

Supporting information

Authors: *Danila Morais de Carvalho, Maarit H. Lahtinen, Martin Lawoko, and Kirsi S. Mikkonen*

Enrichment and identification of lignin-carbohydrate complexes in softwood extract

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Table S1. Mass balance and chemical characterization of galactoglucomannan (GGM) samples^a.

Samples		sdGGM	esGGM	epGGM
Mass balance based on original spruce wood (%)		21.6	4.5 ± 0.2	16.0 ± 0.4
Normalized chemical composition of sugars and lignin (%)	Man	51.1 ± 0.2	38.9 ± 0.4	55.4 ± 0.0
	Glc	14.7 ± 0.3	9.9 ± 0.4	16.3 ± 0.7
	Gal	6.4 ± 0.2	3.7 ± 0.1	7.7 ± 0.2
	Xyl	9.5 ± 0.2	7.9 ± 0.3	10.5 ± 0.5
	GlcA	1.6 ± 0.1	2.7 ± 0.2	1.9 ± 0.2
	Ara	0.6 ± 0.1	3.1 ± 0.1	0.3 ± 0.0
	Fuc	0.2 ± 0.0	0.0 ± 0.0	0.1 ± 0.1
	Rha	0.4 ± 0.1	1.0 ± 0.2	0.3 ± 0.1
	GalA	0.1 ± 0.1	0.0 ± 0.0	0.1 ± 0.1
	Lignin	15.4 ± 2.5	32.9 ± 1.7	7.4 ± 0.6
GGM purity (%) ^b		72.2 ± 0.5	52.5 ± 0.6	79.4 ± 0.5
Acetyl groups	Content (%) ^c	3.8	4.9	3.2
	DS _{Ac} ^d	0.23	0.27	0.27
	Distribution ^e	1.0:0.8	1.0:0.8	1.0:0.8
Molar mass	Mn (kDa)	1.7 ± 0.0	0.7 ± 0.0	2.6 ± 0.0
	Mw (kDa)	7.1 ± 0.1	1.5 ± 0.0	9.3 ± 0.6
	DI	4.3 ± 0.0	2.0 ± 0.1	3.6 ± 0.3

^aIn the table: sdGGM: spray-dried galactoglucomannan; es: ethanol-soluble galactoglucomannan; epGGM: ethanol-precipitated galactoglucomannan; Man: mannose; Glc: glucose; Gal: galactose; Xyl: xylose; GlcA: glucuronic acid; Ara: arabinose; Fuc: fucose; Rha: rhamnose; GalA: galacturonic acid, DS_{Ac}: degree of acetylation of galactoglucomannan samples; Mn: number-average molar mass; Mw: weight-average molar mass; and DI: dispersity index.

^bFor the estimation of the purity of galactoglucomannan were considered the content of mannose, glucose, and galactose in samples. Similarly, arabinoglucuronoxylan content was estimated considering xylose, arabinose, and glucuronic acid moieties and pectin considered the content of fucose, rhamnose, and galacturonic acid.

^cmol acetyl per 100 mol hexose.

^dDS_{Ac} was calculate considering only those sugars present in the galactoglucomannan backbone, *i.e.*, mannose and glucose.

^eAcetylation distribution in mannosyl units (C-2:C-3) obtained by semi-quantitative analysis from NMR spectra.

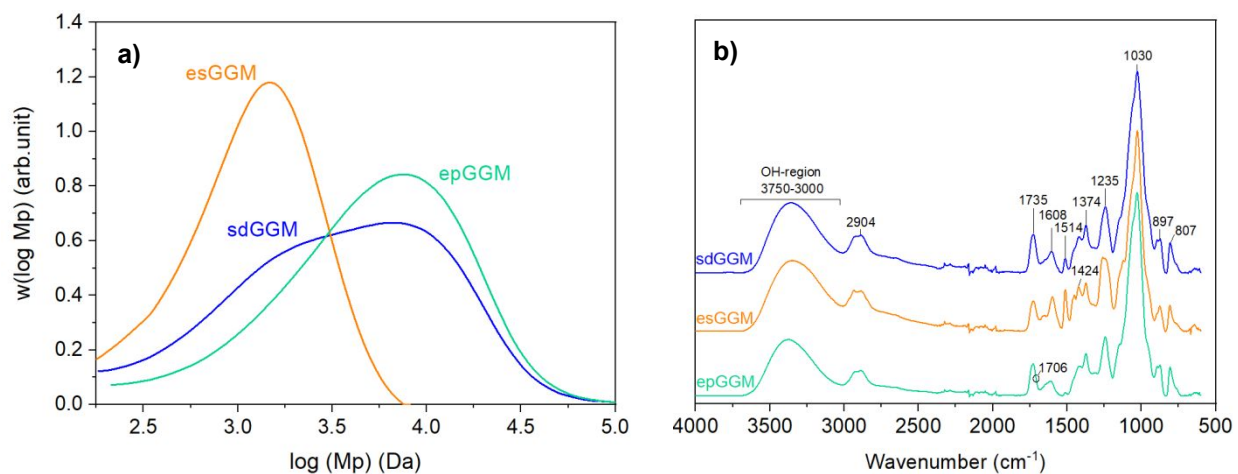


Figure S1. Molar mass distribution (a) and FTIR spectra (b) of sdGGM, esGGM, and epGGM samples.

Table S2. Assignment for the relevant bands observed in FTIR spectra

Absorption band (cm^{-1})	Band assignments ^a
3360	O-H stretching vibration of hydrogen-bonded
2904	Aliphatic C-H stretching
1735	C=O stretching vibration carbonyl groups
1706	Free acetic acid (absent in the spectra)
1608	Water absorption
1514	Aromatic skeletal vibration of lignin rings
1424	Asymmetric aliphatic C-H deformation of methoxyl
1374	Symmetric aliphatic C-H bending of methoxyl in acetyl groups
1235	C-O stretching
1030	C-O and C-C stretching or C-OH bending
897	β -glycosidic linkage between sugar units
807	Aromatic out-of-plane-rings region

^aBand assignments were assessed according to the literature.¹⁻⁸

Table S3. Assignment for the relevant carbon-hydrogen correlation signals of lignin structures in sdGGM, esGGM, and epGGM samples and fractions.

Structural units ^a	δ_C (ppm)	δ_H (ppm)	Assignments ^b	Identified in ^c
β -O-4	59.5	3.70/3.44	C_γ/H_γ in β -O-4	1-27
β -O-4	71.4	4.71	C_α/H_α in γ -hydroxylated β -O-4	1-27
β -O-4 S	85.3	4.16	C_β/H_β in γ -hydroxylated β -O-4 in S-units	2, 5-10, 15, 16, 19, 29, 22-24
β -O-4 G	83.4	4.28	C_β/H_β in γ -hydroxylated β -O-4 in G-units	1-27
β -O-4 BE	84.6	4.34	C_β/H_β in β -O-4 in benzylether structures	2, 6-10, 16, 19, 20, 22-24
β -O-4 GE	81.1	4.64	C_γ/H_γ in γ -acylated β -O-4	1, 2, 4, 6, 7, 16, 20, 22
β -O-4 BE	61.5	3.80	C_γ/H_γ in β -O-4 in benzylether structures	1-17, 19, 20, 22-27
Phenylcoumaran	87.0	5.40	C_α/H_α in phenylcoumaran	1-27
Phenylcoumaran	54.7	3.45	C_β/H_β in phenylcoumaran	1-10, 12, 14-24
Resinol struct.	85.1	4.61	C_α/H_α in resinol structures	1, 2, 4-10, 12, 14, 16-18, 20-24
Resinol struct.	53.6	3.04	C_β/H_β in resinol structures	1-4, 6-10, 12, 16, 17, 20-24
Resinol struct.	71.2	4.14/3.85	C_γ/H_γ in resinol structures	1-20, 22
DBD	83.3	4.80	C_α/H_α in dibenzodioxin	1, 2, 6-10, 12, 16, 17, 20, 22, 23
DBD	85.5	4.01	C_β/H_β in dibenzodioxin	1, 2, 4-11, 16, 19, 20, 22, 23, 27
DBD	60.3	3.42	C_γ/H_γ in dibenzodioxin	1-14, 16-27
OSC (G ₂)	111.8	7.30	C_2/H_2 in (C=O) _a in G-units	1-4, 6-10, 12, 16, 17, 20-23
OSC (G ₆)	122.8	7.64/7.56	C_6/H_6 in (C=O) _a in G-units	2, 3, 6-8, 10, 16, 20, 23
OSC (S _{2,6})	106.4	7.20/7.03	$C_{2,6}/H_{2,6}$ in (C=O) _a in S-units	7, 20
OSC _a	153.8	7.60	C_α/H_α in cinnamyl aldehyde	1-4, 6-9, 11, 12, 16-23
OSC _{β}	126.6	6.79	C_β/H_β in cinnamyl aldehyde	1-4, 6-12, 14, 16, 17, 20-24
Methoxyl	55.4	3.73	C/H in -OMe in G- and S-units	1-27
CA	61.7	4.05	C_γ/H_γ in cinnamyl alcohol	1, 2, 4-12, 14-24, 26, 27
SECO _a	33.6	2.53/2.46	C_α/H_α in secoisolariciresinol	1, 2, 5-17, 20, 22, 23, 25, 26
SECO _{β}	42.3	1.84	C_β/H_β in secoisolariciresinol	1-10, 12, 14, 16-24
DCA _a	31.4	2.52	C_α/H_α in dehydroconiferyl alcohol	1-11, 14, 16-24
DCA _{β}	34.4	1.65	C_β/H_β in dehydroconiferyl alcohol	1-12, 14, 16-27
S _{2,6}	103.9	6.68	C_2/H_2 and C_6/H_6 in S-units	2, 6, 7, 16, 20, 22
G ₂	110.7	6.94	C_2/H_2 in G-units	1-27
G ₅	115.0	6.82	C_5/H_5 in G-units	1-27
G ₆	118.7	6.76	C_6/H_6 in G-units	1-27
H _{2,6}	127.8	7.19	C_2/H_2 and C_6/H_6 in H-units	1-4, 6-10, 12, 14, 16-18, 20-24

^a β -O-4: β -O-4 linkage between lignin including syringyl (S) and guaiacyl (G) units and in lignin involved in benzylether (BE) and gamma-ester (GE) linkages; Phenylcoumaran (β 5); Resinol structures ($\beta\beta$); Dibenzodioxin (DBD); Oxidized side chain (OSC); Methoxyl in S and G units; Cinnamyl alcohol (CA), Secoisolariciresinol (SECO – $\beta\beta$); Dehydro coniferyl alcohol (DCA); and p-hydroxyphenyl (H) unit.

^bContours assignments were assessed according to the literature.^{9, 10}

^cThe numbers refer to the following samples: 1: sdGGM raw; 2: sdGGM SSEH pellet/large; 3: sdGGM SSEH pellet/small; 4: sdGGM SSEH supernatant/large; 5: sdGGM SSEH supernatant/small; 6: esGGM raw; 7: esGGM SSEH pellet/large; 8: esGGM SSEH pellet/small; 9: esGGM SSEH supernatant/large; 10: esGGM SSEH supernatant/small; 11: epGGM raw; 12: epGGM SSEH pellet/large; 13: epGGM SSEH pellet/small; 14: epGGM SSEH supernatant/large; 15: epGGM SSEH supernatant/small; 16: sdGGM reference pellet/large; 17: sdGGM reference pellet/small; 19: sdGGM reference supernatant/large; 19: sdGGM reference supernatant/small; 20: esGGM reference pellet/large; 21: esGGM reference pellet/small; 22: esGGM reference supernatant/large; 23: esGGM reference supernatant/small; 24: epGGM reference pellet/large; 25: epGGM reference pellet/small; 26: epGGM reference supernatant/large; and 27: epGGM reference supernatant/small.

Table S4. Assignment for the relevant lignin substructures in sdGGM, esGGM, and epGGM samples and fractions.

Structural units	δ_C (ppm)	δ_H (ppm)	Assignments ^a	Identified in ^b
β -aryl ether (β -O-4)	71.3	4.90	β -O-4 signal in α	1-10, 12-20, 22-5, 27
β -aryl ether (β -O-4)	83.8	4.26	β -O-4 signal in β of guaiacyl	1- 27
β -aryl ether (β -O-4)	86.0	4.08	β -O-4 signal in β of syringyl	2, 5, 6, 7, 10, 13, 15, 16, 20
β -aryl ether (β -O-4)	61.7	3.40	β -O-4 signal in γ	1-27
Phenyl coumaran (β -5)	87.4	5.46	β -5 signal in α	1-12, 16-24, 26
Phenyl coumaran (β -5)	52.6	3.46	β -5 signal in β	1-26
Phenyl coumaran (β -5)	62.4	3.72	β -5 signal in γ	1-27
Cinnamyl alcohol	128.5	6.42	Cinnamyl alcohol signal in α	1-3, 6-10, 16-23, 25
Cinnamyl alcohol	127.5	6.21	Cinnamyl alcohol signal in β	1, 2, 6-10, 16-23
Cinnamyl alcohol	62.8	3.90	Cinnamyl alcohol signal in γ	1-20, 22-27
Resinol (β - β)	83.4	4.86	β - β signal in α	1, 2, 5- 10, 16, 18, 20, 22
Resinol (β - β)	53.4	3.04	β - β signal in β	1-4, 6-10, 1216, 17, 20-23
Resinol (β - β)	73.1	4.09	β - β signal in γ	1, 4, 9
Dibenzodioxocin (5-5/ β -O-4)	82.9	4.80	5-5/ β -O-4 signal in α	1, 2, 6-8, 10, 16-18, 20, 20, 23
Dibenzodioxocin (5-5/ β -O-4)	86.2	3.90	5-5/ β -O-4 signal in β	2, 6, 7, 16, 20
Spirodienone (β -1)	81.2	5.00	β -1 signal in α	2, 6, 7, 16, 20, 22
Spirodienone (β -1)	59.7	2.70	β -1 signal in β	1, 7, 20
Spirodienone (β -1)	80.1	4.10	β -1 signal in β'	2, 6, 7, 11, 12

^aContours assignments were assessed according to the literature.¹¹

^bThe numbers refer to the following samples: 1: sdGGM raw; 2: sdGGM SSEH pellet/large; 3: sdGGM SSEH pellet/small; 4: sdGGM SSEH supernatant/large; 5: sdGGM SSEH supernatant/small; 6: esGGM raw; 7: esGGM SSEH pellet/large; 8: esGGM SSEH pellet/small; 9: esGGM SSEH supernatant/large; 10: esGGM SSEH supernatant/small; 11: epGGM raw; 12: epGGM SSEH pellet/large; 13: epGGM SSEH pellet/small; 14: epGGM SSEH supernatant/large; 15: epGGM SSEH supernatant/small; 16: sdGGM reference pellet/large; 17: sdGGM reference pellet/small; 19: sdGGM reference supernatant/large; 19: sdGGM reference supernatant/small; 20: esGGM reference pellet/large; 21: esGGM reference pellet/small; 22: esGGM reference supernatant/large; 23: esGGM reference supernatant/small; 24: epGGM reference pellet/large; 25: epGGM reference pellet/small; 26: epGGM reference supernatant/large; and 27: epGGM reference supernatant/small.

Table S5. Assignment for lignin-carbohydrates bonds signals in sdGGM, esGGM, and epGGM samples and fractions.

Structural units ^a	δ_C (ppm)	δ_H (ppm)	Assignments ^b	Identified in ^c
PG	101.1-99.8	5.18-5.00	Phenylglycoside	1, 2, 6-12, 17, 20, 22-24
PG	101.8	4.98-4.90	Phenylglycoside	1, 4, 6-10, 17, 19-23
PG	100.8	4.89-4.84	Phenylglycoside	3, 6, 8-11, 14, 21-23
BE (BE _i)	80.2	4.50	Benzylether	2, 6, 7, 16, 20, 22
GE	62.0-65.0	4.50/4.00	γ -ester	1-27

^aPG: phenylglycoside; BE: benzylether; and GE: γ -ester.

^bContours assignments were assessed according to the literature.^{10, 12, 13}

^cThe numbers refer to the following samples: 1: sdGGM raw; 2: sdGGM SSEH pellet/large; 3: sdGGM SSEH pellet/small; 4: sdGGM SSEH supernatant/large; 5: sdGGM SSEH supernatant/small; 6: esGGM raw; 7: esGGM SSEH pellet/large; 8: esGGM SSEH pellet/small; 9: esGGM SSEH supernatant/large; 10: esGGM SSEH supernatant/small; 11: epGGM raw; 12: epGGM SSEH pellet/large; 13: epGGM SSEH pellet/small; 14: epGGM SSEH supernatant/large; 15: epGGM SSEH supernatant/small; 16: sdGGM reference pellet/large; 17: sdGGM reference pellet/small; 19: sdGGM reference supernatant/large; 19: sdGGM reference supernatant/small; 20: esGGM reference pellet/large; 21: esGGM reference pellet/small; 22: esGGM reference supernatant/large; 23: esGGM reference supernatant/small; 24: epGGM reference pellet/large; 25: epGGM reference pellet/small; 26: epGGM reference supernatant/large; and 27: epGGM reference supernatant/small.

Table S6. Assignment for the relevant carbon-hydrogen correlation signals of polysaccharides linkages in sdGGM, esGGM, and epGGM samples and fractions.

Structural units	δ_C (ppm)	δ_H (ppm)	Assignments ^a	Identified in ^b
Acetyl	20.9	2.01	-CH ₃ in acetyl	1-27
Ara ₁	107.8	4.76	C ₁ /H ₁ in β -L-arabinopyranoside	1-27
Ara ₂	83.2	3.76	C ₂ /H ₂ in β -L-arabinopyranoside	1-23, 25, 27
Ara ₄	84.8	4.00	C ₄ /H ₄ in β -L-arabinopyranoside	2, 6, 7, 9, 10, 11, 16, 20, 22
Ara ₅	62.3	3.42/3.36	C ₅ /H ₅ in β -L-arabinopyranoside	1-25, 27
Glc ₁	102.0	4.32	C ₁ /H ₁ in β -D-glucopyranoside	1-27
Glc ₂	74.8	2.90	C ₂ /H ₂ in β -D-glucopyranoside	1-19, 21-27
Glc ₃	76.7	3.05	C ₃ /H ₃ in β -D-glucopyranoside	1- 27
Glc ₄	79.9	3.38	C ₄ /H ₄ in β -D-glucopyranoside	1-27
Glc ₅	70.2	3.18	C ₅ /H ₅ in β -D-glucopyranoside	1, 3-15, 17-27
Glc ₆ /Man ₆	60.0	3.59	C ₆ /H ₆ in β -D-gluco-/mannopyranoside	1-27
Xyl ₁	101.5	4.38	C ₁ /H ₁ in β -D-xylopyranoside	1-27
Xyl ₂	72.7	3.03	C ₂ /H ₂ in β -D-xylopyranoside	1-27
Xyl ₂ (r _a)	69.7	3.27	C ₂ /H ₂ in β -D-xylopyranoside of xylans reduction end	1-19, 21-27
Xyl ₂ -Ac	73.0	4.40	C ₂ /H ₂ in 2- <i>O</i> -acetyl- β -D-xylopyranoside	2, 6-11, 14-17, 20, 22, 23, 25
Xyl ₁ (₂ Ac)	100.8	4.49	C ₁ /H ₁ in 2- <i>O</i> -acetyl- β -D-xylopyranoside	1-27
Xyl ₃	74.5	3.28	C ₃ /H ₃ in β -D-xylopyranoside	1-23, 25-27
Xyl ₃ -Ac	74.5	4.81	C ₃ /H ₃ in 3- <i>O</i> -acetyl- β -D-xylopyranoside	1-20, 22-27
Xyl _{2,3} -Ac	71.6	4.54	C ₂ /H ₂ in 2,3- <i>O</i> -acetyl- β -D-xylopyranoside	1, 2, 4-11, 12, 15-20, 22-27
Xyl _{2,3} -Ac	72.9	4.90	C ₃ /H ₃ in 2,3- <i>O</i> -acetyl- β -D-xylopyranoside	1-20, 22-27
Xyl ₁ (_{2,3} Ac)	99.3	4.68	C ₁ /H ₁ in 2,3- <i>O</i> -acetyl- β -D-xylopyranoside	1-27
Xyl ₄	75.7	3.51	C ₄ /H ₄ in β -D-xylopyranoside	1-27
Xyl ₅	62.8	3.78/3.17	C ₅ /H ₅ in β -D-xylopyranoside	1-27
Xyl ₄ (nr)	66.0	3.31	C ₄ /H ₄ in β -D-xylopyranoside of xylans non-reducing end	1-15, 17-19, 22-27
Xyl ₅ (nr)	65.3	3.62/2.98	C ₅ /H ₅ in β -D-xylopyranoside of xylans non-reducing end	1-27
Xyl ₄ (nr)	103.4	4.15	C ₁ /H ₁ in β -D-xylopyranoside of xylans non-reducing end	1-6, 8-19, 22-27
Xyl ₄ (r _a)	92.3	4.85	C ₁ /H ₁ in α -D-xylopyranoside of xylans reduction end	1-20, 22-27
Xyl ₄ (r _b)	97.3	4.21	C ₁ /H ₁ in β -D-xylopyranoside of xylans reduction end	1-20, 22-27
Man ₁	100.3	4.51	C ₁ /H ₁ in β -D-mannopyranoside	1-27
Man ₂	74.5	2.90	C ₂ /H ₂ in β -D- mannopyranoside	1-19, 21-27
Man ₂ -Ac	70.9	5.24	C ₂ /H ₂ in 2- <i>O</i> -acetyl- β -D- mannopyranoside	1-27
Man ₃	76.6	3.08	C ₃ /H ₃ in β -D- mannopyranoside	1-20, 22-27
Man ₃ -Ac	73.0	4.82	C ₃ /H ₃ in 3- <i>O</i> -acetyl- β -D- mannopyranoside	1-27
Man ₄	79.3	3.38	C ₄ /H ₄ in β -D- mannopyranoside	1-19, 21-25, 27
Man ₅	76.9	3.63	C ₅ /H ₅ in β -D- mannopyranoside	1-27
Man ₆ -Ac	63.5	4.29/4.03	C ₆ /H ₆ in 6- <i>O</i> -acetyl- β -D- mannopyranoside	1-15, 17-19, 22-27
Man ₄ (r _a)	93.5	4.86	C ₁ /H ₁ in α -D-mannopyranoside of mannan reduction end	1-19, 21-27
Man ₄ (r _b)	93.6	4.59	C ₁ /H ₁ in β -D-mannopyranoside of mannan reduction end	1-15, 17-20, 22-77
Gal ₁	105.0	4.28	C ₁ /H ₁ in β -D-galactopyranoside	1-27
U ₁	97.6	5.10	C ₁ /H ₁ in 4- <i>O</i> -methyl- α -D-glucuronic acid	2, 4, 7, 12, 14, 16, 18, 22, 23, 24, 26
U ₄	81.6	3.09	C ₁ /H ₁ in 4- <i>O</i> -methyl- β -D-glucuronic acid	1-19, 22-27
U*	91.6/94.4	5.19/4.89	C ₁ /H ₁ in galacturonic acid	1-19, 22-27

^aContours assignments were assessed according to the literature.⁹

^bThe numbers refer to the following samples: 1: sdGGM raw; 2: sdGGM SSEH pellet/large; 3: sdGGM SSEH pellet/small; 4: sdGGM SSEH supernatant/large; 5: sdGGM SSEH supernatant/small; 6: esGGM raw; 7: esGGM SSEH pellet/large; 8: esGGM SSEH pellet/small; 9: esGGM SSEH supernatant/large; 10: esGGM SSEH supernatant/small; 11: epGGM raw; 12: epGGM SSEH pellet/large; 13: epGGM SSEH pellet/small; 14: epGGM SSEH supernatant/large; 15: epGGM SSEH supernatant/small; 16: sdGGM reference pellet/large; 17: sdGGM reference pellet/small; 19: sdGGM reference supernatant/large; 19: sdGGM reference supernatant/small; 20: esGGM reference pellet/large; 21: esGGM reference pellet/small; 22: esGGM reference

supernatant/large; 23: esGGM reference supernatant/small; 24: epGGM reference pellet/large; 25: epGGM reference pellet/small; 26: epGGM reference supernatant/large; and 27: epGGM reference supernatant/small.

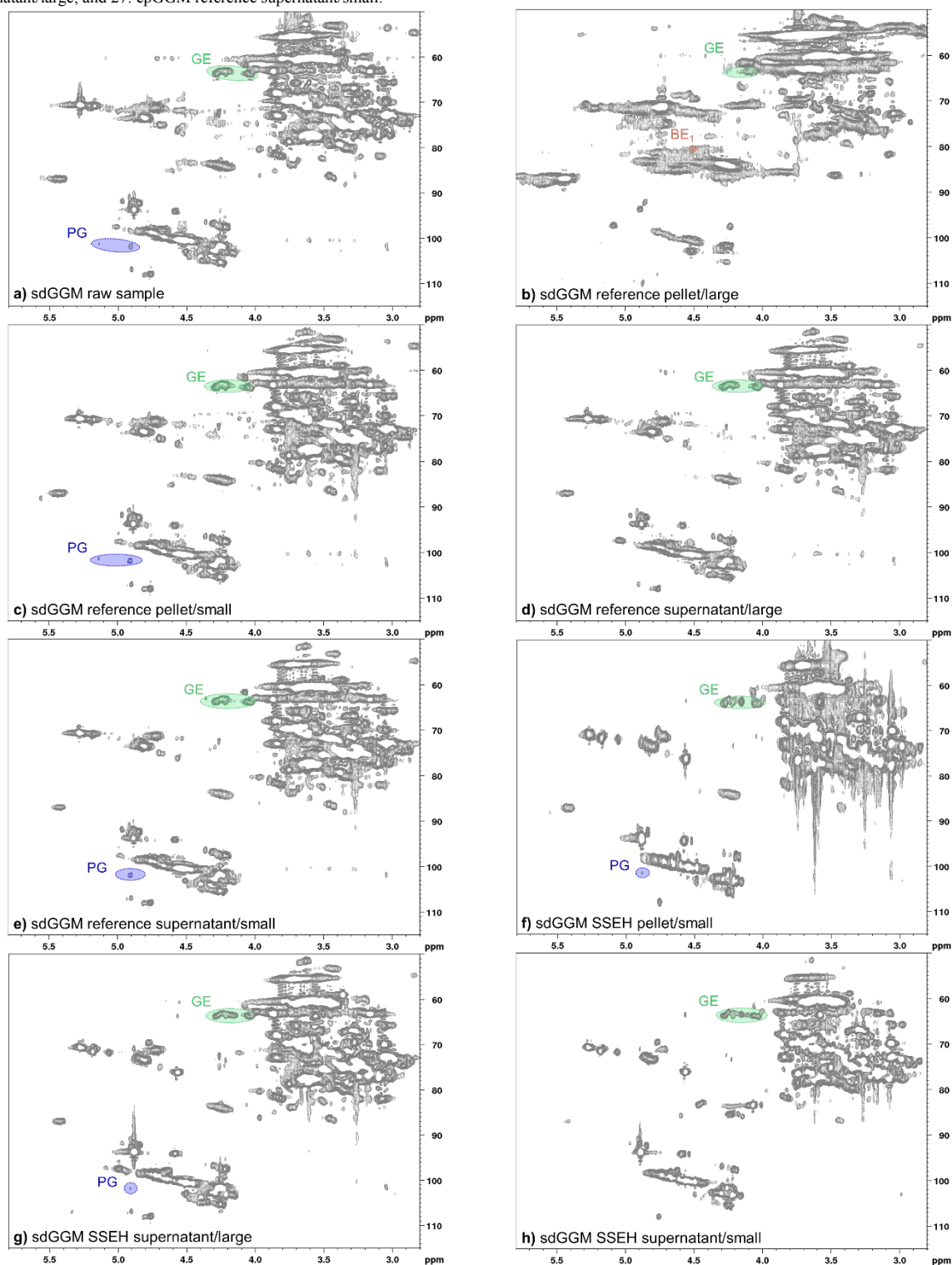


Figure S2. HSQC spectra of sdGGM raw sample (a), sdGGM reference pellet/large (b), sdGGM reference pellet/small (c), sdGGM reference supernatant/large (d), sdGGM reference supernatant/small (e), sdGGM SSEH pellet/small (f), sdGGM SSEH supernatant/large (g), and

sdGGM SSEH supernatant/small (**h**) fractions. Cross-signal for the various LCC bonds identified are highlighted in spectra.

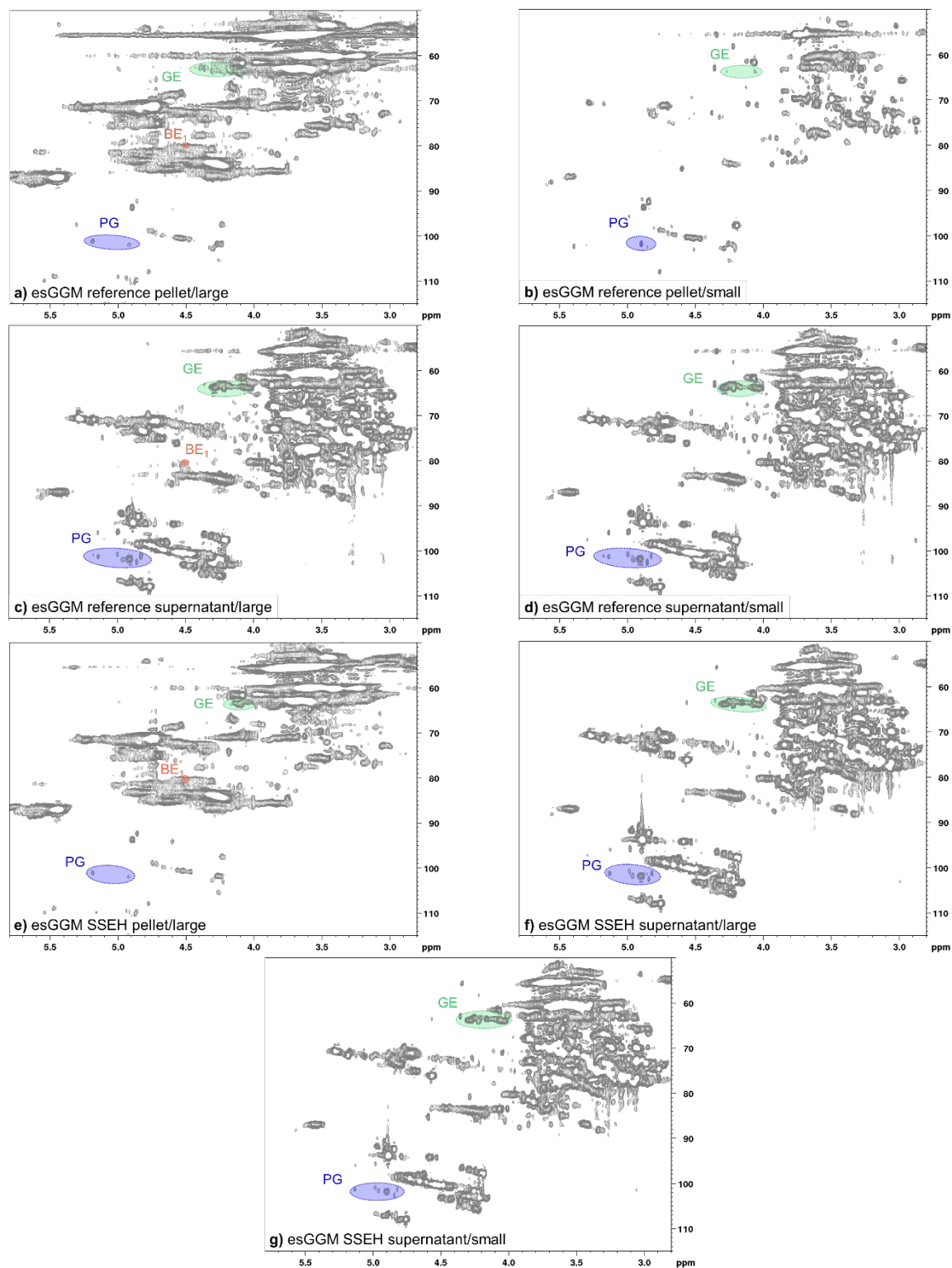


Figure S3. HSQC spectra of esGGM reference pellet/large (**a**), esGGM reference pellet/small (**b**), esGGM reference supernatant/large (**c**), esGGM reference supernatant/small (**d**), esGGM SSEH

pellet/large (e), esGGM SSEH supernatant/large (f), and esGGM SSEH supernatant/small (g) fractions. Cross-signal for the various LCC bonds identified are highlighted in spectra.

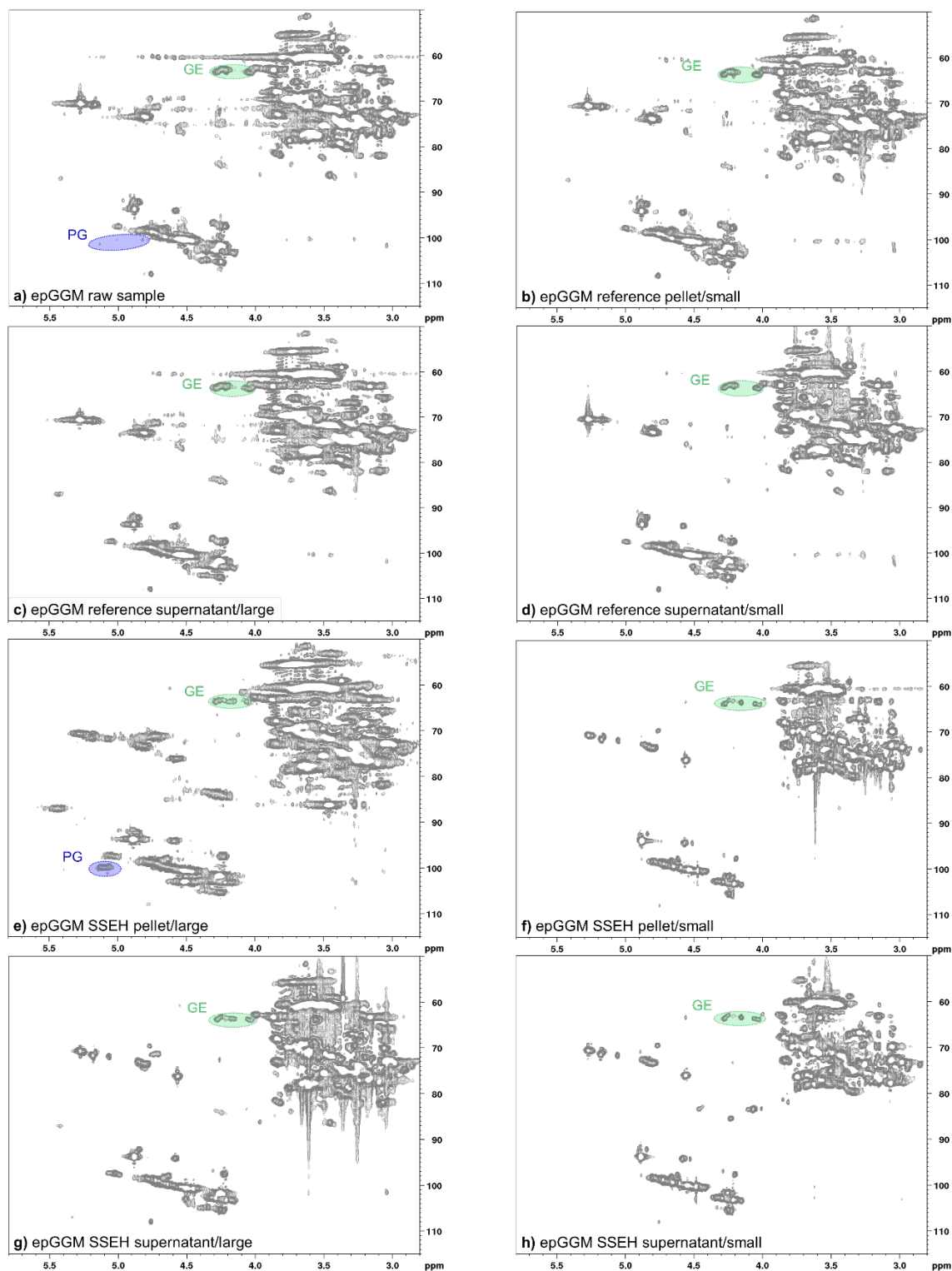


Figure S4. HSQC spectra of epGGM raw sample (a), epGGM reference pellet/small (b), epGGM reference supernatant/large (c), epGGM reference supernatant/small (d), epGGM SSEH

pellet/large (e), epGGM SSEH pellet/small (f), epGGM SSEH supernatant/large (g), and epGGM SSEH supernatant/small (h) fractions. Cross-signal for the various LCC bonds identified are highlighted in spectra.

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