

A polymer of 8-*O*-glucosylated 2-keto-3-deoxy-D-glycero-D-galacto-nonulosonic acid (Kdn) in the cell wall of *Streptomyces* sp. VKM Ac-2090

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The title polymer of Kdn was detected in biological object for the first time.

2-Keto-3-deoxy-D-glycero-D-galacto-nonulosonic acid (Kdn) was first found in the form of an α -2,8-linked oligomer containing 6–7 residues in rainbow trout eggs.¹ Later, Kdn was found in heterooligosaccharides of animal tissues^{2,3} and as a constituent of the capsular heteropolysaccharide of *Klebsiella ozaenae* serotype K4.⁴

We have detected polyKdn in the cell wall of *Streptomyces* sp. VKM Ac-2090 isolated from scab lesions potato.

The polymers were extracted from the cell wall as described previously,⁵ and their structure was examined using NMR spectroscopy.

The ¹³C NMR spectrum of the polymers (Figure 1) displayed the Kdn-containing polysaccharide as the major cell wall anionic polymer, along with several glycerol teichoic acids. The ¹H and ¹³C NMR spectra of the predominant component of the polymer mixture were completely assigned using 2D homonuclear ¹H/¹H COSY, TOCSY and ROESY and heteronuclear ¹H/¹³C HSQC (Figure 2) and HMBC experiments. The residue of nonulosonic acid was identified with Kdn in accordance with the coupling constants in the ¹H NMR spectrum.⁶ The upfield chemical shift of H-3_{eq} (2.205 ppm) was in agreement with the β -configuration of the Kdn residue.⁴ The second sugar residue in the disaccharide repeating unit of the polymer was identified as the terminal β -glucopyranose (β -Glc_p) based on the chemical shifts and coupling constants of the ¹H and ¹³C NMR spectra. The ROESY spectrum (Figure 3) revealed correlation peaks for the anomeric protons of Glc_p and H-8, H-9 and H-9' of Kdn.

The upfield chemical shift of C-9 and the downfield chemical shift of C-8 in the Kdn residue in comparison with that of non-substituted sugar (Table 1) allowed one to conclude that β -Glc_p was bonded with Kdn by the 1 \rightarrow 8 linkage. The presence of a correlation peak of H-1(Glc_p)/C-8(Kdn) in the HMBC spectrum is consistent with this conclusion, too. The downfield shift of C-4 by 2 ppm in the ¹³C NMR spectrum of the β -Kdn residue

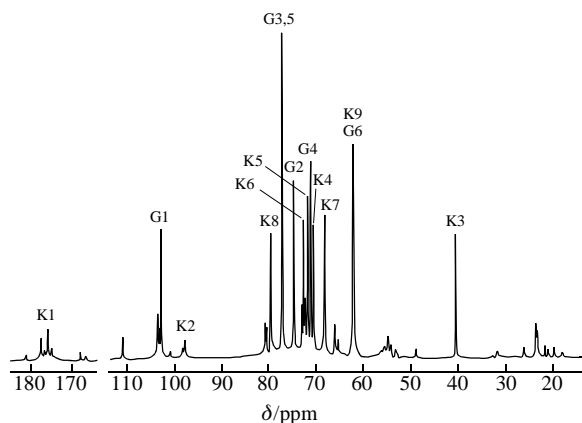


Figure 1 ¹³C NMR spectrum of the polymers of the cell wall of *Streptomyces* sp. VKM Ac-2090. Designations refer to the numbers of carbon atoms in the Kdn (K) and glucopyranose (G) residues.

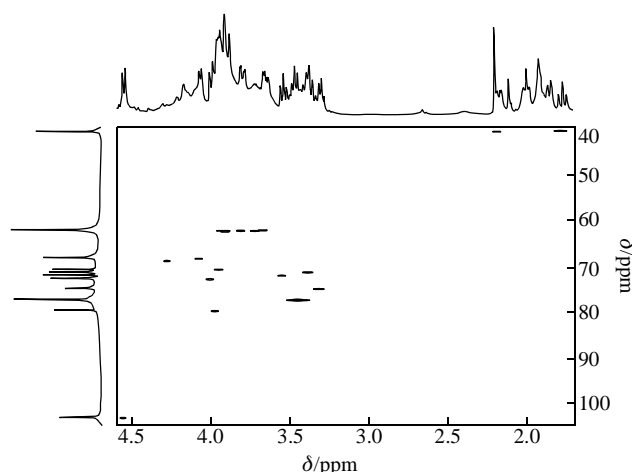


Figure 2 HSQC spectrum of the Kdn-containing polysaccharide of the cell wall of *Streptomyces* sp. VKM Ac-2090. The corresponding parts of the ¹H and ¹³C NMR spectra are displayed along the horizontal and vertical axes, respectively.

Table 1 ¹H NMR data (D₂O, 303 K, acetone: 2.225 ppm) for β -Kdn⁶ and for the polysaccharide of *Streptomyces* sp. VKM Ac-2090 cell wall.

Residue	Chemical shifts, δ /ppm and coupling constants, J/Hz								
	H-3 _{ax}	H-3 _{eq}	H-4	H-5	H-6	H-7	H-8	H-9	H-9'
β -Kdn ⁶	1.80 $J_{3a,3e}$ 12.0	2.23 $J_{3e,4}$ 5.0	4.02 $J_{3a,4}$ 12.0	3.56 $J_{4,5}$ 9.0	4.01 $J_{5,6}$ 9.0	3.88 $J_{6,7}$ 1	3.73 $J_{7,8}$ 8.5	3.63 $J_{9,9'}$ 11	3.88 $J_{8,9'}$ 2
\rightarrow 4)- β -Kdn- (2 \rightarrow 8)	1.80 $J_{3a,3e}$ 13.0	2.20 $J_{3e,4}$ 5.0	3.98 $J_{3a,4}$ 12.3	3.57 $J_{4,5}$ 9.6	4.02 $J_{5,6}$ 9.6	4.09 $J_{6,7}$ 1.2	3.99 $J_{7,8}$ 8.5	3.94 $J_{9,9'}$ 12.5	3.83 $J_{8,9'}$ 3.9
β -Glc _p -(1	H-1 4.58 $J_{1,2}$ 7.9	H-2 3.33 $J_{2,3}$ 8.8	H-3 3.50 $J_{3,4}$ 8.8	H-4 3.38 $J_{4,5}$ 8.7	H-5 3.42 $J_{5,6}$ 1.7	H-6 3.82 $J_{6,6'}$ 12.1	H-6' 3.68 $J_{5,6'}$ 5.7		

Table 2 ¹³C NMR data (D₂O, 303 K, acetone: 31.45 ppm) for β -Kdn⁶ and for the polysaccharide of *Streptomyces* sp. VKM Ac-2090 cell wall.

Residue	Chemical shifts, δ /ppm							
	C-1	C-2	C-3	C-4	C-5	C-6	C-7	C-9
β -Kdn ⁶	174.2	96.02	39.19	68.51	70.67	71.06	69.27	63.86
\rightarrow 4)- β -Kdn- (2 \rightarrow 8)	176.0	97.9	40.5	70.5	71.7	72.6	68.05	61.9
β -Glc _p -(1	102.8	74.7	77.1	71.1	77.1	61.9		

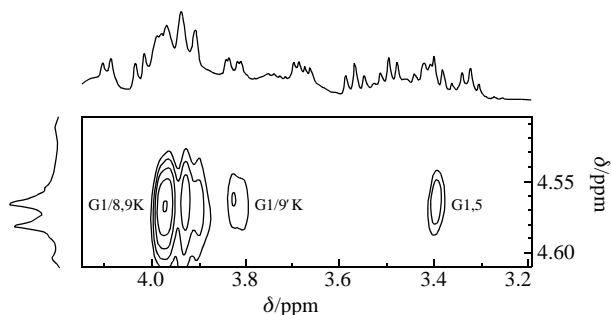
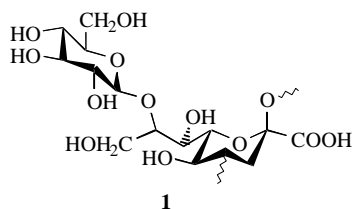


Figure 3 Part of the ROESY spectrum of the Kdn-containing polysaccharide of the cell wall of *Streptomyces* sp. VKM Ac-2090. The corresponding parts of the ^1H NMR spectrum are displayed along the horizontal and vertical axes. Designations refer to the numbers of protons in the Kdn (K) and glucopyranose (G) residues.

compared to that of non-substituted β -Kdn revealed the 2 \rightarrow 4 linkage for the polysaccharide chain. The signals of the terminal monosaccharide residues were not detected. This fact allows us to suggest a high molecular mass of the polymer. Both of the comparable NOE correlation peaks H-1(Glcp)/H-8(Kdn) and H-1(Glcp)/H-9(Kdn) (Figure 3) and relatively large in module negative β -effect of glycosylation for C-9 of the Kdn residue were in agreement with the D-glycero-D-galacto-configuration of Kdn on the assumption of the β -D-configuration of the glucopyranose residue^{7,8} **1**.



Until now, a polymer of Kdn has been found neither in prokaryotic nor eucaryotic cells.

The β -configuration of the glycoside bond in this natural sugar has not been reported previously. This communication is the first report on the polyKdn with the β -configuration of the glycoside bond.

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