

Facile Preparation Method for Inclusion Complexes between Amylose and Polytetrahydrofurans

*Rachmawati Rachmawati, Albert J. J. Woortman, Katja Loos**

Department of Polymer Chemistry, Zernike Institute for Advanced Materials, University of
Groningen, Nijenborgh 4, 9747AG Groningen, The Netherlands



Figure 1. Pressure vessel for preparing amylose inclusion complexes.



Figure 2. Distinctive appearance between amylose and amylose-PTHF complexes after 16 h rotation at 85 °C followed by sedimentation at 85 °C for 1 h (top) and 2 h (bottom).

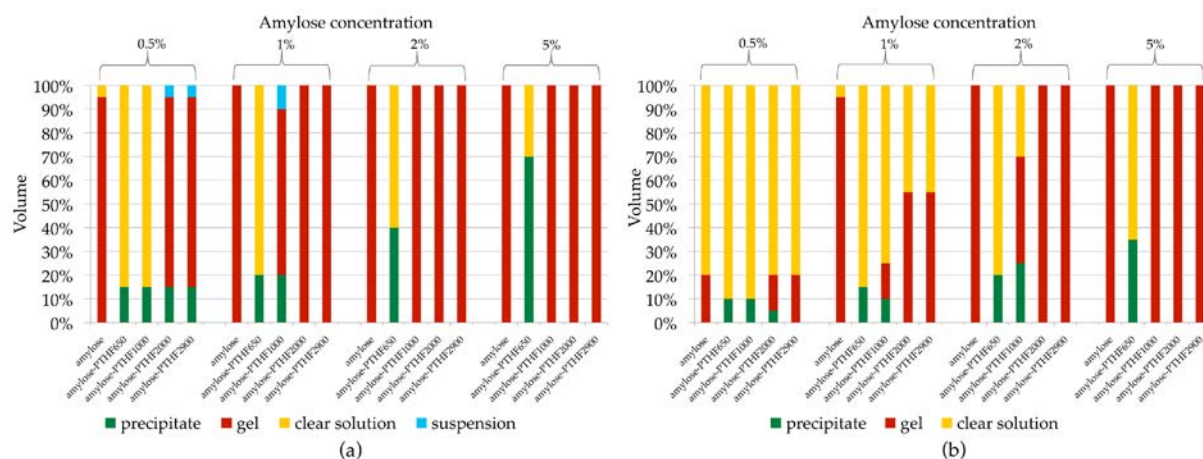


Figure 3. Estimation on gelation and sedimentation behavior of amylose-PTHF complexes after more than 16 h at room temperature (a) and after additional centrifugation for 30 min at 1000 rpm (b).

Table 1. Estimation of uncomplexed amylose for amylose-PTHF complexes.

Guest molecules	Reference (complex) ^a	Reference (amylose) ^a	Gel volume (complex)	Uncomplexed amylose	Uncomplexed amylose
PTHF650	Figure 3(a) (5%) no gel volume	Figure 3(a) (0.5%) gel 95%	0 < 95%	< 10%	< 5.3%
	Figure 2 (5%) precipitate 50%	Figure 3(a) (0.5%) gel 95%	[50/95]x[0.5/5]x 100%	< 5.3%	
PTHF1000	Figure 3(a) (1%) gel volume 70%	Figure 3(a) (0.5%) gel 95%	70% < 95%	< 50%	25-50%
	Figure 3(b) (2%) gel volume 45%	Figure 3(b) (0.5%) gel volume 20% and Figure 3(b) (1%) gel volume 95%	20% < 45% < 95%	25-50%	
PTHF2000 PTHF2900	Figure 3(b) (1%) gel volume 55%	Figure 3(b) (0.5%) gel volume 20%	55% > 20%	> 50%	> 50%

^aFigure 2 and 3 from the Supporting Information section are used as references.

Table 2. DSC data of inclusion complexes between amylose and PTHF.

Sample		Run (°C)	Heating scan		ΔH (J/g)	Cooling scan		ΔH (J/g)
			Onset (°C)	Peak (°C)		Onset (°C)	Peak (°C)	
Method A								
PTHF1100 ^a		1 (120)	21.4	23.3	21.8	10.9	7.7	-19.4
		2 (140)	20.9	23.5	23.4	10.6	7.7	-19.9
		3 (160)	20.1	23.1	22.9	10.3	7.5	-18.6
Potato amylose with								
5% PTHF1100		2 (140)	120.3	128.2	6.9	106.5	101.4	-5.0
		3 (160)	130.0	138.3	5.3	104.4	100.4	-1.6
20% PTHF1100		2 (140)	117.5	124.8	18.2	111.7	105.9	-17.5
		3 (160)	125.6	133.8	21.1	105.5	101.1	-12.8
20% PTHF650		3 (160)	126.0	132.3	17.7	105.0	97.8	-22.3
20% PTHF1000		3 (160)	128.4	133.6	14.2	106.6	100.5	-13.6
20% PTHF2000		3 (160)	130.4	134.6	2.7	108.4	102.2	-3.3
Method B								
Potato amylose with								
20% PTHF650		3 (160)	126.0	133.3	19.9	104.0	95.7	-24.8
20% PTHF1000		3 (160)	126.9	134.1	15.0	106.7	100.5	-16.0
20% PTHF2000		3 (160)	129.5	134.8	4.5	105.1	97.8	-4.8
Synthetic amylose with								
20% PTHF650		3 (160)	122.7	130.8	9.5	107.0	101.2	-14.6
20% PTHF1000		3 (160)	123.8	130.8	8.8	104.6	99.1	-7.6
20% PTHF2000		3 (160)	123.5	130.6	2.9	102.1	97.4	-2.6
20% PTHF1100		3 (160)	124.3	135.2	15.7	109.5	104.3	-9.2
20% PTHF650		4 (160)	127.1	133.4	19.3			
20% PTHF1000		4 (160)	125.3	134.5	14.2			
20% PTHF2000		4 (160)	119.4	134.0	10.0			
20% PTHF1100		4 (160)	122.9	135.5	18.0			
Method C^b		1 (160)				60	40	(19-40)
Potato amylose		1 (160)						
Potato amylose with								
20% PTHF650		1 (160)	129.0	134.2	26.8	105.5	98.6	-17.2
20% PTHF1000		1 (160)	129.7	136.9	10.5	98.3	92.6	-8.7
20% PTHF2000		1 (160)				92.6	85.9	-3.7
20% PTHF2900		1 (160)				93.3	88.1	-1.5

^aMeasured as 2%(w/w) concentration in water. ^bMeasured as 5% (w/w) concentration in water. The rest of the DSC data were obtained as 10% (w/w) concentration in water. The amount of the guest PTHF was calculated based on amylose (w/w).

Table 3. DSC data of first heating scan of inclusion complexes between synthetic amylose and PTHF (method B).

Day	Amylose-PTHF650			Amylose-PTHF1100		
	Onset (°C)	Peak (°C)	ΔH (J/g)	Onset (°C)	Peak (°C)	ΔH (J/g)
0	123.8	131.9	20.2	124.1	136.6	18.7
1	125.4	131.4	24.0	127.3	137.4	22.9
7	126.0	131.5	24.2	128.1	137.3	21.7

Measured for 20% (w/w) PTHF concentration based on amylose. The measurements were conducted to study the stability of amylose-PTHF complexes.

Table 4. XRD data of amylose-PTHF complexes.

Sample	Diffraction peaks							
	2 θ (°)/ <i>d</i> (nm)	Plane (<i>hkl</i>)	2 θ (°)/ <i>d</i> (nm)	Plane (<i>hkl</i>)	2 θ (°)/ <i>d</i> (nm)	Plane (<i>hkl</i>)	2 θ (°)/ <i>d</i> (nm)	Plane (<i>hkl</i>)
Amylose ^a	17.1/0.52		21.7/0.41					
PTHF	19.9/0.45		24.4/0.36					
Amylose inclusion complexes with								
PTHF650	13.1/0.68	200	(18.4)/0.48	221	21.4/0.42	450*	23.7/0.38	550*
	17.3/0.51	A	19.8/0.45	310	22.4/0.40	002 ^A	(25.0)/0.36	340
PTHF- b650	17.2/0.52	A	22.2/0.40	A	(25.9)/0.34	422*		
	19.6/0.45	P	23.9/0.37	222*	(28.6)/0.31	411		
PTHF1000	13.4/0.66	410*	20.0/0.44	310	23.7/0.38	222*	(28.5)/0.31	411
	17.3/0.51	A	21.4/0.42	450*	24.4/0.36	P		
	18.7/0.47	221	22.5/0.40	002 ^A	(27.3)/0.33	222		
PTHF2000	13.3/0.67	111	(18.4)/0.48	221	(21.4)/0.42	450*	24.3/0.37	P
	17.2/0.52	A	19.9/0.45	310 ^P	22.2/0.40	A	28.6/0.31	411
PTHF2900	17.3/0.51	A	20.0/0.44	310 ^P	22.5/0.40	A	24.4/0.36	P
	(18.4)/0.48	221	(21.4)/0.42	450*	(23.8)/0.37	222*		

^aThe amylose was solubilized and freeze-dried. A and P denote amylose and PTHF. **hkl* values of the diffracting planes are determined based on the orthorhombic unit cell of an amylose-*n*-butanol/*n*-pentanol complexes,²¹ while the rest are calculated based on amylose-fatty acid complexes.¹⁵ The *d*-spacing values are calculated based on Bragg's law for *n* = 1. The data in brackets are for shoulder-shaped peaks.