Supplementary information

A novel exopolysaccharide (p-CY01) from the Antarctic bacterium *Pseudoalteromonas* sp. strain CY01 cryopreserves human red blood cells

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Strains

Supplementary Figure S1. Exopolysaccharide (EPS) production and cryoprotective effect on *E. coli*. EPS production was estimated by determining the *E. coli* cell survival ratio after three freeze-thaw cycles in the presence of 0.2% (w/v) crude EPS solutions from the screened strains. The white bar represents EPS production (g/L), and the black bar represents the cryoprotective effect of EPS on *E. coli* (%). The values presented are the mean \pm SD from three experimental repeats.



Supplementary Figure S2. The neighbor-joining phylogenetic tree of 16S rRNA sequences. The neighbor-joining tree illustrates the phylogenetic relationship between strain CY01 and closely related *Pseudoalteromonas* strains based on their 16S rRNA sequences. Numbers on nodes indicate the boostrap analysis with 1,000 sequence replicates.



Supplementary Figure S3. Homogeneity analysis of purified EPS from strain CY01 using an Agilent HPLC system. The EPS was dissolved in 0.1M NaCl solution and prepared as 0.1% (w/v) aqueous solutions. Each run process involved injecting 5 μ L of the sample solution and monitoring it for 70 min at a flow rate of 0.4 mL min⁻¹ at 40°C. The sample was analyzed using an Agilent refractive index detector.



Supplementary Figure S4. 1 H and 13 C NMR (600 and 150 MHz) spectra of p-CY01 in D₂O (a; 1 H NMR, b; 13 C NMR)



Supplementary Figure S5 Selected parts of HSQC (a) and (b), TOCSY (c), COSY (d), and HMBC (e) and (f) NMR (600 MHz, in D₂O) spectra of p-CY01. The ¹H NMR spectrum of p-CY01 showed anomeric proton signals at $\delta_{\rm H}$ 4.54 (d, ${}^{3}J_{1,2} = 8.4$ Hz) and at $\delta_{\rm H}$ 4.95 (br s) to 5.23 (br s) ppm, indicating a β -glycosidical (${}^{3}J_{1,2} \approx 8.0$ Hz) and α -glycosidical (${}^{3}J_{1,2} = < 3.0$ Hz) configurations, respectively. The NMR

and ¹H-¹³C HSQC spectra of p-CY01 displayed five characteristic sugar residues at $\delta_{\rm H}$ 4.54/ $\delta_{\rm C}$ 102.4, 4.96/99.7, 5.12/97.9, 5.21/102.4, and 5.26/97.2 ppm, labeled as **A**, **B**, **C**, **D**, and **E**, respectively (Supplementary Table S1 and Supplementary Fig. S3), along with free sugar units at $\delta_{\rm H}$ 4.66/ $\delta_{\rm C}$ 95.8 and 5.25/91.8, respectively, corresponding to the signals of β -D-Glcp/-D-Galp and α -D-Glcp/-D-Galp. The ¹H-¹³C HSQC, ¹H-¹H TOCSY, including a selective TOCSY method, and ¹H-¹H COSY experiments established their related proton and carbon residues (Supplementary Table 1), assigned as A: \rightarrow 4)- β -D-Glcp-(1 \rightarrow , [^{1,2}] **B**: Terminal- α -D-Glcp-(1 \rightarrow , [³⁻⁵] **C**: \rightarrow 6)- α -D-Galp-(1 \rightarrow , [⁶] **D**: \rightarrow 3,4)- α -D-Galp-(1 \rightarrow , [^{7,8}] and **E**: \rightarrow 4,6)- α -D-Glcp-(1 \rightarrow , [⁹] respectively (Supplementary Table S1 and Supplementary Fig. S4). These were in good agreement with the results of the GC-MS analysis. However, signals for the remaining terminal-galactose unit were hardly distinguishable in the 1D and 2D NMR analyses. The ¹H-¹³C HMBC and ¹H-¹H NOESY analyses support linkages between sugar residues since the methods display clear and strong correlations between sugar units. In the residue **A**, a major backbone of this polysaccharide, a strong correlation peak from the anomeric proton (H-1) to the downfield shifted carbon signal at $\delta_{\rm C}$ 78.6 (C-4) and weak cross peaks with C-2/C-3 were observed in HMBC spectrum, indicating a main repeating unit of p-CY01 as shown in Supplementary Fig. S4 and Fig. S5[.][1,2]



Supplementary Figure S6 Structure of the predominant repeating units and proposed partial structures of the *O*-polysaccharide of p-CY01.



Supplementary Figure S7. Confirmation of interaction between p-CY01 LM and free hemoglobin resulting in precipitated hemoglobin. (A) No decrease in hemoglobin pigment was observed based on p-CY01 LM concentration when compared to PBS. (B) Quantitative measurement of the free hemoglobin ratio using Drabkin's method revealed that even with an increase in hemoglobin concentration, precipitated hemoglobin was not generated when using p-CY01 LM. The value remained comparable to the free hemoglobin ratio in PBS. The presented values represent means \pm standard deviation from three experimental replicates.



Supplementary Figure S8. Biocompatibility of p-CY 01 LM solution (2.5% (w/v) p-CY 01 LM with 1% (v/v) Glycerol and DMSO in ADSOL) on the RBC. RBC integrities after room temperature incubation for an indicated time without freezing. Haemolysis was quantified using Drabkin's assay to determine % RBC integrity (100 - % hemolysis). The values shown are means \pm standard deviation from three experimental replicates.



Supplementary Figure S9. Differential scanning calorimetry (DSC) analysis of p-CY01 LM solutions. Samples were individually cooled from +5 °C to -78 °C at a rate of 40 °C min⁻¹ and thawing at 2°C min⁻¹. (A) 1% Glycerol (v/v) and 1% DMSO (v/v) were used as a negative control. (B) 1% Glycerol (v/v), 1% DMSO (v/v) and 0.5% p-CY01 LM (w/v). (C) 1% Glycerol (v/v), 1% DMSO (v/v) and 2.5% p-CY01 LM (w/v). The supercooling points of (A) 1% Glycerol (v/v) and 1% DMSO (v/v) were used as a negative control, (B) 1% Glycerol (v/v), 1% DMSO (v/v) and 0.5% p-CY01 LM (w/v) and 0.5% p-CY01 LM (w/v) and 1% DMSO (v/v) and 1% DMSO (v/v) and 2.5% p-CY01 LM (w/v) and 1% DMSO (v/v) and 1% DMSO (v/v) and 0.5% p-CY01 LM (w/v) and

Residues	H-1	H-2	H-3	H-4	H-5	H-6
	C-1	C-2	C-3	C-4	C-5	C-6
A : →4)-β-D-Glcp-	4.54	3.35	3.64	3.66	3.64	3.67
(1→	102.4	73.7	75.0	78.6	77.0	60.0
B : T-α-D-Glc <i>p</i> - $(1 \rightarrow$	4.96	3.61	3.69	3.87	3.93	3.62
	99.7	73.2	70.6	71.2	69.0	60.0
C: →6)- α -D-Galp-(1→	5.12	4.38	3.96	4.18	3.98	3.83
	97.9	74.4	68.7	69.3	69.2	67.6
D : →3,4)- α -D-Gal <i>p</i> -(1→	5.21	4.16	4.48	3.84	4.44	-
	102.4	70.5	81.5	75.8	70.5	-
E: →4,6)-α-D-	5.26	3.60	3.98	3.80	3.96	3.84
Glc <i>p</i> -(1→	97.2	71.2	72.6	75.2	70.2	68.9

Supplementary Table 1. ¹H and ¹³C NMR (600 and 150 MHz, in D₂O) data for the residues **A** to **E** from p-CY01.

Variables	Effect	S.E.	t-statistics	P-value
Glycerol	2.321	0.9096	1.28	0.224
DMSO	2.621	0.9096	1.44	0.173
ADSOL	5.826	0.9096	3.20	0.007
p-CY01 LM	51.694	0.9096	28.41	0.000

Supplementary Table 2. Statistical analysis of cryo-additives using Plackett-Burman design.

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