

SUPPLEMENTARY INFORMATION

Screening ionic liquids for dissolving hemicellulose by COSMO-RS based on the selective model

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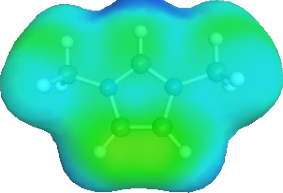
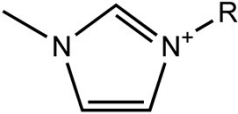
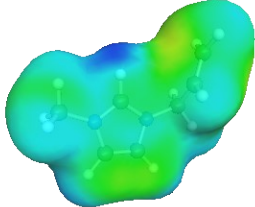
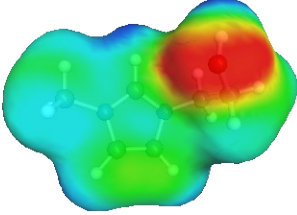
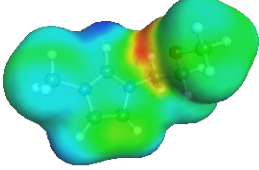
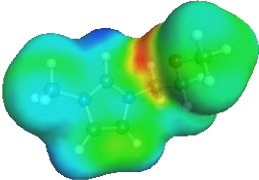
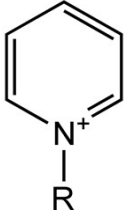
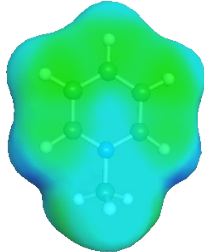
Table S1. The molecular model and conformation of hemicellulose and ions

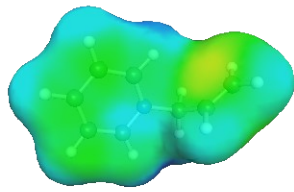
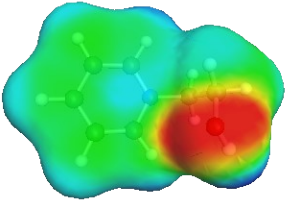
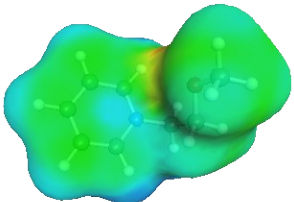
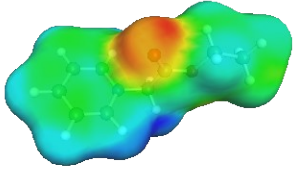
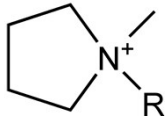
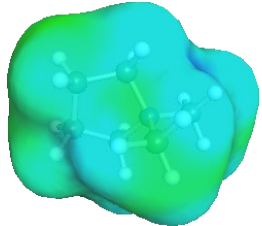
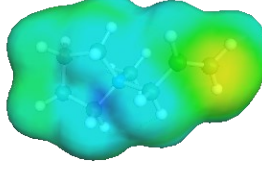
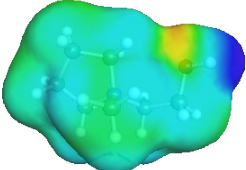
Table S2. R^2 , RSS and Optimization time of the six hemicellulose models

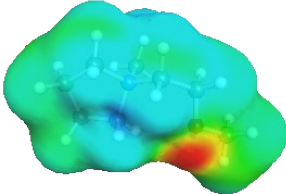
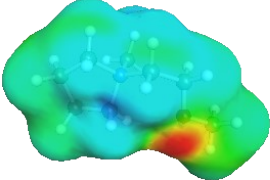
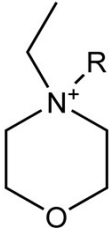
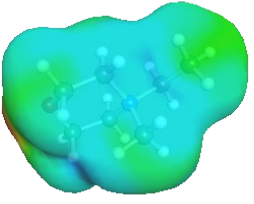
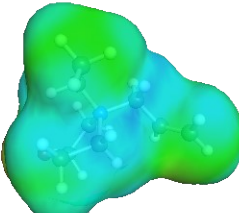
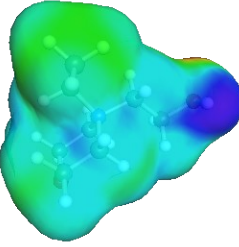
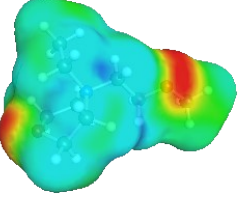
Table S3. The R^2 of the integral area and $\ln\gamma$ of 38 kind anions in four intervals
considered

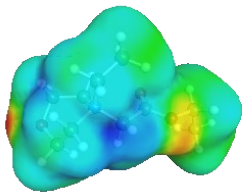
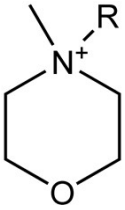
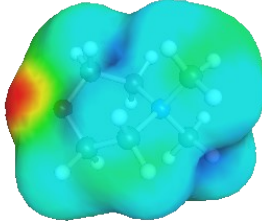
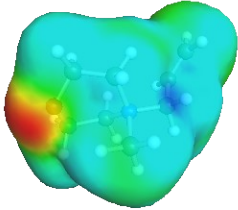
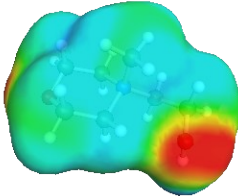
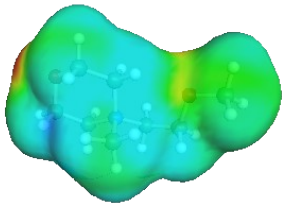
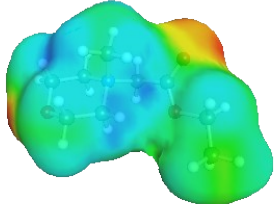
Figure S1. σ -profile curves of hemicellulose and 38 anions.

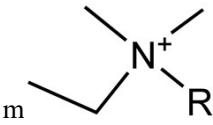
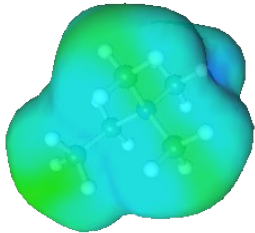
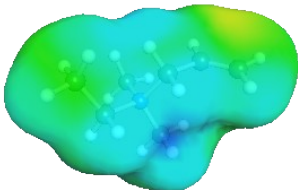
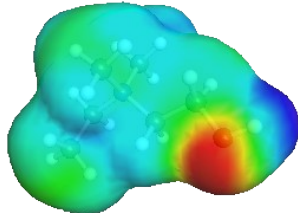
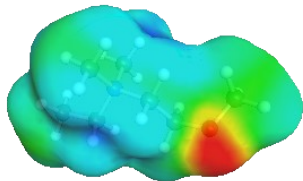
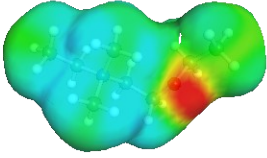
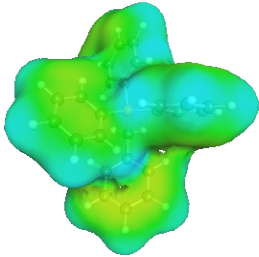
Table S1. The molecular model and conformation of hemicellulose and ions used in this work

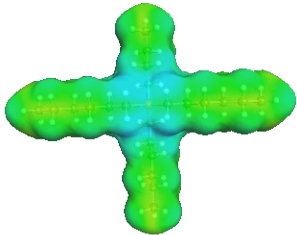
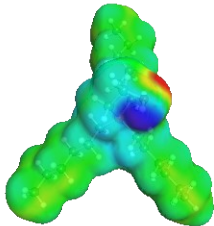
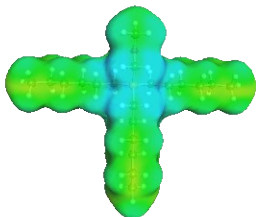
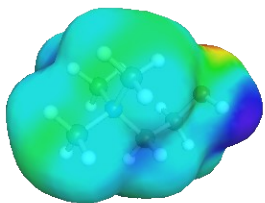
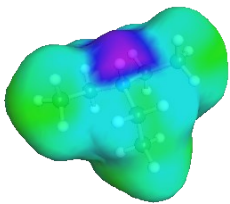
No.	Name of cation	Acronym	Conformation
1-5	1-R-3-Methylimidazolium	1: [Mmim], R=methyl	
		2: [Amim], R=allyl	
		3: [HOEmim], R=2-hydroxyethyl	
		4: [EtOMmim], R=2-methoxyethyl	
		5: [EOEmim], R=2-ethoxy-2oxoethyl	
6-10	N-R-Pyridinium 	6: [Mpy], R=methyl	

No.	Name of cation	Acronym	Conformation
		7: [Apy], R=allyl	
		8: [HOEtpy], R=2-hydroxyethyl	
		9: [EtOMpy], R=2-methoxyethyl	
		10: [EOEpy], R=2-ethoxy-2oxoethyl	
11-15	1R-1-Methylpyrrolidiniu m 	11: [Mpy], R=methyl	
		12: [Apy], R=allyl	
		13: [HOEtpy], R=2-hydroxyethyl	

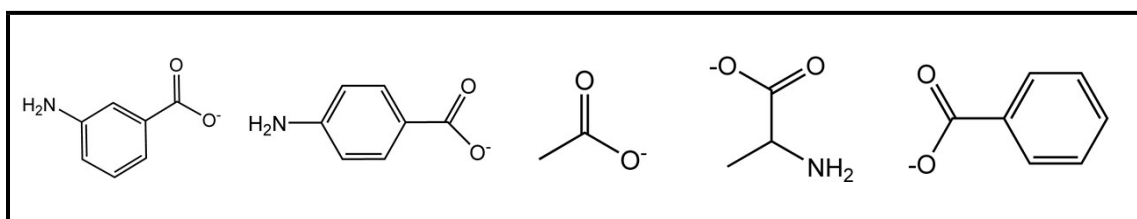
No.	Name of cation	Acronym	Conformation
		14: [EtOMpy], R=2-methoxyethyl	
		15: [EOEpy], R=2-ethoxy-2oxoethyl	
16-20	4-R-4-Ethylmorpholinium 	16: [Memor], R=methyl	
		17: [Aemor], R=allyl	
		18: [HOEtemor], R=2-hydroxyethyl	
		19: [EtOMemor], R=2-methoxyethyl	

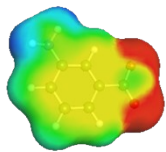
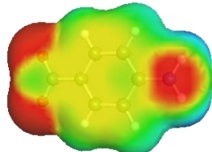
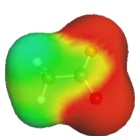
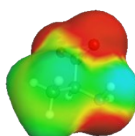
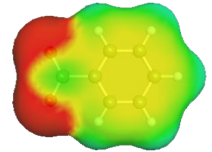

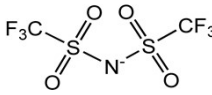


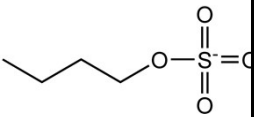
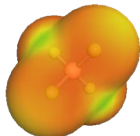
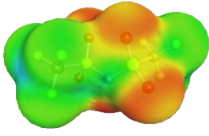


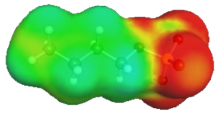
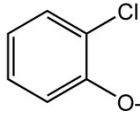
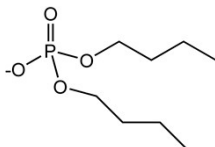
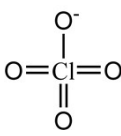
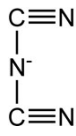
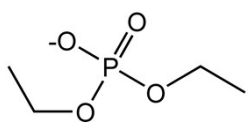
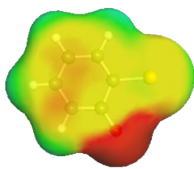
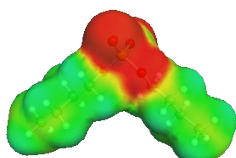
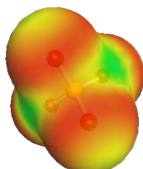
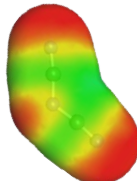
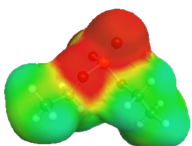
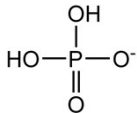
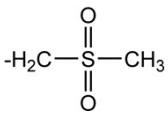
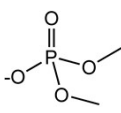
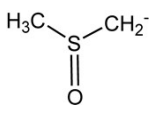
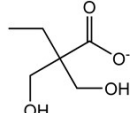
No.	Name of cation	Acronym	Conformation
		20:[EOEemor], R=2-ethoxy-2oxoethyl	
21-25	4-R-4-Methylmorpholinium 	21: [Mmmor], R=methyl	
		22: [Ammor], R=allyl	
		23: [HOEmmor], R=2-hydroxyethyl	
		24: [EtOMmmor], R=2-methoxyethyl	
		25:[EOEmmor], R=2-ethoxy-2oxoethyl	

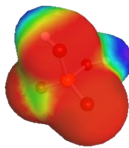
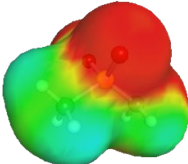
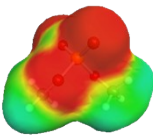
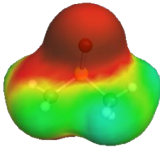
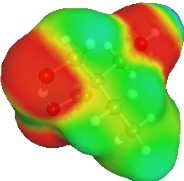
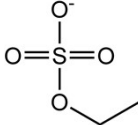
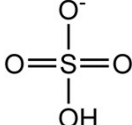
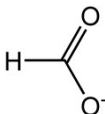
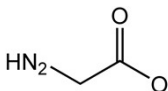
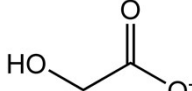
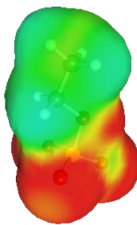
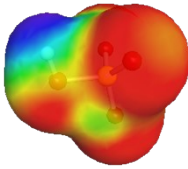
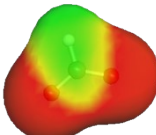
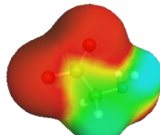
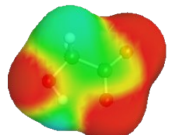
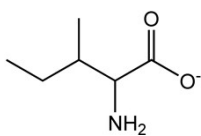
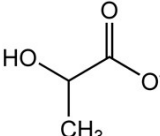

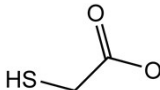
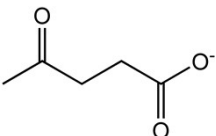
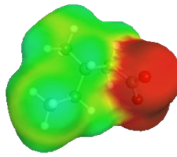
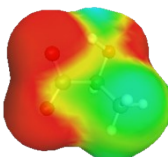

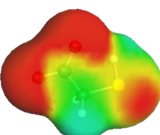
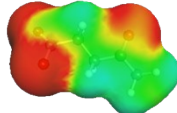
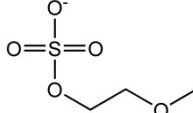
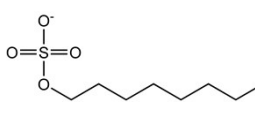
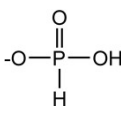

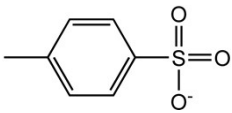
No.	Name of cation	Acronym	Conformation
26-30	R-ethyl-dimethylammoniu 	26: [Medmam], R=methyl	
		27: [Aedmam], R=allyl	
		28:[HOEtedmam], R=2-hydroxyethyl	
		29:[EtOMedmam], R=2-methoxyethyl	
		30:[EOEedmam], R=2-ethoxy-2oxoethyl	
31	Benzyl-triphenyl-phosphonium	[Bentpho]	

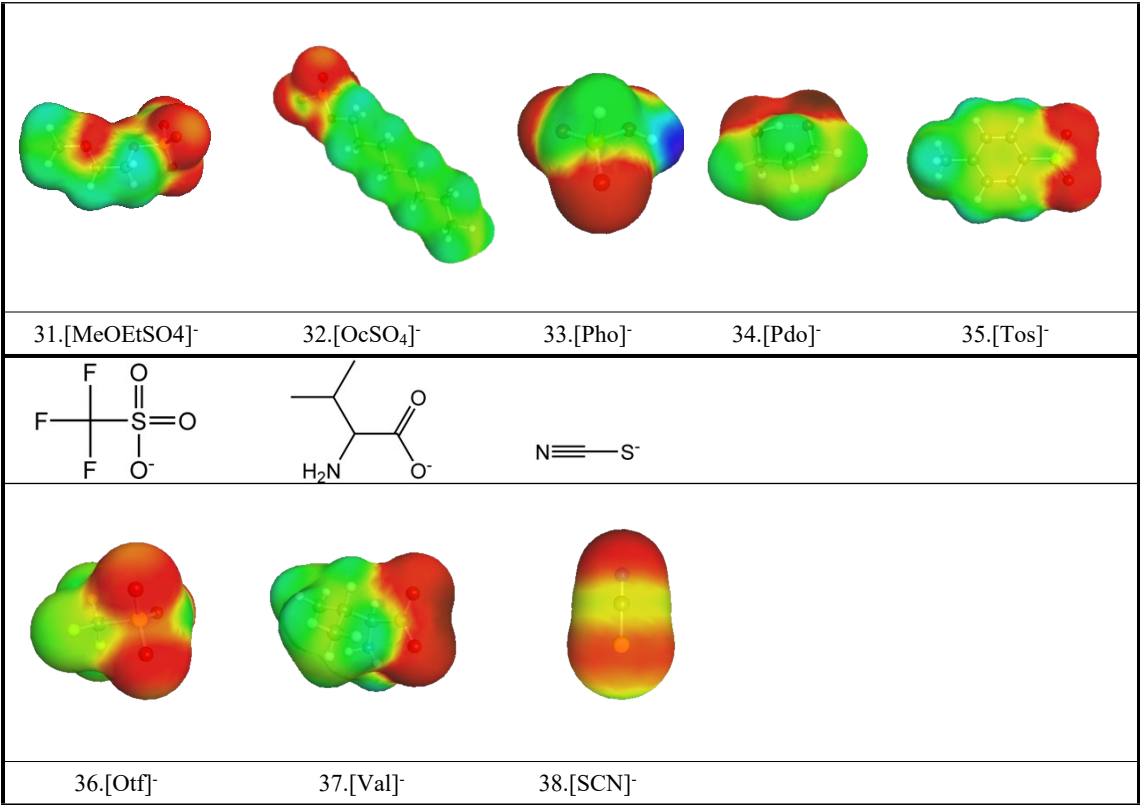
No.	Name of cation	Acronym	Conformation
32	Butyl-trihexyl-phosphonium	[Butpho]	
33	Trihexyl(2-hydroxyethyl)phosphonium	[Thhpho]	
34	Trihexyl(methoxymethyl)-phosphonium	[Thmompho]	
35	Choline	[Choline]	
36	Triethylamine	[Triam]	

The conformation of anions studied in this work



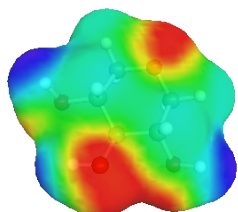
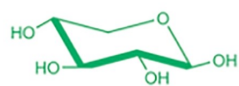
				
1.[Oabc]⁻	2.[Pabc]⁻	3.[Ac]⁻	4.[Ala]⁻	5.[BEN]⁻
				
				
6.[BF ₄] ⁻	7.[TF ₂ N] ⁻	8.[Br] ⁻	9.[Cl] ⁻	10.[BuSO ₄] ⁻
				
				
11.[Clpol]⁻	12.[DBP]⁻	13.[ClO ₄] ⁻	14.[N(CN) ₂] ⁻	15.[DEP]⁻
				

				
16.[DHP] ⁻	17.[DMSO ₂] ⁻	18.[DMPO ₄] ⁻	19.[DMSO] ⁻	20.[DMBA] ⁻
				
				
21.[MeSO ₄] ⁻	22.[HSO ₄] ⁻	23.[HCOO] ⁻	24.[Gly] ⁻	25.[MeHOCOO] ⁻
				
				
26.[Isoec] ⁻	27.[EtHOCOO] ⁻	28.[I] ⁻	29.[MeHSCOO] ⁻	30.[lev] ⁻
				

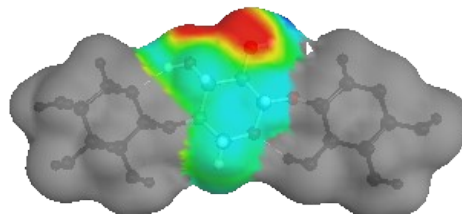
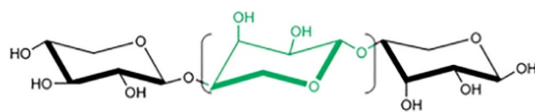


The conformation of hemicellulose models studied in this work

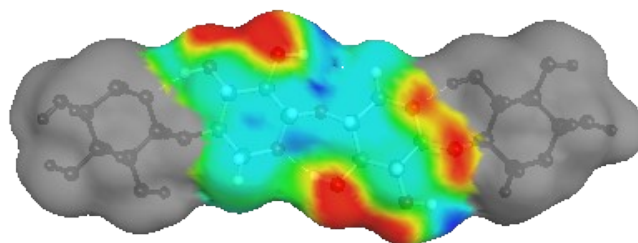
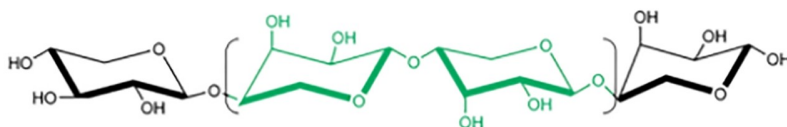
Model 1



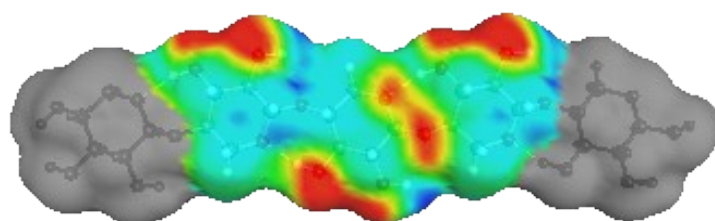
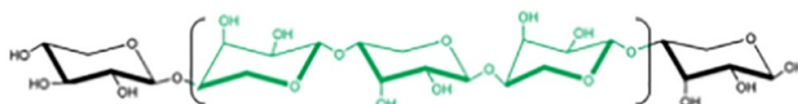
Model 2



Model 3



Model 4



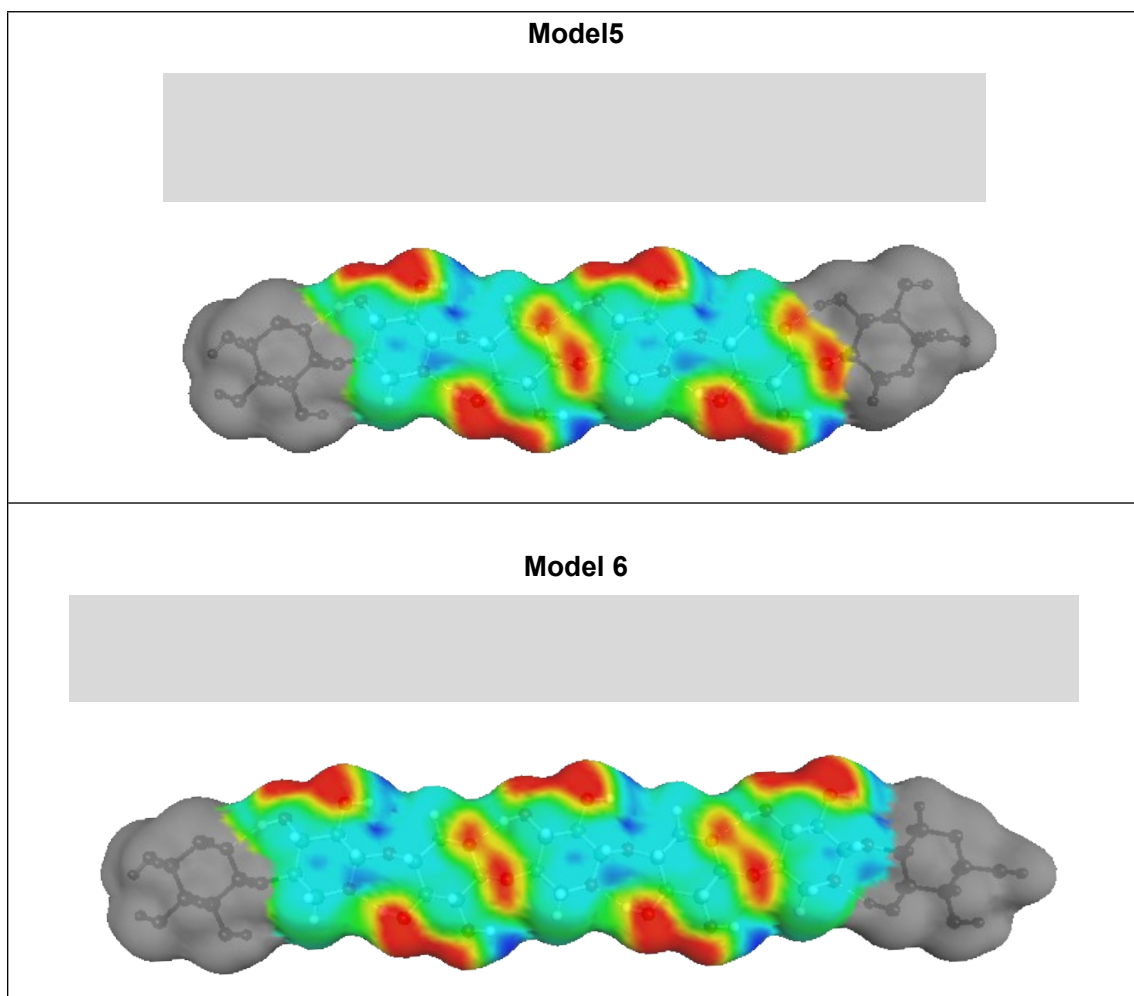


Table S2. R², RSS and Optimization time of the six hemicellulose models

	Model 1	Model 2	Model 3	Model 4	Model 5	Model 6
R ²	0.162	0.648	0.824	0.827	0.828	0.828
RSS	15.43	0.09	0.20	0.51	0.96	1.57
Optimization time(min)	6	56	176	434	678	790

Table S3. The R² of the integral area and lny of 38 kind anions in four intervals considered.

Region	+0.015~+0.021	+0.016~+0.021	+0.015~+0.022	+0.016~+0.022
R ²	0.703	0.779	0.622	0.630

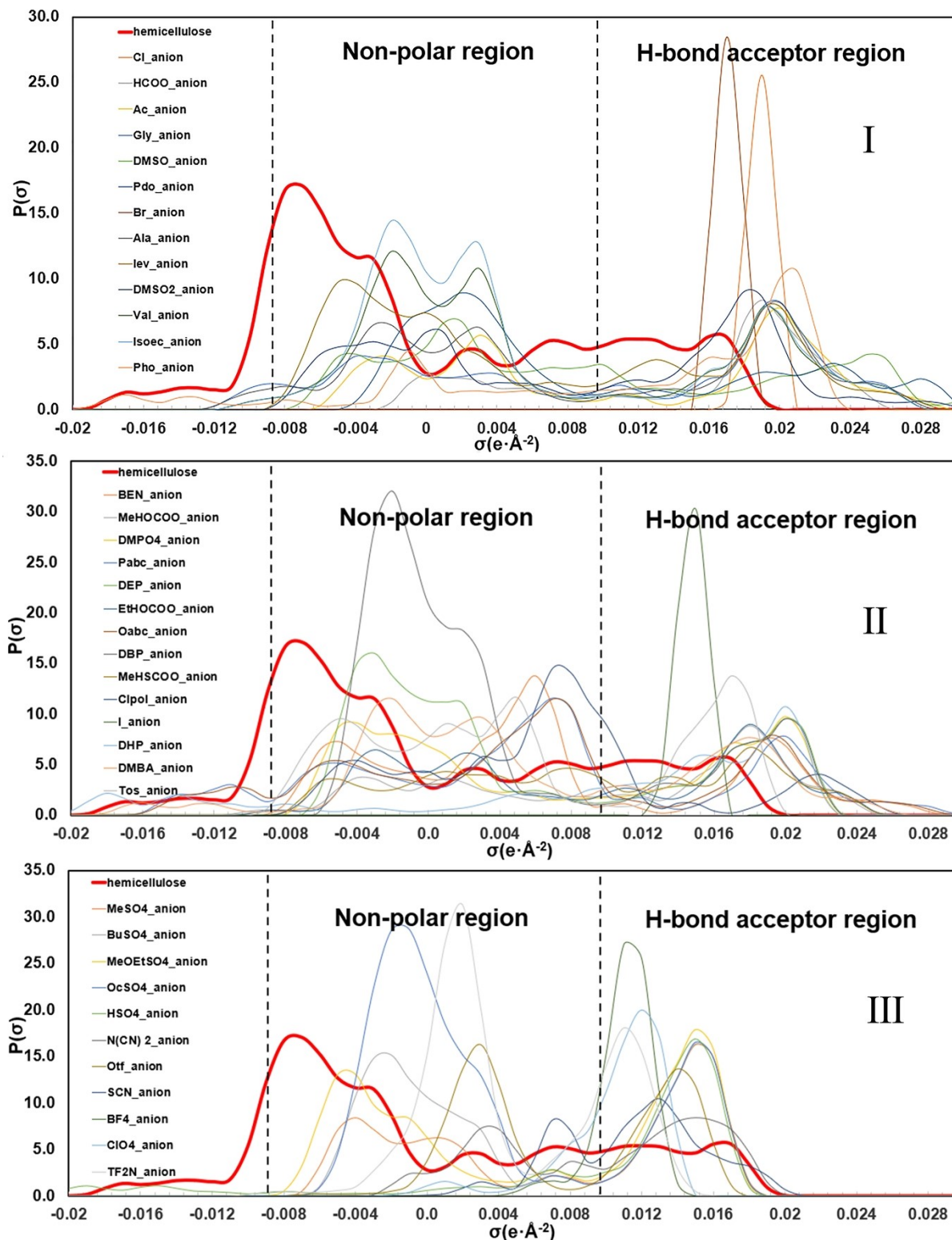


Figure S1. σ -profile curves of hemicellulose and 38 anions. The red curve is the σ -profile of the hemicellulose model. The anions are divided into three groups according to the order of activity coefficients.