Synthesis of 2'-C- β -difluoromethylribonucleosides and enzymatic incorporation into oligonucleotides

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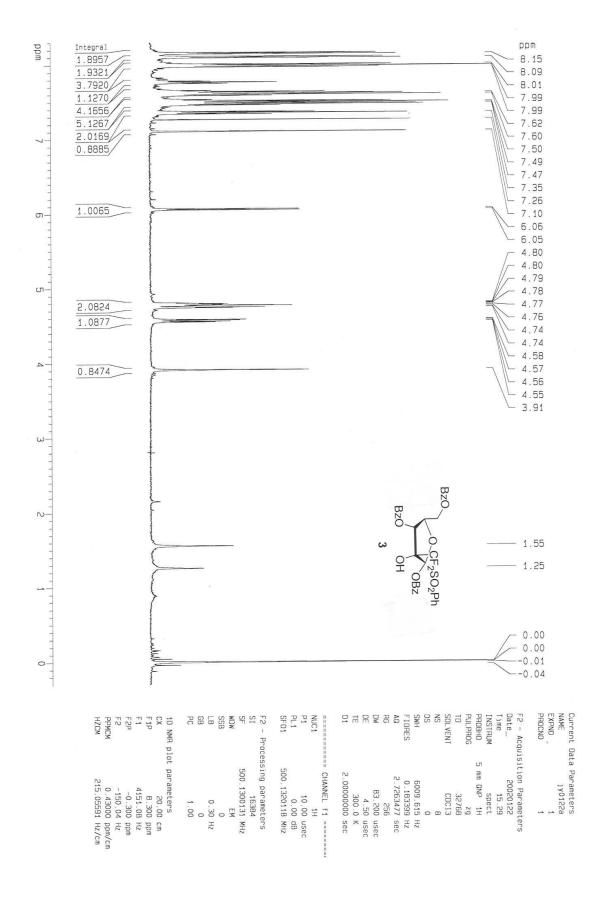
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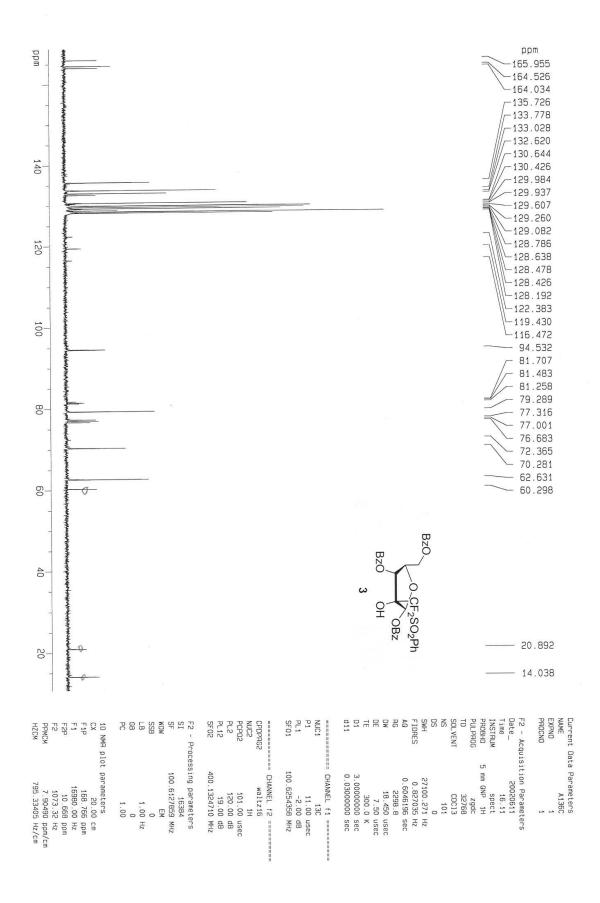
Supporting Information

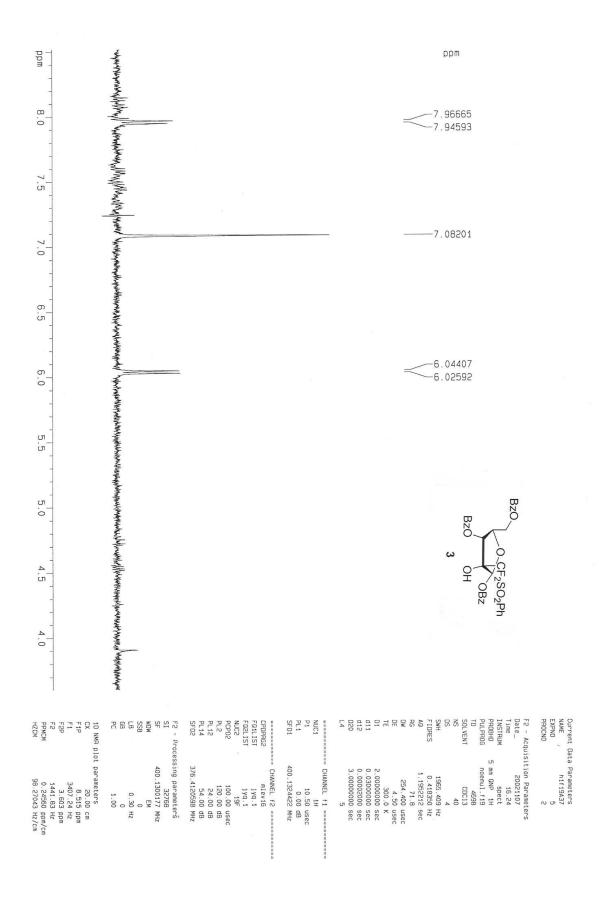
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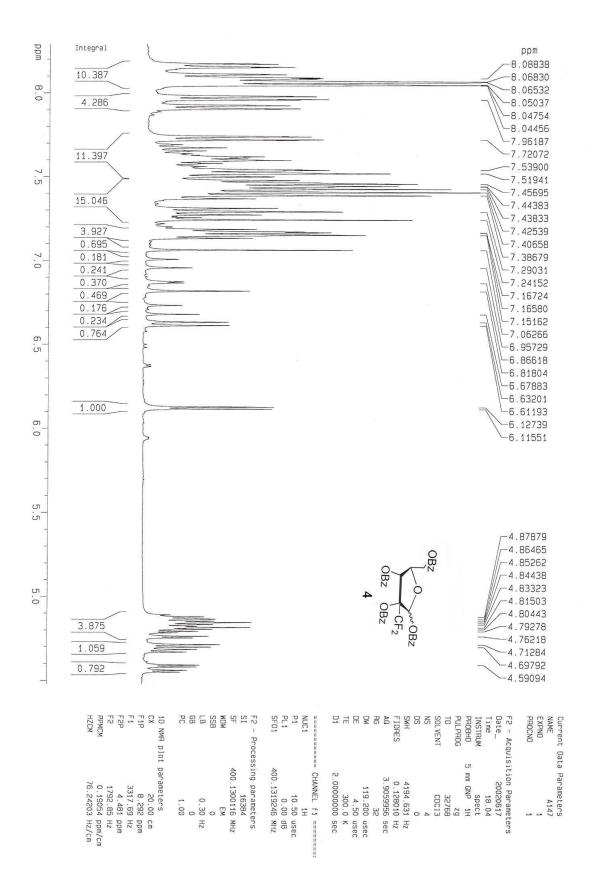
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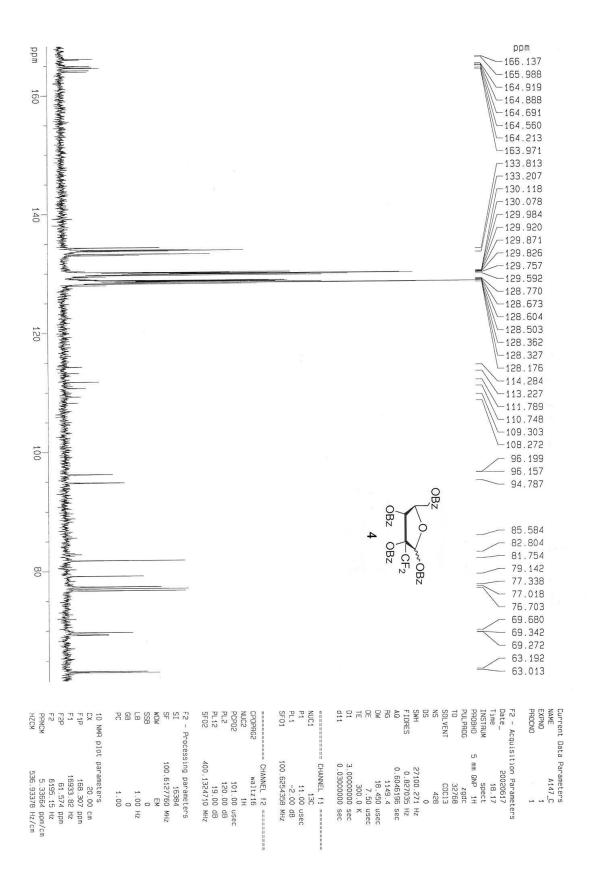
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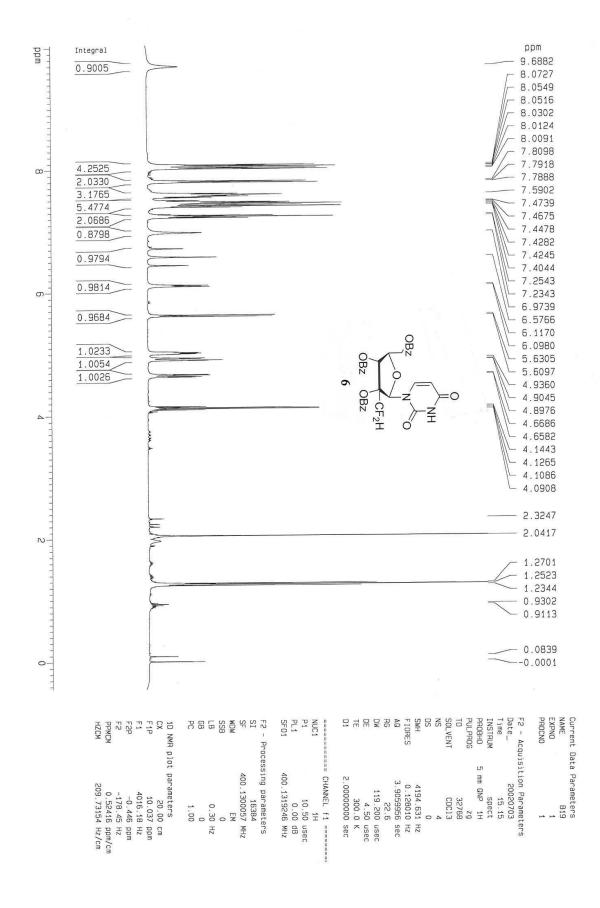


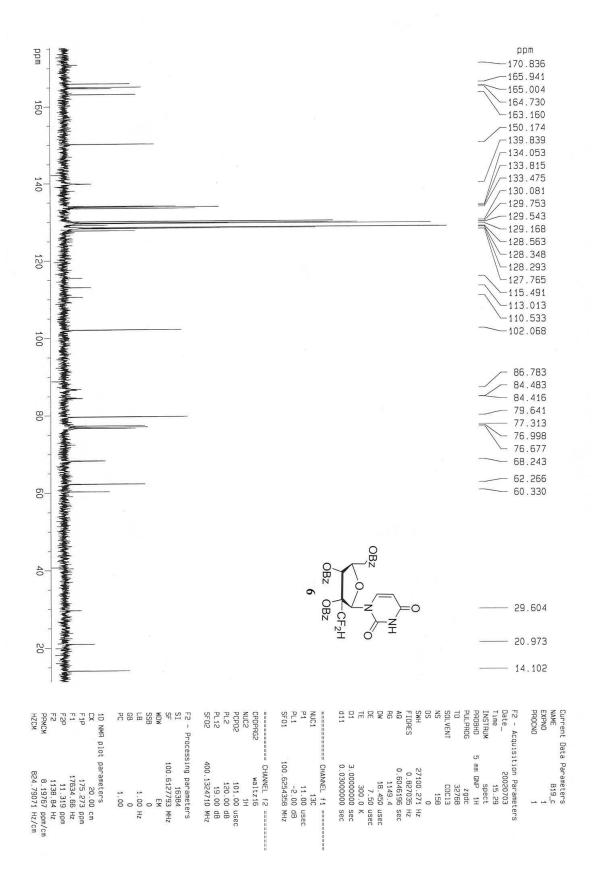


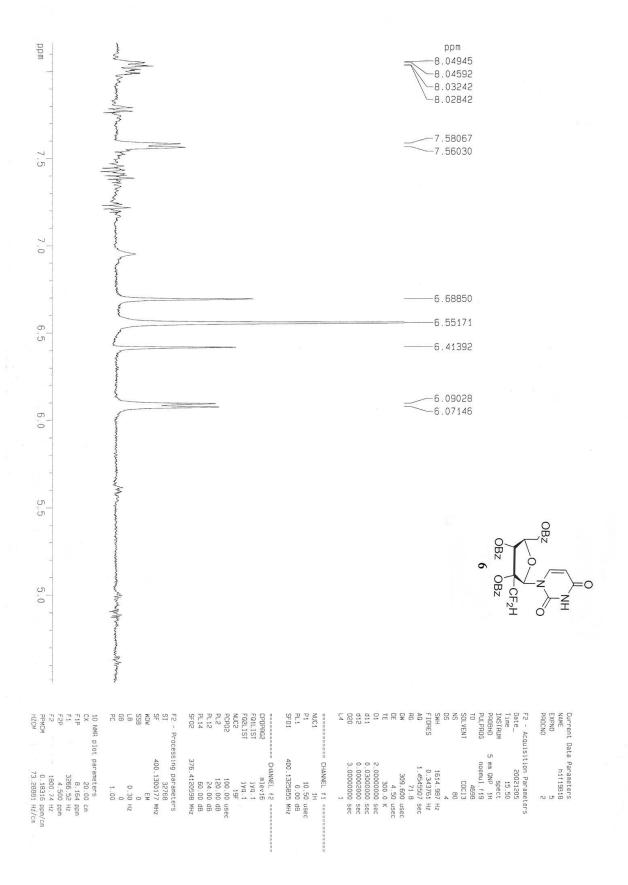


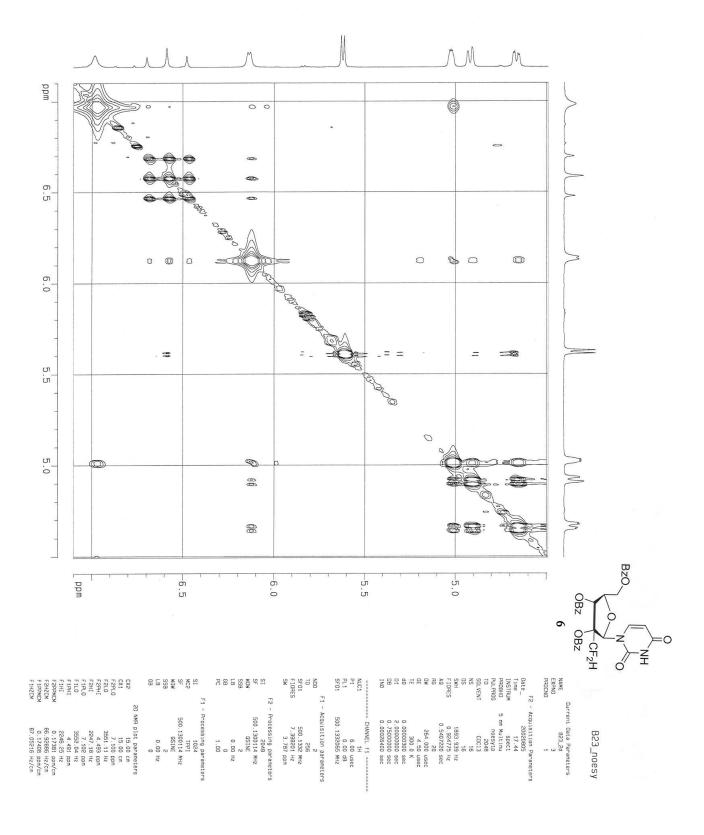


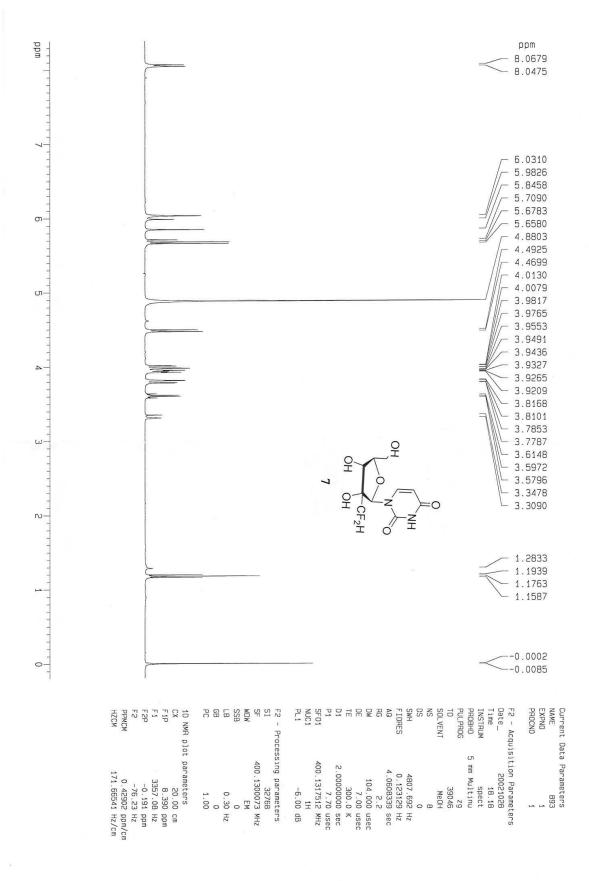


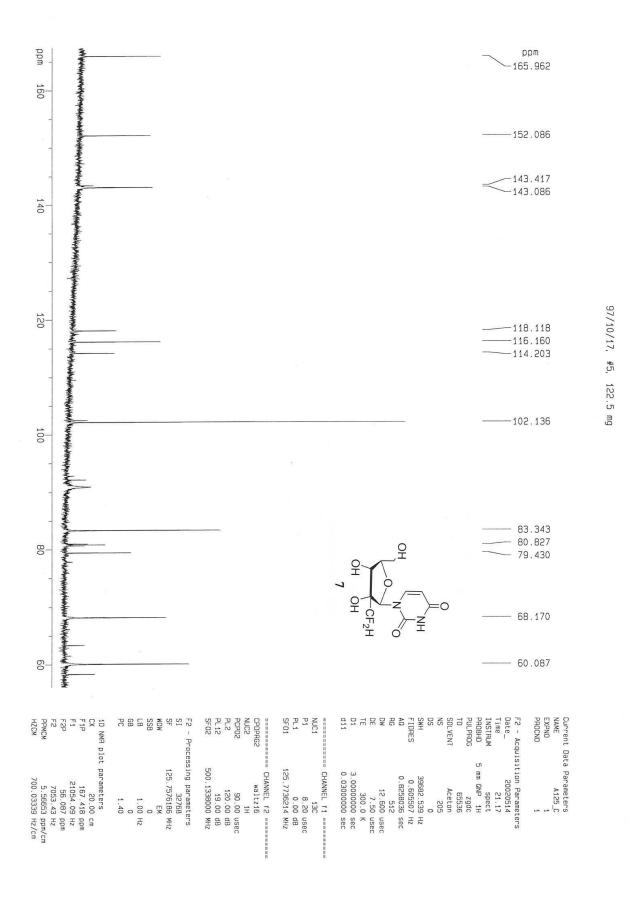


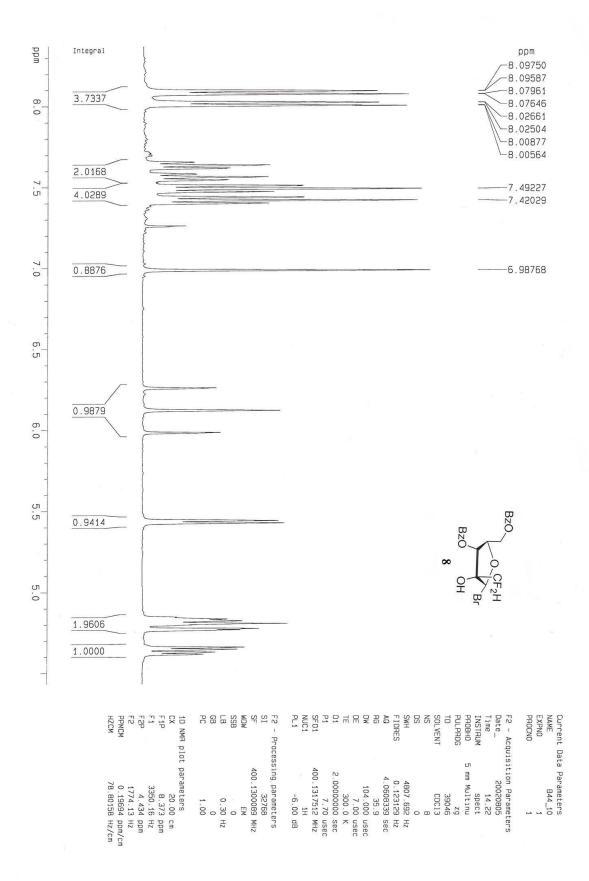


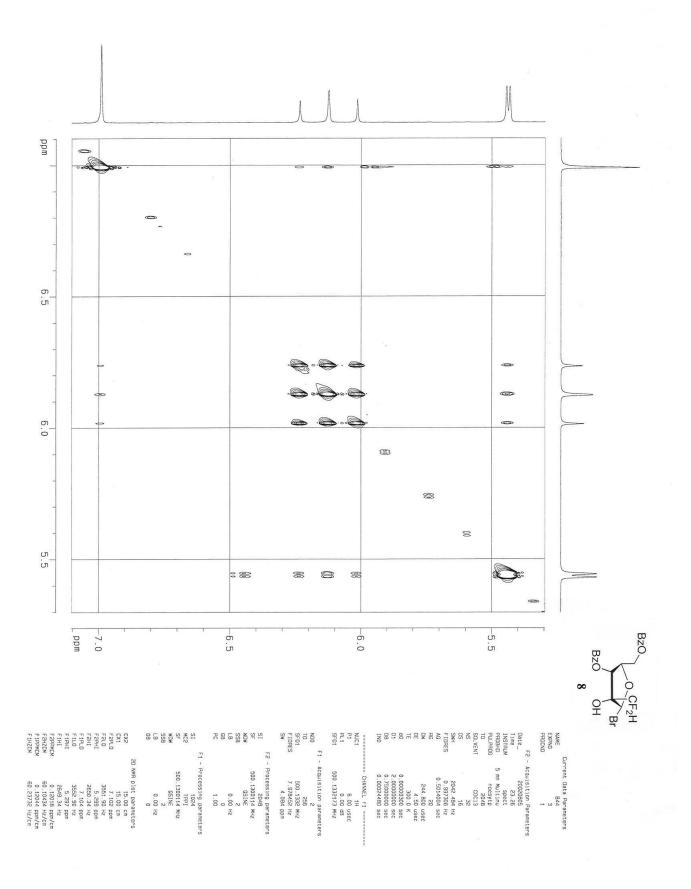


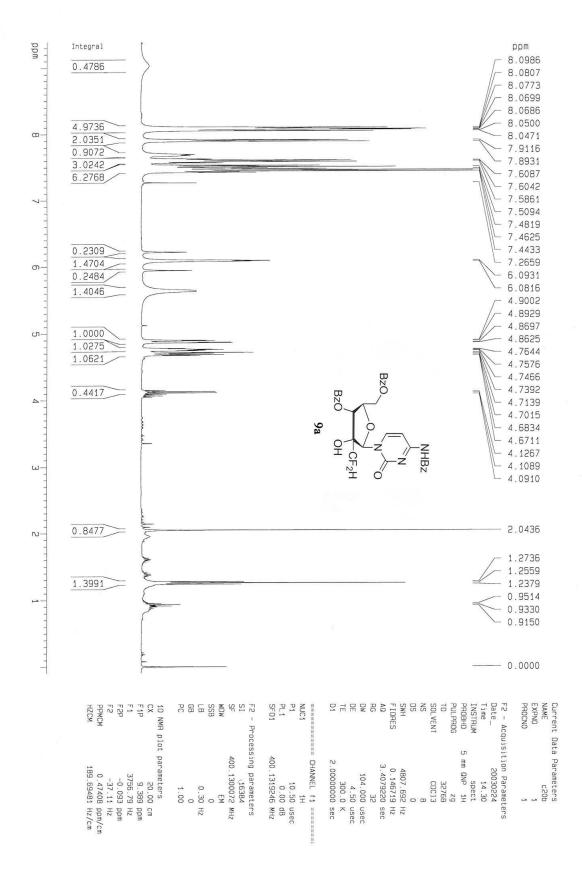


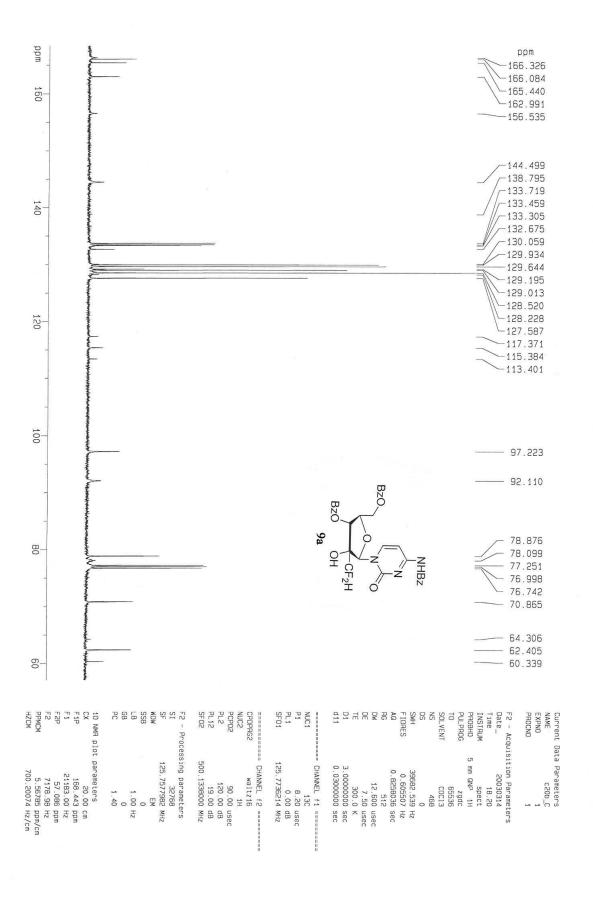








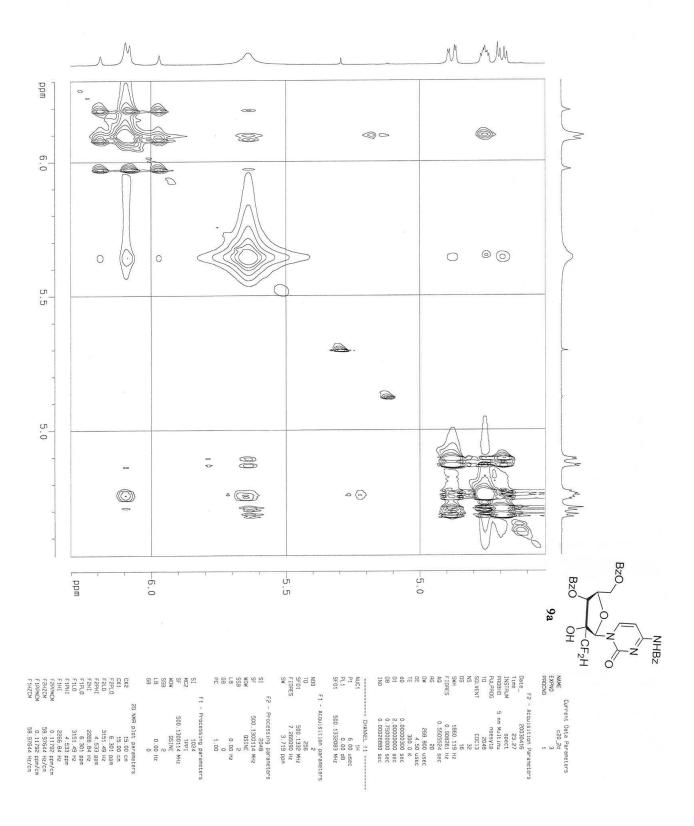


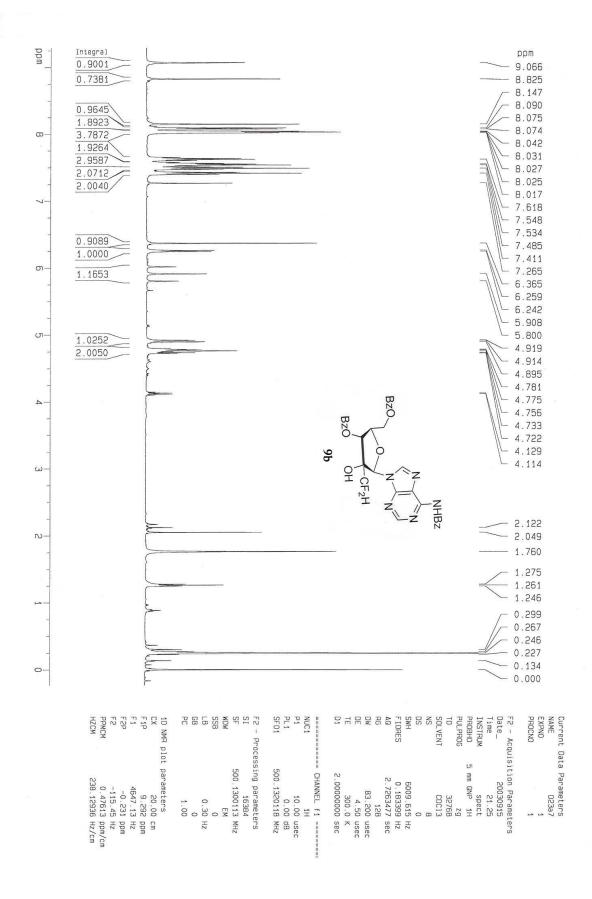


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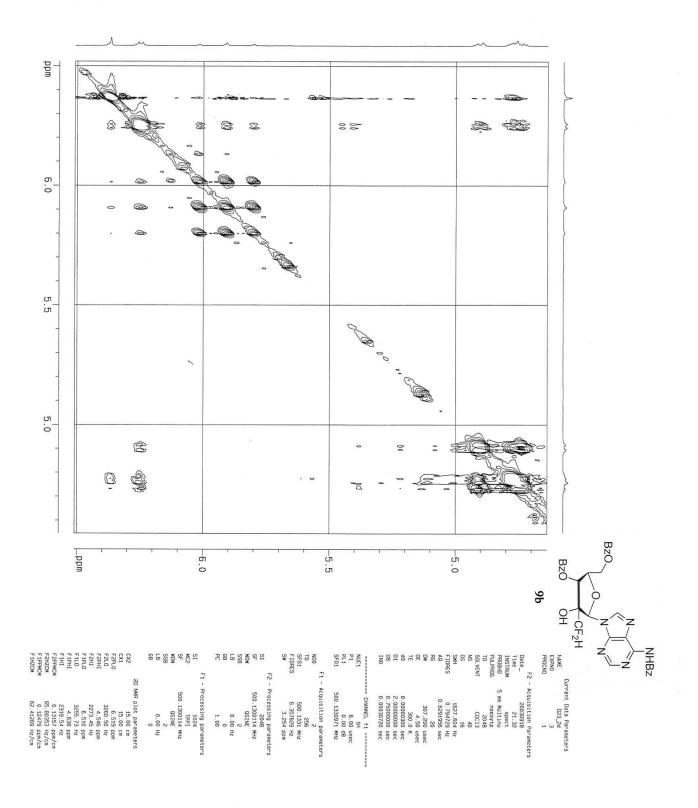
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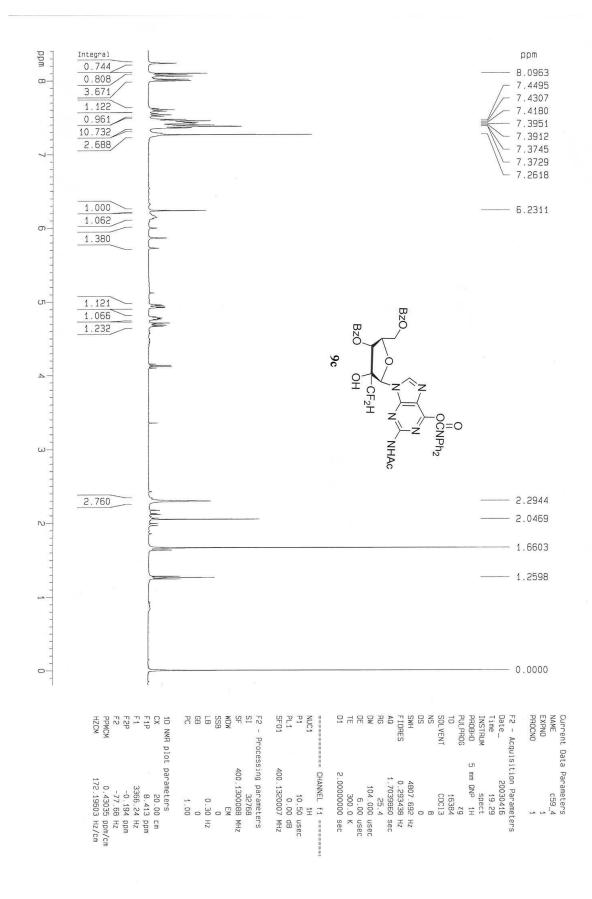
122.5 mg

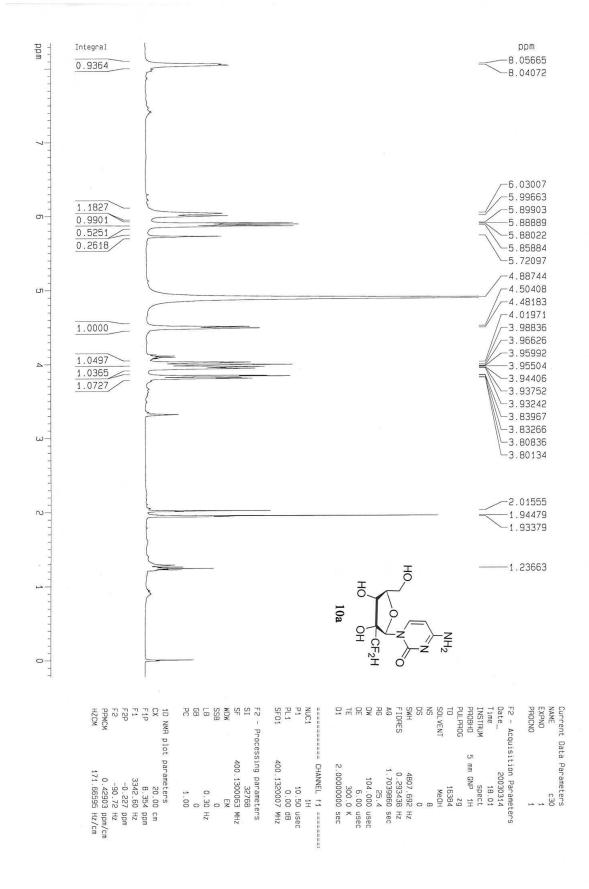


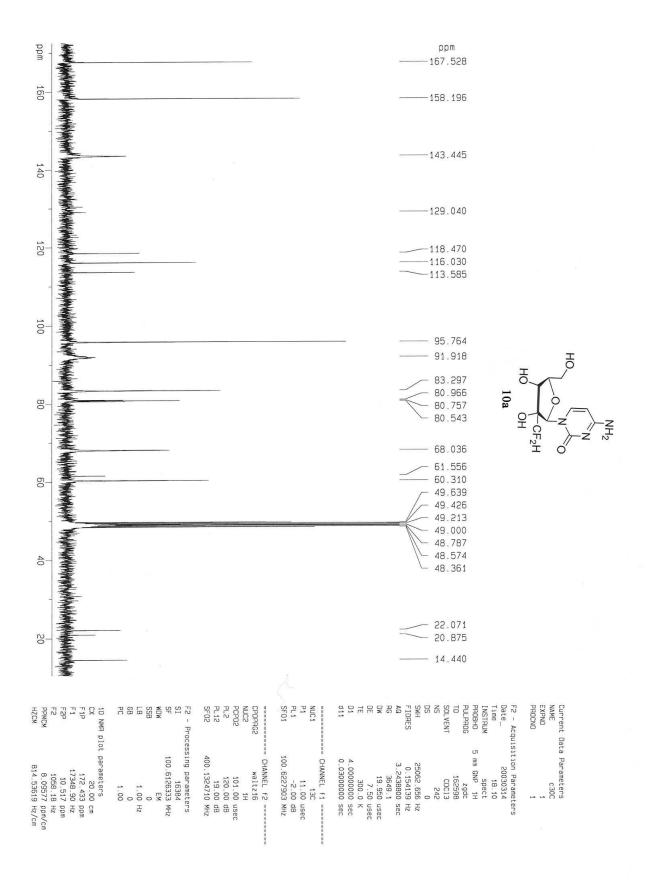


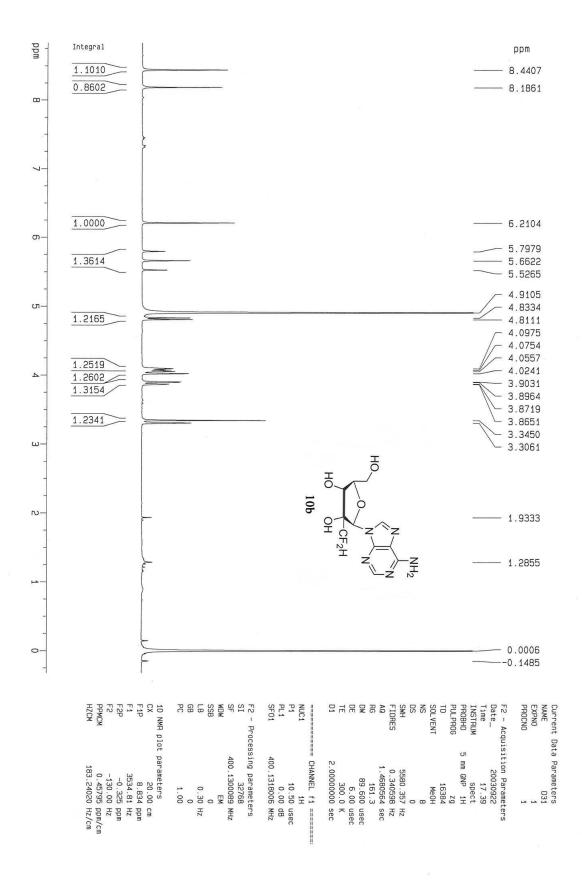
97/10/17, #5, 122.5 mg

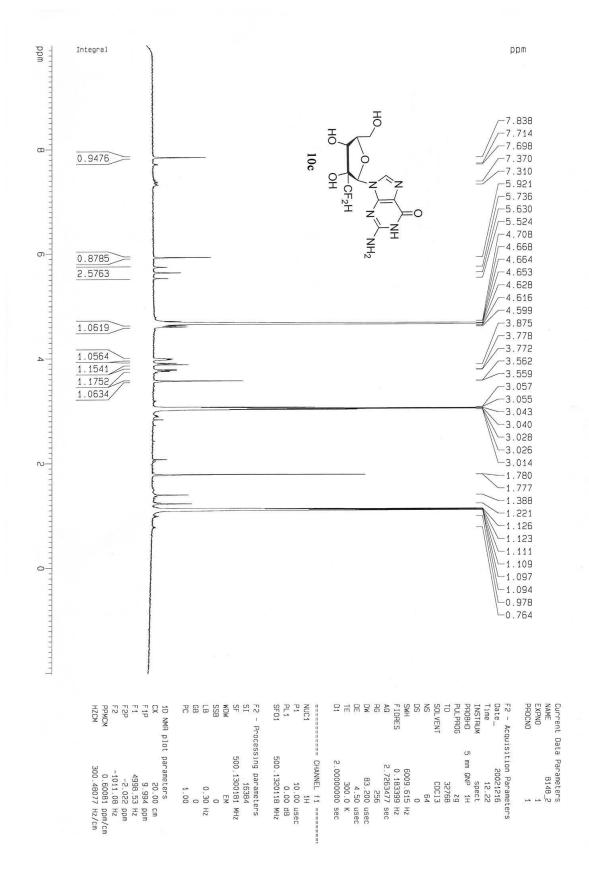


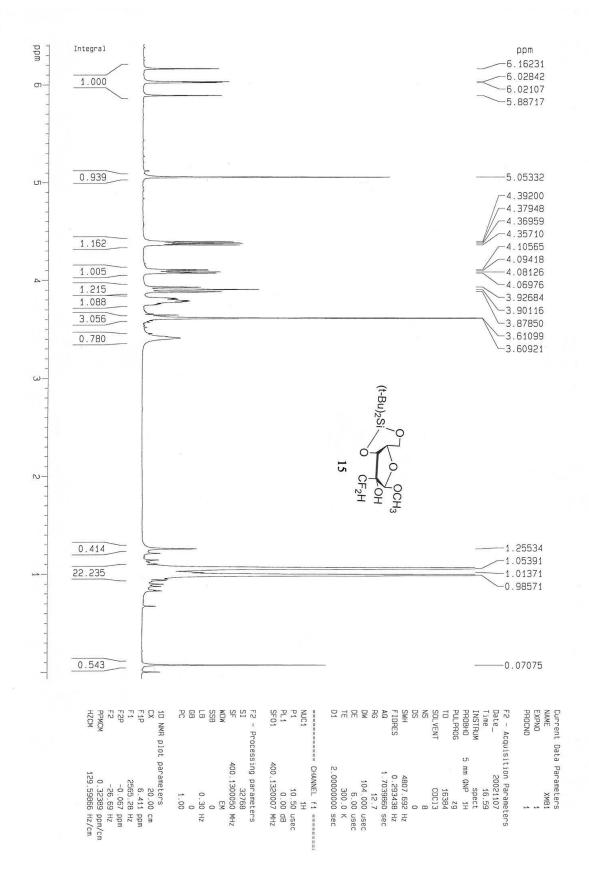


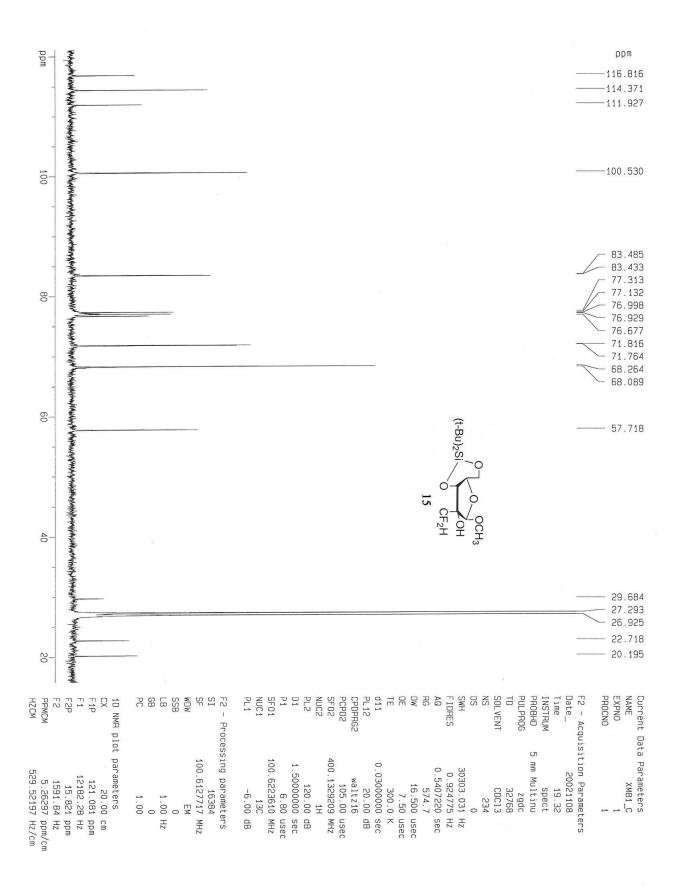


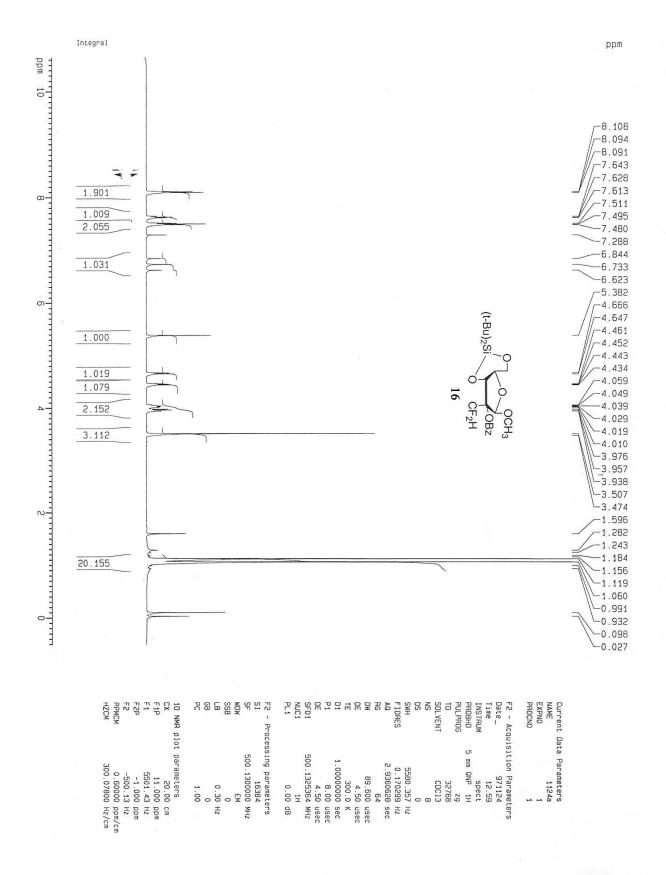


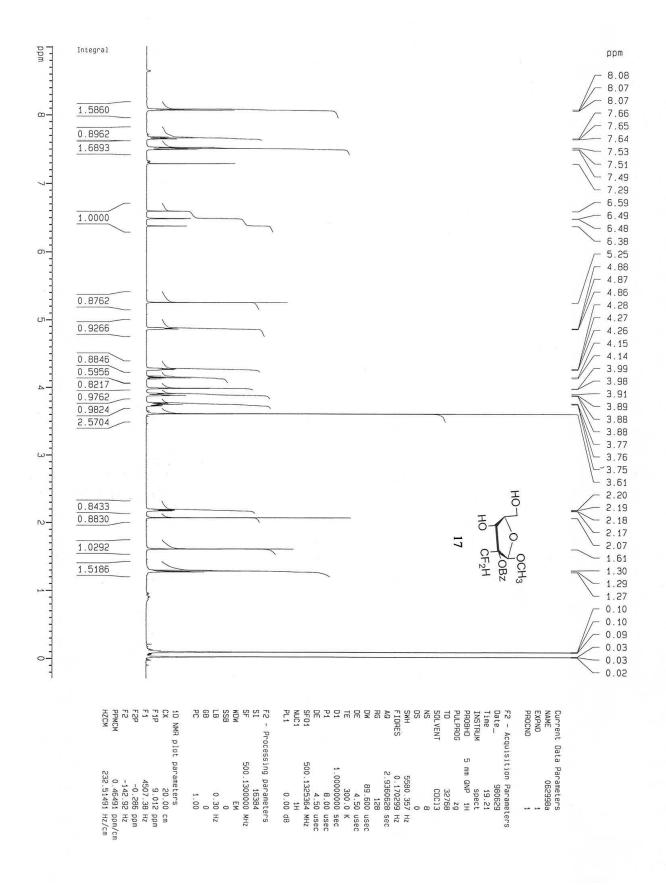


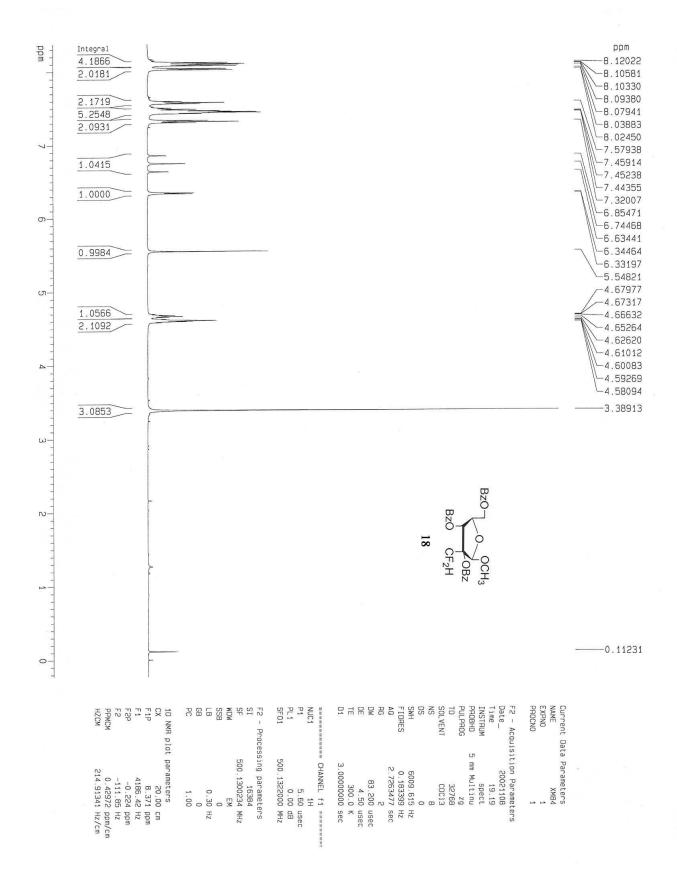


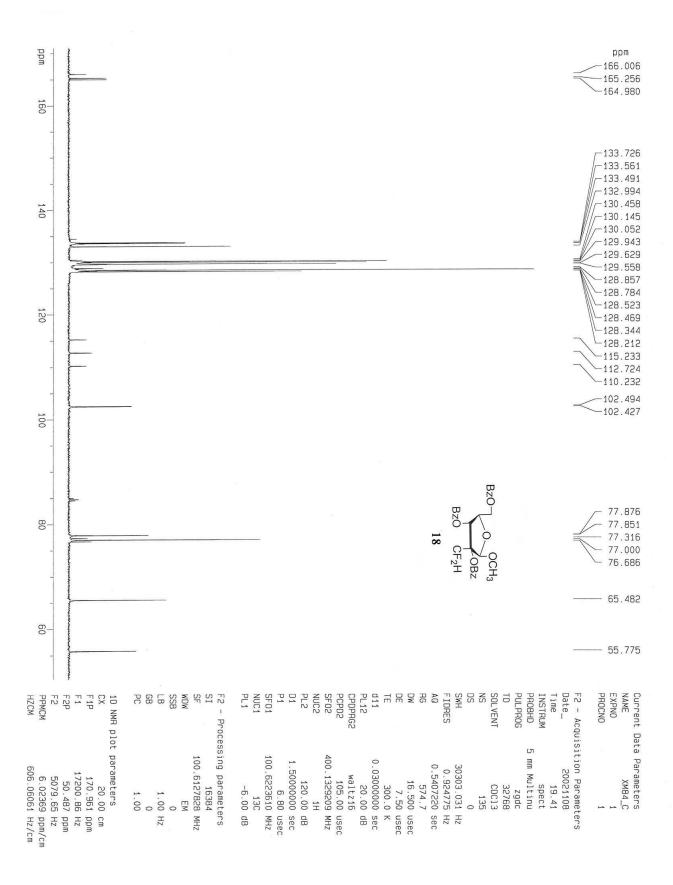


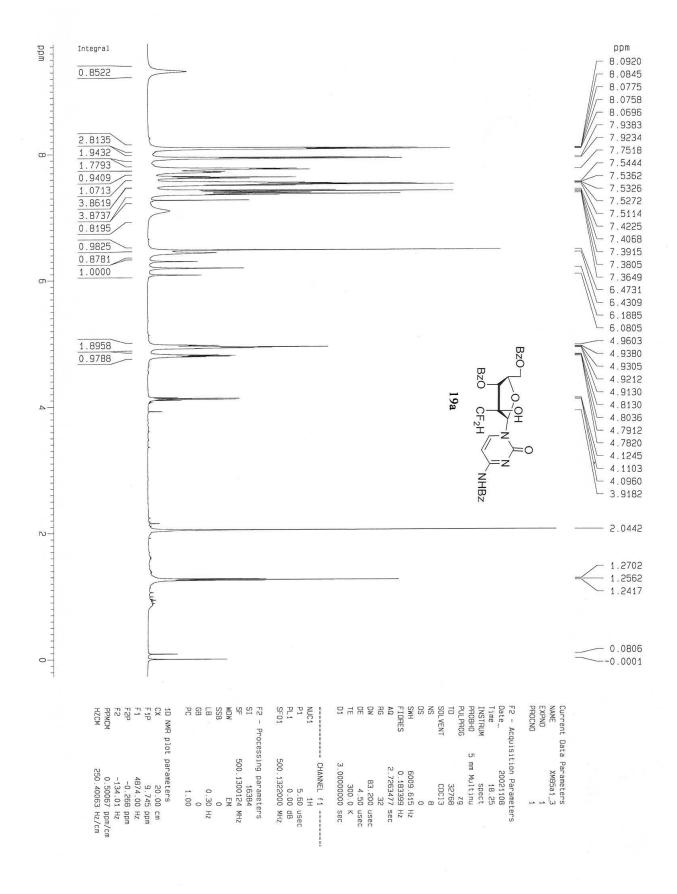


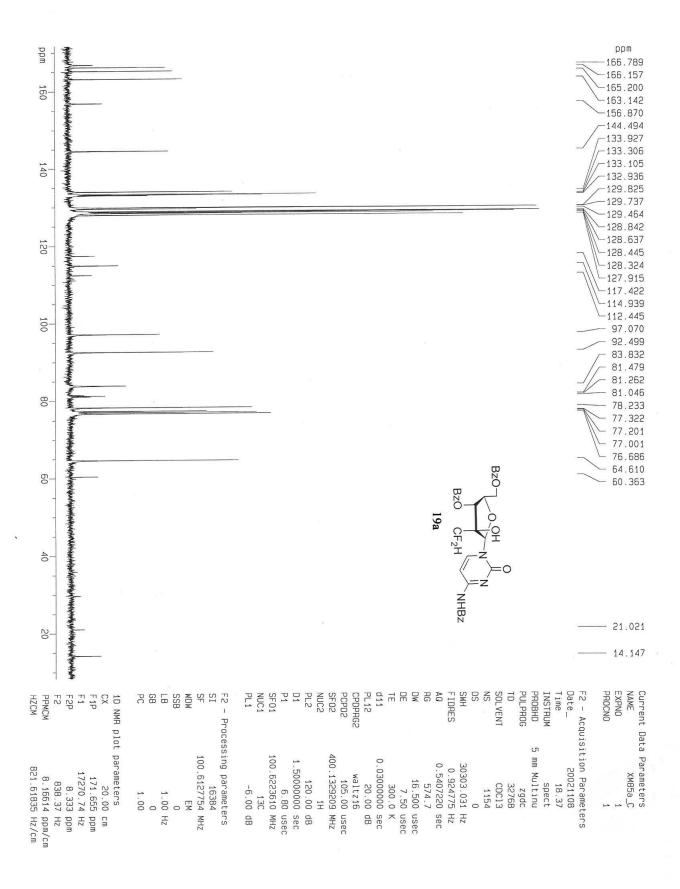


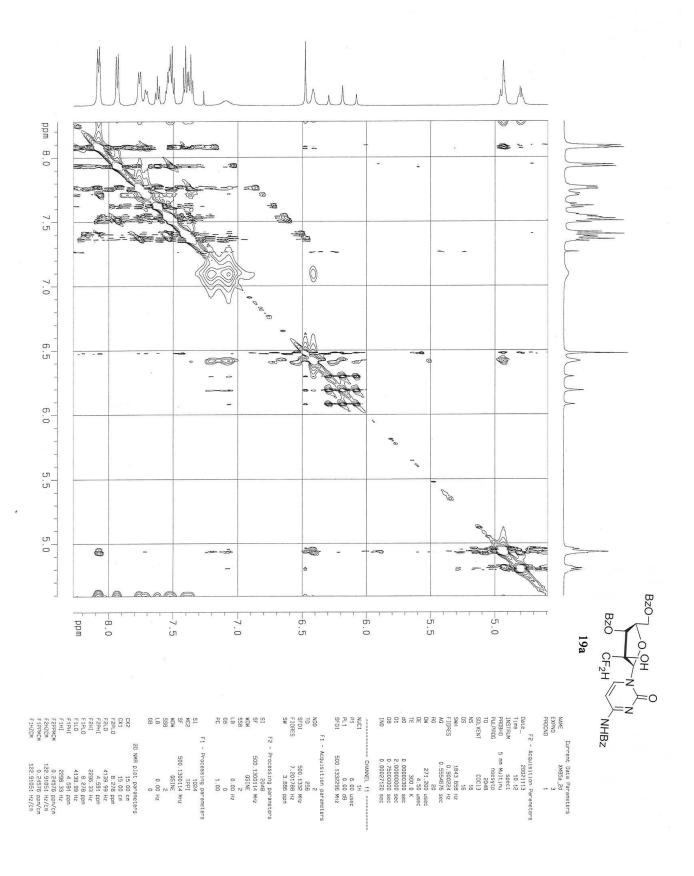


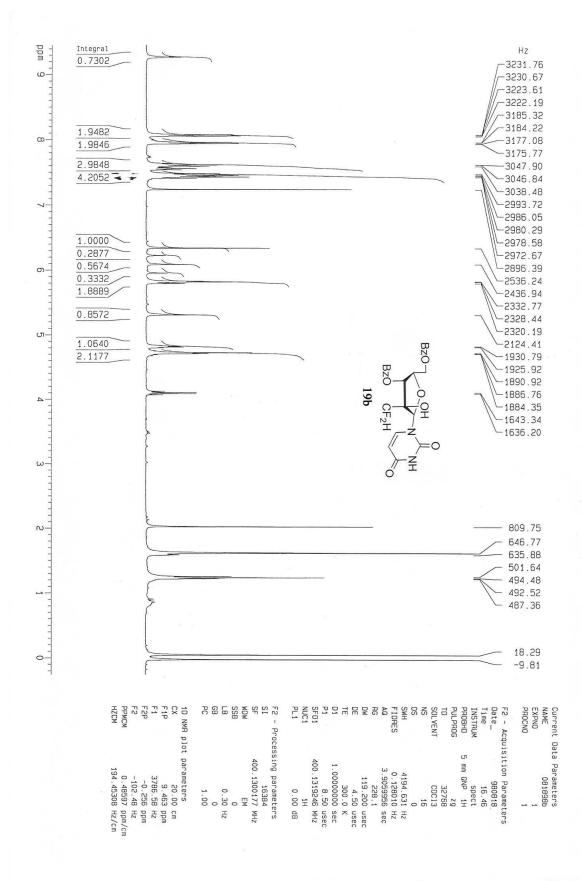


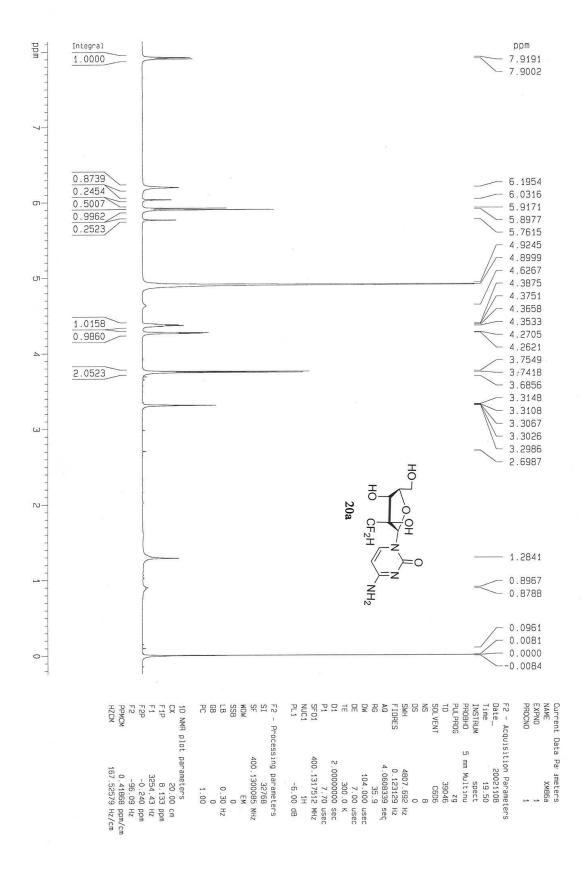


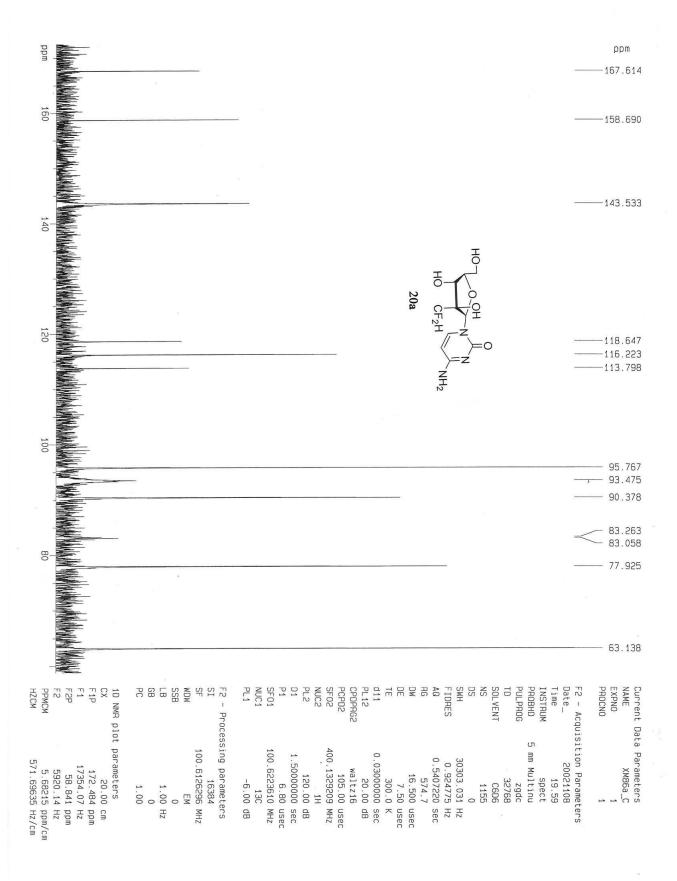


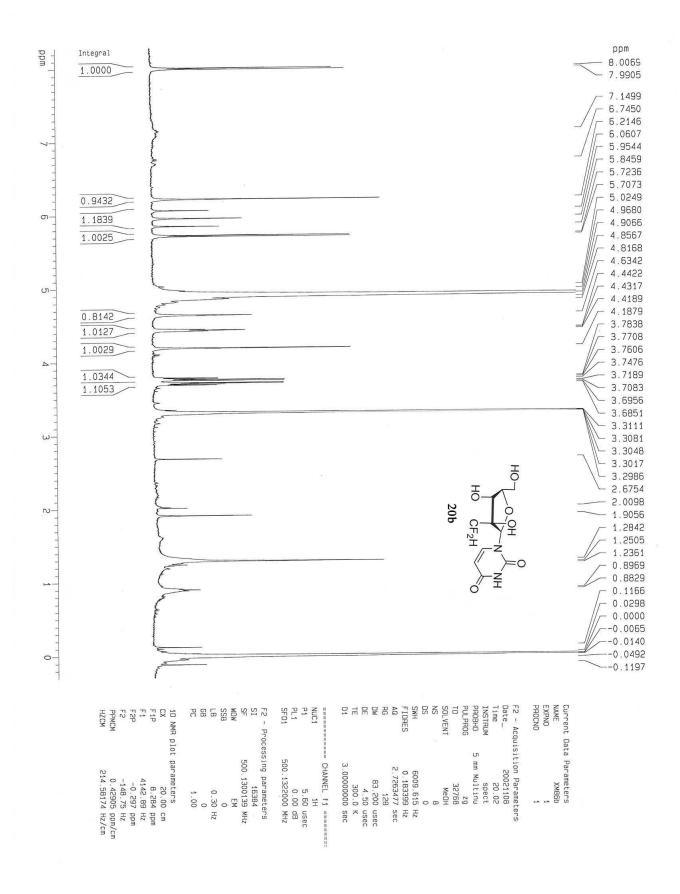


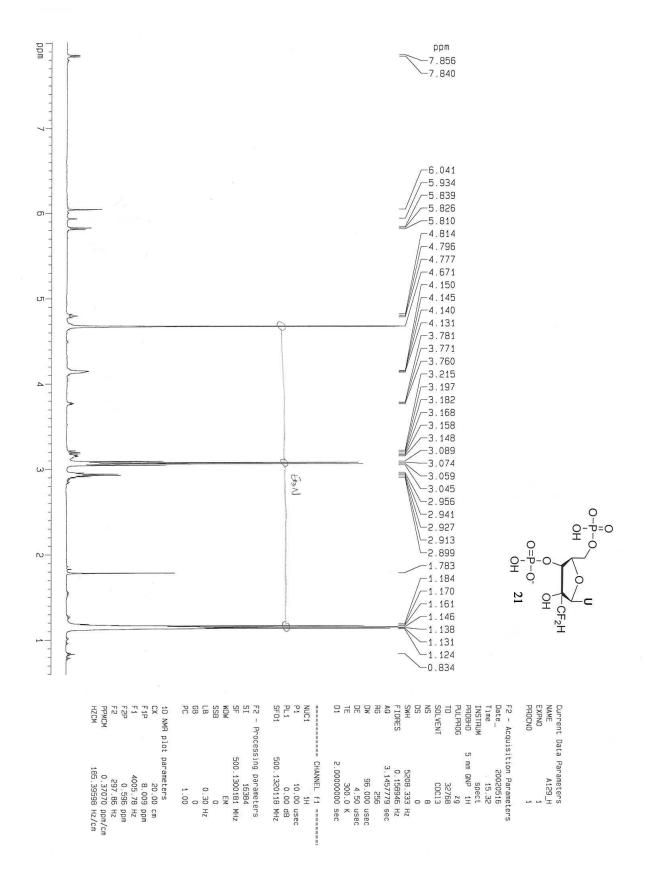


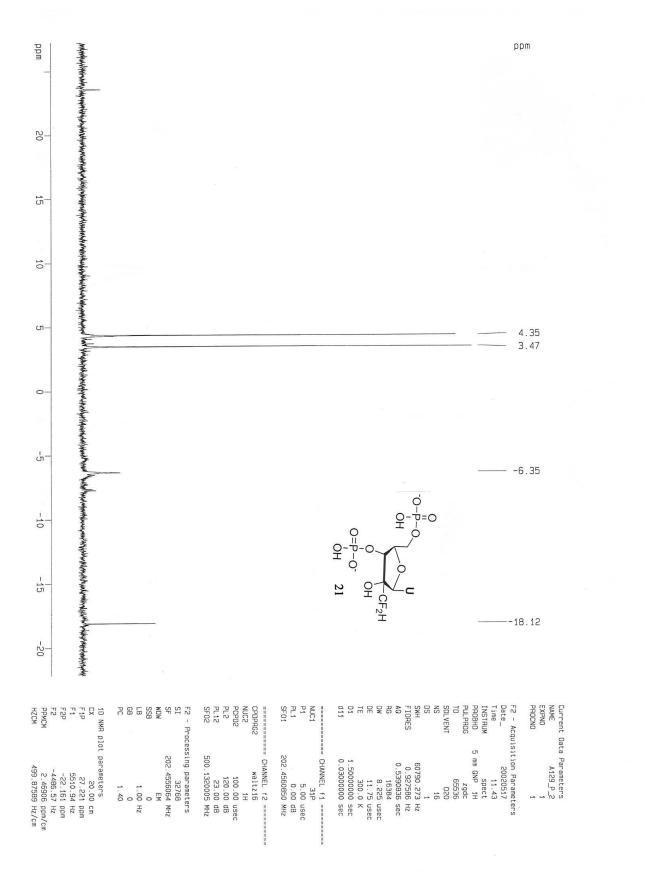






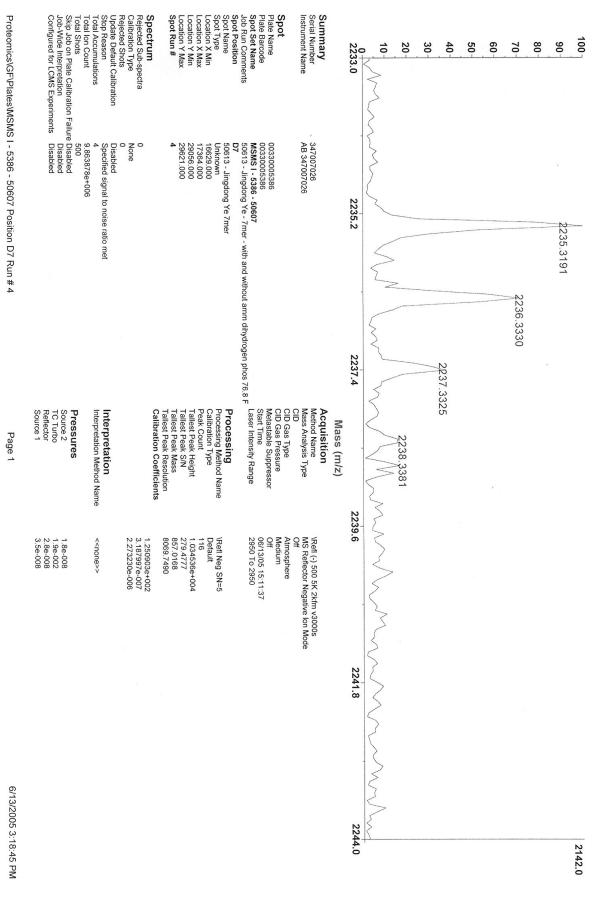






Spectrum Report

Final - Total Shots 500 - Position D7



P1 nuclease digestion. $*p(dN)_{12}rU_{2'CF2H}(dN)_{12}$ (or $*p(dN)_{12}rU(dN)_{12}$) was incubated in P1 nuclease buffer in the presence of 0.01 mg/ml P1 nuclease (USB) for 10 min at 37 °C. The mixture was then heated at 90 °C for 1 minute, combined with gel loading buffer (final concentration: 50 mM Tris-HCl, pH 8, 3.5 M urea and 0.0025% (w/v) each of bromophenol blue and xylene cyanol) and separated by 20% sequencing PAGE. Figure S1 shows the digestion pattern. The shorter P1 digestion products of the CF₂ oligonucleotide, arising from cleavage upstream of the modification (12 nucleotides and less), comigrate with their wild-type counterparts during PAGE. This is expected, as these products lack the modification and therefore are identical to those produced from wild-type oligonuleotide. The 12mer was assigned using the oligonucleotide from which the modified CF₂ oligonucleotide was generated (Figure S1, lane 5). We observe no product corresponding to P1 cleavage at the site bearing the difluoromethyl group (13mer), suggesting that the modification hinders P1 activity. The longer P1 digestion products of the CF2 oligonucleotide, arising from cleavage downstream of the modification (14 nucleotides and greater), migrate modestly slower than their wild-type counterparts, presumably due to the presence of the difluoromethyl modification. The species labeled as A and B in lanes 3 and 4 respectively, are present in the undigested controls (lanes 1 and 2, respectively) and most likely correspond to cleavage products arising from internal transphosphorylation at the ribose linkage.

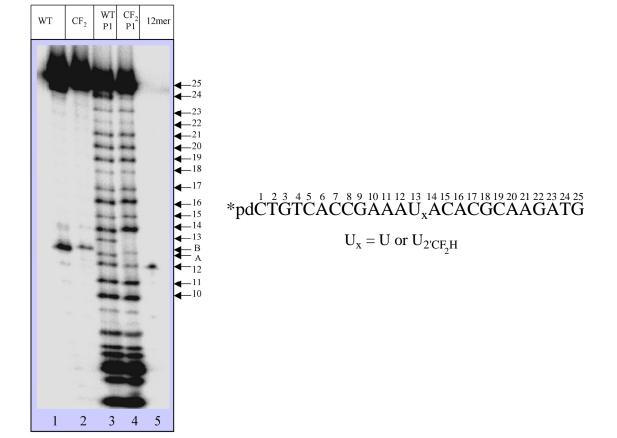


Figure S1. Nuclease P1 digestion of chimeric oligonucleotides containing uridine or 2'-C- β -difluoromethyluridine. Lane 1, WT, *pd(CTGTCACCGAAA)rUd(ACACGCAAGATG), input; Lane 2, CF₂, *pd(CTGTCACCGAAA)rU_{2'CF2H}d(ACACGCAAGATG), input; Lane 3, WT, with P1 nuclease; Lane 4, CF₂, with P1 nuclease; Lane 5, 12mer, *pd(CTGTCACCGAAA).