity of guanosine in LJM-6247, it is therefore likely that previous results identifying consecutive guanosine residues as preferred sequences for RNA phenoxy radical reactivity simply reflect the high number of guanosines at these sites.

Having established that RNA biotinylation occurs on guanosine independent of sequence context, we next investigated radical biotinylation of DNA oligonucleotides under identical conditions. Oligodeoxyribonucleotide LJM-6131 has the same sequence as RNA LJM-6133. Remarkably, LJM-6131 was not detectably biotinylated by HRP under identical conditions (Fig. 2A, Fig. S11–S12). Previous reports suggest that deoxyguanosine can react with phenoxy radicals under some conditions, but this reaction is much more efficient with catalysts other than peroxidases. For example, Fe(III) or Cu(II) catalyzed reactions between deoxyguanosine and ochratoxin A, a mycotoxin with a single phenoxy group, up to 5-fold faster than with HRP/H₂O₂ at 24 h.^[1b] Several reports describe heme-dependent self-biotinylation of G-quadruplex DNAs by BT. ^[4]

Similar reactions show that heme is capable of oxidizing a library of phenolic substrates 7to 68-fold more efficiently than HRP.^[6] There is a report of double-stranded DNA biotinylation by HRP under conditions different from those presented here, complicating comparison of results.^[7] In order to assess double-stranded DNA biotinylation under the present conditions, oligodeoxyribonucleotide LJM-6349 was hybridized with LJM-6133 to form a duplex. No biotinylation of the duplex was observed (Fig. S13). We therefore conclude that DNA is intrinsically much less reactive with phenoxy radicals than RNA, limiting practical HRP-catalyzed BT conjugation. To eliminate the possibility that DNA biotinylation was undetectable because biotinylated DNA, but not RNA, suffers rapid depurination, we exposed reacted DNA oligonucleotides to hot piperidine treatment to cleave apurinic sites. No evidence of detectable DNA depurination was observed under these conditions (Fig. S14).

Serendipitously, we also studied DNA LJM-6132, a version of LJM-6131 differing only by 5' conjugation of fluorescein (FAM). Remarkably, whereas LJM-6131 resisted biotinylation, LJM-6132 was efficiently biotinylated in vitro by HRP in the presence of BT and H₂O₂ (Fig. 2B, lane 7; Fig. S9). This surprising result was confirmed in both native and denaturing polyacrylamide gels under conditions where biotin-streptavidin complexes remain stable (Fig. S11). Thus, fluorescein conjugation is necessary and sufficient for DNA phenoxy radical biotinylation. The multiple streptavidin-dependent bands observed in lane 7 of Fig. 2B presumably result from different numbers of biotinylated oligonucleotides binding to streptavidin tetramers.

Chembiochem. Author manuscript; available in PMC 2022 April 16.

Inspection of gel images also revealed unexpected shifted products in the absence of BT (Fig. 2B, lane 4). These products are of lower molecular weights than those observed for streptavidin complexes (Fig. 2B, lane 7). We hypothesized that these biotinylation-independent products resulted from phenoxy radical reactions between LJM-6132 molecules and between LJM-6132 and tyrosine residues of HRP. We therefore quenched reactions with proteinase K digestion for 1 h at 56 °C, to remove proteins. Indeed, most biotinylation-independent products were eliminated by proteinase K digestion, confirming that they reflect direct reaction of LJM-6132 with HRP (Fig. S2). Interestingly, several reaction products formed in the absence of BT resisted proteinase K digestion for 24 h (Fig. S3). We hypothesize that these proteinase K-resistant species are DNA-DNA crosslinked products. The presence of both DNA-protein and DNA-DNA products in radical reactions lacking BT suggests that fluorescein itself is activated by H_2O_2 similarly to BT, at phenolic functional groups present in both compounds. Given that DNA-DNA and DNA-HRP products are suppressed in the presence of excess BT, we conclude that radical activation of BT by HRP is efficient.

To confirm that electrophoretic mobility shifts detected for BT-dependent products indeed reflect noncovalent biotin-streptavidin complexes, we demonstrated that such products are sensitive to competition by addition of free biotin. Streptavidin was incubated for 30 min with $0-200 \mu$ M free biotin before the addition of LJM-6132 reaction products. Competition was observed as expected (Fig. 3, lanes 3–8). Maximum competition was observed at free biotin concentrations near equivalence with total biotin binding sites on streptavidin (Fig. 3, lanes 7–8). This result confirms that fluorescein-modified LJM-6132 undergoes phenoxy radical biotinylation in a manner dependent on the simultaneous presence of FAM, HRP, BT, and H₂O₂. It is striking that FAM reaction products remain fluorescent (Fig. 3, lane 3) while proteinase-K resistant FAM-FAM cross-linked products do not (Fig. S3A–B).

These results show that the resistance of unmodified DNA oligonucleotides to radical biotinylation can be practically overcome by FAM conjugation, a particularly convenient and accessible modification. We hypothesized that the fluorescein-BT reaction involves the phenolic functional groups of fluorescein, analogous to BT. To test this, we exploited the structural similarity of rhodamine (TAMRA) and fluorescein, where TAMRA lacks phenolic functional groups (Fig. 4A). DNA oligonucleotide conjugates to FAM (LJM-6132) and TAMRA (LJM-6223; Table 1) were subjected to radical biotinylation reactions. Whereas LJM-6132 was detectably biotinylated, TAMRA-conjugated LJM-6223 was not (Fig. 4B). Lower molecular weight BT-independent DNA-HRP products are also not observed with TAMRA-conjugated DNA. We deduce that phenoxy radical biotinylation of FAM-conjugated DNA involves the phenolic functional group of FAM. While results of preliminary mass spectrometric analyses to confirm the FAM biotinylation product were indeterminate, we propose the reaction mechanism shown in Scheme 1. This is, to our knowledge, the first report of peroxide-mediated activation of BT allowing biotinylation of fluorescein by phenoxy radical chemistry.

In summary, we present evidence that RNA is significantly more reactive to phenoxy radical biotinylation than DNA, and that RNA biotinylation occurs at guanosine residues. Radical biotinylation is at the C8 position of guanosine independent of sequence context, and is

Chembiochem. Author manuscript; available in PMC 2022 April 16.

therefore more likely for G-rich RNA sequences. In contrast, DNA biotinylation of the same sequence under identical conditions is so low as to be undetectable by our streptavidin binding assay. Importantly, phenoxy radical biotinylation of DNA oligonucleotides is conveniently facilitated by fluorescein conjugation. Fluorescein is directly activated by HRP and H_2O_2 , allowing radical biotinylation by BT. This insight may be helpful for future chemical biology applications as it enables proximity biotinylation of FAM-conjugated DNA by peroxides in cases where biotinylation of unmodified DNA is undetectable. Further study will be required to understand the relative susceptibility of DNA vs. RNA to phenoxy radical chemistry.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

Acknowledgements

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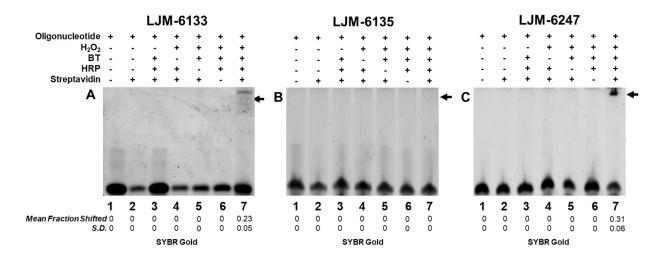


Figure 1. Oligoribonucleotide radical biotinylation is catalyzed by HRP in the presence of $\rm H_2O_2$ and biotin tyramide.

Biotinylation is detected by a streptavidin-dependent gel shift under denaturing conditions, followed by staining with SYBR Gold. A. Biotinylation of LJM-6133 containing G₃ motifs. B. Biotinylation of LJM-6135 lacking G₃ motifs. C. Biotinylation of LJM-6247 containing the same number of G residues as LJM-6133, without G₃ motifs. Statistics are calculated from independent experiments performed in triplicate; S.D. is standard deviation.

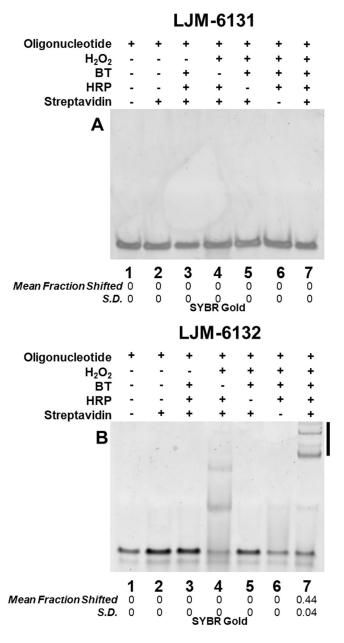


Figure 2. Oligodeoxyribonucleotide 5' FAM modification is necessary and sufficient for efficient radical biotinylation catalyzed by HRP in the presence of H_2O_2 and biotin tyramide. Biotinylation of the indicated DNA oligonucleotides is detected by a streptavidin-dependent gel mobility shift under native conditions. Statistics are calculated from independent experiments performed in triplicate; S.D. is standard deviation.

Page 8

Wilbanks et al.

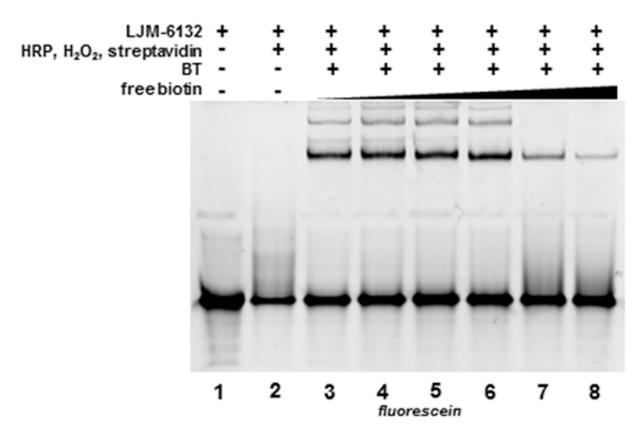


Figure 3. Biotinylation confirmed by free biotin competition for streptavidin binding to LJM-6132.

Biotin excess of $0 - 200 \,\mu\text{M}$ (lanes 3–8) in a 30-min pre-incubation inhibits LJM-6132 binding by streptavidin under native gel conditions.

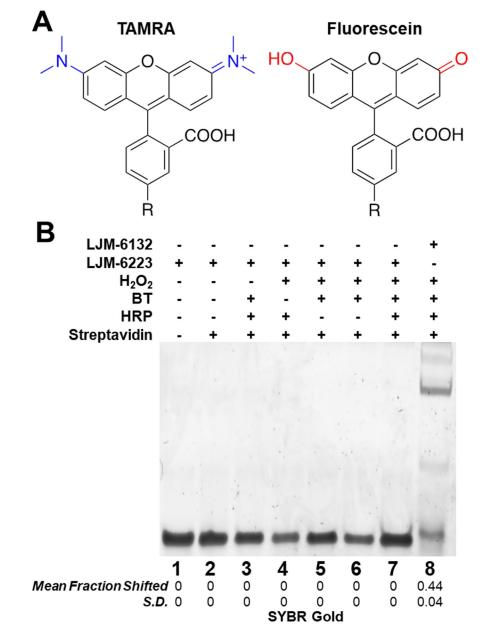
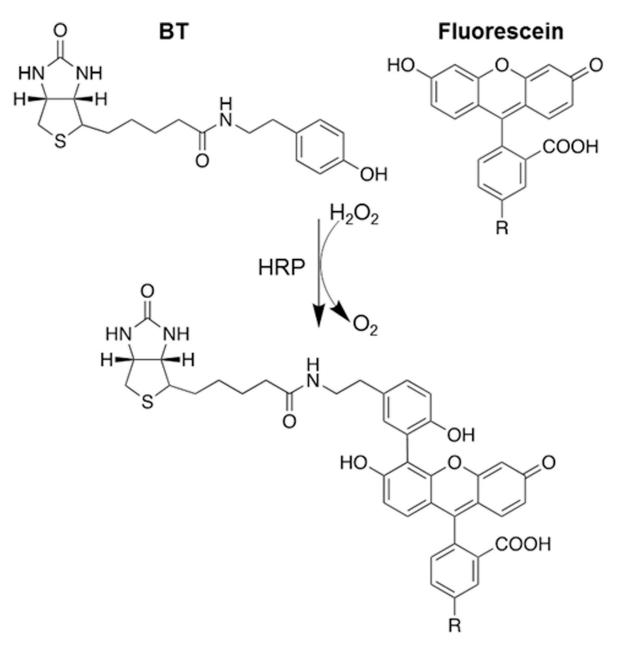


Figure 4. Radical biotinylation of FAM but not TAMRA.

A. Structures of FAM and TAMRA highlighting phenolic group of FAM (red) vs. amino groups of TAMRA (blue). R: oligonucleotide. B. Absence of detectable radical biotinylation under native gel conditions of 5' TAMRA-modified LJM-6223 relative to 5' FAM-modified LJM-6132. Statistics are calculated from independent experiments performed in triplicate; S.D. is standard deviation.



Scheme 1. Proposed mechanism of HRP-catalyzed phenoxy radical biotinylation of FAM by biotin tyramide in the presence of H₂O₂. R: oligonucleotide.

Chembiochem. Author manuscript; available in PMC 2022 April 16.

Table 1:

Oligonucleotides studied in this work

Sequence (5'-3')
AGGGCTAAGGTCTGACTGGGAATGGGA
FAM-AGGGCTAAGGTCTGACTGGGAATGGGA
AGGGCUAAGGUCUGACUGGGAAUGGGA
AGCGCUAAGGUCUGACUGAGAAUGUGA
AGAGCUGAGGUCUGACGGUGAGUGAGA
TAMRA-AGGGCTAAGGTCTGACTGGGAATGGGA
AGUGAUGAGGAACGUUGGAGCGCGUGC
UGCGUCGUGGACCGUCGGAGCGUGCGU
CGCGUAGUGGUUAGAAGGCGAGCGUGA
TCCCATTCCCAGTCAGACCTTAGCCCT