### **Efficient Access to Peptidyl-RNA Conjugates for Picomolar Inhibition of** Non-ribosomal FemX<sub>wv</sub> Aminoacyl Transferase

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Dedicated to Professor Michel Therisod on the occasion of his 65th birthday

Abstract: Peptidyl-RNA conjugates have various applications in studying the ribosome and enzymes participating in tRNA-dependent pathways such as Fem transferases in peptidoglycan synthesis. Herein a convergent synthesis of peptidyl-RNAs based on Huisgen-Sharpless cycloaddition for the final ligation step is developed. Azides and alkynes are introduced into tRNA and UDP-MurNAc-pentapeptide, respectively. Synthesis of 2'-azido RNA helix starts from 2'-azido-2'-deoxyadenosine that is coupled to deoxycytidine by phosphoramidite chemistry. The resulting dinucleotide is deprotected and ligated to a 22-nt RNA helix mimicking the acceptor arm of Ala-tRNAAla by T4 RNA ligase. For alkyne UDP-MurNAc-pentapeptide, meso-cystine is enzymatically incorporated into the peptidoglycan precursor and reduced, and L-Cys is converted to dehydroalanine with O-(mesitylenesulfonyl)hydroxylamine. Reaction of but-3-yne-1-

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### Introduction

Peptidyl-RNA conjugates have various applications in functional and structural studies of the ribosome and for improv-

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tapeptide. The CuI-catalyzed azide alkyne cycloaddition reaction in the presence of tris[(1-hydroxypropyl-1H-1,2,3-triazol-4-yl)methyl]amine provided the peptidyl-RNA conjugate, which was tested as an inhibitor of non-ribosomal  $Fem X_{Wv}$  aminoacyl transferase. The bi-substrate analogue was found to inhibit FemX<sub>wv</sub> with an IC<sub>50</sub> of  $(89 \pm$ 9) рм, as both moieties of the peptidyl-RNA conjugate contribute to high-affinity binding.

thiol with dehydroalanine affords the

alkyne-containing UDP-MurNAc-pen-

ing stability and cell permeability of short interfering RNAs.<sup>[1]</sup> Two strategies are currently explored for synthesis of peptidyl-RNA conjugates. In the divergent method, the peptide and oligonucleotide fragments are sequentially assembled by solid-phase synthesis on the same support. In the convergent method, the peptide and oligonucleotide fragments are independently synthesized prior to ligation. The first method is limited by the lack of a suitable solid support to assemble the peptide and oligonucleotide moieties and the lack of a uniform protection strategy suitable for any peptide sequence. A promising approach relies on polymerization of 3'-aminoacylamino-3'-deoxyadenosine linked to the solid support at the 2' position.<sup>[2]</sup> The modified adenosine provides the C-terminal extremity of the peptide chain and the 3' terminus of the oligonucleotides. Automated synthesis proceeds by Fmoc solid-phase synthesis followed by standard automated DMT/phosphoramidite synthesis. Successful protection of amino acid side chains is currently limited to Ser, Thr, Tyr, Asp and Glu.<sup>[3]</sup> To incorporate Arg in peptidyl-RNA conjugates, a convergent approach was developed which relies on a final native chemical ligation step.<sup>[4]</sup> This requires synthesis of a cysteinyl 3'-adenosine analogue resulting in the obligate presence of a cysteinyl residue at the C terminus of the peptide moiety of the conjugate. Here we develop a convergent synthetic route to peptidyl-RNA conjugates based on the Huisgen-Sharpless Cu<sup>I</sup>-

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catalyzed azide-alkyne cycloaddition reaction (Cu-AAC) and apply the method to the synthesis of a picomolar inhibitor of non-ribosomal FemX<sub>Wy</sub> aminoacyl transferase.

Aminoacyl-tRNAs participate in various biosynthetic pathways as donors of activated aminoacyl residues in addition to their role in protein synthesis by the ribosome.<sup>[5]</sup> Among non-ribosomal aminoacyl transferases, FemX<sub>Wv</sub> from Weissella viridescens<sup>[6]</sup> catalyzes transfer of Ala from Ala-tRNA<sup>Ala</sup> to the amino group of L-Lys at the 3-position of peptidoglycan precursors<sup>[7]</sup> (Figure 1A). Enzymes of the Fem family are attractive targets for the development of antibiotics active against resistant bacteria, since the residues incorporated by the enzymes are essential for the last crosslinking step of peptidoglycan polymerization in several important β-lactam-resistant pathogens such as staphylococci, pneumococci and streptococci.<sup>[8]</sup> Blocking Fem activity would therefore result in production of incomplete precursors acting as chain terminators that block formation of the essential stress-bearing peptidoglycan layer of the bacterial cell wall.

The alanyl-tRNA substrate of  $Fem X_{Wy}$  is produced by the alanyl-tRNA synthetase that aminoacylates the 3'-position of the terminal adenosine residue (A76) of tRNA<sup>Ala</sup> (Figure 1 A).<sup>[7]</sup> The resulting 3'-acylated tRNA binds to the acceptor site and delivers Ala to the ribosome. The same 3' Ala-tRNA<sup>Ala</sup> regioisomer binds to FemX<sub>wv</sub> with high affinity, although the enzyme catalyzes aminoacyl transfer only from the 2'-position following trans-esterification within the active site.<sup>[9]</sup> This enables  $\text{Fem}X_{Wv}$  to efficiently compete with the translation machinery for a common pool of AlatRNA<sup>Ala [9]</sup> The aminoacyl-transfer reaction catalyzed by FemX<sub>wv</sub> may proceed by a substrate-assisted catalytic mechanism involving direct attack of the amino group of the peptidoglycan precursor on the carbonyl group of Ala attached to the 2' position of the terminal ribose.<sup>[9]</sup> Our target (Figure 1B) is a bi-substrate composed of a peptidoglycan precursor analogue covalently linked to an RNA helix mimicking the tRNAAla acceptor arm. The convergent synthesis of the bi-substrate (Scheme 1) is based on a Cu-AAC reaction between alkyne UDP-MurNAc-pentapeptide 9 and 2'-azido RNA helix 11.

### **Results and Discussion**

For synthesis of 2'-azido RNA helix 11, we started from the 2'-azido-2'-deoxyadenosine 1 (Scheme 1).<sup>[9,10]</sup> Coupling of 1 with commercially available Ac-dC-PCNE afforded dinucleotide 2, which was phosphorylated and deprotected to obtain 3. An additional deprotection step afforded dinucleotide 4, which was ligated to 22-nt RNA helix 10 by T4 RNA ligase to obtain 24-nt azido RNA helix 11.

Alkyne UDP-MurNAc-pentapeptides 9a and 9b were obtained by semisynthesis (Scheme 1). Briefly, UDP-MurNAcpentapeptide analogue 6, which contained meso-cystine instead of L-Lys, was obtained enzymatically by sequential addition of L-Ala, D-Glu, meso-cystine and D-Ala-D-Ala to UDP-MurNAc by Mur synthetases MurC, D, E and F from

Escherichia coli, respectively.[11] The natural substrate of E. coli MurE, meso-diaminopimelic acid (meso-DAP), differs from meso-cystine only by replacement of a methylene by a disulfide group. This account for successful incorporation of meso-cystine instead of meso-DAP by MurE, albeit at the cost of 28-fold reduction in enzyme catalytic efficiency (data not shown). The mesocystine-containing UDP-MurNAc-pentapeptide 6 was reduced by dithiothreitol to afford L-cysteine-containing UDP-MurNAc-pentapeptide 7. Transformation of L-Cys of 7 into dehydroalanine of 8 was achieved with O-(mesitylenesulfonyl)hydroxylamine

(MSH) by using procedures proteins,<sup>[12]</sup> for developed which proved compatible with the peptidoglycan precursor. In the last step, reaction of but-3-yne-1-thiol with dehy-



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### CCGGGG NHAc NH<sub>2</sub> H VHB2 NHBz dC-PCNE NH<sub>2</sub> RÖ TBDMSO Ňз твомо 3 R = OTBDMS 1 2 11 С нo 4 R= OH R = UDPMurNAc NH =0 OH OH HOO HN g 13a (L-Cys) 13b (D-Cys) ΗÓ ΗÔ HC R 9a (L-Cys) 9b (D-Cys)

Scheme 1. Convergent synthesis of peptidyl-RNA conjugates 13a and 13b. a) 1. Tetrazole, CH<sub>2</sub>Cl<sub>2</sub>, RT, 1 h; 2. I<sub>2</sub>, 30 min, RT; 3. TCA, CH<sub>2</sub>Cl<sub>2</sub>, RT, 30 min; b) 1. bis(2-cyanoethyl)diisopropylphosphoramidite, tetrazole, CH<sub>2</sub>Cl<sub>2</sub>, RT, 1 h; 2. I<sub>2</sub>, RT, 30 min; 3. CH<sub>3</sub>NH<sub>2</sub>, RT, 24 h; c) HCl (6 m)/THF/CH<sub>3</sub>OH, 24 h; d) 22 nt RNA helix<sup>Ala</sup>, T4 RNA ligase; e) 1. L-Ala, MurC; D-Glu, MurD; 2. meso-Cystine, MurE; 3. D-Ala-D-Ala, MurF; f) DTT, Tris-HCl buffer, pH 8.0, 2 h, RT; g) MSH, DMF, 2 h, RT; h) (S)-But-3-ynyl ethanethioate, DMF, 2 h, 37°C; i) CuSO4, THPTA, Na ascorbate, 4°C, 24 h.

droalanine afforded UDP-MurNAc-pentapeptide analogues 9a and 9b, which contained the desired alkyne. Diastereoisomers 9a and 9b, which resulted from non-stereoselective thiol conjugate addition<sup>[13]</sup> to dehydroalanine 8, were purified by HPLC.

Optimization of the cycloaddition reaction<sup>[14]</sup> was performed with 24-nt azido RNA helix 11 and commercially available 1,8-nonadiyne, which afforded compound 12 (Figure 2A). The reaction conditions for this critical step were azido-RNA helix 11 (0.1 mм), nonadiyne (5 mм), copper sulfate (0.5 mm), sodium ascorbate (5 mm) and the tris[(1-hydroxypropyl-1*H*-1,2,3-triazol-4-yl)methyl]amine (THPTA) ligand (3.5 mm) in water for 24 h at 7°C. The THPTA ligand<sup>[15]</sup> was used to stabilize Cu<sup>I</sup> in aqueous buffer and avoid RNA degradation. These conditions provided the expected triazole analogue 12 in 70% yield, as detected by gel electrophoresis (Figure 2A) and mass spectrometry (calculated and observed masses of 7771.8 and 7772.1, respectively). The same approach was used to connect alkyne UDP-MurNAc-pentapeptides 9a and 9b (100 µM) to azido RNA helix 11 (50 µm; Scheme 1). The resulting bi-substrates 13a and 13b were purified by denaturing polyacrylamide gel electrophoresis (Figure 2A). These initial conditions were optimized by increasing the concentrations of UDP-MurNAc-pentapeptide 9a (20 mм) and RNA helix 11 (1 mM), which increased the yield from 4 to 36%.

Interaction of 13a with FemX<sub>Wy</sub> was analyzed by size-exclusion chromatography (Figure 3A). In the presence of a threefold molar excess of  $Fem X_{Wv}$  13a was quantitatively eluted as a complex. Highly purified bi-substrate 13a was recovered from the corresponding fractions following phenol extraction to remove the protein (Figure 3B) and analyzed by mass spectrometry (calculated and observed masses of 8828.6 and 8828.4, respectively).

The RNA helices were tested as inhibitors of  $Fem X_{Wv}$  by using a coupled assay<sup>[16]</sup> that relies on aminoacylation of tRNA<sup>Ala</sup> by aminoacyl-tRNA synthetase AlaRS and transfer of Ala from the resulting Ala-tRNAAla to UDP-MurNAcpentapeptide by  $\text{Fem}X_{Wv}$  (Figure 2B). RNA helix 10 inhibited Fem $X_{Wv}$  with an IC<sub>50</sub> of (89 ± 4)  $\mu$ M.<sup>[9]</sup> Low affinity is accounted for by the presence of vicinal hydroxyl groups in the terminal adenosine, which are also present in the





Figure 2. Inhibition of FemX<sub>Wv</sub>. Structure of tRNA analogues and analysis by denaturing PAGE (A).  $IC_{50}$  of the tRNA analogues (B).



Figure 3. Isolation of the complex containing **13a** and FemX<sub>Wv</sub>. Size-exclusion chromatography of the complex (A) and denaturing PAGE analysis of purified **13a** (B).

tRNA<sup>Ala</sup> product of the FemX<sub>wv</sub>-catalyzed reaction.<sup>[9]</sup> In agreement, substitution of 2'-hydroxyl by an azido group (helix **11**) led to a 56-fold decrease in the IC<sub>50</sub> ((1.6± 0.3) µM). Addition of nonadiyne to **11** by the Cu-AAC reaction (compound **12**) led to a further eightfold decrease in the IC<sub>50</sub> ((0.20±0.02) µM). Thus, the nonadiyne carbon chain may mimic the L-Lys side chain of UDP-MurNAcpentapeptide. Bi-substrate **13a** inhibited FemX<sub>wv</sub> with an



**General reagents and materials:** Solvents were dried by standard methods and distilled before use. Unless otherwise specified, materials were purchased from commercial suppliers and used without further purification. TLC: precoated silica gel thin layer sheets 60  $F_{254}$  (Merck). Flash chromatography: silica gel, 180–240 mesh (Merck). Optical rotations were measured on a PerkinElmer 341 digital polarimeter. Spectra were

 $IC_{50}$  of  $89\pm9\,$ рм, whereas diastereoisomer 13b inhibited FemX<sub>Wv</sub> with an IC<sub>50</sub> of  $2.3 \pm$ 0.2 nm. The L and D configurations of Cys were assigned to diastereoisomers 13a and 13b, respectively, based on the higher affinity of FemX<sub>wv</sub> for 13a and the presence of L-Lys at the analogous position of the natural substrate. Together, these results indicate that the RNA helix and UDP-MurNAc-peptide moieties of bi-substrate 13a both contributed to high-affinity binding to FemX<sub>wy</sub>.

### Conclusion

We have developed a convergent method to synthesize peptidyl-RNA conjugates based on Huisgen-Sharpless cycloaddition for the final ligation step. The reaction conditions reported here are fully compatible with two types of biomolecules, RNA and the peptidoglycan precursor. We also show that introduction of dehydroalanine as electrophilic site by using the MSH reagent is compatible with the peptidoglycan precursor containing nucleotide (UDP), sugar (MurNAc) and peptide (L-Alaγ-D-Glu-L-Cys-D-Ala-D-Ala) moieties. Finally, association of both partners of the enzymatic acyl-transfer reaction in peptidyl-RNA 13a induces an 18000-fold increase in  $\text{Fem}X_{W_{V}}$ inhibition efficiency and paves the way for the design of antibacterial agents acting on this new target.

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#### recorded on Bruker AVANCE 400 spectrometer for <sup>1</sup>H (400 MHz), <sup>13</sup>C (101 MHz) and <sup>31</sup>P (122 or 162 MHz) in CDCl<sub>3</sub>, D<sub>2</sub>O or DMSO. Chemicals shifts are expressed in parts per million (ppm) relative to residual CHCl<sub>3</sub> ( $\delta$ = 7.26) or HDO ( $\delta$ = 4.79) for <sup>1</sup>H and CDCl<sub>3</sub> ( $\delta$ = 77.16) for <sup>13</sup>C as internal references. Signals were assigned on the basis of DEPT 135, COSY, HSQC and HMBC. High-resolution mass spectra were carried recorded on a Bruker micrOTOF spectrometer. Unless otherwise specified, HPLC was performed on a HPLC system with C18 reverse-phase columns (analytical column, 250×4.6 mm, HYPERSIL-100 C18; semipreparative column, 250×21.2 mm, HYPERSIL HS C18; Thermoelectron Corporation). The solvent system consisted of 50 mm aqueous NH<sub>4</sub>OAc:CH<sub>3</sub>CN with a linear gradient (100:0 to 67:33) applied from 5 to 50 min at a flow rate of 1 (analytical) or 10 mLmin<sup>-1</sup> (semipreparative) and UV detection at 260 nm. For separation of UDP-MurNAc-peptides, fast protein liquid chromatography (FPLC) was performed on an Äkta purifier system (Amersham Pharmacia Biotech) with an analytical C18 reverse-phase column (Nucleosil 100-3 C18, 250×4.6 mm, Macherey-Nagel). The solvent system consisted of 50 mm aqueous NH<sub>4</sub>OAc/CH<sub>3</sub>CN with a linear gradient (100:0 to 95:5) applied from 7 to 37 min at a flow rate of 1 mLmin<sup>-1</sup> and UV detection at 260 nm. Analytical denaturing polyacrylamide gel electrophoresis (PAGE) was performed in gels (20× 20×0.1 cm) containing acrylamide (13%, w/v), bis-acrylamide (4.4%, w/ v), urea (8M), tetramethylethylenediamine (TEMED, 0.04% v/v), ammonium persulfate (0.08% w/v) and 1× TBE buffer (pH 8.2) [Tris (89 mm), borate (89 mm), EDTA (2 mm)]. Gels were loaded (lane width 6 mm) with RNA helices (100 pmol) in a volume of 10 µL containing bromophenol blue (0.01 % w/v) and glycerol (25 % v/v). Electrophoresis was performed for 120 min at 600 V, and gels stained with ethidium bromide (0.5 µg mL<sup>-1</sup>) were imaged with a Herolab E.A.S.Y 429K camera. Preparative denaturing PAGE was performed under the same conditions, except that 1 nmol of RNA helices was loaded per lane and the concentration of ethidium bromide used for staining was reduced to 0.25 µgmL<sup>-1</sup>. RNA bands were cut off from gels with a sterile scalpel and electroeluted for 2 h at 100 V in dialysis bags (cut-off 1 kDa; Spectra/Por 7 Dialysis Membrane, Spectrum Labs). RNA helices were concentrated to a final volume of 250 µL by ultrafiltration (Microcon, cut-off 3 kDa, Millipore), purified by gel filtration (Sephadex G-75 column 10/ 300 GL, GE Healthcare) equilibrated in Tris-HCl buffer (25 mm, pH 7.5) containing NaCl (100 mм) and MgCl<sub>2</sub> (5 mм).

# $(2R,3S,5R)\mbox{-}5\mbox{-}[4\mbox{-}Acetamido\mbox{-}2\mbox{-}oxopyrimidin\mbox{-}1(2H)\mbox{-}yl]\mbox{-}2\mbox{-}(hydroxymethyl)tetrahydrofuran\mbox{-}3\mbox{-}yl[(2R,3S,4R,5R)\mbox{-}4\mbox{-}azido\mbox{-}5\mbox{-}(6\mbox{-}benzamido\mbox{-}9H\mbox{-}hyl)tetrahydrofuran\mbox{-}3\mbox{-}yl[(2R,3S,4R,5R)\mbox{-}4\mbox{-}azido\mbox{-}5\mbox{-}(6\mbox{-}benzamido\mbox{-}9H\mbox{-}hyl)tetrahydrofuran\mbox{-}3\mbox{-}yl[(2R,3S,4R,5R)\mbox{-}4\mbox{-}azido\mbox{-}5\mbox{-}(6\mbox{-}benzamido\mbox{-}9H\mbox{-}hyl)tetrahydrofuran\mbox{-}3\mbox{-}yl[(2R,3S,4R,5R)\mbox{-}4\mbox{-}azido\mbox{-}5\mbox{-}(6\mbox{-}benzamido\mbox{-}9H\mbox{-}hyl)tetrahydrofuran\mbox{-}3\mbox{-}yl[(2R,3S,4R,5R)\mbox{-}4\mbox{-}azido\mbox{-}5\mbox{-}(6\mbox{-}benzamido\mbox{-}9H\mbox{-}hyl)tetrahydrofuran\mbox{-}3\mbox{-}yl[(2R,3S,4R,5R)\mbox{-}4\mbox{-}azido\mbox{-}5\mbox{-}(6\mbox{-}benzamido\mbox{-}9H\mbox{-}hyl)tetrahydrofuran\mbox{-}3\mbox{-}yl[(2R,3S,4R,5R)\mbox{-}4\mbox{-}azido\mbox{-}5\mbox{-}(6\mbox{-}benzamido\mbox{-}9H\mbox{-}hyl)tetrahydrofuran\mbox{-}3\mbox{-}yl[(2R,3S,4R,5R)\mbox{-}4\mbox{-}4\mbox{-}2\mbox{-}4\mbox{-}4\mbox{-}4\mbox{-}4\mbox{-}3\mbox{-}4\mbox{-}3\mbox{-}4\mbox{$

purin-9-yl)-3-(tert-butyldimethylsilyloxy)tetrahydrofuran-2-yl]methyl 2cyanoethyl phosphate (2): A 0.45 M solution of tetrazole in MeCN (2.9 mL, 1.29 mmol) and Ac-dC-PCNE (250 mg, 0.324 mmol) were added to a solution of  $\mathbf{1}^{[9,10]}$  (66 mg, 0.130 mmol) in CH<sub>2</sub>Cl<sub>2</sub> (700 µL) at room temperature under argon atmosphere. After stirring for 1 h at room temperature, 0.1 m iodine in THF:H2O:pyridine (75:2:20, 3.3 mL) was added. After 30 min, the reaction mixture was diluted with EtOAc (10 mL), washed with water (10 mL), aqueous saturated Na<sub>2</sub>S<sub>2</sub>O<sub>3</sub> solution (10 mL) and brine (10 mL), dried over anhydrous MgSO<sub>4</sub>, filtered and concentrated at reduced pressure. The residue was then stirred with 0.18 M trichloroacetic acid solution in dichloromethane (7.2 mL) at room temperature for 30 min. The reaction mixture was diluted with CH2Cl2 (5 mL) and the organic phase was washed with water (5 mL), aqueous saturated NaHCO3 solution at 0°C (5 mL) and brine (5 mL), dried over anhydrous MgSO4, filtered and concentrated. The crude product was purified by preparative TLC with CH<sub>2</sub>Cl<sub>2</sub>:MeOH (90:10) as eluent to give 2 (two diastereoisomers, ratio 1:1, 95 mg, 82 %). <sup>1</sup>H NMR (400 MHz, CDCl<sub>3</sub>):  $\delta =$ 9.44 (br, 1H, NHAc), 8.75 (s, 1H, H2Ad or H8Ad), 8.31 (s, 1H, H2Ad or H8<sup>Ad</sup>), 8.16 (dd, J = 7.5, 1.9 Hz, 1 H, H6<sup>Cyt</sup>), 8.02 (d, J = 1.3 Hz, 2 H, Bz), 7.58 (ddd, J=7.2, 5.1, 1.6 Hz, 1 H, Bz), 7.49 (ddd, J=8.3, 6.9, 2.5 Hz, 2 H, Bz), 7.31 (dd, J=7.5, 2.8 Hz, 1 H, H5<sup>Cyt</sup>), 6.15-6.05 (m, 2 H, H1'<sup>Ad</sup>, H1'<sup>Cyt</sup>), 5.07 (ddd, J=8.6, 5.9, 2.6 Hz, 1H, H3'Cyt), 4.97-4.93 (m, 1H, H3'Ad), 4.80-4.75 (m, 1H,  $H2'^{Ad}$ ), 4.27–4.15 (m, 5H,  $OCH_2$ ,  $H4'^{Ad}$ ,  $H5'^{Cyt}$ ), 3.77 (dd, J=8.7, 2.7 Hz, 2H, H5'<sup>Ad</sup>), 3.47 (s, 3H, Me<sup>Ac</sup>), 2.76–2.71 (m, 2H, CH2CN), 2.68-2.57 (m, 1H, H2'aCyt), 2.36-2.29 (m, 1H, H2'bCyt), 0.97 (s, 9H, tBu<sup>TBS</sup>), 0.21 (s, 3H, Me<sup>TBS</sup>), 0.20 ppm (s, 3H, Me<sup>TBS</sup>); <sup>13</sup>C NMR (101 MHz, CDCl<sub>3</sub>):  $\delta = 170.7$  (C=O<sup>Bz</sup>), 162.6 (C=O<sup>Bz</sup>), 155.5 (C2<sup>Ad</sup> or C8<sup>Ad</sup>), 145.3 (C6<sup>Cyt</sup>), 142.5, (C2<sup>Ad</sup> or C8<sup>Ad</sup>), 133.0, 128.9, 128.3 (Bz), 116.5 (Cq), 96.8 (C5<sup>Cyt</sup>), 87.8 (C1<sup>'Ad</sup>), 87.3 (C1<sup>'Cyt</sup>), 77.5 (C3<sup>'Cyt</sup>), 71.9 (C3<sup>'Ad</sup>), 62.6 (C2<sup>'Ad</sup>), 61.5 (C5<sup>'Cyt</sup>), 61.3 (C5<sup>'Ad</sup>), 39.7 (C2<sup>'Cyt</sup>), 25.8 (*t*Bu<sup>TBS</sup>), 25.0 (CH<sub>3</sub><sup>Ac</sup>), 18.5 (CH<sub>2</sub>CN), 18.1 (Cq<sup>TBS</sup>), -4.5, -4.8 ppm (2 × Me<sup>TBS</sup>); <sup>31</sup>P NMR (<sup>1</sup>H-decoupled, 122 MHz, DMSO):  $\delta$  = -2.60, -2.66 (2s); ESI<sup>+</sup> MS: *m*/z calcd for C<sub>37</sub>H<sub>47</sub>N<sub>12</sub>O<sub>11</sub>PNaSi<sup>+</sup> [*M*+Na]<sup>+</sup>: 917.2886; found: 917.3253.

[(2R,3S,5R)-5-[4-Amino-2-oxopyrimidin-1(2H)-yl]-3-({[(2R,3S,4R,5R)-5-(6-amino-9H-purin-9-yl)-4-azido-3-(tert-butyldimethylsilyloxy)tetrahydrofuran-2-yl]methoxy}oxidophosphoryloxy)tetrahydrofuran-2-yl]methyl hydrogenphosphate (3): Bis(2-cyanoethyl)diisopropylphosphoramidite (88 mg, 320 µmol, 2.5 equiv) was added to phosphotriester 2 (115 mg, 130 µmol). Ultradry CH2Cl2 (400 µL) was added, followed by tetrazole in CH<sub>3</sub>CN (2.8 mL, 1.3 mmol). The mixture was stirred at room temperature for 1 h, and I<sub>2</sub> in THF:H<sub>2</sub>O:pyridine, (75:2:20, 3.2 mL, 320 µmol) was added. After being stirred at room temperature for 30 min, the mixture was diluted with EtOAc (5 mL) and washed successively with saturated aqueous Na<sub>2</sub>S<sub>2</sub>O<sub>3</sub> solution (5 mL) and brine (5 mL). The organic layer was dried over anhydrous MgSO4 and concentrated to dryness. The crude intermediate was dissolved in 15 mL of aqueous 5м MeNH<sub>2</sub>. The reaction was stirred for 12 h at room temperature and concentrated under reduced pressure. The ammonium salt of dinucleotide 3 was purified by HPLC, lyophilized and recovered as a white solid (35 mg, 33 %).  $R_{\rm f} = 0.2$  (CH<sub>2</sub>Cl<sub>2</sub>:MeOH 94:6); <sup>1</sup>H NMR (400 MHz, D<sub>2</sub>O):  $\delta = 8.47$  (s, 1H, H2<sup>Ad</sup> or H8<sup>Ad</sup>), 8.30 (s, 1H, H2<sup>Ad</sup> or H8<sup>Ad</sup>), 7.86 (d, J=7.7 Hz, 2H, H6<sup>Cyt</sup>), 6.19 (d, J = 4.8 Hz, 1H, H5<sup>Cyt</sup>), 6.13–6.06 (m, 2H, H1'<sup>Ad</sup>, H1'<sup>Cyt</sup>), 4.97 (t, J = 5.5 Hz, 1H, H3'<sup>Ad</sup>), 4.89–4.84 (m, 1H, H3'<sup>Cyt</sup>), 4.34–4.28 (m, 2H, H4'Ad, H4'Cyt), 4.24-4.10 (m, 2H, H5'Ad), 4.03 (s, 2H, H5'Cyt), 2.42 (ddd, J=7.9, 5.4, 2.7 Hz, 1H, H2'<sup>Cyt</sup>), 1.93-1.85 (m, 1H, H2'<sup>Cyt</sup>), 1.00 (s, 9H, tBu<sup>TBS</sup>), 0.28 (s, 3H, Me<sup>TBS</sup>), 0.27 ppm (s, 3H, Me<sup>TBS</sup>); HPLC retention time: 24.7 min (MeCN:NH4OAc 0:100 to 50:50); ESI- MS: m/z calcd for C<sub>25</sub>H<sub>37</sub>N<sub>11</sub>O<sub>12</sub>P<sub>2</sub>Si<sup>-</sup> [M-H]<sup>-</sup>: 774.1951; found: 774.1959.

[(2R,3S,5R)-5-[4-Amino-2-oxopyrimidin-1(2H)-yl]-3-({[(2R,3S,4R,5R)-5-(6-amino-9H-purin-9-yl)-4-azido-3-hydroxytetrahydrofuran-2-yl]methoxy}oxidophosphoryloxy)tetrahydrofuran-2-yl]methyl hydrogenphosphate (4): Partially protected dinucleotide 3 (34 mg, 44 µmol) was treated with aqueous 6M HCI:THF:MeOH (1:2:1, 2.8 mL) at room temperature for 24 h. The reaction mixture was concentrated in vacuo, diluted with water (3.5 mL) and washed with CH2Cl2 (3.5 mL). The aqueous layer was evaporated under reduced pressure, and the ammonium salt of dinucleotide 4 was purified by HPLC, lyophilized and recovered as a white solid (13.9 mg, 44%). <sup>1</sup>H NMR (400 MHz,  $D_2O$ ):  $\delta = 8.49$  (s, 1 H,  $H2^{Ad}$  or H8<sup>Ad</sup>), 8.25 (s, 1 H, H2<sup>Ad</sup> or H8<sup>Ad</sup>), 7.81 (d, J = 7.7 Hz, 1 H, H6<sup>Ad</sup>), 6.13– 6.10 (m, 2H, H5<sup>Cyt</sup>, H1'<sup>Ad</sup>), 6.08 (d, J = 8.0 Hz, 1H, H1'<sup>Cyt</sup>), 4.87–4.84 (m, 2H, H2'Ad, H3'Cyt), 4.35 (br, 1H, H3'Ad), 4.29 (br, 2H, H4'Cyt, H4'Ad), 4.22–4.12 (m, 2H, H5'<sup>Ad</sup>), 4.02 (dd, J = 4.6, 2.6 Hz, 2H, H5'<sup>Cyt</sup>), 2.40 (ddd,  $J=7.7, 5.7, 1.9 \text{ Hz}, 1 \text{ H}, \text{H2}'_{a}^{\text{Cyt}}), 1.92 -1.85 \text{ ppm} (m, 1 \text{ H}, \text{H2}'_{b}^{\text{Cyt}});$ <sup>13</sup>C NMR (101 MHz, D<sub>2</sub>O):  $\delta = 155.0$  (C2<sup>Ad</sup> or C8<sup>Ad</sup>), 152.5 (C2<sup>Ad</sup> or C8<sup>Ad</sup>), 141.8 (C6<sup>Cyt</sup>), 139.7(Cq), 96.0 (C1'<sup>Cyt</sup>), 85.7 (C1'<sup>Ad</sup>), 85.5 (C5<sup>Cyt</sup>), 84.7 (C4<sup>'Cyt</sup>, C4<sup>'Ad</sup>), 83.5 (C3<sup>'Ad</sup>), 70.4 (C2<sup>'Ad</sup>), 64.9 (C3<sup>'Cyt</sup>), 64.5 (C5<sup>'Cyt</sup>), 64.2 (C5'<sup>Ad</sup>), 38.1 ppm (C2'<sup>Cyt</sup>); <sup>31</sup>P NMR (<sup>1</sup>H-decoupled, 162 MHz, D<sub>2</sub>O):  $\delta = 0.19$ , -1.12 (2s); HPLC retention time: 19.2 min (MeCN:NH<sub>4</sub>OAc 0:100 to 25:75 from 7 to 37 min); ESI- MS: m/z calcd for  $C_{19}H_{24}N_{11}O_{12}P_2^{-}[M-H]^{-}: 660.1087; found: 660.1116.$ 

**UDP-MurNAc-L-Ala-γ-D-Glu-meso-cystine (5):** The reaction mixture (100 mL) contained Tris-HCl (100 mM, pH 8.6), MgCl<sub>2</sub> (40 mM), potassium phosphate (20 mM), ATP (5 mM), UDP-MurNAc-L-Ala-D-Glu (0.1 mM),<sup>[11]</sup> meso-cystine (0.5 mM) and His-tagged MurE from *E. coli* (8.7 mg).<sup>[17]</sup> After 18 h at 37 °C under gentle stirring, the magnesium phosphate precipitate was removed by centrifugation and the supernatant was lyophilized. Product **5** was purified by gel filtration on a Sephadex G-25 column (115×2 cm) in water.<sup>[18]</sup> Yield, 8.4 µmol (80%). The reaction product was used without further purification, although analytical HPLC revealed the presence of UDP-MurNAc-L-Ala-D-Glu (ca. 10%) in addition to **5**.

**UDP-MurNAc-L-Ala-γ-D-Glu-***meso***-cystine-D-Ala-D-Ala (6)**: The reaction mixture (8 mL) contained Tris-HCl (100 mM, pH 8.6), MgCl<sub>2</sub> (30 mM), ATP (5 mM), UDP-MurNAc-L-Ala-γ-D-Glu-*meso*-cystine **5** (1 mM), D-Ala-D-Ala (3 mM) and His-tagged MurF from *E. coli* 

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## **FULL PAPER**

(90 µg).<sup>[19]</sup> After 3 h of incubation at 37 °C, the reaction mixture was lyophilized and product **6** was purified on a Sephadex G-25 column (115×2 cm) in water. Yield, 7 µmol (90%). ESI MS: m/z calcd for C<sub>40</sub>H<sub>62</sub>N<sub>9</sub>O<sub>28</sub>P<sub>2</sub>S<sub>2</sub><sup>-</sup> 1242.2626  $[M-H]^-$ ; found 1242.1827. The reaction product was used without further purification although analytical HPLC revealed the presence of UDP-MurNAc-L-Ala-D-Glu (ca. 10%) in addition to **6**.

**UDP-MurNAc-L-Ala**-γ-**D**-**Glu-L-Cys-D-Ala**-**D**-**Ala** (7): Dithiothreitol (15.43 mg, 100 μmol) was added to 6 (3.3 mM, 3 μmol) in Tris-HCl buffer (50 mM, pH 8.0). After 2 h of incubation at room temperature, the product was purified by size-exclusion chromatography on a Sephadex G-25 column ( $30 \times 1$  cm) in water, lyophilized, dissolved in water and UV-quantified ( $\varepsilon_{250 \text{ nm}} = 10000 \text{ M}^{-1} \text{ cm}^{-1}$ , 2.7 μmol, 90% yield). Purity was assessed by analytical FPLC on a C18 column and product **7** was used without further purification. Retention time: 22.0 min; ESI MS: *m/z* calcd for C<sub>37</sub>H<sub>57</sub>N<sub>8</sub>O<sub>26</sub>P<sub>2</sub>O<sup>-</sup> [*M*-H]<sup>-</sup>: 1123.2585; found 1123.1970.

**UDP-MurNAc-L-Ala-γ-D-Glu-dehydroalanine-D-Ala-D-Ala (8)**: *O*-Mesitylenesulfonylhydroxylamine (MSH)<sup>[12]</sup> in DMF (7 mm, 7.04 μmol) was added to a solution of **7** (1.4 mm, 704 nmol) in phosphate buffer (50 mm, pH 8.0) at room temperature. Total conversion of Cys to dehydroalanine was typically obtained after 2 h of incubation at room temperature, as assessed by determining the sulfhydryl concentration with Ellman's reagent. Product **8** was purified by size-exclusion chromatography on a Sephadex G-25 column (30×1 cm) in water followed by FPLC on a C18 column. Product **8** was lyophilized, dissolved in water and UV-quantified ( $\varepsilon_{260 nm} = 10000 \text{ M}^{-1} \text{ cm}^{-1}$ , 200 nmol, 28 % yield). Retention time: 20.2 min; ESI MS: m/z calcd for  $C_{37}H_{55}N_8O_{26}P_2^{-1}$  [M-H]<sup>-</sup>: 1089.2708; found: 1089.2710.

UDP-MurNAc-L-Ala-γ-D-Glu-L-Cys(S-but-3-ynyl)-D-Ala-D-Ala (9a) and UDP-MurNAc-L-Ala-γ-D-Glu-D-Cys(S-but-3-ynyl)-D-Ala-D-Ala (9b): NaOH (1 M, 10 µmol) was added at room temperature to (S)-but-3-ynyl ethanethioate in DMF (1 M, 10 µmol). After 10 min phosphate buffer (80 µL, 250 mM, pH 8.0) and compound 8 in water (1 mM, 100 nmol) were sequentially added and the mixture was incubated for 2 h at 37°C. The mixture of 9a and 9b was purified by size-exclusion chromatography on a Sephadex G-25 column (30×1 cm), lyophilized and dissolved in water. Diastereoisomers 9a and 9b were purified by FPLC on a C18 column, lyophilized and dissolved in RNAse-free water (Sigma) ( $\varepsilon_{260 nm}$ =10 000 m<sup>-1</sup> cm<sup>-1</sup>; 9a: 25 nmol, 25%; 9b:25 nmol, 25%). Retention times: 36.5 and 33.9 min for 9a and 9b, respectively; ESI MS: *m/z* calcd for C<sub>41</sub>H<sub>61</sub>N<sub>8</sub>O<sub>26</sub>P<sub>2</sub>S<sup>-</sup> 1175.2898 [*M*-H]<sup>-</sup>; found: 1175.2820 and 1175.2893 for 9a and 9b, respectively.

2'-Azido-RNA helix (11): 2'-Azido-dinucleotide 4 was ligated to RNAhelix 10 (5'-GGGGCCUUAGCUCAGGCUCCAC-3') mimicking the acceptor arm of the tRNA<sup>Ala[6c]</sup> with purified T4 RNA ligase.<sup>[6c]</sup> RNA-helix 10 was synthesized by the phosphoramidite method and purified by polyacrylamide gel electrophoresis (Eurogentec). The ligation reaction was performed at 37°C for 120 min in 500 µL of HEPES buffer (50 mm, pH 7.5) containing RNA helix 10 (40 nmol), 2'-azido-dinucleotide 4 (400 nmol), T4 RNA ligase (0.6 mg), DMSO (10%), ATP (1 mm) and MgCl<sub>2</sub> (15 mm). The product of the ligation reaction, 11, was purified by anion-exchange chromatography (DNAPac PA100, 250×4 mm; Dionex) with a linear ammonium acetate gradient (25 to 2500 mm, pH 8.0) containing 0.5% acetonitrile, which was applied for 30 min at a flow rate of 1 mLmin<sup>-1</sup>.<sup>[6b]</sup> Fractions containing the ligation product were identified by PAGE, lyophilized, dissolved in RNAse-free water and stored at  $-20^{\circ}$ C ( $\varepsilon_{260 \text{ nm}} = 2.26 \ 10^5 \text{ M}^{-1} \text{ cm}^{-1}$ ; 15.4 nmol, 46%); ESI MS: m/z calcd for  $C_{227}H_{284}N_{93}O_{164}P_{23}$  [*M*]: 7651.6; found: 7651.5.

**Nonadiyne derivative of 2'-azido-RNA-helix (12):** 2'-Azido-RNA-helix **11** was coupled to 1,8-nonadiyne by Cu<sup>1</sup>-catalyzed azido–alkyne cycloaddition (Cu-AAC) reaction. The mixture ( $50 \ \mu$ L in water) contained 2'azido-RNA-helix **11** (2 nmol, 100  $\mu$ M), 1,8-nonadiyne ( $5 \ m$ M), CuSO<sub>4</sub> (0.5 mM), Na ascorbate ( $5 \ m$ M) and the Cu<sup>1</sup>-stabilizing ligand THPTA ( $3.5 \ m$ M).<sup>[15a]</sup> After 24 h of incubation at 4 °C, Cu–THPTA complex and Na ascorbate were removed by gel filtration on a Sephadex G-25 column ( $30 \times 1 \ m$ ) in water. Product **12** was purified by denaturing PAGE followed by gel filtration. Concentration was determined spectrophotometrically ( $\varepsilon_{260 \text{ nm}} = 2.26 \text{ } 10^8 \text{ M}^{-1} \text{ cm}^{-1}$ ; 1.4 nmol, 70%). ESI MS: *m*/*z* calcd for C<sub>236</sub>H<sub>296</sub>N<sub>93</sub>O<sub>164</sub>P<sub>23</sub> [*M*]: 7771.8; found: 7772.1.

RNA-helix-UDP-MurNAc-pentapeptide bi-substrate (13a and 13b): 2'-Azido-RNA-helix 11 was coupled to 9a or 9b by Cu-AAC reaction. The mixture (200 µL in phosphate buffer, 25 mM, pH 8.0), which contained 2'azido-RNA-helix 11 (5 nmol, 25 µм), 9a or 9b (20 nmol, 50 µм), CuSO<sub>4</sub> (0.5 mm), Na ascorbate (5 mm) and THPTA (3.5 mm), was incubated at 4°C for 24 h. Compounds 13a and 13b were purified by denaturing PAGE and gel filtration as described above for compound 12 ( $\varepsilon = 2.36 \times$  $10^5 \text{ m}^{-1} \text{ cm}^{-1}$  at 260 nm, 200 pmol; 4%). The procedure was optimized by increasing the concentration of 9a (20 mm, 200 nmol) and 11 (1 mm, 10 nmol) in the ligation step and by purifying 13a by anion-exchange chromatography (DNAPac PA-100, Dionex) with a solvent system consisting of a 30 min linear gradient from 25 to 2500 mM of NH<sub>4</sub>OAc (pH 8.0) containing 0.5% CH<sub>3</sub>CN at a flow rate of 1 mL min<sup>-1</sup>. Compound 13a was identified by denaturing PAGE and its concentration was determined spectrophotometrically ( $\varepsilon_{260 \text{ nm}} = 2.36 \times 10^5 \text{ M}^{-1} \text{ cm}^{-1}$ ; 3.6 nmol, 36%). ESI MS: m/z calcd for  $C_{268}H_{346}N_{101}O_{190}P_{25}S$  [M]: 8828.6; found: 8828.4.

Analysis of the interaction of 13a with FemX<sub>wv</sub> by size-exclusion chro**matography**: Bi-substrate **13a** (200 pmol) was incubated with Fem $X_{Wy}$ (1200 pmol) for 10 min at 4°C in 500  $\mu L$  of buffer A (25 mm Tris-HCl, pH 7.5; 100 mм NaCl; 5 mм MgCl<sub>2</sub>). The FemX<sub>wv</sub>-13a complex was loaded onto a Sephadex G-75 column (10/300 GL, GE Healthcare) equilibrated with buffer A and elution was performed at a flow rate of 0.8 mLmin<sup>-1</sup> at room temperature. Relevant fractions (5 mL) were pooled, concentrated to a final volume of 500 µL by ultrafiltration (3 kDa, Microcon, Millipore) and incubated for 10 min at 65 °C followed by 5 min at 0°C. Compound 13a was extracted twice with phenol (500  $\mu L;$  phenol, water-saturated, stabilized, pH 4.1; Eurobio) and once with a phenol:chloroform solution (5:1 v:v, 500 µL, Sigma). The product was purified on a Sephadex 75 10/300 GL column in buffer A. Compound 13a was concentrated to a final volume of 250 µL by ultrafiltration. The overall yield of the size-exclusion chromatography purification step was 43 % (85 pmol).

IC<sub>50</sub> determination: W. viridescens FemX<sub>Wv</sub><sup>[16]</sup> E. faecalis alanyl-tRNA synthetase (AlaRS)<sup>[7a]</sup> and T7 RNA polymerase<sup>[6c]</sup> were purified as previously described. Full-length tRNAAla (5'-GGGGCCUUAGCUCAG-CUGGGAGAGCGCCUGCUUUGCACGCAGGAGGUCAGCGGU-UCGAUCCCGCUAGGCUCCACCA-3') was obtained by in vitro transcription using T7 RNA polymerase.<sup>[6c]</sup> Inhibition of FemX<sub>Wv</sub> was tested in a radioactive coupled assay<sup>[8a]</sup> involving acylation of tRNA<sup>Ala</sup> by [<sup>14</sup>C]Ala and transfer of [<sup>14</sup>C]Ala from the resulting [<sup>14</sup>C]Ala-tRNA<sup>Ala</sup> to UDP-MurNAc-L-Ala-y-D-Glu-L-Lys-D-Ala-D-Ala (UDP-MurNAc-pentapeptide) to form UDP-MurNAc-L-Ala-y-D-Glu-L-Lys(L-[14C]Ala)-D-Ala-D-Ala ([14C]UDP-MurNAc-hexapeptide). The standard assay mixture (10 µL) contained Tris-HCl (50 mM, pH 7.5), alanyl-tRNA synthetase (800 nm), ATP (7.5 mm), β-mercaptoethanol (2 mm), MgCl<sub>2</sub> (12.5 mm), L- $[^{14}C]$ Ala (50 μм, 3700 Bq nmol<sup>-1</sup>, ICN, Orsay, France), tRNA<sup>Ala</sup> (0.4 μм), UDP-MurNAc-pentapeptide (50 μм), FemX<sub>Wv</sub> (5 nм) and various concentrations of compounds 10, 11 and 12. Reactions were performed at 30°C for 10 min and stopped at 95°C for 10 min. Inhibition of FemX<sub>Wv</sub> by 13a and 13b was tested under the same conditions, except for the volume of the mixture (100 µL and 10 µL, respectively), FemX<sub>Wy</sub> concentration (0.05 µm and 0.5 µm, respectively) and incubation time (100 min for both). The reaction mixture was lyophilized and dissolved in water (10 µL). L-[14C]Ala and [14C]UDP-MurNAc-hexapeptide were separated by descending paper chromatography (Whatman 4MM, Elancourt, France) with isobutyric acid:1 M ammonium hydroxide (5:3). Radioactive spots were identified by autoradiography, cut out and counted by liquid scintillation. IC50 were determined by plotting residual activity versus inhibitor concentration. Curves represent fits of a four-parameter logistic function of experimental values (Sigma Plot 9.0).

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### Peptidyl-RNA Conjugates -

- M. Fonvielle, D. Mellal, D. Patin, M. Lecerf, D. Blanot, A. Bouhss, M. Santarem, D. Mengin-Lecreulx, M. Sollogoub, M. Arthur,\*
- *M. Ethève-Quelquejeu*\*..... **IIII**-**IIII**

Efficient Access to Peptidyl-RNA Conjugates for Picomolar Inhibition of Non-ribosomal FemX<sub>wv</sub> Aminoacyl Transferase



**Conjugation of complex biomolecules:** Peptidyl–RNA conjugates have various applications to study the ribosome and enzymes participating in tRNAdependent pathways such as Fem transferases in peptidoglycan synthesis. Here, a convergent route for synthesis of peptidyl–RNAs based on Huisgen– Sharpless cycloaddition for the final ligation step is developed (see figure). Both moieties of the peptidyl–RNA conjugate contribute to high-affinity binding to FemX, providing a picomolar inhibitor.