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Antithrombin Stabilisation by Sulfated Carbohydrates Correlates with Anticoagulant Activity

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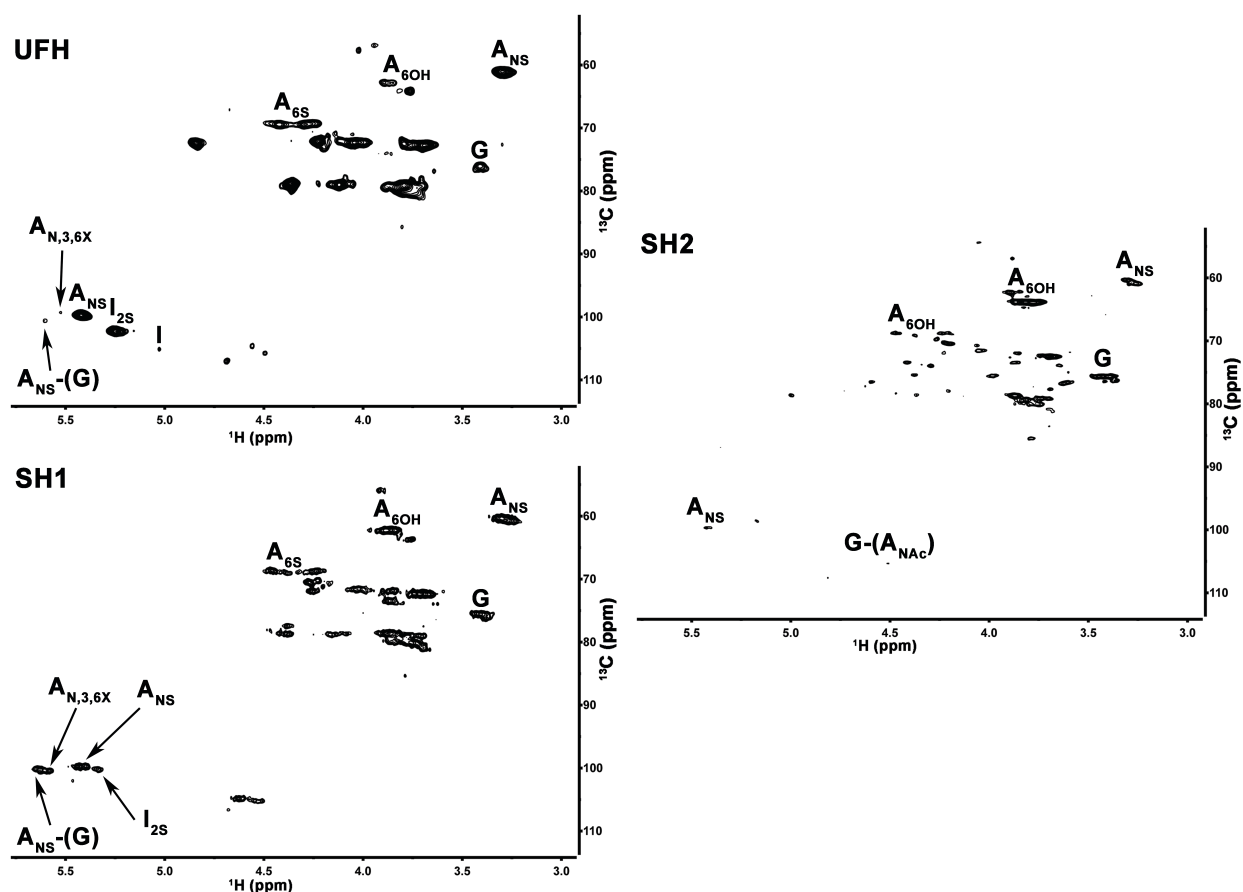
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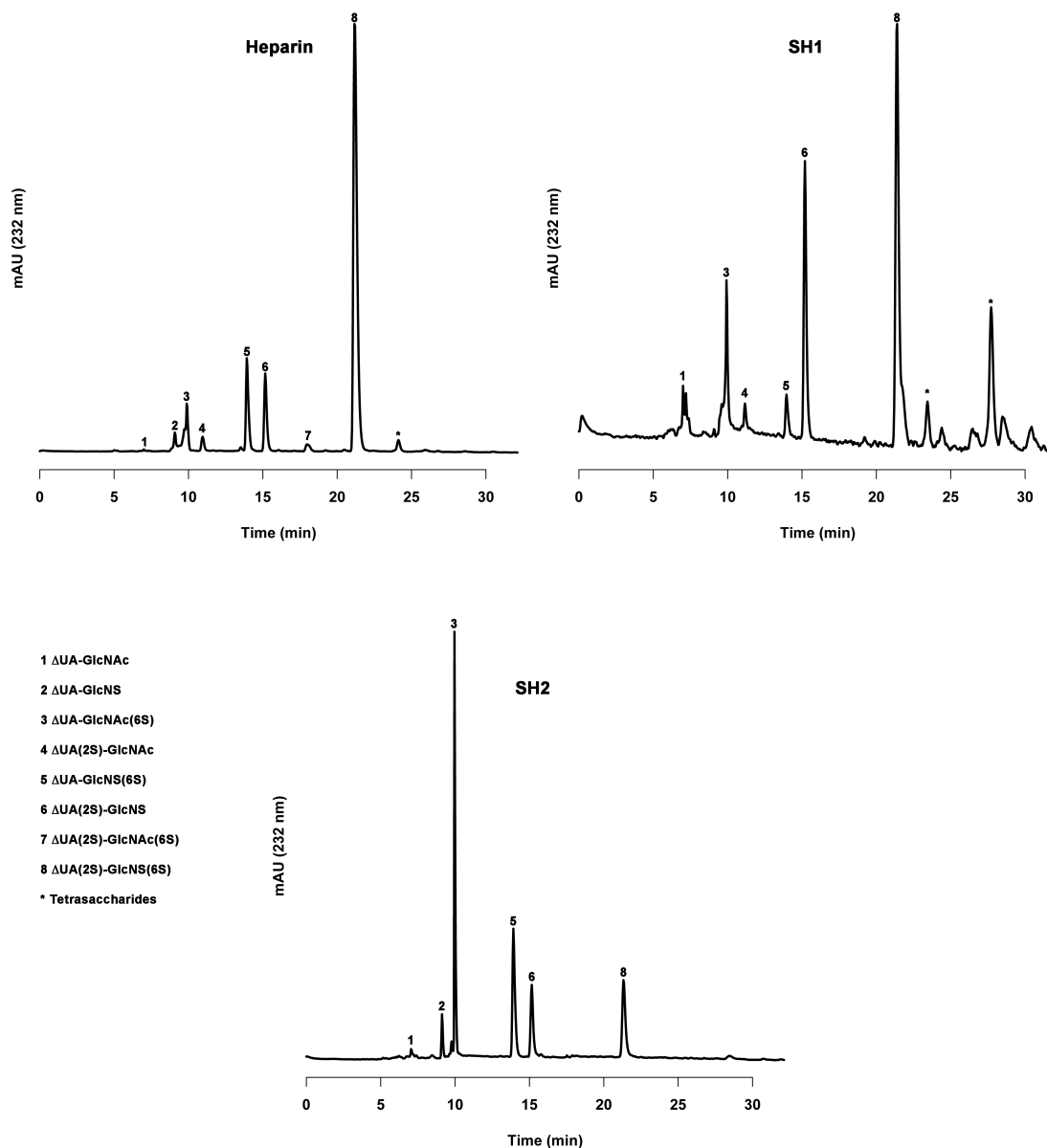
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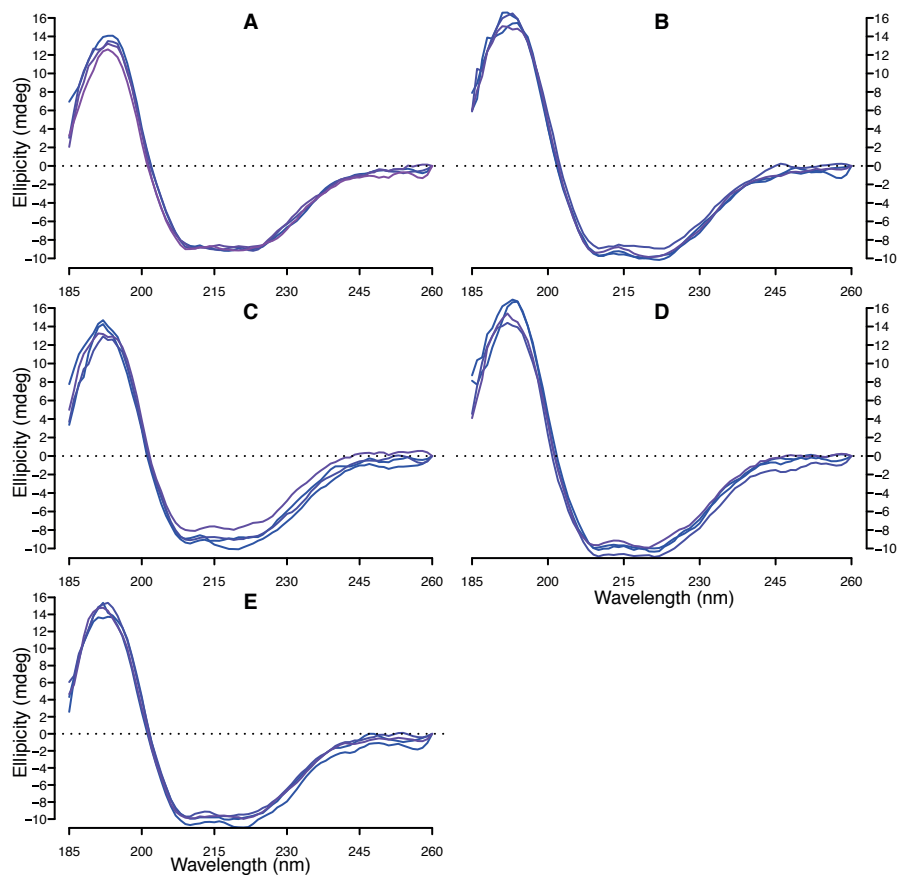


Supplementary figure 1. ^1H - ^{13}C HSQC spectra (600 MHz) of unfractionated heparin (UFH, upper left), and Shrimp heparinoids-1 (SH-1, upper right) and -2 (SH-2, bottom) from the Pacific White Shrimp, *L.vannemei*. Signals from selected structural features are labelled; A_{NS} indicates 2-deoxy-2-sulfamino α -D-glucopyranose; $\text{I}_{2\text{S}}$, 2-O-sulfo- α -L-iduronic acid; G, β -D-glucuronic acid; I, α -L-Iduronic acid; $\text{A}_{3\text{S}}$, 2-deoxy-3-O-sulfo-2-amino α -D-glucopyranose; A_{NAC} , 2-deoxy-2-acetyl amino α -D-glucopyranose. $\text{A}_{\text{NS},3\text{S},6\text{X}}$ The substitution at position -6 has not been determined.

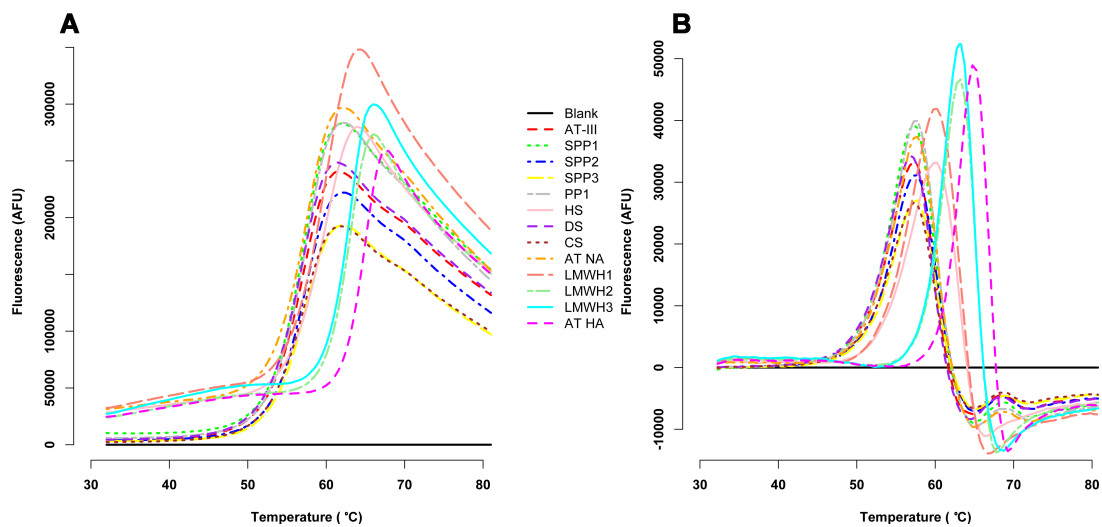


Polysaccharide	Disaccharides (%)							
	Δ UA-NAc	Δ UA-NS	Δ UA-NAc(6S)	Δ UA(2S)-NAc	Δ UA-NS(6S)	Δ UA(2S)-NS	Δ UA(2S)-NAc(6S)	Δ UA(2S)-NS(6S)
Heparin	0.14	2.07	6.65	1.69	10.33	8.73	1.33	69.07
SH1	9.31	0.00	13.04	1.86	3.45	21.79	0.00	50.56
SH2	1.59	4.83	35.02	0.00	24.75	14.42	0.00	19.39

Supplementary figure 2. SAX-HPLC disaccharide analysis of unfractionated heparin (heparin, upper left), Shrimp heparinoid-1 (SH1, upper right) and Shrimp heparinoid-2 (SH2, bottom) from the Pacific White Shrimp, *L.vannemei*, and their disaccharide composition analyses (Table). 100 mg of each sample was digested using a mixture of heparin lyases (2.5 mIU each) and analysed on a PhenoSphere™ 5 μ m SAX 80 Å LC Column 150 x 4.6 mm.



Supplementary figure 3. SRCD spectra of AT and AT complexes individual scans used for Figure 1 and 2. A, AT; B, AT:Pentasaccharide; C, AT:SH1; D, AT:SH2 and E, AT:UFH.



Supplementary figure 4. Determination of the melting temperature of AT alone and AT bound to a range of polysaccharides via the Fluorescence shift assay. **A.** Raw data and **B.** First derivative. These melting temperatures are compared to the anticoagulant activity of the polysaccharide in supplementary table 2.

Supplementary table 1. Statistics on secondary structure determination for AT:pentasaccharise and AT:SH2 (Supplementary Figure 4 B and D), illustrating their similar secondary structure content. Numbers in brackets are standard deviations.

	AT:Pentasacchride	AT:SH2
α -helix	85.8 (1.3)	87.3 (1.6)
Antiparallel β -sheet	0.1 (0.1)	0.1 (0.1)
Parallel β -sheet	1.8 (0.2)	1.5 (0.1)
β -turn	8.7 (0.2)	8.8 (0.4)
Unordered	6.3 (0.1)	4.8 (0.6)

Supplementary table 2. List of polysaccharides employed in Figure 2, their melting temperature changes relative to AT and their associated literature references. ^amanufacturers values of activity.

Polysaccharide	Anti-Xa (IU/mg)	T _m (°C) change (from AT)	Reference
SPP 1	≤5	None	1,2
SPP 2	≤5	None	1,2
SPP 3 ^a	≤5	None	
PP 1 ^a	≤5	None	
HS	~10	+2.6	3
DS	≤5	-1.3	3
CS	≤5	None	
AT NA	≤5	None	4,5,6
LMWH1 ^a	100-120	+6.6	
LMWH2 ^a	80-120	+8.0	
LMWH3 ^a	140-160	+9.3	
AT HA	~400	+12.0	4,5,6

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