

Commercially available carrageenans show broad variation in their structure, composition, and functionality

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Tab. S1: Primer sequences used for gene amplification for ligation-independent cloning.

Gene locus	Primer	Primer sequence (5' → 3')
ZGAL_236	forward	TACTTCCAATCCATGCAACAACCTACGAAGACTTCAAATCCG AAC
	reverse	TATCCACCTTTACTGTTACTCCACGAGTATCTTTTTTGAAACC TCTCC
ZGAL_4265	forward	TACTTCCAATCCATGGTTCCAACCTGAATTGAGGGCCG
	reverse	TATCCACCTTTACTGTCAGTTACACGAAGTAATACTACCTAG ATTTTGGTTAC

Tab. S2: Molecular weight and polydispersity (determined by HPSEC-RI/MALLS) of the carrageenans used in this study.

Sample	M _w , kDa	Polydispersity index
KC1	1013 ± 35	1.9
KC2	965 ± 23	2.3
KC3	925 ± 25	1.7
IC1	999 ± 16	1.9
IC2	621 ± 25	2.3
IC3	523 ± 61	1.7
IC4	866 ± 5	1.8
IC5	915 ± 3	2.0
LC1	923 ± 8	2.0
LC2	641 ± 9	1.8
LC3	1787 ± 18	2.0
LC4	1499 ± 2	2.2
LC5	821 ± 13	1.7
LC6	1081 ± 26	2.7
C1	1158 ± 21	2.1
C2	886 ± 30	2.0

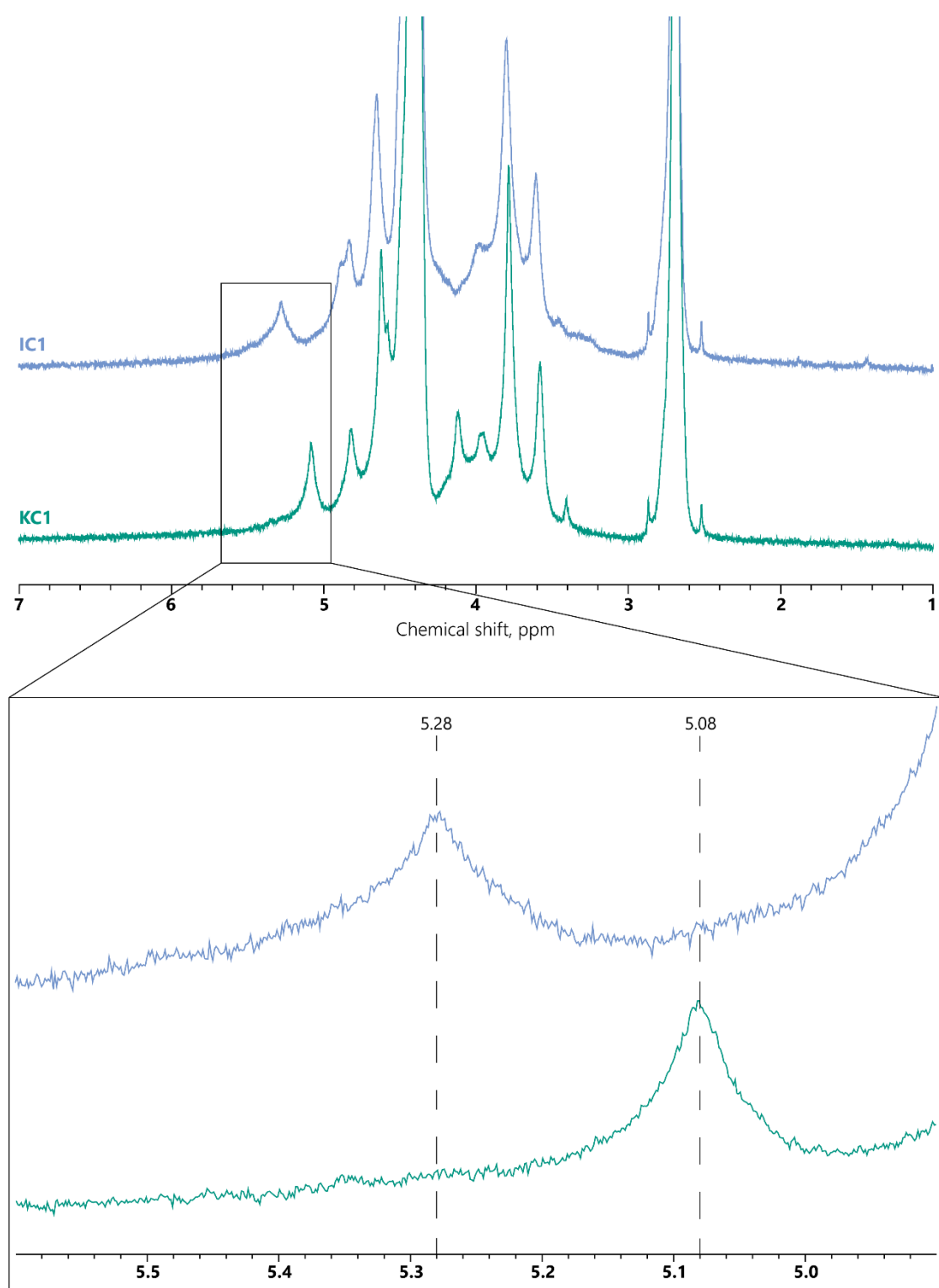


Fig. S1: ^1H NMR spectra of the commercial carrageenans KC1 and IC1.

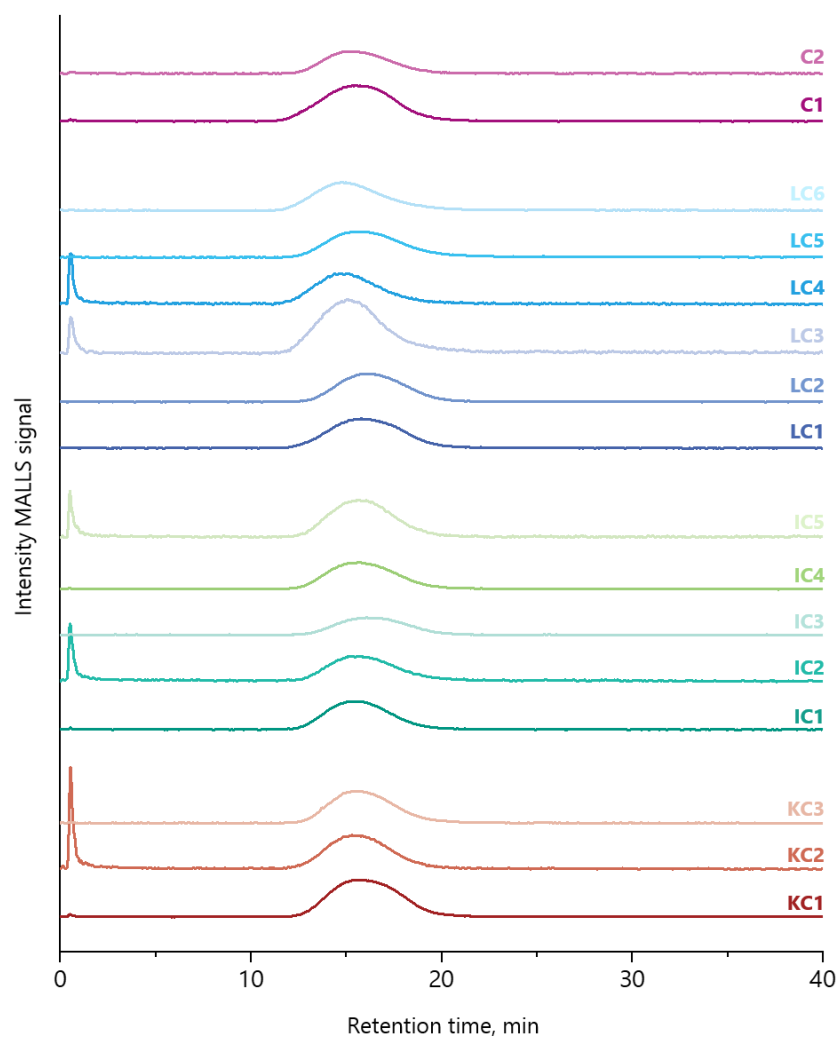


Fig. S2: HPSEC-MALLS chromatograms of the commercial carrageenans used in this study.

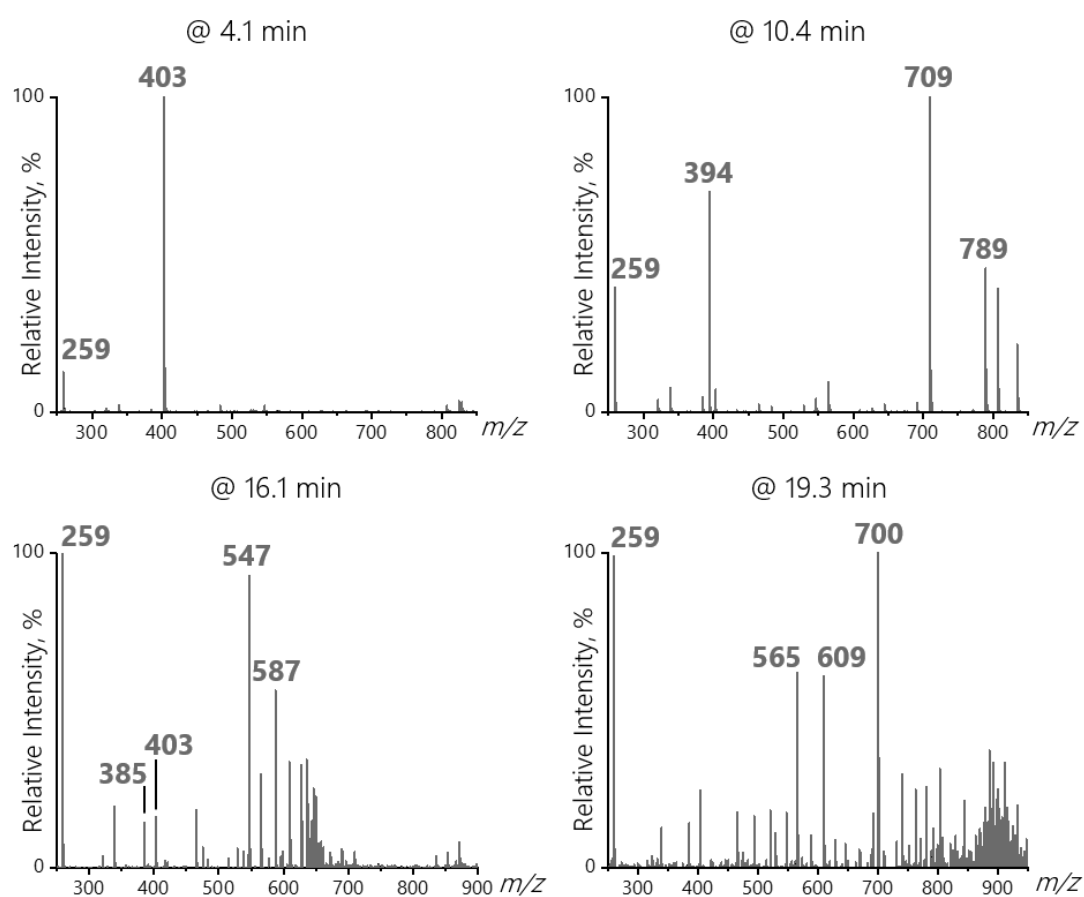


Fig. S3: Mass spectra of the oligosaccharides enzymatically released from KC1 after κ -carrageenase treatment. The analysis was conducted using HPLC-MS in negative full scan mode. The retention times of the oligosaccharides are indicated above the mass spectra.

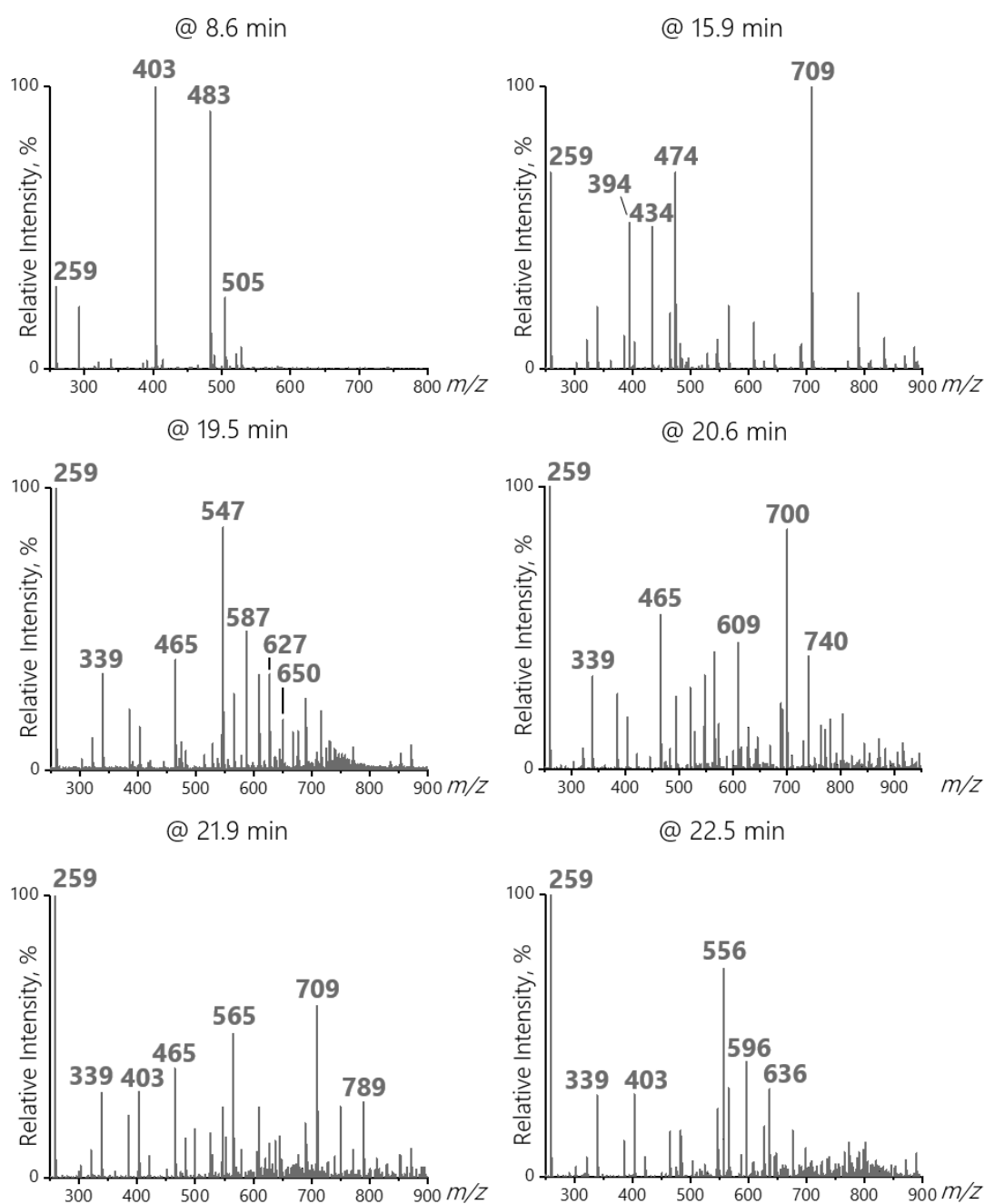


Fig. S4: Mass spectra of the oligosaccharides enzymatically released from IC1 after α -carrageenase treatment. The analysis was conducted using HPLC-MS in negative full scan mode. The retention times of the oligosaccharides are indicated above the mass spectra.

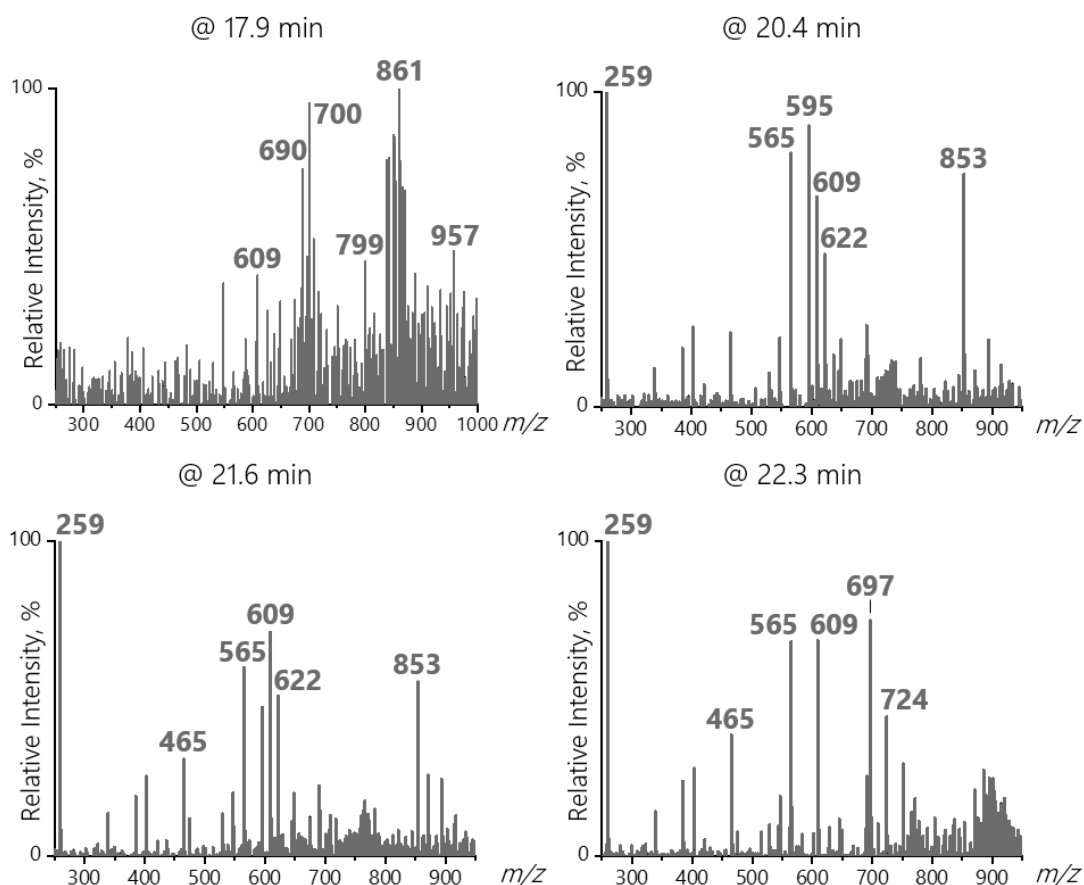


Fig. S5: Mass spectra of the oligosaccharides eluting between 17.9 min and 22.3 min, enzymatically released from LC2 after κ -carrageenase treatment. The analysis was conducted using HPLC-MS in negative full scan mode. The retention times of the oligosaccharides are indicated above the mass spectra.

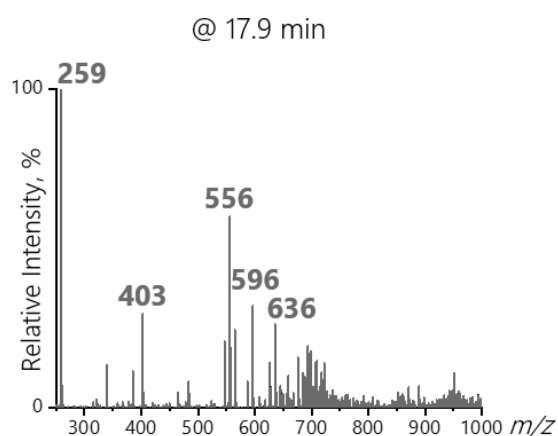


Fig. S6: Mass spectra of the oligosaccharide eluting at 17.9 min, enzymatically released from LC1 after κ -carrageenase treatment. The analysis was conducted using HPLC-MS in negative full scan mode. The retention times of the oligosaccharides are indicated above the mass spectra.

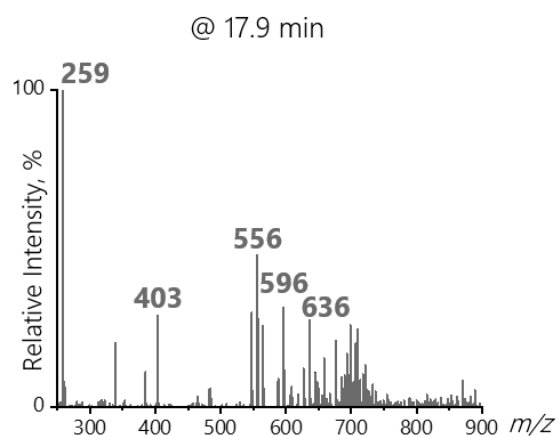


Fig. S7: Mass spectra of the oligosaccharide eluting at 17.9 min, enzymatically released from C1 after κ -carrageenase treatment. The analysis was conducted using HPLC-MS in negative full scan mode. The retention times of the oligosaccharides are indicated above the mass spectra.

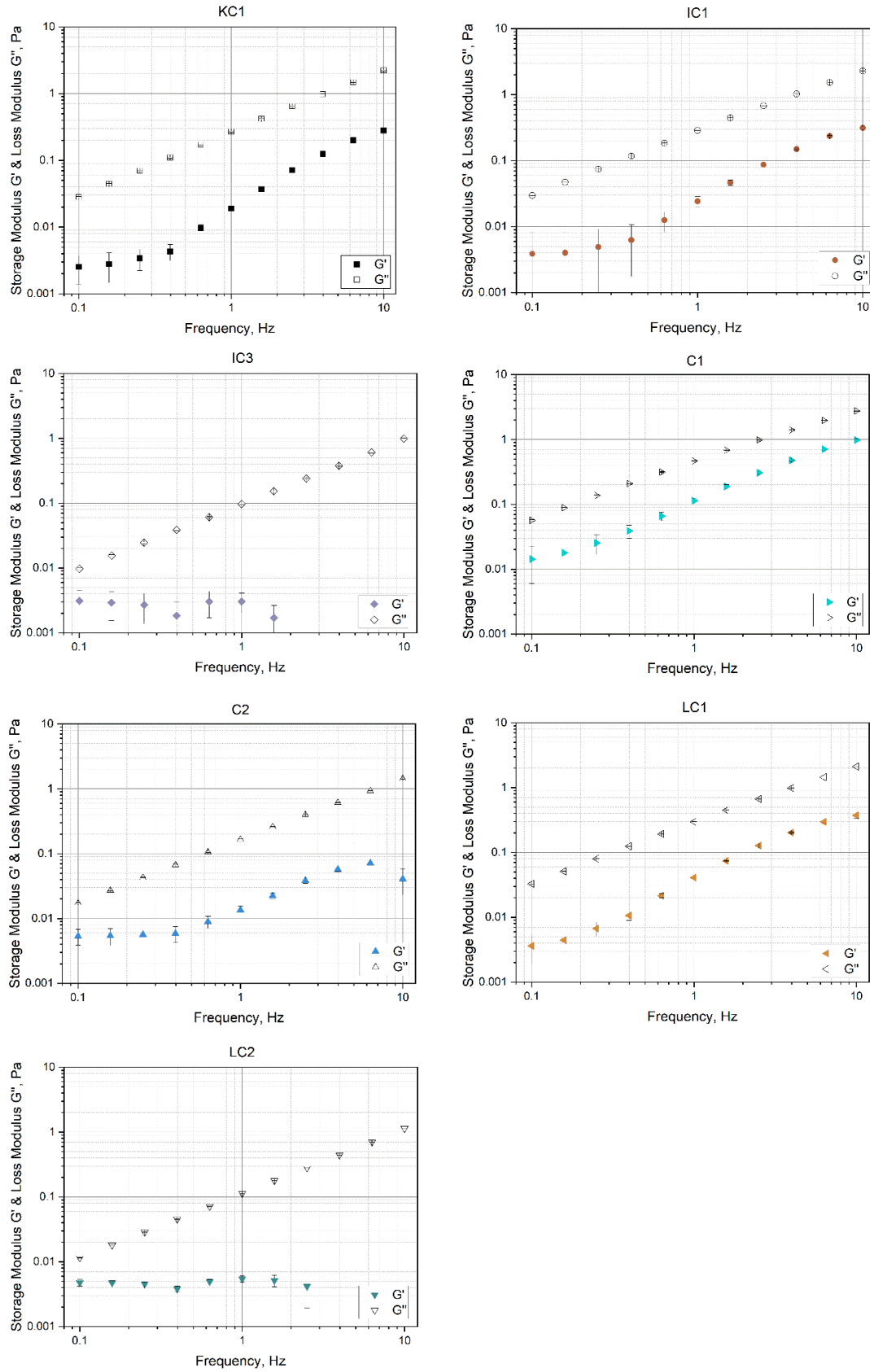


Fig. S8: Storage Modulus G' and loss modulus G'' of the seven rheologically analyzed commercial carrageenan samples. The filled symbols correspond to the storage modulus G' , while the empty symbols represent the loss modulus G'' . The samples were prepared without CaCl_2 addition and exhibit a predominantly viscous behavior.

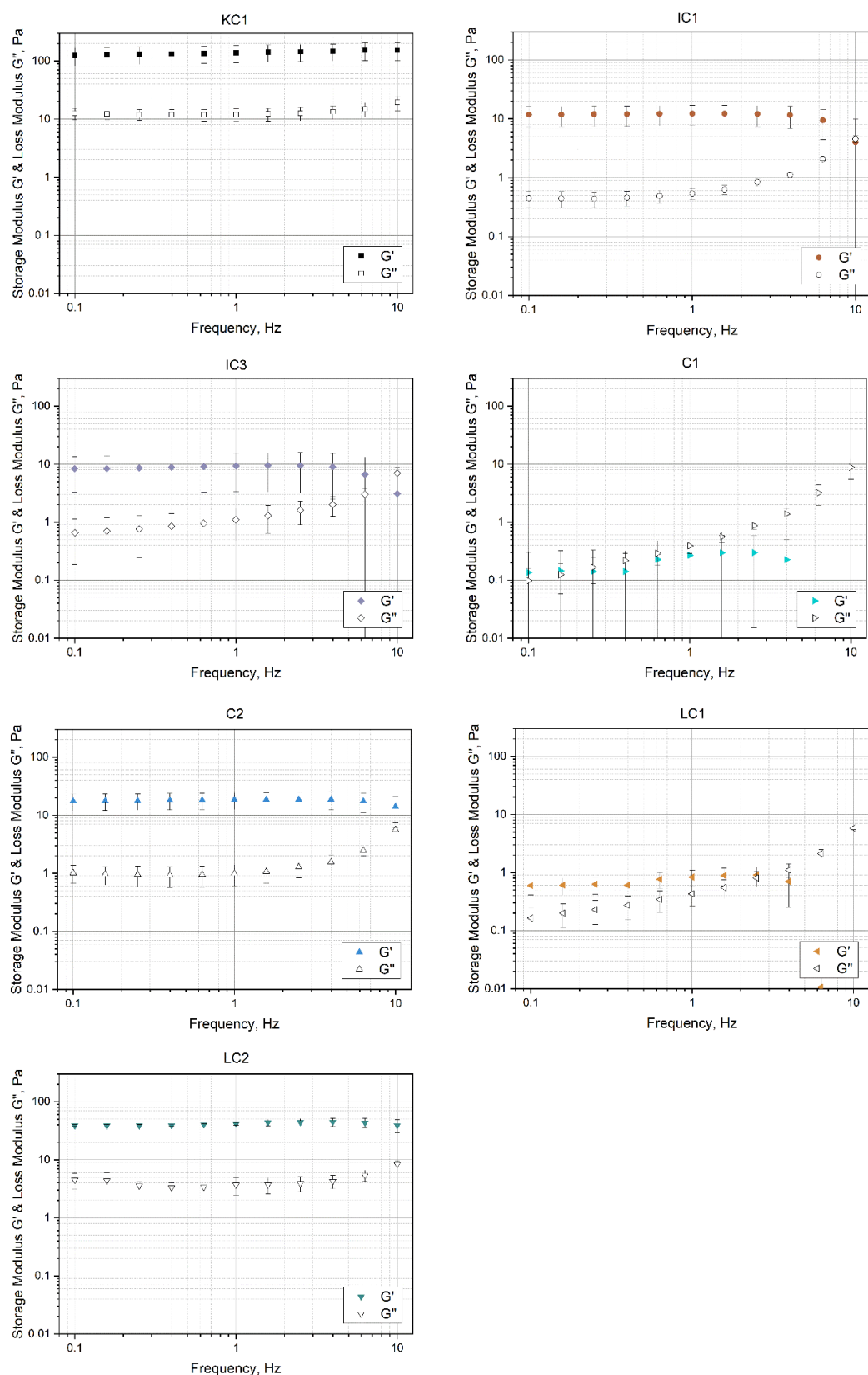


Fig. S9: Storage Modulus G' and loss modulus G'' of the seven commercially available carrageenan samples analyzed rheologically with the addition of CaCl_2 . The filled symbols correspond to the storage modulus G' , while the empty symbols represent the loss modulus G'' . With exception of samples C1 and LC1, all other measured samples exhibit gel formation, although the gel strengths vary.

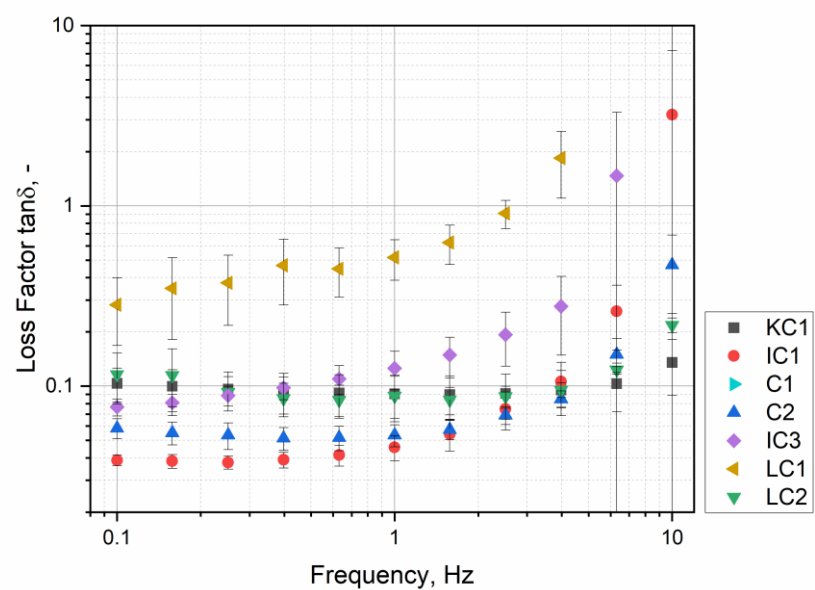


Fig. S10: Loss factors of the different commercial carrageenan samples in 0.1 M CaCl_2 solution measured at a frequency range between 0.1 and 10 Hz at an amplitude γ of 2 %.