



**Figure S1.**  $^1\text{H}$   $^{13}\text{C}$  NMR HSQC-DEPT spectra of the OS of *B. holmesii* ATCC 51541 and *B. pertussis* 606 [overlay spectrum]; \*Resonances placed in frames indicate variants of the residue F  $\rightarrow$ 3,4)-L- $\alpha$ -D-Hepp-(1 $\rightarrow$  in different environment; The remaining resonances of the initially identified residue G ( $\delta_{\text{H1}}/\delta_{\text{C1}}$  4.95/96.9) and ( $\delta_{\text{H2}}/\delta_{\text{C2}}$  3.85/73.4) were not resolved from those of residue F.

**Table S1.**  $^1\text{H}$   $^{13}\text{C}$  NMR chemical shifts of the core oligosaccharides of *B. pertussis* 606 and *B. holmesii* ATCC 51541 \*

Residue	Strain	Chemical Shifts (ppm)							
		H-1	H-2	H-3	H-4	H-5	H-6, H-6'	H-7	H-8, H-8'
		C-1	C-2	C-3	C-4	C-5	C-6	C-7	C-8
<b>Kdo</b> $\rightarrow$ 4,7)-anhKdof	Bp606			3.12	4.38	4.02	4.13	3.89	3.66, 3.60
			203.1	42.9	77.4	83.8	75.4	83.6	61.1
	Bhol			3.12	4.39	4.02	4.14	3.89	3.66, 3.60
			202.9	42.8	77.4	83.8	75.4	83.7	61.1
<b>Kdo'</b> $\rightarrow$ 4,7)-anhKdof	Bp606			3.11	4.50	4.12	4.08	3.74	3.67, 3.58
			203.3	39.1	75.7	80.2	74.7	84.8	61.8
	Bhol			3.09	4.51	4.12	4.08	3.74	3.67, 3.58
			203.2	38.8	75.8	80.2	74.7	84.8	61.8
<b>A</b> $\rightarrow$ 4)- $\alpha$ -Glc <sub>p</sub> N-(1 $\rightarrow$	Bp606	5.47	3.31	3.94	3.63	3.73	3.78, 3.73		
		97.4	54.5	70.3	74.9	71.9	60.5		
	Bhol	5.48	3.33	3.94	3.63	3.74	3.78	3.73	
		97.3	54.5	70.3	74.9	71.9	60.5		
<b>B</b> $\rightarrow$ 2,7)-L- $\alpha$ -D-Hepp-(1 $\rightarrow$	Bp606	5.38	3.86	3.90	3.84	3.46	4.18	3.70	
		99.7	79.9	70.5	66.6	72.2	68.0	70.3	
	Bhol	5.39	3.85	3.91	3.84	3.45	4.17	3.70	
		99.7	79.8	70.4	66.5	72.1	67.9	70.3	
<b>C</b> L- $\alpha$ -D-Hepp-(1 $\rightarrow$	Bp606	5.27	3.96	3.64	3.78	3.55	3.90	3.65, 3.62	
		101.2	70.1	70.3	65.8	72.5	68.5	62.5	
	Bhol	5.27	3.96	3.74	3.81	3.54	3.94	3.62	
		101.2	70.1	70.5	65.9	72.5	68.5	62.7	
<b>D</b> $\alpha$ -Gal <sub>p</sub> NA-(1 $\rightarrow$	Bp606	5.16	3.44	4.03	4.15	4.33			
		94.5	50.5	66.7	69.7	72.3	175.1		
	Bhol	5.17	3.47	4.04	4.16	4.33			
		94.4	50.5	66.7	69.7	72.3	175.1		
<b>E</b> $\alpha$ -Glc <sub>p</sub> N-(1 $\rightarrow$	Bp606	5.12	3.28	3.85	3.45	3.70	3.78, 3.72		
		96.0	54.1	69.8	69.4	72.2	60.2		
	Bhol	5.12	3.28	3.86	3.44	3.70	3.78, 3.72		
		96.0	54.1	69.8	69.4	72.2	60.2		
<b>F</b> $\rightarrow$ 3,4)-L- $\alpha$ -D-Hepp-(1 $\rightarrow$	Bp606	5.07	3.90	3.84	4.23	3.54	3.95	3.67, 3.65	
		97.7	73.5	79.9	71.7	71.3	68.9	62.7	
	Bhol	5.07	3.90	3.84	4.23	3.53	3.93	3.67, 3.65	
		97.8	73.3	76.7	71.7	71.4	68.9	62.7	
<b>H</b> $\alpha$ -Glc <sub>p</sub> A-(1 $\rightarrow$	Bp606	4.97	3.52	3.69	3.39	4.04			
		101.0	71.9	71.6	72.3	73.8	176.6		
	Bhol	4.99	3.52	3.68	3.39	4.03			
		101.0	71.9	71.6	72.3	73.8	176.6		
<b>I*</b> $\alpha$ -Glc <sub>p</sub> A-(1 $\rightarrow$	Bp606	4.96	3.51	3.67	3.37	4.01			
		101.1	72.1	71.5	72.2	73.8	176.7		
	Bhol	4.95	3.52	3.67	3.38	4.01			
		101.2	72.1	71.6	72.2	73.9	176.7		

Residue	Strain	Chemical Shifts (ppm)							
		H-1	H-2	H-3	H-4	H-5	H-6, H-6'	H-7	H-8, H-8'
		C-1	C-2	C-3	C-4	C-5	C-6	C-7	C-8
J →4,6)-β-Glcp-(1→	Bp606	4.39	3.27	3.56	3.44	3.67	3.76, 3.89		
		101.4	73.4	76.4	78.9	72.9	67.2		
	Bhol	4.40	3.27	3.55	3.44	3.66	3.76, 3.89		
		101.4	73.4	76.4	79.0	73.0	67.3		

\* The additional CH<sub>2</sub>- signals at δ<sub>H1</sub>/δ<sub>C1</sub> (3.13/40.5 ppm), δ<sub>H1'</sub>/δ<sub>C1'</sub> (3.18/39.8 ppm) and δ<sub>H2</sub>/δ<sub>C2</sub> (3.9/60.2 ppm), δ<sub>H2'</sub>/δ<sub>C2'</sub> (4.09/62.0 ppm) for the *B. pertussis* 606 OS and at δ<sub>H1</sub>/δ<sub>C1</sub> (3.17/40.0 ppm) and δ<sub>H2</sub>/δ<sub>C2</sub> (4.07/62.2 ppm) for the *B. holmesii* OS V indicate the presence of phosphoethanolamine in the intact core oligosaccharides of these strains. P-EtN correlations were not resolved and the substitution position is only tentative.

**Table S2.** Selected inter-residue NOE and <sup>3</sup>J<sub>H,C</sub>-connectivities from the anomeric atoms of the core oligosaccharide of *B. pertussis* 606

Residue	Atom H-1/C-1 Connectivities to Inter-Residue			
	(ppm)	δ <sub>C</sub>	δ <sub>H</sub>	Atom/Residue
A →4)-α-GlcpN-(1→	97.4/5.47	78.9	3.44	C-4, H-4 of J
B →2,7)-L-α-D-Hepp-(1→	99.7/5.38	79.9	3.84	C-3, H-3 of F
C L-α-D-Hepp-(1→	101.2/5.27	74.9	3.63	C-6, H-6/6' of A
D α-GalpNA-(1→	94.5/5.15	67.2	3.76/3.89	C-6, H-6/6' of J
E α-GlcpN-(1→	96.0/5.12	70.3	3.70	C-7, H-7 of B
F →3,4)-L-α-D-Hepp-(1→	97.7/5.07	83.7	4.02	C-5, H-5 of Kdo
H α-GlcpA-(1→	101.0/4.97	79.9	3.86	C-2, H-2 of B
I α-GlcpA-(1→	101.1/4.96	79.9	3.85	C-2, H-2 of B
J →4,6)-β-Glcp-(1→	101.4/4.39	71.7	4.23	C-4, H-4 of F