Supplemental data

GENERAL MATERIALS AND PROCEDURE FOR SYNTHESIS OF GalNAc/GlcNAc-PP-LIPID

All solvents were dried with a solvent-purification system from Innovative Technology, Inc. All reagents were obtained from commercial sources and used without further purification. The 200-400 mesh silica gel 60 RP-18 was utilized for purification. ¹H, ¹³C and ³¹P NMR spectra were recorded at the indicated field strengths. Mass spectral data was collected using a Shimadzu LCMS-2010A Liquid Chromatography Mass Spectrometer or a Bruker micrOTOF Instrument.

The general procedure for chemical synthesis of GalNAc/GlcNAc-PP-lipid compounds is shown as Fig. S1. Specifically, the N,N'-Carbonyldiimidazole (CDI) (23 mg, 0.14 mmol) was added to a reaction vessel containing the diisopropylethylammonium salt of GalNAc/GlcNAc-1- $PO_4^{2^2}$ (23 mg, 0.034 mmol). The vessel was flushed with argon, after which anhydrous THF was added (3 mL). The reaction was stirred at room temperature for 2 h. Addition of dry CH₃OH (42 µL) followed by stirring for an additional 1 h at room temperature served to remove excess CDI. The reaction mixture was concentrated and then anhydrous THF (3 mL) was once again added. To a separate reaction vessel, lipid phosphate (0.026 mmol) was added, and the vessel was then flushed with argon. The activated GalNAc/GlcNAc-1- $PO_4^{2^2}$ was transferred into this vessel via syringe. The reaction was stirred at room temperature for 3 days, after which the reaction mixture was concentrated. Dissolution of the residue in water, followed by filtration through a short bed of C-18 reverse phase silica gel (water then 1:1 Water/Isopropanol) afforded the crude product.

The crude product was treated with a 0.1% NaOCH₃ in CH₃OH solution (3 mL). The reaction was stirred at room temperature for 40 min. The reaction mixture was neutralized with Amberlyst-15 ion exchange resin until pH around 7 and then concentrated, after which the residue was dissolved in water. Purification via C-18 reverse phase column chromatography (Solvent A: Isopropanol (Lipids: Undecaprenyl, Solanesyl,) or CH₃CN (Lipids: Heptaprenyl, Pentaprenyl, MS-Pentaprenyl, cis-Farnesyl, Undecyl, Geranyl); Solvent B = 1.7% NH₄HCO₃; A/B = 0%, 5%, 10%, 15%, 20%, 25%, 30%, 35%, 40%, 45% and 50% (10 mL each)) provided GalNAc/GlcNAc-PP-lipid as a white solid following lyophilization.

SYNTHESIS AND CHARACTERIZATION OF EACH OF GalNAc/GlcNAc-PP-LIPID COMPOUNDS

Chemical synthesis and characterization of some GalNAc/GlcNAc-PP-lipid compounds have been reported in our previously published paper (Woodward R, et al. (2010) In vitro

bacterial polysaccharide biosynthesis: defining the functions of Wzy and Wzz. *Nat Chem Biol* 6:418-423.) and those data as well as the data for the newly-synthesized compounds in this study were all listed below for reference.

Chemical synthesis of GalNAc-PP-Geranyl-GalNAc-PP-Geranyl was prepared using the general procedure as described above. Following the deprotection step, the crude product was purified via C-18 reverse phase column chromatography to afford GalNAc-PP-Geranyl as a white solid (10 mg, 11% over two steps). ¹H NMR (500 MHz, D₂O): δ 5.57 (dd, *J* = 3.4 Hz, *J* = 7.0 Hz, 1H), 5.48 (td, *J* = 1.1 Hz, *J* = 7.1 Hz, 1H), 5.28-5.22 (m, 1H), 4.53 (t, *J* = 6.9 Hz, 2H), 4.29 (dt, *J* = 3.0 Hz, *J* = 10.9 Hz, 1H), 4.23 (t, *J* = 6.2 Hz, 1H), 4.08 (d, *J* = 2.6 Hz, 1H), 4.01 (dd, *J* = 3.2 Hz, *J* = 10.9 Hz, 1H), 3.85-3.78 (m, 2H), 2.23-2.17 (m, 2H), 2.16-2.12 (m, 2H), 2.11 (s, 3H), 1.76 (s, 3H), 1.73 (s, 3H), 1.67 (s, 3H); ¹³C NMR (125 MHz, D₂O): δ 160.3, 143.0, 133.7, 124.1, 121.6, 94.7, 72.1, 68.4, 67.8, 62.5, 61.1, 49.8, 38.8, 25.6, 24.8, 22.2, 17.0, 15.6; ³¹P NMR (162 MHz, D₂O): δ -10.2, -12.5; LRMS (*m*/*z*): [M-H]⁻ calcd. for C₁₈H₃₂NO₁₂P₂, 516.1; found 516.1.

Chemical synthesis of GalNAc-PP-cis-Farnesyl-GalNAc-PP-cis-Farnesyl was prepared using the general procedure as described above. Following the deprotection step, the crude product was purified via C-18 reverse phase column chromatography to afford GalNAc-PP-cis-Farnesyl as a white solid (11 mg, 6% over two steps). ¹H NMR (500 MHz, D₂O): δ 5.58 (dd, *J* = 3.4 Hz, *J* = 7.0 Hz, 1H), 5.51 (t, *J* = 6.1 Hz, 1H), 5.32-5.22 (m, 2H), 4.52 (t, *J* = 7.0 Hz, 2H), 4.31 (dt, *J* = 3.0 Hz, *J* = 10.9 Hz, 1H), 4.25 (t, *J* = 6.2 Hz, 1H), 4.09 (d, *J* = 3.0 Hz, 1H), 4.01 (dd, *J* = 3.2 Hz, *J* = 10.9 Hz, 1H), 3.88-3.75 (m, 2H), 2.24-2.13 (m, 8H), 2.12 (s, 3H), 1.83 (s, 3H), 1.75 (s, 6H), 1.69 (s, 3H); ¹³C NMR (125 MHz, D₂O): δ 175.0, 142.8, 137.2, 133.7, 124.8, 124.4, 120.5, 94.7, 72.1, 68.5, 67.9, 62.9, 61.1, 49.8, 32.1, 31.6, 31.2, 25.9, 24.9, 22.7, 22.5, 22.2, 17.0; ³¹P NMR (162 MHz, D₂O): δ -10.2, -12.5; LRMS (*m*/*z*): [M-H]⁻ calcd. for C₂₃H₄₀NO₁₂P₂, 584.2; found 584.2.

Chemical synthesis of GalNAc-PP-MS-Pentaprenyl-GalNAc-PP-MS-Pentaprenyl was prepared using the general procedure as described above. Following the deprotection step, the crude product was purified via C-18 reverse phase column chromatography to afford GalNAc-PP-MS-Pentaprenyl as a white solid (27 mg from 65 mg lipid phosphate, 26% over 2 steps). ¹H NMR (500 MHz, D₂O): δ 5.55 (dd, J = 3.3 Hz, J = 7.1 Hz, 1H), 5.22-5.05 (m, 4H), 4.26 (dt, J = 2.9 Hz, J = 11.0 Hz, 1H), 4.19 (dd, J = 5.2 Hz, J = 7.1 Hz, 1H), 4.06-3.93 (m, 4H), 3.84-3.72 (m, 2H), 2.11-2.02 (m, 11H), 2.01-1.92 (m, 6H), 1.69 (s, 3H), 1.66 (s, 3H) 1.61 (s, 3H), 1.58 (s, 6H), 1.54-1.24 (m, 4H), 1.22-1.10 (m, 1H), 0.92 (d, J = 6.5 Hz, 3H); ¹³C NMR (125 MHz, D₂O): δ 174.5, 135.0, 134.8, 134.5, 130.8, 125.7, 124.4, 124.2, 94.7, 72.1, 68.6, 67.7, 64.9, 61.3, 49.9, 49.8, 39.6, 37.3, 37.1, 31.8, 29.0, 26.6, 26.6, 26.5, 25.4, 24.9, 23.2, 22.3, 18.8, 18.8, 17.4, 15.7, 15.7; ³¹P

NMR (162 MHz, D₂O): δ -9.9, -11.8; LRMS (*m*/*z*): [M-H]⁻ calcd. for C₃₃H₅₈NO₁₂P₂, 722.3; found, 722.3.

Chemical synthesis of GalNAc-PP-Pentaprenyl-GalNAc-PP-Pentaprenyl was prepared using the general procedure as described above. Following the deprotection step, the crude product was purified via C-18 reverse phase column chromatography to afford GalNAc-PP-Pentaprenyl as a white solid (22 mg from 28 mg lipid phosphate, 49% over 2 steps). ¹H NMR (500 MHz, CD₃OD): δ 5.67 (dd, *J* = 3.1 Hz, *J* = 6.8 Hz, 1H), 5.56 (t, *J* = 6.6 Hz, 1H), 5.30-5.16 (m, 4H), 4.58 (t, *J* = 6.5 Hz, 2H), 4.36 (d, *J* = 10.9 Hz, 1H), 4.30 (dd, *J* = 6.0 Hz, 1H), 4.14 (d, *J* = 3.0 Hz, 1H), 4.07 (dd, *J* = 2.9 Hz, *J* = 10.9 Hz, 1H), 3.94-3.84 (m, 2H), 2.30-2.23 (m, 2H), 2.22-2.11 (m, 13H), 2.10-2.02 (m, 4H), 1.85 (s, 3H), 1.80 (s, 3H), 1.75 (s, 3H), 1.71 (s, 3H), 1.68 (s, 6H); ¹³C NMR (125 MHz, D₂O): δ 174.7, 160.3, 141.2, 135.8, 134.9, 134.4, 130.6, 124.6, 124.5, 124.3, 121.2, 94.7, 72.2, 68.7, 67.7, 62.7, 61.3, 49.9, 48.9, 39.7, 39.7, 32.0, 31.9, 26.7, 26.6, 26.1, 25.4, 23.1, 22.9, 22.3, 17.4, 15.8, 15.7; ³¹P NMR (162 MHz, D₂O): δ -10.4, -13.1; LRMS (*m*/*z*): [M-H]⁻ calcd. for C₃₃H₅₆NO₁₂P₂, 720.3; found, 720.3.

Chemical synthesis of GalNAc-PP-Heptaprenyl-GalNAc-PP-Heptaprenyl was prepared using the general procedure as described above. Following the deprotection step, the crude product was purified via C-18 reverse phase column chromatography to afford GalNAc-PP-Heptaprenyl as a white solid (22 mg from 42 mg lipid phosphate, 36% over 2 steps). ¹H NMR (500 MHz, D₂O): δ 5.61 (dd, *J* = 2.7 Hz, *J* = 5.9 Hz, 1H), 5.50 (t, *J* = 5.5 Hz, 1H), 5.24-5.08 (m, 6H), 4.51 (t, J = 5.7 Hz, 2H), 4.29 (d, J = 10.8 Hz, 1H), 4.23 (t, J = 5.5 Hz, 1H), 4.07 (d, J = 1.4Hz, 1H), 4.00 (dd, J = 2.2 Hz, J = 10.7 Hz, 1H), 3.88-3.75 (m, 2H), 2.20-1.96 (m, 27H), 1.79 (s, 3H), 1.72 (s, 3H), 1.70 (s, 3H), 1.68 (s, 3H), 1.62 (s, 3H), 1.60 (s, 9H); ¹³C NMR (125 MHz, D₂O): δ 174.5, 172.4, 172.2, 171.9, 140.8, 136.3, 135.9, 135.9, 135.9, 135.8, 132.1, 125.7, 125.7, 125.6, 125.4, 122.7, 122.6, 96.5, 70.6, 68.7, 64.0, 64.0, 62.3, 41.0, 41.0, 41.0, 40.9, 40.8, 34.5, 33.2, 30.9, 30.8, 30.7, 30.6, 30.5, 28.0, 27.9, 27.8, 27.7, 27.7, 27.7, 26.0, 24.0, 23.8, 23.0, 20.8, 20.7, 20.7, 17.9, 16.8, 16.2, 15.7, 14.5; 174.7, 165.3, 134.8, 134.6, 134.3, 130.4, 125.2, 125.0, 124.8, 124.5, 124.3, 124.3, 121.2, 94.6, 72.1, 68.7, 67.7, 62.6, 61.3, 49.9, 39.7, 39.7, 32.0, 31.9, 31.9, 29.9, 29.8, 29.7, 29.5, 26.7, 26.6, 26.2, 25.4, 23.3, 23.2, 23.0, 23.0, 22.3, 17.3, 15.7; ³¹P NMR (162 MHz, D₂O): δ -9.7, -11.4; LRMS (*m*/*z*): [M-H]⁻ calcd. for C₄₃H₇₂NO₁₂P₂, 856.5; found, 856.7.

Chemical synthesis of GalNAc-PP-Solanesyl-GalNAc-PP-Solanesyl was prepared using the general procedure as described above. However, the two purification steps were reversed as the final product proved too insoluble to obtain suitable spectral data. The crude product was thus purified via C-18 reverse phase column chromatography to afford Peracetylated GalNAc-PP-

Solanesyl as a white solid (35 mg from 45 mg lipid phosphate, 50%). ¹H NMR (500 MHz, CD₃OD): δ 5.61 (dd, *J* = 3.3 Hz, *J* = 7.3 Hz, 1H), 5.43 (t, *J* = 6.6 Hz, 2H), 5.22 (dd, *J* = 3.1 Hz, *J* = 11.3 Hz, 1H), 5.14-5.04 (m, 8H), 4.59 (t, *J* = 7.0 Hz, 1H), 4.53 (t, *J* = 6.3 Hz, 3H), 4.21 (dd, *J* = 8.2 Hz, *J* = 10.8 Hz, 1H), 4.04 (dd, *J* = 5.8 Hz, *J* = 10.8 Hz, 1H), 2.11 (s, 3H), 2.10-2.02 (m, 18H), 1.99 (s, 3H), 1.99-1.94 (m, 17H), 1.90 (s, 3H), 1.69 (s, 3H), 1.65 (s, 3H), 1.59 (s, 3H), 1.58 (s, 21H); ¹³C NMR (125 MHz, CD₃OD): δ 174.5, 172.4, 172.2, 171.9, 140.8, 136.3, 135.9, 135.9, 135.9, 135.8, 132.1, 125.7, 125.7, 125.6, 125.4, 122.7, 122.6, 96.5, 70.6, 68.7, 64.0, 64.0, 62.3, 41.0, 41.0, 40.9, 40.8, 34.5, 33.2, 30.9, 30.8, 30.7, 30.6, 30.5, 28.0, 27.9, 27.8, 27.7, 27.7, 26.0, 24.0, 23.8, 23.0, 20.8, 20.7, 20.7, 17.9, 16.8, 16.2, 15.7, 14.5; ³¹P NMR (162 MHz, CD₃OD): δ -9.6, -12.5; LRMS (*m*/z): [M-H]⁻ calcd. for C₅₉H₉₄NO₁₅P₂, 1118.6; found, 1118.5.

Final deprotection with 0.1% NaOCH₃ in CH₃OH and filtration through a short bed of C-18 reverse phase silica gel afforded GalNAc-PP-Solanesyl (31 mg, 99%). LRMS (m/z): [M-H]⁻ calcd. for C₅₃H₈₈NO₁₂P₂, 992.6; found 992.6.

Chemical synthesis of GalNAc-PP-Undecaprenyl-GalNAc-PP-Undecaprenyl was prepared using the general procedure as described above. However, the two purification steps were reversed as the final product proved too insoluble to obtain suitable spectral data. The crude product was thus purified via C-18 reverse phase column chromatography to afford Peracetylated GalNAc-PP-Undecaprenyl as a white solid (20 mg from 23 mg lipid phosphate, 59%). ¹H NMR (500 MHz, CD₃OD): δ 5.60 (dd, J = 3.2 Hz, J = 7.3 Hz, 1H), 5.42 (t, J = 6.1 Hz, 2H), 5.21 (dd, J = 3.2 Hz, J = 11.3 Hz, 1H), 5.14-5.05 (m, 10H), 4.58 (t, J = 6.9 Hz, 1H), 4.54-4.45 (m, 3H), 4.21 (dd, J = 8.1 Hz, J = 10.9 Hz, 1H), 4.04 (dd, J = 5.8 Hz, J = 10.8 Hz, 1H), 2.10 (s, 3H), 2.08-2.00 (m, 34H), 1.98 (s, 3H), 1.97 (s, 3H), 1.96-1.92 (m, 6H), 1.89 (s, 3H), 1.71 (s, 3H), 1.66-1.62 (m, 21H), 1.58 (s, 3H), 1.57 (s, 9H); ¹³C NMR (125 MHz, CD₃OD): δ 174.4, 172.3, 172.2, 171.9, 140.6, 136.5, 136.4, 136.3, 136.3, 136.1, 135.9, 132.1, 126.3, 126.3, 126.3, 126.0, 125.6, 125.6, 125.6, 123.6, 123.6, 96.5, 70.4, 68.8, 68.7, 63.8, 63.8, 62.3, 41.0, 41.0, 40.9, 34.4, 33.4, 33.4, 33.4, 33.2, 33.0, 30.8, 30.8, 30.7, 30.6, 30.5, 30.5, 30.4, 28.0, 27.8, 27.8, 27.8, 27.7, 27.7, 26.1, 24.0, 23.9, 23.8, 23.0, 20.8, 20.7, 17.9, 16.3, 14.5, 14.0; ³¹P NMR (162 MHz, CD₃OD): δ -9.7, -12.6; LRMS (*m*/z): [M-H]⁻ calcd. for C₆₉H₁₁₀NO₁₅P₂, 1254.7; found, 1254.7.

Final deprotection with 0.1% NaOCH₃ in CH₃OH and filtration through a short bed of C-18 reverse phase silica gel afforded GalNAc-PP-Undecaprenyl (18 mg, 99%). LRMS (m/z): [M-H]⁻ calcd. for C₆₃H₁₀₄NO₁₂P₂,1128.7; found, 1128.7.

*Chemical synthesis of GlcNAc-PP-Pentaprenyl-*GlcNAc-PP-Pentaprenyl was prepared in a pure form, and directly used without any purification after deacetylation of AcGlcNAc-PP-Pentaprenyl, which was purified by C-18 reverse phase column chromatography [Solvent A:

CH₃CN; Solvent B = 1.7% NH₄HCO₃; A/B = 0%→60% with 5% increment each time in concentration of A (10 mL each)]. Fraction containing product was lyophilized to afford pure AcGlcNAc-PP- Pentaprenyl as a white solid (14 mg from 40 mg lipid phosphate 21% yield). ¹H NMR (500 MHz, MeOD): δ 5.60 (br s, 1H), 5.48 (t, *J* = 6.5 Hz, 1H), 5.34 (t, *J* = 10.0 Hz, 1H), 5.17-5.09 (m, 5H), 4.56 (t, *J* = 6.5 Hz, 2H), 4.40-4.31 (m, 3H), 4.18 (dd, *J* = 1.5, 12.5 Hz, 1H), 2.16-1.97 (m, 8H), 2.07 (s, 3H), 2.01 (s, 3H), 2.00 (s, 3H), 1.97 (s, 3H), 1.76 (s, 3H), 1.70 (s, 3H), 1.68 (s, 3H), 1.63 (s, 3H), 1.61 (s, 3H); ¹³C NMR (125 MHz, MeOD): δ 172.6, 171.1, 170.5, 169.9, 139.7, 135.2, 134.7, 134.5, 130.7, 124.4, 124.1, 121.7, 121.6, 94.5, 71.5, 68.6, 68.3, 62.6, 61.4, 39.4, 31.9, 31.6, 26.4, 26.21, 26.19, 24.5, 22.3, 21.4, 19.30, 19.26, 19.25, 16.4, 14.8; ³¹P NMR (162 MHz, MeOD): δ -10.5, -13.4; LRMS (*m*/*z*): [M-H]⁻ calcd. for C₃₉H₆₂NO₁₅P₂, 846.3, found 846.3.

Final deprotection with catalytic amount of 0.1% NaOCH₃ in CH₃OH and subsequent addition of Amberlyst-15 ion exchange resin to the reaction mixture until around neutral pH (pH 7.0) afforded the GlcNAc-PP-Pentaprenyl in quantitative yield. LRMS (*m/z*): [M-H]⁻ calcd. for $C_{33}H_{56}NO_{12}P_2$, 720.2, found 720.2.

Chemical synthesis of GalNAc-PP-Undecyl-GalNAc-PP-Undecyl was prepared in a pure form, and directly used without any purification after deacetylation of AcGalNAc-PP-Undecyl, which was purified by C-18 reverse phase column chromatography [Solvent A: CH₃CN; Solvent B = 1.7% NH₄HCO₃; A/B = 0% \rightarrow 10% with 1% increment in concentration of A (10 mL each)]. Fraction containing product was lyophilized to afford pure AcGalNAc-PP-Undecyl as a white solid (19 mg from 30 mg lipid phosphate 30% yield). ¹H NMR (500 MHz, MeOD): δ 7.58 (s, 1H), 5.67-5.65 (m, 1H), 5.46 (d, *J* = 3.0 Hz, 1H), 5.24 (dd, *J* = 3.5, 11.0 Hz, 1H), 4.61-4.55 (m, 2H), 4.23 (dd, J = 3.0, 11.0 Hz, 1H), 4.09-4.06 (m, 1H), 4.03-3.99 (m, 2H), 2.15 (s, 3H), 2.023 (s, 3H), 2.021 (s, 3H), 1.95 (s, 3H), 1.70-1.64 (m, 2H), 1.41-1.31 (m, 16H), 0.92 (t, *J* = 7.0 Hz, 3H); ¹³C NMR (125 MHz, MeOD): δ 172.8, 170.8, 170.7, 170.5, 119.0, 95.1(d, *J* = 5.0 Hz), 68.8, 67.3, 67.2, 66.2, 66.1, 60.8, 31.7, 30.4, 30.3, 29.4, 29.12, 29.07, 25.5, 22.3, 21.5, 19.3, 19.22, 19.18, 13.0; ³¹P NMR (162 MHz, MeOD): δ -10.4 (d, *J* = 16.7 Hz), -13.2 (d, *J* = 16.2 Hz); LRMS (*m*/z): [M-H]⁻ calcd. for C₂₅H₄₅NO₁₅P₂, 660.5; found, 660.5.

Final deprotection with catalytic amount of 0.1% NaOCH₃ in CH₃OH and subsequent addition of Amberlyst-15 ion exchange resin to the reaction mixture until around neutral pH (pH 7.0) afforded the GalNAc-PP-Undecyl in quantitative yield. LRMS (m/z): [M-H]⁻ calcd. for C₁₉H₃₈NO₁₂P₂, 534.2; found 534.2.

ESI-MS ANALYSIS OF O-UNIT-PP-SUBSTRATES

Following the chemical synthesis of GalNAc-PP-lipid compounds, four glycosyl residues are successively ligated at the non-reducing end by well-characterized glycosyltransferases (WbnH, WbnJ, WbnK and WbnI), allowing for the synthesis of various lipids appended with the same oligosaccharide glycan moiety. ESI-MS was used to monitor the reaction for each of the enzymatic glycosylation steps. The ESI-MS spectra for all of the O-unit-PP-lipid substrates are shown in Fig. S2.

Primer name Sequ	lence		
WaaL _{R3} -NcoI-F	5-ATGCCCATGGCTACCTCAACATTATTTTTCTTTCTCGAG-3		
WaaL _{R3} -HindIII-R	5- ATGC <u>AAGCTT</u> TTAGTGGTGGTGGTGGTGGTGGTGGTGGTGGTGCTTGTTTT TCATCGCTAATA-3		
WaaL _{R3} -R209A-F	5'-TTTCTGATATTAGGAACTTTATCCGCCAGGGGCCTGGCTTTCC-3'		
WaaL _{R3} -R209A-R	5'-GGAAAGCCAGGCCCCTCCGGATAAAGTTCCTAATATCAGAAA-3'		
WaaL _{R3} -R268A-F	5- CTAACGTATAAACTTCAACAAACTAATAGTTCTTAT <u>GCA</u> TATGCAAATGGT ACTCAAGGC-3		
WaaL _{R3} -R268A-R	5- GCCTTGAGTACCATTTGCATA <u>TGC</u> ATAAGAACTATTAGTTTGTTGAAGTTT ATACGTTAG-3		
WaaL _{R3} -H319A-F	5- GACCTTTAGACAATCAATAGGGCCA <u>GCA</u> AATTTTGCGCTATTCATCTGGTT TG-3		
WaaL _{R3} -H319A-R	5- CAAACCAGATGAATAGCGCAAAATT <u>TGC</u> TGGCCCTATTGATTGTCTAAAG GTC-3		

Table S1. The oligonucleotide primers used for cloning and mutagenesis of waaL



Fig. S1. The general scheme for chemical synthesis of GalNAc/GlcNAc-pp-lipid compounds.



unit-PP-Undecyl. C, O-unit-PP-cis-Farnesyl. D, O-unit-PP-MS-Pentaprenyl. E, O-unit-PP-

Pentaprenyl. F, O-unit-PP-Heptaprenyl. G, O-unit-PP-Solanesyl. H, O-Unit-PP-Undecaprenyl.



Fig. S3. Western blotting analyses of in vitro ligation reaction samples using O-unit-PP-

Heptaprenyl donor.



 $\alpha\text{-}D\text{-}PerNAcp\text{-}(1\text{-}3)\text{-}\alpha\text{-}L\text{-}Fucp\text{-}(1\text{-}4)\text{-}\beta\text{-}D\text{-}Glcp\text{-}(1\text{-}3)\text{-}\alpha\text{-}D\text{-}GalNAc\text{-}1\text{-}PP\text{-}Und$



Fig. S4. Structures of O-units of E. coli O157 and O86

re_nneroozo	LSFGLSF IGIRTRITLI LSVVAAIVAITLILIINGAILILLEFVF ILFF IVNVKKIKFKGILV FIL	217
Te_ADU91200	GSLLSLFHLHGLKN-KKIFIGLGILGFVMGIFASILSGS <mark>R</mark> GGWIYVPMGFVLVSLFTFKSLKHSLLLIFSSFV	229
Hp_EGC72795	FSFVIAFHLLDVNKYKQKLVALVILFGIFGVLASLLSFARGGWIGVPILLLILIFLYRHLLSKKLLLGGITF	233
As_YP_001140043	LNWFVFQQRQLLRWIRYGALVALVAALFALYLSQSRGGWLALGGIVGYVICYKALFKPWKYIAIAM	226
Vc_AAL77359	LVLSCIYLLLNSQSKVLKLLASIGGVLSLAAIIMTDVRGVILFLPVVIIYLVITTIKLRWKYYVAL-TLS	226
Pd_ZP_06156631	AIMSLSFALASTDKRVKVITTVCFSFSCLALIIGESRGLWLALALTLFTFAIYSMIRFYQPKYWLYIMG	218
Br_CBW75858	LAFASLERDRKIWLGEWALKIIALLCGCYASYLSGARGAWIALPVLVWASMAGRHWLSNRWTAAGLV	233
Ec AAC69671	PSLLASILILKSDFRHKTTLYTINFMLSLCAVIVTETRAAILVFPFFALILIVMDSYINKRINYKL-YCF	242
Ec AAC69682	TTI YSSIVETKWKSKYKWEI EYANFAL SEAAVMMTGTRAALETEPI MTMVII EI OHRDOKVEI EKG-I SG	245
Pa CAC73082	TCAL SADAL I KI DSAVKVVI VICHELI VITATE TETPAATEVVETVCATILI SEVENKELETKACHELI VITATE TETPAATEVVETVCATI	255
Cm 7R 00055745	TALLAGATINI DI DITUNINA UNITATI CIA UNITATI ANA	200
Cy_2F_06355745.	IALLASQATINERERTIVSETEEMPELSEAVIIIIQITAALLVIPVESIGE-PEIMIKANKAMELKA-ELA	252
Ec_AAB18599(K12)) IGIVSGVAILYIKKNHPFLFLLNSCAVLYVLALIQI <mark>R</mark> AILLLFPIICVAALIAYYNKSPKKFISS-IVL	246
Ea_CBA19033	ISLVMMQSILMLKIRYRLGFYIAAFILSYSAIILTGTRAAMIAYPALILLTVIVTKNIITVRHKIAV-ILL	236
Ec_AAC69648	PALLNLWLIKKTSYKIAFVIFSAVFLFLLLGTLSRGAWLAVFIVTLLWLILNRQWKLLMLT-SIV	238
Se_AAS22256	PALLNIWLFRKTSLKLAFFALSAVYLFLMLGTLSRGAWLAVLVVGVLWAILNRQWKLMGIG-AAI	237
Se_NP_462613	PALLNIWLFRKNAIKLVFLVLSAIYLFFILGTLSRGAWLAVLIVGVLWAILNRQWKLIGVG-AIL	237
Ec AAC69661 (R3)	PALLNLWLIKSAKYRISEVVLSVIFIFLILGTLSRGAWLSVLVIGLIWILMFKOWKLLLVG-VMV	238
Xh VP 003470214	PVIICTWAIWKKHSIISWIIIITASAISIFIIIGTISRCAWVAIVVATVETTVINRFWKIIAA-TIC	242
Pm CAR46213	PITI ALWALKKNHSTINWILLS	240
Dm EFE51ACA	DI DI WU I DEVI TEVI	224
FF_EFE51464		234
Pa_AAG08384	TALWLAYWMQSKPILAPLPLISLALLGGLIAIGSRIPLVGLIAARMWL-VLAGDKKKALIALALA	241
Kp_BAH65745	GILICGMLLKEKASHWLYLPVVIMLVMLLLTQSRGPIIALVLAVGCILHLHVFIRRNLLIAAALA	206
VC_AAL77364	SIIFTLPFLLDRDSSKLFKLLFILFISSSVICLILSGSRAPILAIVISSFILILFTRPKVLLFSFFSL	226
Lb_YP_797859	PVFLFLRSLFDRNFKKAGIQGIILLSFLYVVFLNNARSSMLGAIFSSITAFLVLGIVRKELPSAKILLFASGIL	262
	*	
Vc AAL 76923		288
T- ADU01000		200
1e_AD091200		300
Hp_EGC72795	LCIISVVLIVNNKFVNRLSEAQYELKIYL-SGDNKVSSIGERLDMWIIGSKAFLEHPISGWSLKELDYY	301
As_YP_001140043	LCIASIGITYHTNQLVQLRVADAVSDLNFAEKGSYNSSWGL <mark>R</mark> VVAWQSAWLGFLDAPLTGVGTNGFDAL	295
Vc_AAL77359	VTVLSGVFYATFQSDINARIAQTQDEIALIKQGDLSSSIGI <mark>R</mark> LDLWMHGVEIIAQNPLFGVGDSGLQGS	295
Pd_ZP_06156631	LVILIGSMGIVLKPQLEARWQQTEQEYQAIQNGNMCTSIGL <mark>R</mark> LQMWQAASILVEDDPILGTGDAHIQK	286
Br_CBW75858	LLLIAALGALSTTNMVQERVSALESDIRKLSAGNTESSTGIRLELWKASLELYAEQPVFGVGKGRLKEG	302
Ec AAC69671	ITIALLAGVFSFKDTLLMRMNDLNNDLVNYSHDNTRTSVGARLAMYEVGLKTYSPI-GOSLEKRAEK	308
Ec_AAC69682	VETILLACCI TENKETDREINSI KADVISYA-TKNNSOSSVCAREAMVNACIKCSPDCENWOSI FORAFK	314
Pa_CAC73082	SSATULCE FLOCETHORANDI INDUSSY-SMNNSETSVCARIAMVOSCTRACFRALLCSARORSER	323
Cr 7P 06255745		220
Cy_2F_06355745.	PVLLVGLASIFLKSTIENKIQMMVDLNST SQNNSNISTGALAMQRAGIEAGKVNLVGQSLEQ-KGAE	320
EC_AAB18599 (K12)	LIAILASIVIIFNKPIQNRYNEALNDLNSYINANSVISLGA <mark>R</mark> LAMYEIGLNIFIKSPFSFRSAESRAES	315
Ea CBA19033	VPVLLTSTSLVEKDLVMSRTHDEERNTTATNDTEAENSVESRVWMOVVATRTCSEAPL-COSAEORASE	304
	THEETOTOETINDE ANTINI ENTINI INDIEAENOTION WAR TAINTO	001
Ec_AAC69648	ISVAAVGVFTYKGDHAGKDRLIYKLQQTDSSYRVTNGTQGTAWTLIMENPLKGYGYGDDIYHAI	302
Ec_AAC69648 Se_AAS22256	ISVAAVGVFTYKDQHAGKDRLIYKLQQTDSSYRVINGTQGTAWILIMENPLKGYGYGDDIYHAI LVIAGALVITQQIHKPNQDRLLYKLQQTDSSYRVINGTQGTAWILIQENPFKGYGYSNEVYDSI	302 301
Ec_AAC69648 Se_AAS22256 Se_NP_462613	ISVAAVGVFTYKIQDHAGKDRLIYKLQQTDSSYRVINGTQGTAWILIMENPLKGYGYGDDIYHAI LVIAGALVITQQIHKFNQDRLLYKLQQTDSSYRVINGTQGTAWILIQENPFKGYGYSNEVYDSI LAIIGALVITQHNNKPDPEHLLYKLQQTDSSYRVINGTQGTAWILIQENPIKGYGYGNDVYDGV	302 301 301
Ec_AAC69648 Se_AAS22256 Se_NP_462613 Ec_AAC69661(R3)	ISVAAVGVFTYKGDHAGKDRLIYKLQQTDSSYRVTNGTQGTAWILIMENPLKGYGYGDDIYHAI LVIAGALVITQQIHKPNQDRLLYKLQQTDSSYRVTNGTQGTAWILIQENPFKGYGYSNEVYDSI LAIIGALVITQHNNKPDPEHLLYKLQQTDSSYRVTNGTQGTAWILIQENPIKGYGYGNDVYDGV AIIALSVIFTHENPIKGYGYGNVAYKDV	302 301 301 209
Ec_AAC69648 Se_AAS22256 Se_NP_462613 Ec_AAC69661(R3) Xb_YP_003470214	ISVAAVGVFTYKGDHAGKDRLIYKLQQTDSSYRYTNGTQGTAWTLIMENPLKGYGYGDDIYHAI LVIAGALVITQQIHKPNQDRLLYKLQQTDSSYRYTNGTQGTAWILIQENPFKGYGYSNEVYDSI LAIIGALVITQHNNKPDPEHLLYKLQQTDSSYRYTNGTQGTAWILIQENPIKGYGYGNDVYDGV AIIALSVIFTHKEMTAKLTYKLQQTNSSYRYANGTQGSALDLILENPVIGYGGNVAYKDV LGLIAGSIHWSAVSNPATKILIHKLTQTDSSYRYSNGTQGSAWSLIMENPVKGYGFGNVYHOV	302 301 301 209 306
Ec_AAC69648 Se_AAS22256 Se_NP_462613 Ec_AAC69661(R3) Xb_YP_003470214 Pm_C4R46213	ISVAAVGVFTYKODHAGKDRLIYKLQQTDSSYRVINGTQGTAWILIMENPLKGYGYGDDIYHAI LVIAGALVITQQIHKPNQDRLLYKLQQTDSSYRVINGTQGTAWILIQENPFKGYGYSNEVYDSI LAIIGALVITQHNNKPDPEHLLYKLQQTDSSYRVINGTQGTAWILIQENPFKGYGYGNDVYDGV AIIALSVIFTH	302 301 301 209 306 303
Ec_AAC69648 Se_AAS22256 Se_NP_462613 Ec_AAC69661(R3) Xb_YP_003470214 Pm_CAR46213 Px_FEF51464	ISVAAVGVFTYKODHAGKDRLIYKLQQTDSSYRVINGTQGTAWILIMENPLKGYGYGDDIYHAI LVIAGALVITQQIHKPNQDRLLYKLQQTDSSYRVINGTQGTAWILIQENPFKGYGYSNEVYDSI LAIIGALVITQRPTKGYGYGNUYDGY AIIALSVIFTH	302 301 301 209 306 303 297
Ec_AAC69648 Se_AAS22256 Se_NP_462613 Ec_AAC69661(R3) Xb_YP_003470214 Pm_CAR46213 Pr_EFE51464 Be_AAC69224	ISVAAVGVFTYKEQDHAGKDRLIYKLQQTDSSYRVINGTQGTAWILIMENPLKGYGYGDDIYHAI LVIAGALVITQQIHKPNQDRLLYKLQQTDSSYRVINGTQGTAWILIQENPFKGYGYSNEVYDSI LAIIGALVITQREMTAKLTYKLQQTDSSYRVINGTQGTAWILIQENPIKGYGYGNDVYDGV AIIALSVIFTHKEMTAKLTYKLQQTNSSYRVANGTQGSALDLILENPVIGYGYGNVAYKDV LGLLAGSIHWSAVSNPATKLLLHKLTQTDSSYRVSNGTQGSALVSLIMENPVKGYGFGNKVYHQV VALFFGVTQLPNQFINAKLKHKLSQTNSGLRFDGGTQGSALDLILAPIKGYGYGNQLYHNV LSLVSLISIKL	302 301 301 209 306 303 297
Ec_AAC69648 Se_AAS22256 Se_NP_462613 Ec_AAC69661(R3) Xb_YP_003470214 Pm_CAR46213 Pr_EFE51464 Pa_AAG08384	ISVAAVGVFTYKIQDHAGKDRLIYKLQQTDSSYRVINGTQGTAWILIMENPLKGYGYGDDIYHAI LVIAGALVITQQIHKPNQDRLLYKLQQTDSSYRVINGTQGTAWILIQENPLKGYGYGNEVYDSI LAIIGALVITQNNKPDPEHLLYKLQQTDSSYRVINGTQGTAWILIQENPIKGYGYGNDYDGV AIIALSVIFTHKEMTAKLTYKLQQTNSSYRVANGTQGSALDLILENPVIGYGYGNVAYKDV LGLLAGSIHWSKEMTAKLTYKLQQTNSSYRVSNGTQGSALDLILENPVIGYGYGNVAYKDV LGLLAGSIHWS	302 301 301 209 306 303 297 293
Ec_AAC69648 Se_AAS22256 Se_NP_462613 Ec_AAC69661(R3) Xb_YP_003470214 Pm_CAR46213 Pr_EFE51464 Pa_AAG08384 Kp_BAH65745	ISVAAVGVFTYKODHAGKDRLIYKLQQTDSSYRVINGTQGTAWILIMENPLKGYGYGDDIYHAI LVIAGALVITQQIHKPNQDRLLYKLQQTDSSYRVINGTQGTAWILIQENPFKGYGYGNDVYDSI LAIIGALVITQRDPFKGYGYGNDVYDGV AIIALSVIFTH	302 301 301 209 306 303 297 293 263
Ec_AAC69648 Se_AAS22256 Se_NP_462613 Ec_AAC69661(R3) Xb_YP_003470214 Pm_CAR46213 Pr_EFE51464 Pa_AAC08384 Kp_BAH65745 VC_AAL77364	ISVAAVGVFTYKODHAGKDRLIYKLQQTDSSYRVINGTQGTAWILIMENPLKGYGYGDDIYHAI LVIAGALVITQQIHKPNQDRLLYKLQQTDSSYRVINGTQGTAWILIQENPFKGYGYGNDVYDSI LAIIGALVITQRPIKGYGYGNDVYDGV AIIALSVIFTH	302 301 301 209 306 303 297 293 263 295
Ec_AAC69648 Se_AAS22256 Se_NP_462613 Ec_AAC69661(R3) Xb_YP_003470214 Pm_CAR46213 Pr_EFE51464 Pa_AAG08384 Kp_BAH65745 VC_AAL77364 Lb_YP_797859	ISVAAVGVFTYKEQHAGKDRLIYKLQQTDSSYRVINGTQGTAWILIMENPLKGYGYGDDIYHAI LVIAGALVITQQIHKPNQDRLLYKLQQTDSSYRVINGTQGTAWILIQENPFKGYGYSNEVDSI LAIIGALVITQREMTAKLTYKLQQTDSSYRVINGTQGTAWILIQENPIKGYGYGNDVYDGV AIIALSVIFTHKEMTAKLTYKLQQTDSSYRVINGTQGSALDLILENPIKGYGYGNVAYKDV LGLLAGSIHWSAVSNPATKLLLHKLTQTDSSYRVINGTQGSALDLILENPVKGYGFGNKVYHQV VALFFGVTQLPNNQFINAKLKHKLSQTNSGLRFDGGTQGSALDLILAQPIKGYGYGNQLYHNV LSLVSLISIKL	302 301 301 209 306 303 297 293 263 295 323
Ec_AAC69648 Se_AAS22256 Se_NP_462613 Ec_AAC69661(R3) Xb_YP_003470214 Pm_CAR46213 Pr_EFE51464 Pa_AAG08384 Kp_BAH65745 VC_AAL77364 Lb_YP_797859	ISVAAVGVFTYKODHAGKDRLIYKLQQTDSSYRVINGTQGTAWTLIMENPLKGYGYGDDIYHAI LVIAGALVITQQIHKPNQDRLLYKLQQTDSSYRVINGTQGTAWILIQENPFKGYGYGNDVYDGV AIIALSVIFTH	302 301 301 209 306 303 297 293 263 295 323
Ec_AAC69648 Se_AAS22256 Se_NP_462613 Ec_AAC69661(R3) Xb_YP_003470214 Pm_CAR46213 Pr_EFE51464 Pa_AAG08384 Kp_BAH65745 VC_AAL77364 Lb_YP_797859	ISVAAVGVFTYKODHAGKDRLIYKLQQTDSSYRVINGTQGTAWILIMENPLKGYGYGDDIYHAI LVIAGALVITQQIHKPNQDRLLYKLQQTDSSYRVINGTQGTAWILIQENPFKGYGYGNDVYDGV AIIALSVIFTH	302 301 301 209 306 303 297 293 263 295 323
Ec_AAC69648 Se_AAS22256 Se_NP_462613 Ec_AAC69661(R3) Xb_YP_003470214 Pm_CAR46213 Pr_EFE51464 Pa_AAC08384 Kp_BAH65745 VC_AAL77364 Lb_YP_797859	ISVAAVGVFTYKDQHAGKDRLIYKLQQTDSSYRVINGTQGTAWLIIMENPLKGYGYGDDIYHAI LVIAGALVITQQIHKPNQDRLLYKLQQTDSSYRVINGTQGTAWLLIQENPFKGYGYGNDVYDSI LAIIGALVITQQIHKPNQDRLLYKLQQTDSSYRVINGTQGTAWLLIQENPFKGYGYGNDVYDGV AIIALSVIFTH	302 301 301 209 306 303 297 293 263 295 323
Ec_AAC69648 Se_AAS22256 Se_NP_462613 Ec_AAC69661(R3) Xb_YP_003470214 Pm_CAR46213 Pr_EFE51464 Pa_AAC08384 Kp_BAH65745 VC_AAL77364 Lb_YP_797859 Vc_AAL76923 Te_ADI91200	ISVAAVGVFTYKODHAGKDRLIYKLQQTDSSYRVINGTQGTAWTLIMENPLKGYGYGDDIYHAI LVIAGALVITQQIHKPNQDRLIYKLQQTDSSYRVINGTQGTAWTLIQENPFKGYGYSNEVYDSI LAIIGALVITQQIHKPNQDRLLYKLQQTDSSYRVINGTQGTAWILIQENPFKGYGYGNDVYDGV AIIALSVIFTH	302 301 301 209 306 303 297 293 263 295 323 359 373
Ec_AAC69648 Se_AAS22256 Se_NP_462613 Ec_AAC69661 (R3) Xb_YP_003470214 Pm_CAR46213 Pr_EFE51464 Pa_AAG08384 Kp_BAH65745 VC_AAL77364 Lb_YP_797859 Vc_AAL76923 Te_ADU91200 H= E6C72705	ISVAAVCVFTYKODHAGKDRLIYKLQQTDSSYRVTNGTQGTAWTLIMENPLKGYCGYGDDIYHAI LVIAGALVITQQIHKPNQDRLLYKLQQTDSSYRVTNGTQGTAWTLIQENPLKGYCGYGDDIYHAI LVIAGALVITQQIHKPNQDRLLYKLQQTDSSYRVTNGTQGTAWILIQENPLKGYCGYGNDVYDGV AIIALSVIFTH	302 301 301 209 306 303 297 293 263 295 323 359 373
Ec_AAC69648 Se_AAS22256 Se_NP_462613 Ec_AAC69661(R3) Xb_YP_003470214 Pm_CAR46213 Pr_EFE51464 Pa_AAG08384 Kp_BAH65745 VC_AAL77364 Lb_YP_797859 Vc_AAL76923 Te_ADU91200 Hp_EGC72795	NIELGYGVGVFTYKODHAGKDRLIYKLQQTDSSYRVINGTQGTAWTLIMENPLKGYGYGDDIYHAI LVIAGALVITQQIHKPNQDRLLYKLQQTDSSYRVINGTQGTAWILIQENPFKGYGYGNDVYDGV AIIALSVIFTH	301 301 209 306 303 297 293 263 295 323 359 373 374
Ec_AAC69648 Se_AAS22256 Se_NP_462613 Ec_AAC69661(R3) Xb_YP_003470214 Pm_CAR46213 Pr_EFE51464 Pa_AAG08384 Kp_BAH65745 VC_AAL77364 Lb_YP_797859 Vc_AAL76923 Te_ADU91200 Hp_EGC72795 As_YP_001140043	NIELFKEGKVDEWTSTVPRGHALSQYFEAIASNTLGILAIFAMLILPFGVFLNDYRKTGSPISQTGYLFA	301 301 209 306 303 297 293 263 295 323 359 373 374 361
Ec_AAC69648 Se_AAS22256 Se_NP_462613 Ec_AAC69661(R3) Xb_YP_003470214 Pm_CAR46213 Pr_EFE51464 Pa_AAC08384 Kp_BAH65745 VC_AAL77364 Lb_YP_797859 Vc_AAL76923 Te_ADU91200 Hp_EGC72795 As_YP_001140043 Vc_AAL77359	NIELGIGUINELEVADINELTALINITATIONALISE AND ALARS AND ALAR	301 301 209 306 303 297 293 263 295 323 359 373 374 361 357
Ec_AAC69648 Se_AAS22256 Se_NP_462613 Ec_AAC69661(R3) Xb_YP_003470214 Pm_CAR46213 Pr_EFE51464 Pa_AAG08384 Kp_BAH65745 VC_AAL77364 Lb_YP_797859 Vc_AAL76923 Te_ADU91200 Hp_EGC72795 As_YP_001140043 Vc_AAL77359 Pd_ZP_06156631	ISVAAVGVFTYKDDHAGKDRLIYKLQQTDSSYRVTNGTQGTAWTLIMENPLKGYGYGDDIYHAI LVIAGALVITQQIHKPNQDRLLYKLQQTDSSYRVTNGTQGTAWTLIMENPLKGYGYGDDYYHAI LVIAGALVITQQIHKPNQDRLLY	301 301 209 306 303 297 293 263 295 323 359 373 374 361 357 353
Ec_AAC69648 Se_AAS22256 Se_NP_462613 Ec_AAC69661(R3) Xb_YP_003470214 Pm_CAR46213 Pr_EFE51464 Pa_AAG08384 Kp_BAH65745 VC_AAL77364 Lb_YP_797859 Vc_AAL76923 Te_ADU91200 Hp_EGC72795 As_YP_001140043 Vc_AAL77359 Pd_ZP_06156631 Br_CBW75858	NIELFKEGKVDEWTSTVPRGHARSQYFEAIASNUTLGILAIFAMLILPFGVFLNDYRKTGSPISQTGYLFA NIELFKEGKVDEWTSTVPRGHARSQYFEAIASNUTLGILAIFAMLILPFGVFLNDYRKTGSPSQTGYLFA NIELFKCGRUPELAINAQFINSYFEAIASNUTLGILAIFAMLILPFGVFLNDYRKTGSPSQTGYLFA NIELFKCGRUPELAINAQFINSYFEAIASNUTLGILAIFAMLILPFGVFLNDYRKTGSP	302 301 301 209 306 303 297 293 263 295 323 373 374 353 374 361 353 373
Ec_AAC69648 Se_AAS22256 Se_NP_462613 Ec_AAC69661(R3) Xb_YP_003470214 Pm_CAR46213 Pr_EFE51464 Pa_AAG08384 Kp_BAH65745 VC_AAL77364 Lb_YP_797859 Vc_AAL76923 Te_ADU91200 Hp_EGC72795 As_YP_001140043 Vc_AAL77359 Pd_ZP_06156631 Br_CBW75858 Ec_AAC69671	NIELFKEGKVDEWTSTVPRGHALSQYFEAIASNTLGILAIFAMLILPFGVFLNDYRKTGSPSQTGYLFA MLLVLGIVLSFTQGHSQVFRGHALSQYFEAIASNTLGILAIFAMLILPFGVFLNDYRKTGSPSQTGYLFA MLLVLGIVLSFT	302 301 301 209 306 303 297 293 263 295 323 373 374 357 353 373 373
Ec_AAC69648 Se_AAS22256 Se_NP_462613 Ec_AAC69661 (R3) Xb_YP_003470214 Pm_CAR46213 Pr_EFE51464 Pa_AAG08384 Kp_BAH65745 VC_AAL77364 Lb_YP_797859 Vc_AAL76923 Te_ADU91200 Hp_EGC72795 As_YP_001140043 Vc_AAL77359 Pd_ZP_06156631 Br_CBW75858 Ec_AAC69662	NIELFKEGKVDEWTSTVPRGHAFSQYFEAIASNTILGILAIFAMLILPFGVFLNDYRKTGSPSQFWLGRGFSY-ELDF CLLFFSLQKTDFGITAQNRLESIVEIHFGQYFEAIASNTILGILAIFAMLILPFGVFLNDYRKTGSPSQTGYLFA MLLVLGIVLSFTQVGQKITDPLFGKEKHTDSGRTFIWDSTFPLIRDNFFAGVGPGN	302 301 301 209 306 303 297 293 263 295 323 373 374 361 357 373 373 373 373 373
Ec_AAC69648 Se_AAS22256 Se_NP_462613 Ec_AAC69661 (R3) Xb_YP_003470214 Pm_CAR46213 Pr_EFE51464 Pa_AAG08384 Kp_BAH65745 VC_AAL77364 Lb_YP_797859 Vc_AAL77364 Lb_YP_797859 Vc_AAL77359 Pd_ZP_001140043 Vc_AAL77359 Pd_ZP_06156631 Br_CBW75858 Ec_AAC69671 Ec_AAC6962 Pa_C4C73082	NIELEKEGKVDEWTSTVPRGHALSQYFEAIASN TLGILAIFAMLILPFGVFLNDYRKTGSPISQTGYLFA NIELFKCHISQKVYKHRHPINEYLNEFVKNGIGGIALSIFINWGSGLEFFKFKNARSNYSSEVKLPAGIGAIFV KKDADRGLVPPLALAAAAASQYMQNLVIRGIGGIALSIFINWGSGLACVLIP SKMTN	302 301 301 209 306 303 297 293 293 293 295 323 373 373 374 351 357 353 373 373 373 373
Ec_AAC69648 Se_AAS22256 Se_NP_462613 Ec_AAC69661(R3) Xb_YP_003470214 Pm_CAR46213 Pr_EFE51464 Pa_AAG08384 Kp_BAH65745 VC_AAL77364 Lb_YP_797859 Vc_AAL77364 Lb_YP_797859 Vc_AAL76923 Te_ADU91200 Hp_EGC72795 As_YP_001140043 Vc_AAL77359 Pd_ZP_06156631 Br_CBW75858 Ec_AAC69671 Ec_AAC69682 Pa_CAG73082 Cv_ZP_06355745	NIELFKBGKVDEWTSTVPRGHAKSQYFEAIASNTILGILAIFAMLILPFGVFLNDYRKTGSPISQTGYLFA NIELFK	302 301 301 209 306 297 293 263 295 323 373 374 361 357 353 373 373 378 388 384
Ec_AAC69648 Se_AAS22256 Se_NP_462613 Ec_AAC69661(R3) Xb_YP_003470214 Pm_CAR46213 Pr_EFE51464 Pa_AAG08384 Kp_BAH65745 VC_AAL77364 Lb_YP_797859 Vc_AAL76923 Te_ADU91200 Hp_EGC72795 As_YP_001140043 Vc_AAL77359 Pd_ZP_06156631 Br_CBW75858 Ec_AAC69671 Ec_AAC69682 Pa_CAG73082 Cy_ZP_06355745.	NIELFKEGKVDEWTSTVPRGHALSQYFEAIASNTLGILAIFAMLILPFGVFLNDYRKTGSPSQTGYLFA MLLVLGIVLSFTQGRIKCLYQQDESYFAIASNTLGILAIFAMLILPFGVFLNDYRKTGSP	302 301 301 209 303 297 293 263 295 323 373 374 361 357 353 373 373 373 373 373 373 373 373
Ec_AAC69648 Se_AAS22256 Se_NP_462613 Ec_AAC69661 (R3) Xb_YP_003470214 Pm_CAR46213 Pr_EFE51464 Pa_AAG08384 Kp_BAH65745 VC_AAL77364 Lb_YP_797859 Vc_AAL77364 Lb_YP_797859 Vc_AAL77359 Pd_ZP_001140043 Vc_AAL77359 Pd_ZP_06156631 Br_CBW75858 Ec_AAC69682 Pa_CAG73082 Cy_ZP_06355745. Ec_AAB18599 (K12)	NIELFKCRUPLAGKDELIYKLQQTDSSYRVINGTQGTAWILIQENPLKGYGYGDDIYHAI LVIAGALVITQQIHKPNQDRLLYKLQQTDSSYRVINGTQGTAWILIQENPLKGYGYGDDIYHAI LVIAGALVITQ	302 301 301 209 306 303 297 293 263 295 323 373 374 361 357 373 373 373 373 373 373 373 378 388 38
Ec_AAC69648 Se_AAS22256 Se_NP_462613 Ec_AAC69661 (R3) Xb_YP_003470214 Pm_CAR46213 Pr_EFE51464 Pa_AAG08384 Kp_BAH65745 VC_AAL77364 Lb_YP_797859 Vc_AAL77364 Lb_YP_797859 Vc_AAL77359 Pd_ZP_001140043 Vc_AAL77359 Pd_ZP_06156631 Br_CBW75858 Ec_AAC69671 Ec_AAC69671 Ec_AAC6962 Pa_CAG73082 Cy_ZP_06355745. Ec_AAB18599 (K12) Ea_CBA19033	NIVELIGISUARGURING LEMENTAL TARTAL TARTAL AND THE AND	302 301 301 209 306 297 293 263 295 323 295 323 373 374 361 357 357 357 357 357 357 357 357 357 357
Ec_AAC69648 Se_AAS22256 Se_NP_462613 Ec_AAC69661 (R3) Xb_YP_003470214 Pm_CAR46213 Pr_EFE51464 Pa_AAG08384 Kp_BAH65745 VC_AAL77364 Lb_YP_797859 Vc_AAL77364 Lb_YP_797859 Vc_AAL77359 Vc_AAL77359 Pd_ZP_06156631 Br_CBW75858 Ec_AAC69671 Ec_AAC69682 Pa_CAG73082 Cy_ZP_06355745. Ec_AAB18559 (K12) Ea_CBA19033 Ec_AAC69648	<pre>NINDINGLY AND AND AND AND AND AND AND AND AND AND</pre>	302 301 301 209 303 297 293 263 295 323 295 323 373 374 361 357 353 373 373 373 373 378 388 384 379 369 371
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Fig. S5. Multiple sequence alignment analysis of WaaL homologues.