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APPENDICES

Appendix A**Culture media****1. Malt Extract Agar (MEA)**

Malt extract	20.0	g
Peptone	1.0	g
Dextrose	20.0	g
Agar	25.0	g

Dissolved in distilled water to final volume 1 liter.

Note that: sterile dextrose should be prepared separately and added after autoclaving to prevent caramelization.

2. Yeast Malt Agar (YMA)

Yeast extract	3.0	g
Malt extract	3.0	g
Bacto-peptone	5.0	g
Dextrose	10.0	g
Agar	20.0	g

Dissolved in distilled water to final volume 1 liter.

3. Pullulan Production (PM) medium

Sucrose	50.0	g
Bacto-peptone	0.6	g
K ₂ HPO ₄	5.0	g
MgSO ₄ ·7H ₂ O	0.4	g
NaCl	1.0	g
Yeast extract	0.4	g

Dissolved in 800 ml of distilled water and adjusted pH to 6.5 with HCl.

Added distilled water to final volume 1 liter.

4. Starch Agar medium

Soluble starch	10.0	g
NaNO ₃	2.0	g
MgSO ₄ ·7H ₂ O	0.5	g
NaCl	0.5	g
FeSO ₄ ·7H ₂ O	0.01	g
Yeast extract	0.4	g
Agar	20.0	g

Dissolved in distilled water to final volume 1 liter.

5. Crypto-Basol (CB) medium

Yeast nitrogenous base	6.7	g
Asparagine	2.0	g
KH ₂ PO ₄	5.0	g
Oat spelt xylan	10.0	g

Dissolved in distilled water to final volume 1 liter.

6. LB (Luria-Bertani) medium

Bacto-tryptone	10.0	g
Yeast extract	5.0	g
NaCl	10.0	g
Agar	10.0	g

Dissolved in 800 ml of distilled water and adjusted pH to 7.5 with NaOH.

Added distilled water to final volume 1 liter.

Note that: added antibiotics to final concentration 50 µg/ml, if needed.

Appendix B

PCR reactions and condition

1. DNA Amplification of DNA walking in GenomeWalker library

Reaction

28 μ l	sterile distilled water
10 μ l	5 x amplifying buffer
2 μ l	dNTPs (10 mM each)
2 μ l	AP1 primer (10 μ M)
2 μ l	GSP primer (10 μ M)
1 μ l	Phusion Hot Start DNA polymerase (2 U/ μ l) (Finnzymes)
<u>5 μl</u>	DNA library
50 μ l	Total volume

Condition

98°C	30 sec	
98°C	10 sec	} 35 cycles
55°C	30 sec	
72°C	1 min	
72°C	7 min	

2. DNA Amplification of DIG-labeling probe

Reaction (Probe 3:1 ratio (unlabeled/labeled))

1 μ l	DNA (~0.1 ng)
6 μ l	primer AP-GSP3f (3.2 pmol/ μ l)
6 μ l	primer AP-GSP5r (3.2 pmol/ μ l)
5 μ l	10 x amplifying buffer
1 μ l	dATP (10 mM)
1 μ l	dCTP (10 mM)
1 μ l	dGTP (10 mM)

0.75 μ l dTTP (10 mM)
 2.5 μ l DIG UTP (1 mM)
 23.5 μ l sterile distilled water
 2 μ l *Taq* DNA polymerase (1 U/ μ l) (Roche)
 50 μ l Total volume

Condition

96°C 1 min
 96°C 30 sec
 55°C 1 min
 72°C 1 min
 72°C 7 min

} 35 cycles

3. DNA Amplification of α -amylase mRNA detection and large subunit (26S)

Reaction

5 μ l 10 x amplifying buffer
 2 μ l Primer F (10 μ M)
 2 μ l Primer R (10 μ M)
 5 μ l dNTPs (2 mM each)
 32 μ l sterile distilled water
 2 μ l Red *Taq* DNA polymerase (1 U/ μ l) (Sigma)
 2 μ l DNA template
 50 μ l Total volume

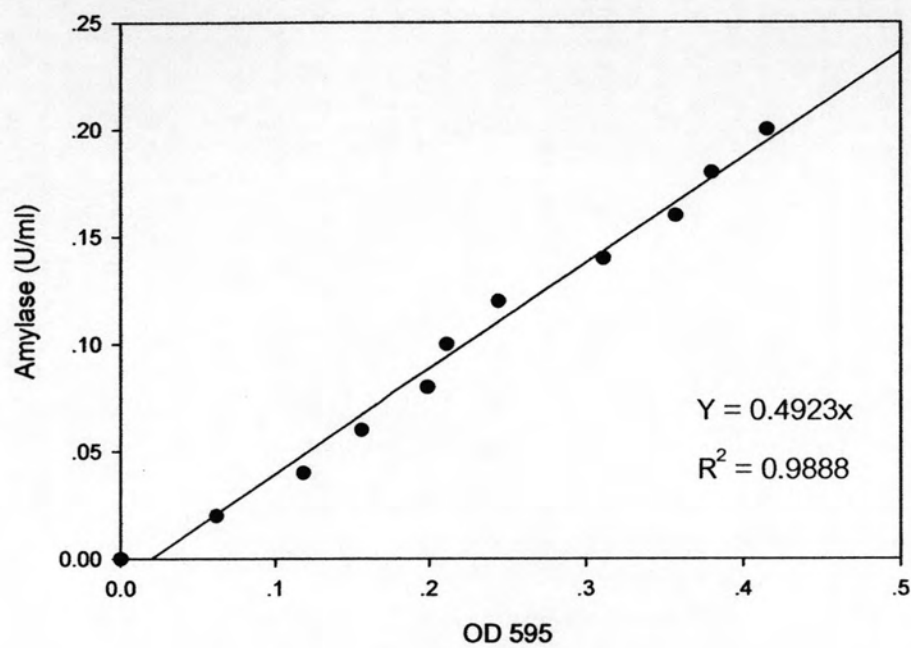
Condition

96°C 1 min
 96°C 30 sec
 52°C 1 min
 72°C 1 min
 72°C 7 min

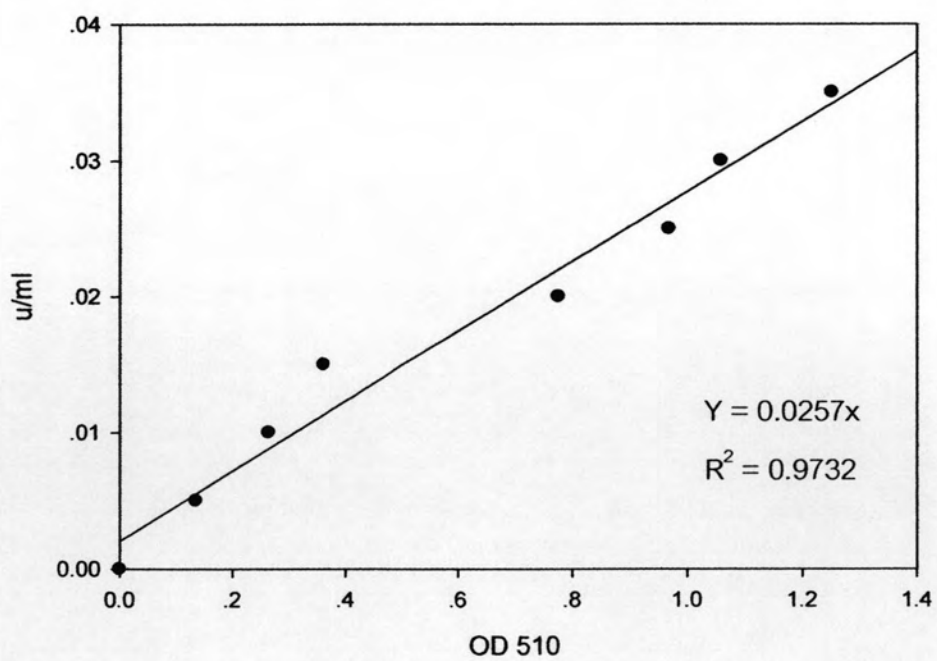
} 35 cycles

Appendix C

Standard curve

A. Standard curve for α -amylase

B. Standard curve for pullulanase



Appendix D

Data tables of Figure 4.15 – 4.26 and semi-log growth curves

Table 1 Cell growth (OD_{600}) of *A. pullulans* cultured in standard PM (5.0% sucrose and 0.1% N-sources) at day 0, 2, 4, 6, and 8 (Numbers were average \pm standard error)

Day	CU 3	CU 20	CU 36	NRM2	NRRL Y-12974
0	0.0 \pm 0.0	0.0 \pm 0.0	0.0 \pm 0.0	0.0 \pm 0.0	0.0 \pm 0.0
2	1.9 \pm 0.1	1.2 \pm 0.0	1.9 \pm 0.1	2.8 \pm 0.3	2.1 \pm 0.4
4	4.8 \pm 1.0	2.7 \pm 0.1	3.6 \pm 1.0	3.3 \pm 0.1	4.8 \pm 0.8
6	5.4 \pm 0.4	3.0 \pm 0.1	4.1 \pm 1.3	3.7 \pm 0.3	4.3 \pm 0.2
8	5.6 \pm 0.5	3.6 \pm 0.1	4.4 \pm 0.1	4.1 \pm 0.1	2.4 \pm 0.2

Table 2 Cell growth (OD_{600}) of *A. pullulans* NRRL Y-12974 cultured in modified PM at day 0, 2, 4, 6, and 8 (Numbers were average \pm standard error)

Day	5.0% Sucrose, 0.1% N-sources	5.0% Sucrose, 0.3% N-sources	5.0% Starch, 0.1% N-sources	5.0% Starch, 0.3% N-sources
0	0.0 \pm 0.0	0.0 \pm 0.0	0.0 \pm 0.0	0.0 \pm 0.0
2	2.1 \pm 0.4	2.0 \pm 0.1	2.6 \pm 0.0	3.0 \pm 0.1
4	4.8 \pm 0.8	3.7 \pm 0.4	4.0 \pm 0.5	4.6 \pm 0.6
6	4.3 \pm 0.2	4.3 \pm 0.5	4.3 \pm 0.2	6.2 \pm 1.2
8	2.4 \pm 0.2	5.7 \pm 0.7	4.5 \pm 0.2	6.9 \pm 0.4

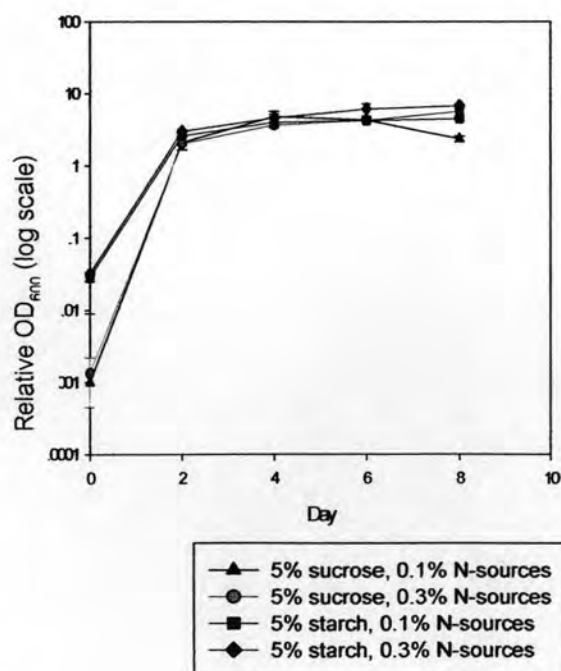
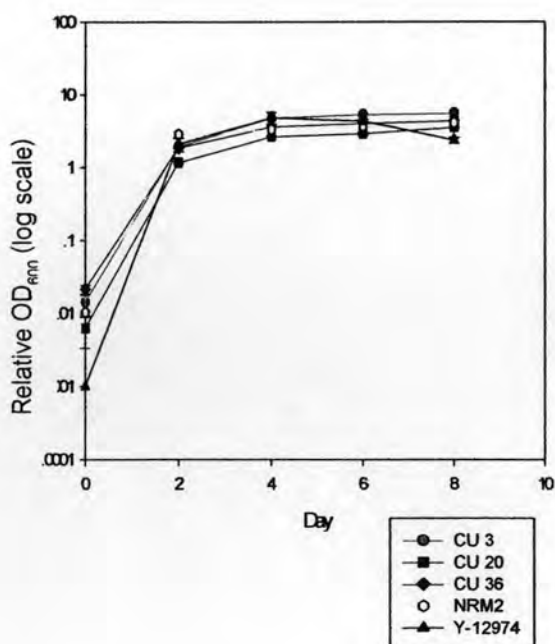


Table 3 pH measurement of culture supernatants of *A. pullulans* grown in standard PM (5.0% sucrose and 0.1% N-sources) at day 0, 2, 4, 6, and 8 (Numbers were average \pm standard error)

Day	CU 3	CU 20	CU 36	NRM2	NRRL Y-12974
0	6.5 \pm 0.0	6.5 \pm 0.0	6.5 \pm 0.0	6.5 \pm 0.0	6.5 \pm 0.0
2	4.0 \pm 0.0	4.4 \pm 0.0	4.8 \pm 0.1	3.9 \pm 0.0	5.5 \pm 0.4
4	3.6 \pm 0.0	3.7 \pm 0.1	3.9 \pm 0.0	3.7 \pm 0.0	4.3 \pm 0.1
6	3.5 \pm 0.0	3.7 \pm 0.0	3.7 \pm 0.1	3.6 \pm 0.0	4.2 \pm 0.0
8	3.5 \pm 0.0	3.7 \pm 0.1	3.5 \pm 0.0	3.6 \pm 0.0	4.2 \pm 0.0

Table 4 pH measurement of culture supernatants of *A. pullulans* NRRL Y-12974 grown in modified PM at day 0, 2, 4, 6, and 8 (Numbers were average \pm standard error)

Day	5.0% Sucrose, 0.1% N-sources	5.0% Sucrose, 0.3% N-sources	5.0% Starch, 0.1% N-sources	5.0% Starch, 0.3% N-sources
0	6.5 \pm 0.0	6.5 \pm 0.0	6.5 \pm 0.0	6.5 \pm 0.0
2	5.5 \pm 0.4	5.8 \pm 0.0	6.0 \pm 0.1	6.0 \pm 0.0
4	4.3 \pm 0.1	5.2 \pm 0.0	5.4 \pm 0.0	5.5 \pm 0.0
6	4.2 \pm 0.0	5.0 \pm 0.0	5.3 \pm 0.1	5.5 \pm 0.0
8	4.2 \pm 0.0	5.6 \pm 0.4	5.8 \pm 0.0	5.7 \pm 0.0

Table 5 EPS yield (g/L) of *A. pullulans* cultured in standard PM (5.0% sucrose and 0.1% N-sources) at day 2, 4, 6, and 8 (Numbers were average \pm standard error)

Day	CU 3	CU 20	CU 36	NRM2	NRRL Y-12974
2	9.8 \pm 0.1	3.8 \pm 0.1	3.1 \pm 0.1	5.8 \pm 0.0	12.0 \pm 0.1
4	17.6 \pm 0.2	14.1 \pm 1.0	6.2 \pm 0.1	10.0 \pm 0.1	31.3 \pm 1.1
6	21.6 \pm 0.8	22.2 \pm 0.8	7.1 \pm 1.0	11.6 \pm 0.2	39.0 \pm 0.1
8	24.6 \pm 0.4	29.0 \pm 1.4	6.7 \pm 0.3	17.0 \pm 0.3	39.8 \pm 0.2

Table 6 EPS yield (g/L) of *A. pullulans* NRRL Y-12974 cultured in modified PM at day 2, 4, 6, and 8 (Numbers were average \pm standard error)

Day	5.0% Sucrose, 0.1% N-sources	5.0% Sucrose, 0.3% N-sources	5.0% Starch, 0.1% N-sources	5.0% Starch, 0.3% N-sources
2	12.0 \pm 0.1	6.3 \pm 0.1	28.6 \pm 0.6	30.3 \pm 0.9
4	31.3 \pm 1.1	15.0 \pm 1.9	31.6 \pm 1.0	36.0 \pm 0.4
6	39.0 \pm 0.1	23.8 \pm 0.9	30.8 \pm 0.6	22.5 \pm 0.7
8	39.8 \pm 0.2	16.8 \pm 1.0	29.6 \pm 1.3	18.0 \pm 2.0

Table 7 EPS yield (g EPS dry weight / g cell dry weight) of *A. pullulans* cultured in standard PM (5.0% sucrose and 0.1% N-sources) at day 2, 4, 6, and 8 (Numbers were average \pm standard error)

Day	CU 3	CU 20	CU 36	NRM2	NRRL Y-12974
2	3.2	1.9	1.2	1.7	3.2
4	3.4	4.5	1.0	2.3	5.5
6	2.9	5.8	0.9	2.3	6.7
8	3.1	7.1	0.7	2.9	6.5

Table 8 EPS yield (g EPS dry weight / g cell dry weight) of *A. pullulans* NRRL Y-12974 cultured in modified PM at day 2, 4, 6, and 8 (Numbers were average \pm standard error)

Day	5.0% Sucrose, 0.1% N-sources	5.0% Sucrose, 0.3% N-sources	5.0% Starch, 0.1% N-sources	5.0% Starch, 0.3% N-sources
2	3.2	1.7	7.7	8.1
4	5.5	2.3	7.3	4.0
6	6.7	2.9	5.9	2.6
8	6.5	2.1	4.7	1.7

Table 9 Molecular weight (kD) of EPS precipitated from culture supernatants of *A. pullulans* grown in standard PM (5.0% sucrose and 0.1% N-sources) at day 2, 4, 6, and 8 (Numbers were average \pm standard error)

Day	CU 3	CU 20	CU 36	NRM2	NRRL Y-12974
2	722 \pm 45	1,070 \pm 90	664 \pm 97	512 \pm 18	128 \pm 118
4	327 \pm 30	302 \pm 71	415 \pm 131	89 \pm 11	170 \pm 40
6	500 \pm 51	114 \pm 8	186 \pm 15	33 \pm 3	52 \pm 3
8	459 \pm 9	43 \pm 5	76 \pm 11	18 \pm 4	26 \pm 1

Table 10 Molecular weight (kD) of EPS precipitated from culture supernatants of *A. pullulans* NRRL Y-12974 grown in modified PM at day 2, 4, 6, and 8 (Numbers were average \pm standard error)

Day	5.0% Sucrose, 0.1% N-sources	5.0% Sucrose, 0.3% N-sources	5.0% Starch, 0.1% N-sources	5.0% Starch, 0.3% N-sources
2	128 \pm 118	857 \pm 67	2 \pm 0	2 \pm 0
4	170 \pm 40	425 \pm 56	3 \pm 0	3 \pm 0
6	52 \pm 3	223 \pm 60	3 \pm 0	3 \pm 0
8	26 \pm 1	356 \pm 23	3 \pm 0	2 \pm 0

Table 11 Viscosity (cP) of EPS precipitated from culture supernatants of *A. pullulans* grown in standard PM (5.0% sucrose and 0.1% N-sources) at day 2, 4, 6, and 8. Measured at 25°C with a rotation of 30 rpm (shear rate of 39.6 1/S) (Numbers were average \pm standard error)

Day	CU 3	CU 20	CU 36	NRM2	NRRL Y-12974
2	4.3 \pm 0.1	9.9 \pm 0.5	5.8 \pm 1.1	3.7 \pm 0.3	6.4 \pm 1.0
4	5.0 \pm 0.1	7.0 \pm 0.2	32.1 \pm 2.5	2.5 \pm 0.2	3.4 \pm 0.1
6	8.3 \pm 0.8	5.6 \pm 0.4	38.5 \pm 5.0	2.1 \pm 0.0	2.5 \pm 0.1
8	8.5 \pm 0.7	3.4 \pm 0.4	66.1 \pm 3.9	2.4 \pm 0.1	2.3 \pm 0.2

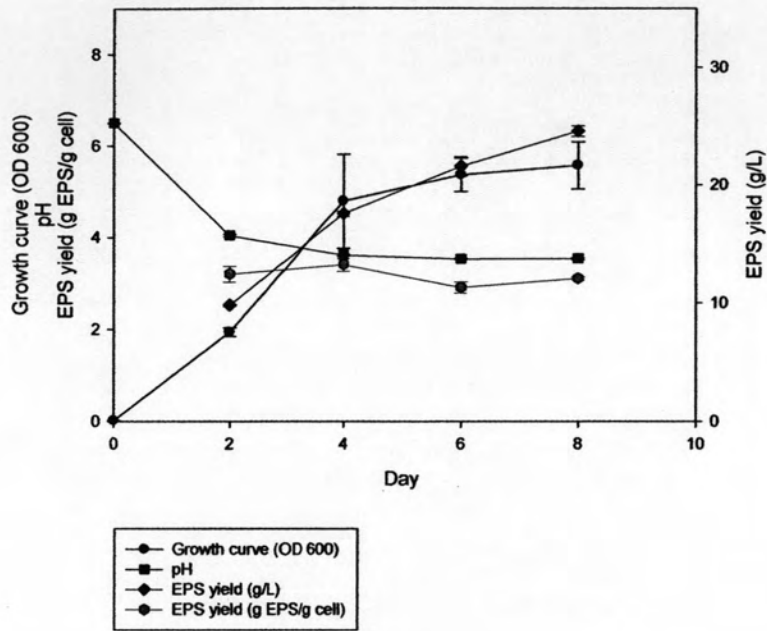
Table 12 Viscosity (cP) of EPS precipitated from culture supernatants of *A. pullulans* NRRL Y-12974 grown in modified PM at day 2, 4, 6, and 8. Measured at 25°C with a rotation of 30 rpm (shear rate of 39.6 1/S) (Numbers were average \pm standard error)

Day	5.0% Sucrose, 0.1% N-sources	5.0% Sucrose, 0.3% N-sources	5.0% Starch, 0.1% N-sources	5.0% Starch, 0.3% N-sources
2	6.4 \pm 1.0	6.0 \pm 0.6	2.4 \pm 0.3	1.5 \pm 0.1
4	3.4 \pm 0.1	7.5 \pm 0.9	2.5 \pm 0.1	2.1 \pm 0.1
6	2.5 \pm 0.1	5.0 \pm 1.5	2.1 \pm 0.0	1.7 \pm 0.1
8	2.3 \pm 0.2	5.8 \pm 0.2	2.4 \pm 0.2	3.3 \pm 0.3

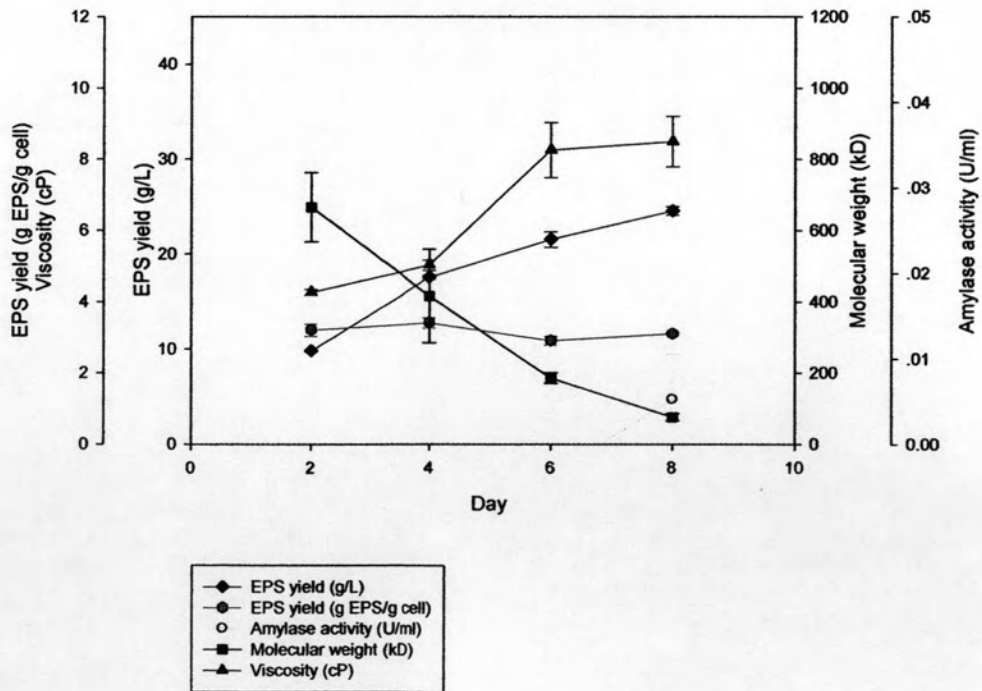
Appendix E

Graphs of combine data of each strain from Result 4.2.1

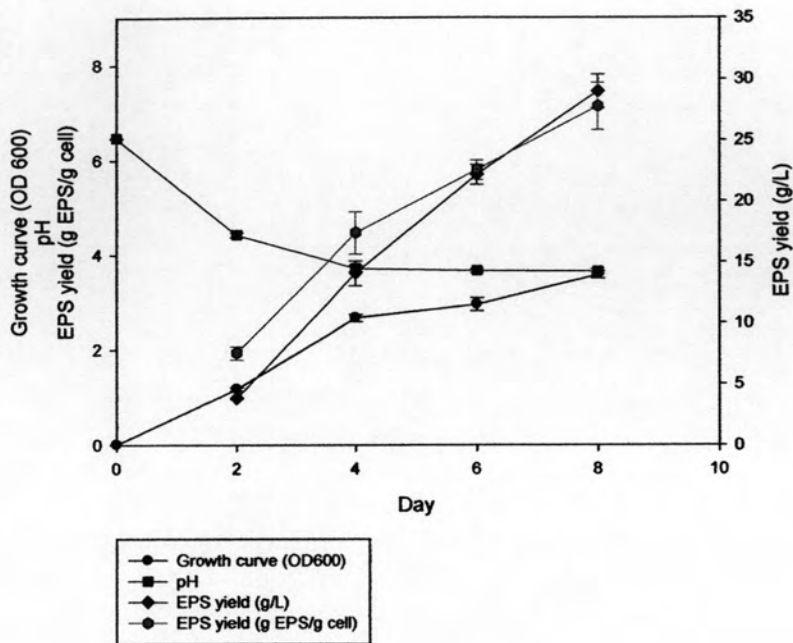
CU 3



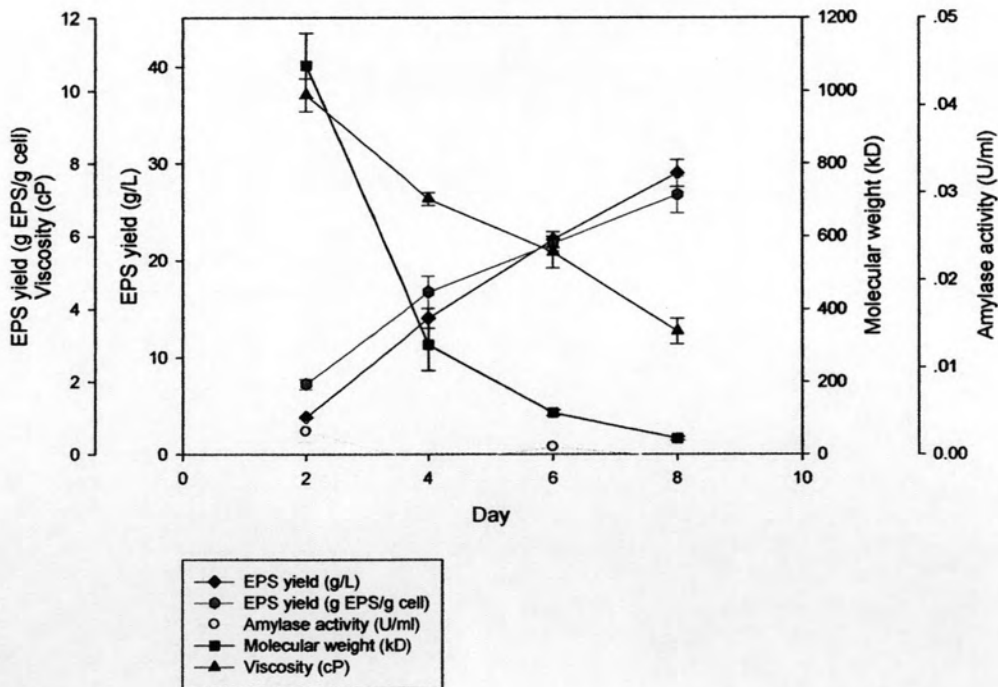
CU 3



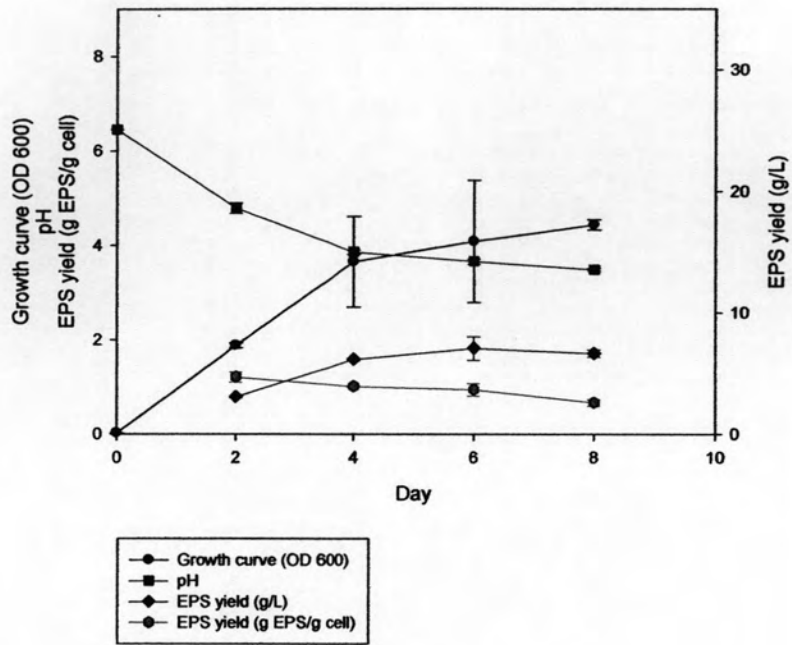
CU 20



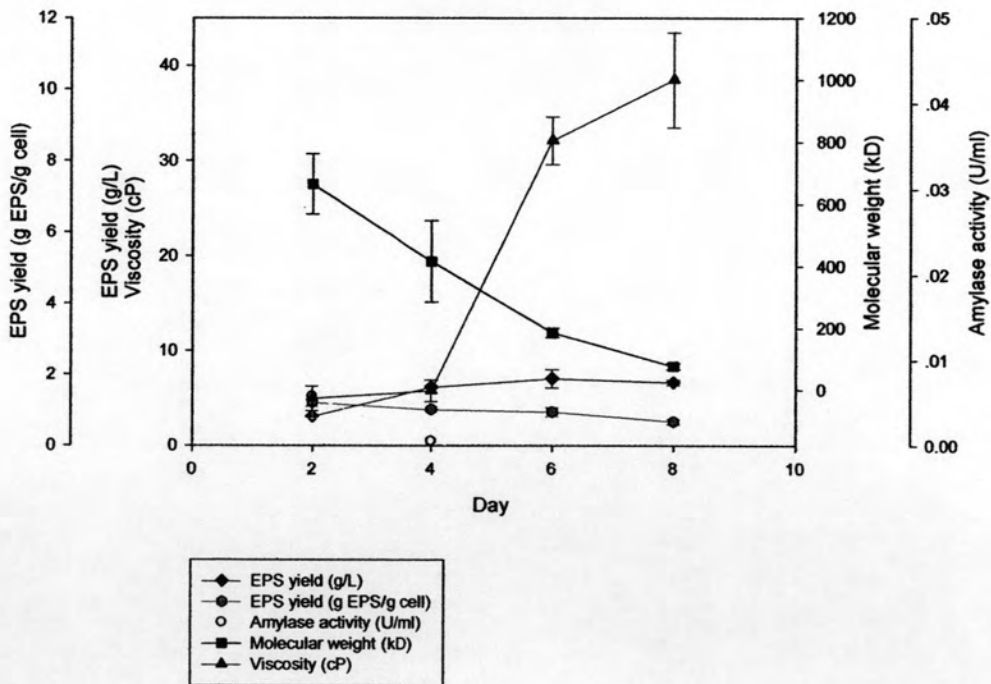
CU 20



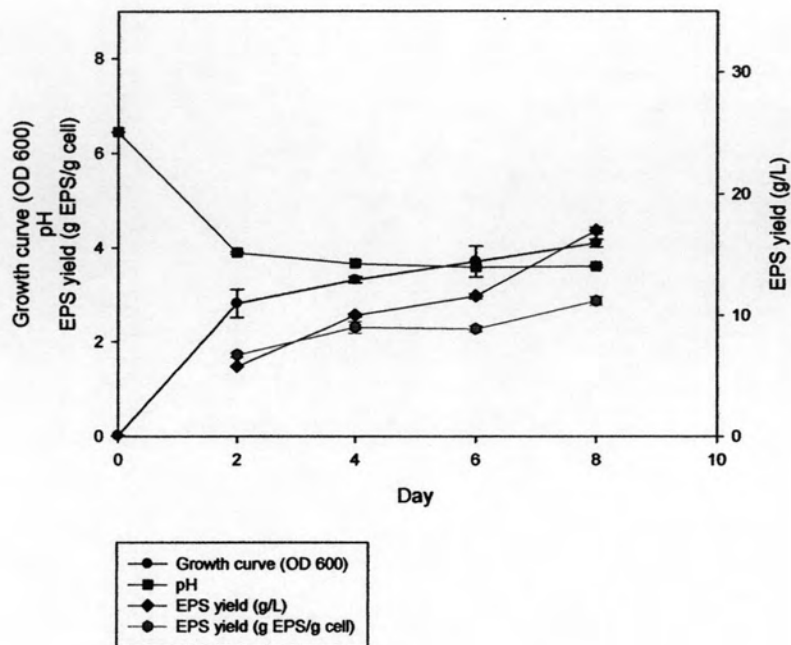
CU 36



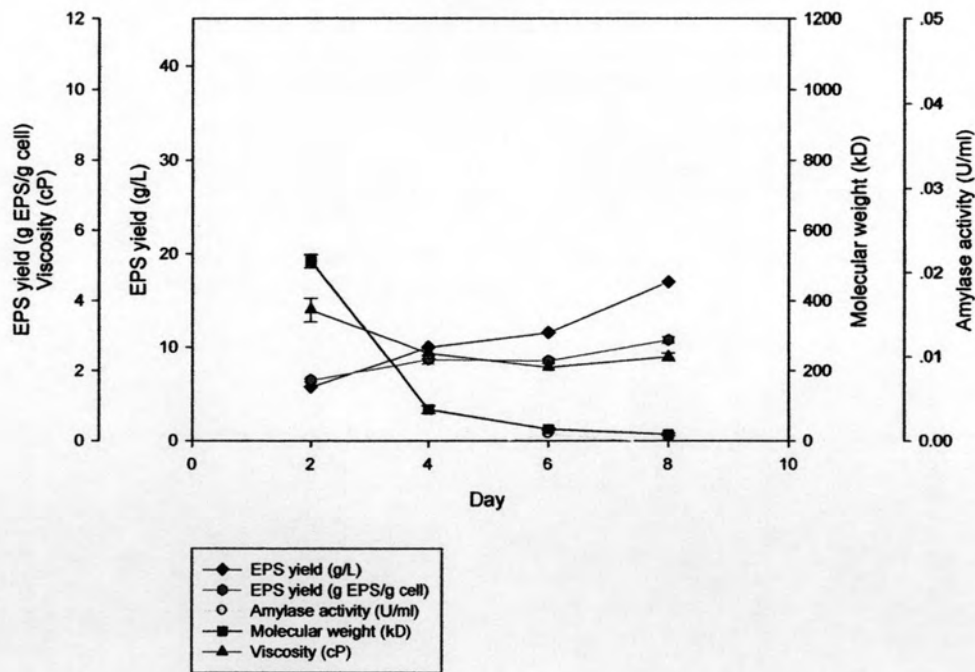
CU 36



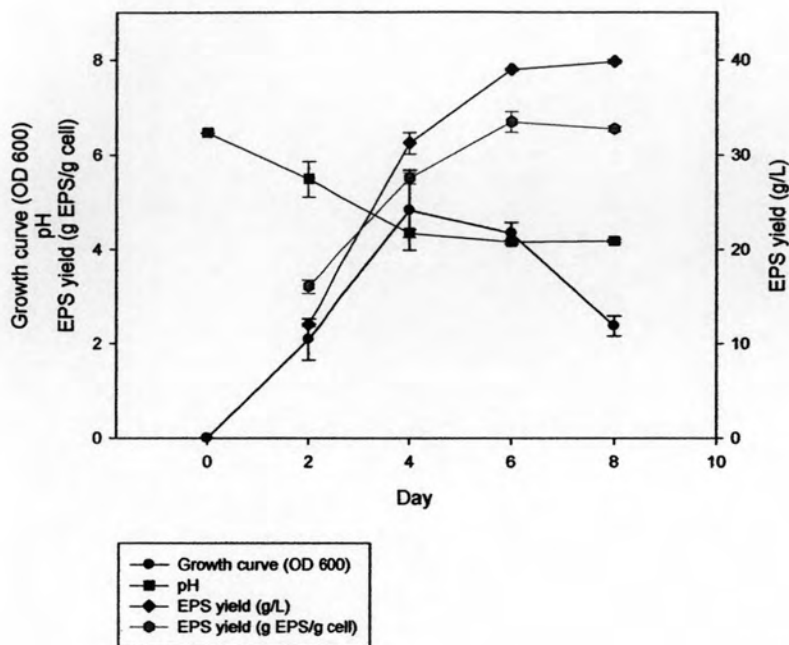
NRM2



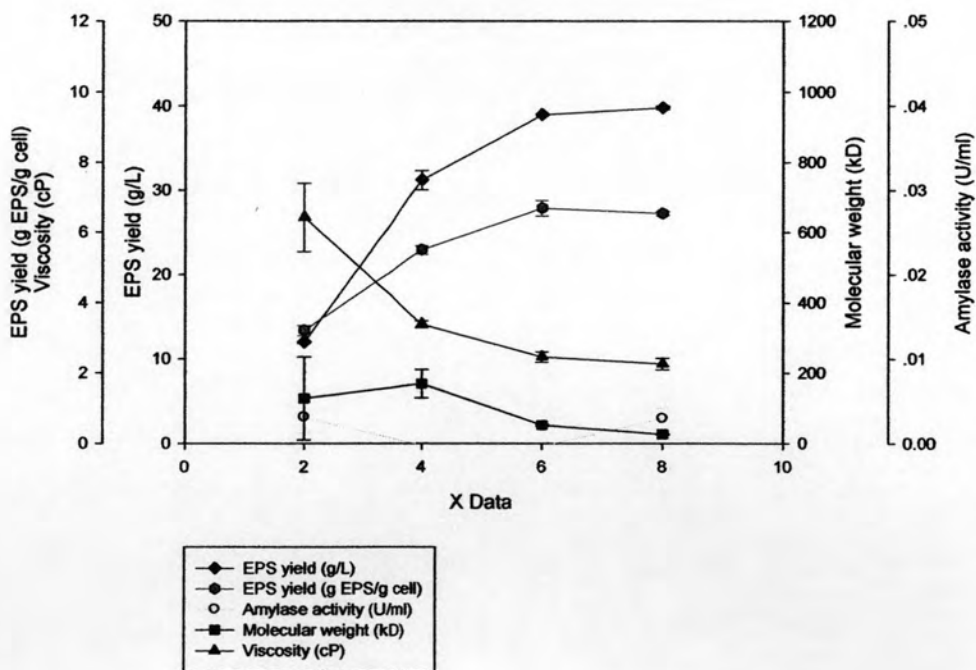
NRM2



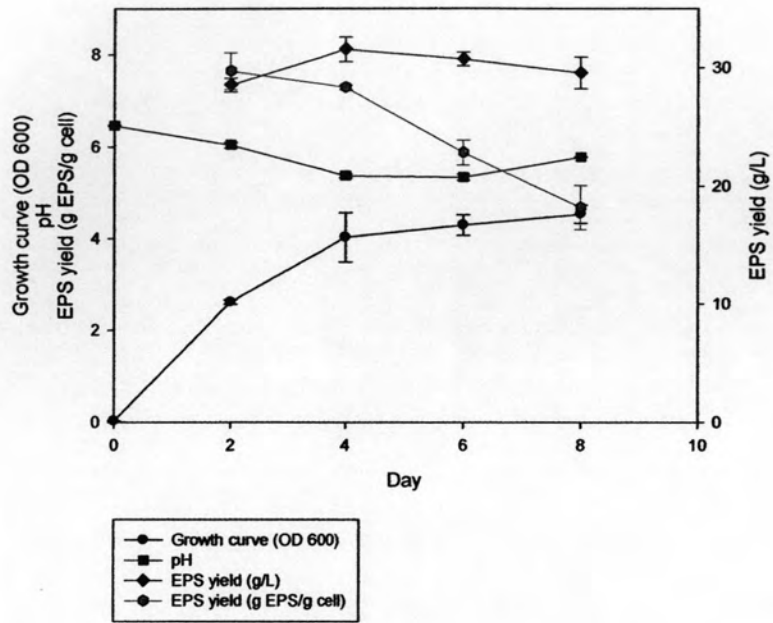
Y-12974 (PM; 5% sucrose, 0.1% N-sources)



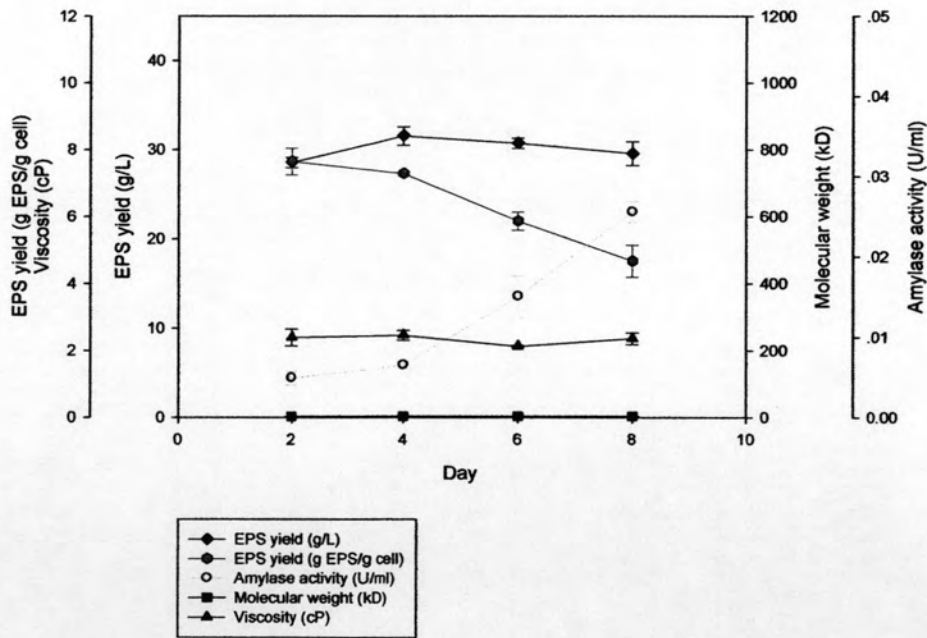
Y-12974 (PM; 5% sucrose, 0.1% N-sources)



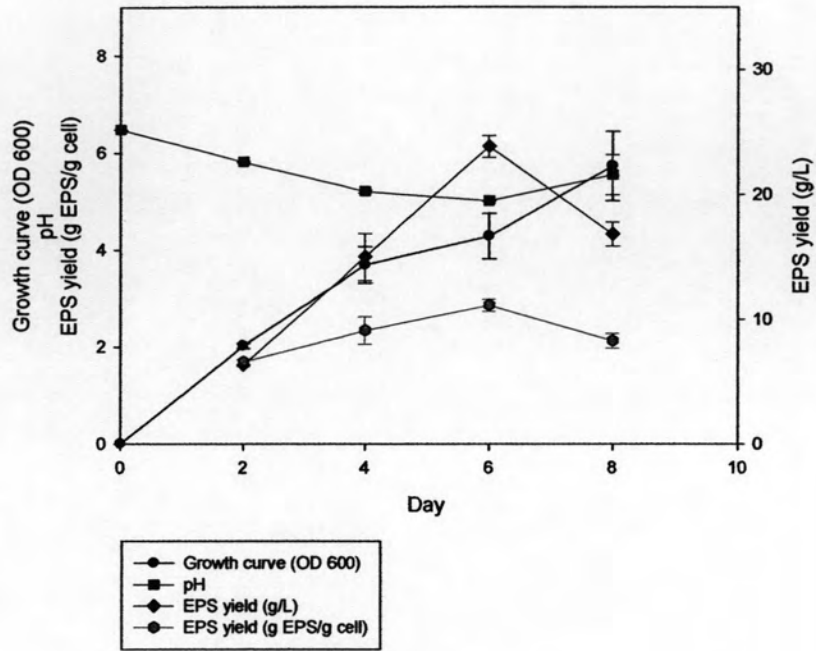
Y-12974 (PM; 5% starch, 0.1% N-sources)



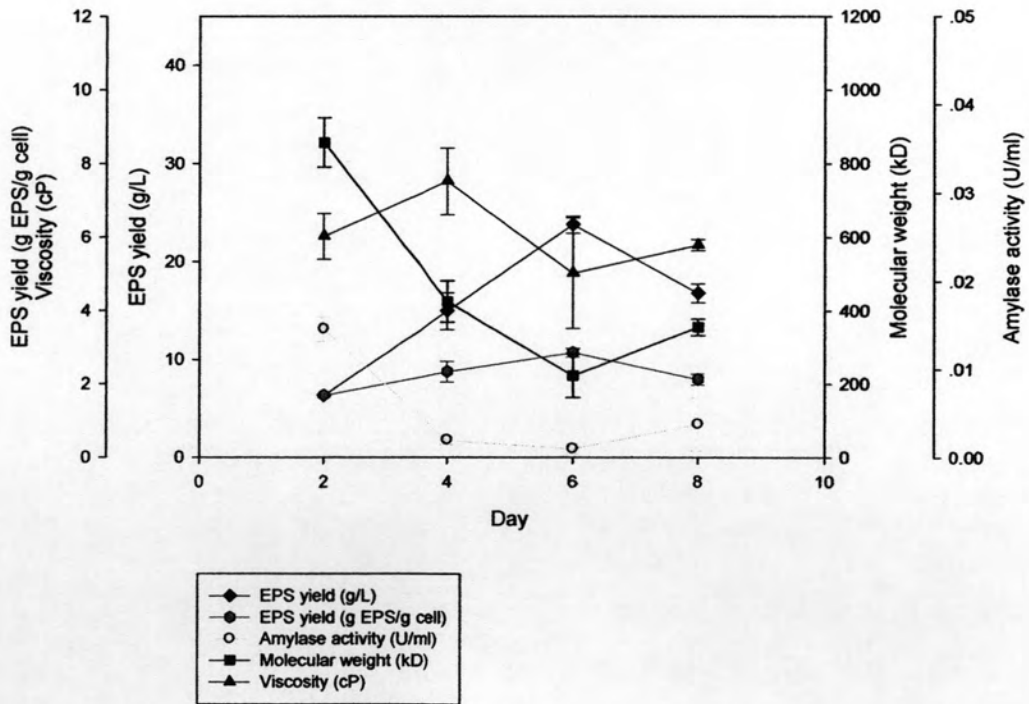
Y-12974 (PM; 5% starch, 0.1% N-sources)



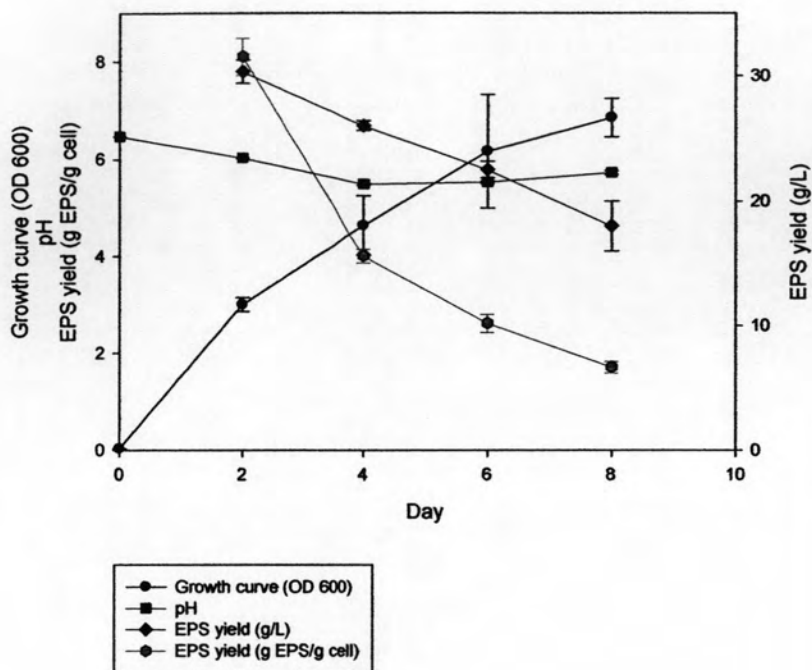
Y-12974 (PM; 5% sucrose, 0.3% N-sources)



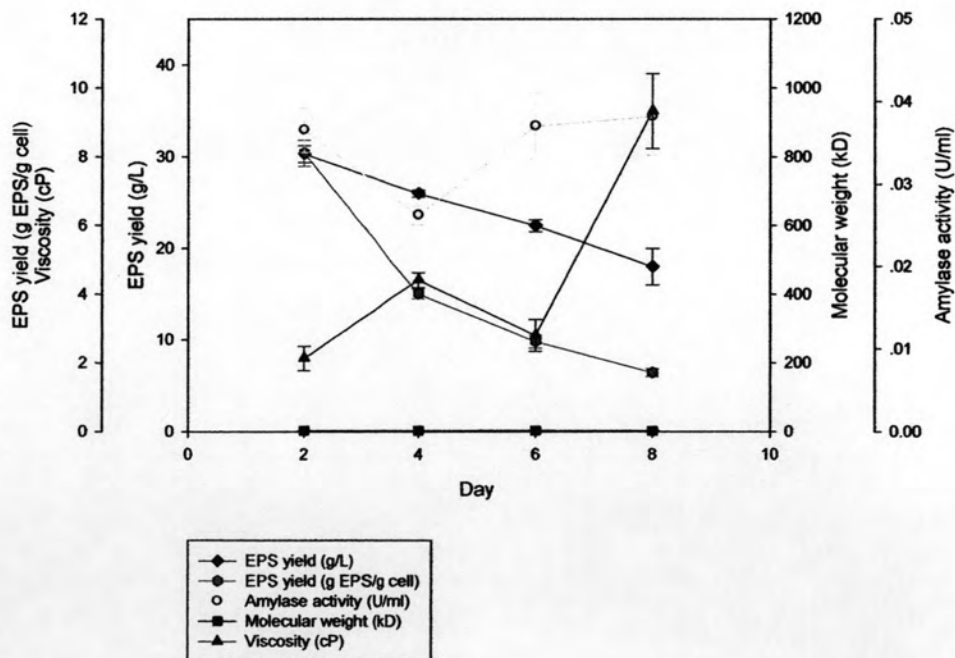
Y-12974 (PM; 5% sucrose, 0.3% N-sources)



Y-12974 (PM; 5% starch, 0.3% N-sources)



Y-12974 (PM; 5% starch, 0.3% N-sources)



Appendix F

NMR

Pullulan Repeat Unit: Proton NMR

Pullulan has a repeat unit of maltotriose (Glc- α -1,4-Glc- α -1,4-Glc) connected by an α -1,6-linkage. The NMR shows two α -1,4 proton signals (5.32 ppm and 5.36 ppm) and one α -1,6 proton (4.94 ppm). Equal intensity of three signal indicates a 2:1 ratio of α -1,4 : α -1,6. Chemical shifts of protons around the glucose rings (H2, H3, H4, H5, and H6) are shown in table 1 (courtesy of Neil P.J. Price, USDA, Peoria, IL, USA).

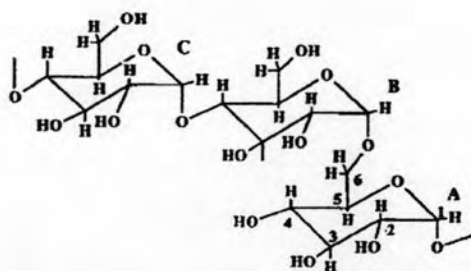
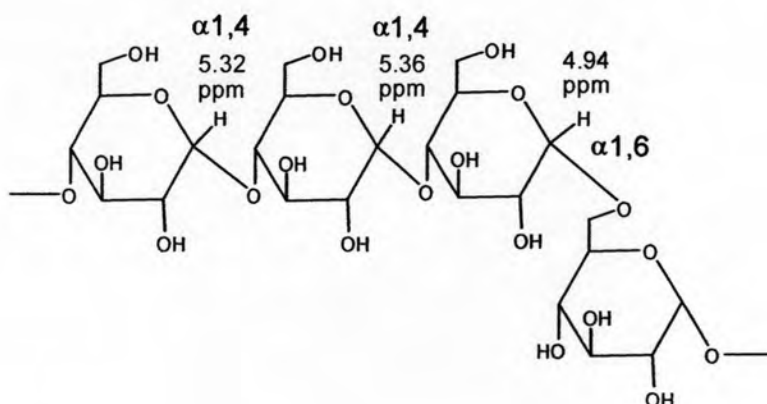
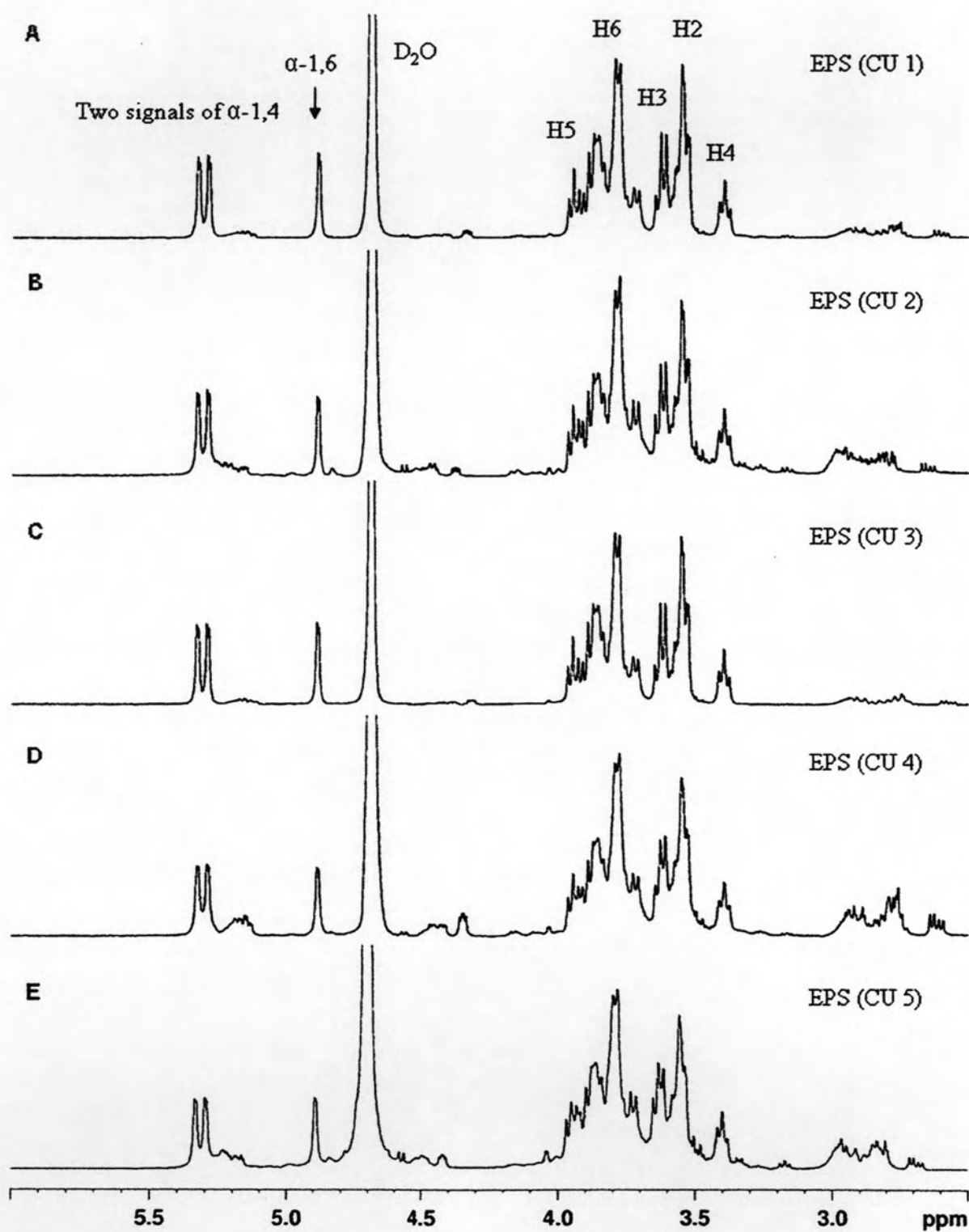
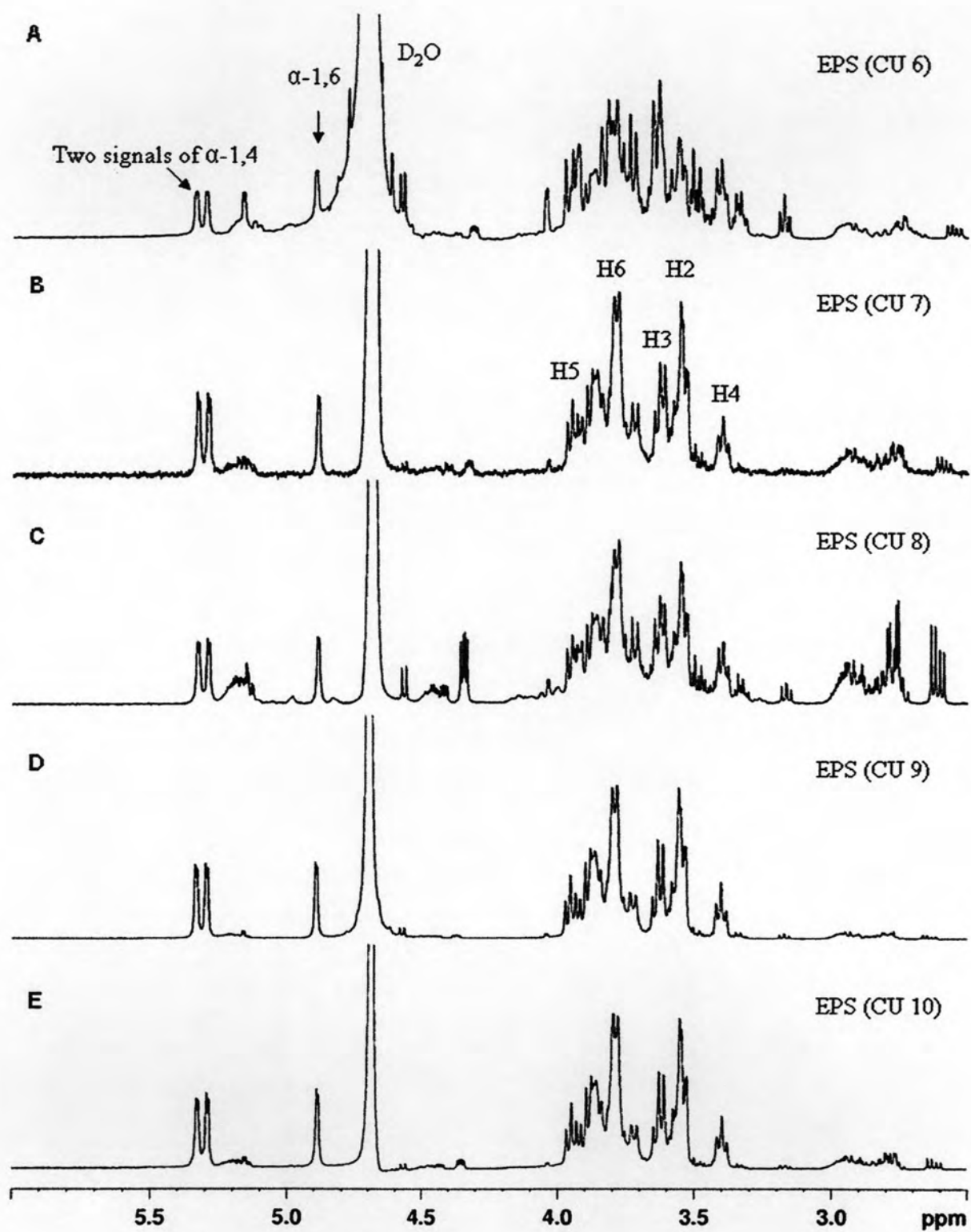


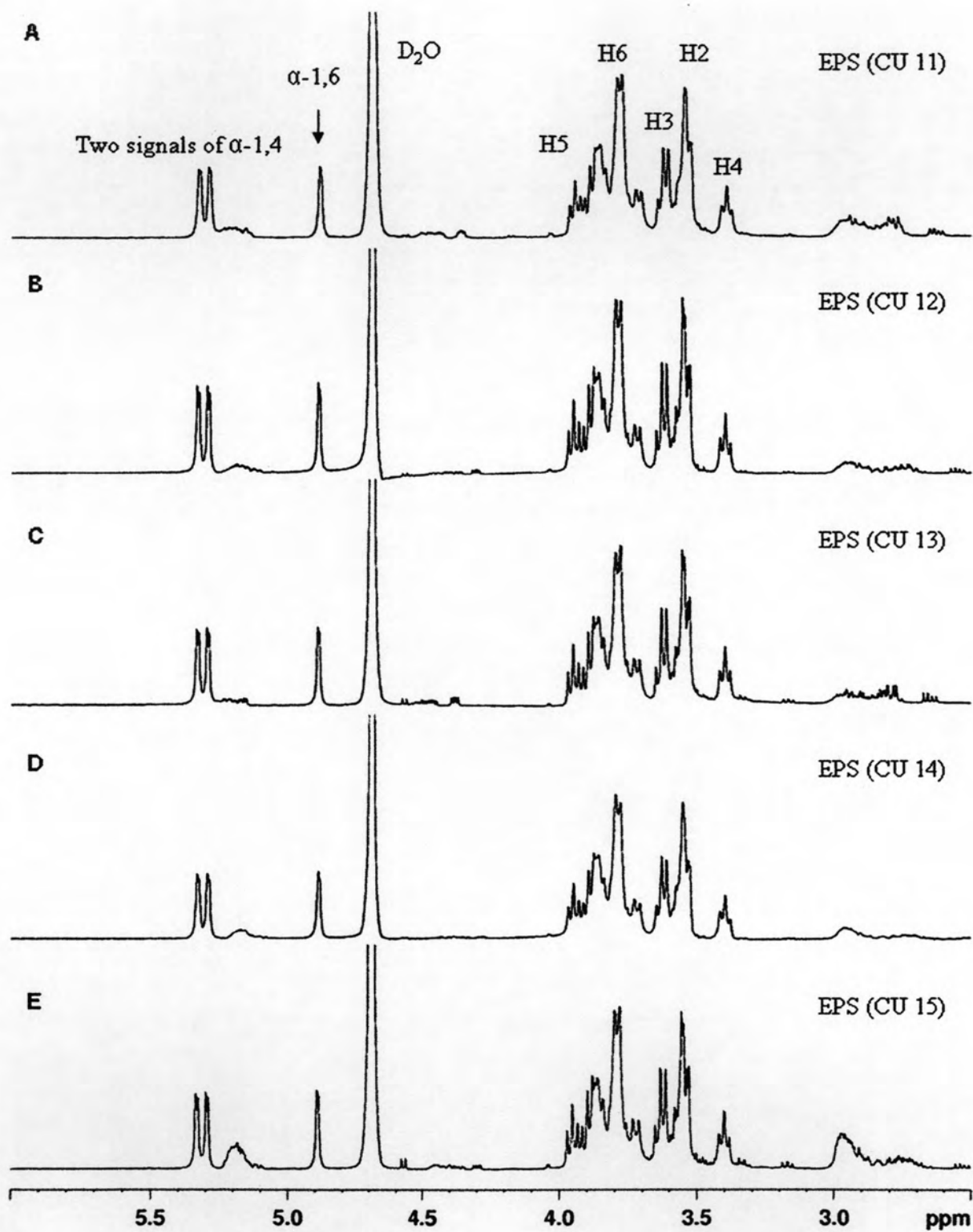
Table 1.
Chemical Shifts (ppm) of Pullulan Protons at 55°C (DSS reference).

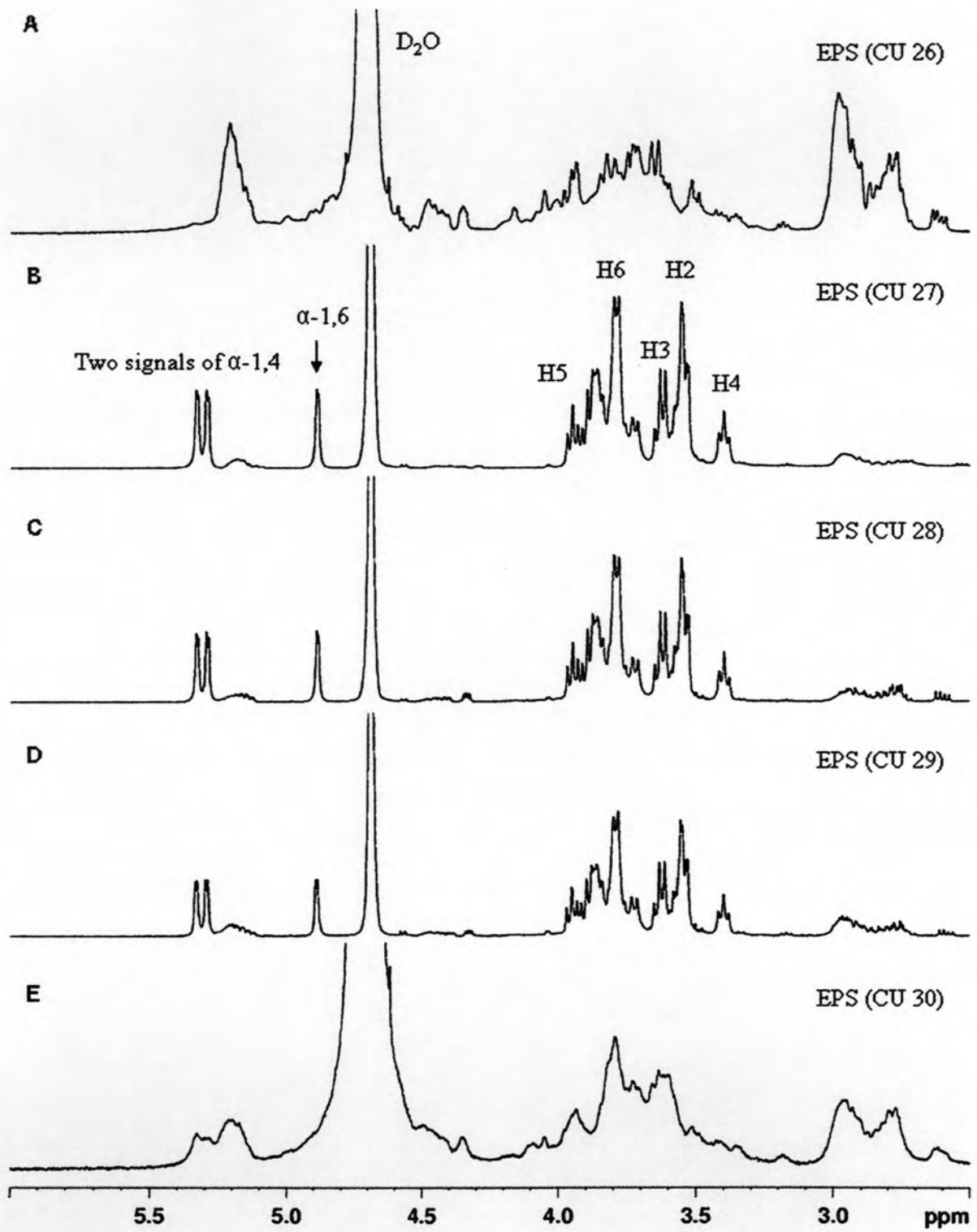
Sigma, Pfanstiehl	H-1	H-2	H-3	H-4	H-5	H-6
(1 \rightarrow 4)-(1 \rightarrow 6)-(1 \rightarrow 4) Glc	4.94	3.59	3.99	3.65	3.84	3.82, 3.91
(1 \rightarrow 4)-(1 \rightarrow 4)-(1 \rightarrow 6) Glc	5.36	3.63	3.94	3.58	3.91	3.82, 3.84
(1 \rightarrow 6)-(1 \rightarrow 4)-(1 \rightarrow 4) Glc	5.32	3.60	3.69	3.45	3.91	3.79, 3.91
AVEBE						
(1 \rightarrow 4)-(1 \rightarrow 6)-(1 \rightarrow 4) Glc	4.94	3.59	3.99	3.65	3.84	3.82, 3.91
(1 \rightarrow 4)-(1 \rightarrow 4)-(1 \rightarrow 6) Glc	5.36	3.63	3.94	3.58	3.91	3.82, 3.84
(1 \rightarrow 6)-(1 \rightarrow 4)-(1 \rightarrow 4) Glc	5.32	3.60	3.69	3.45	3.91	3.79, 3.91
(1 \rightarrow 4)-(1 \rightarrow 4)-(1 \rightarrow 4) Glc	5.31	3.62	3.98	3.65	3.91	n.d.
α Reducing Glc	5.23	3.56	3.97	3.65	n.d.	n.d.
β Reducing Glc	4.63	3.27	3.77	3.65	n.d.	n.d.

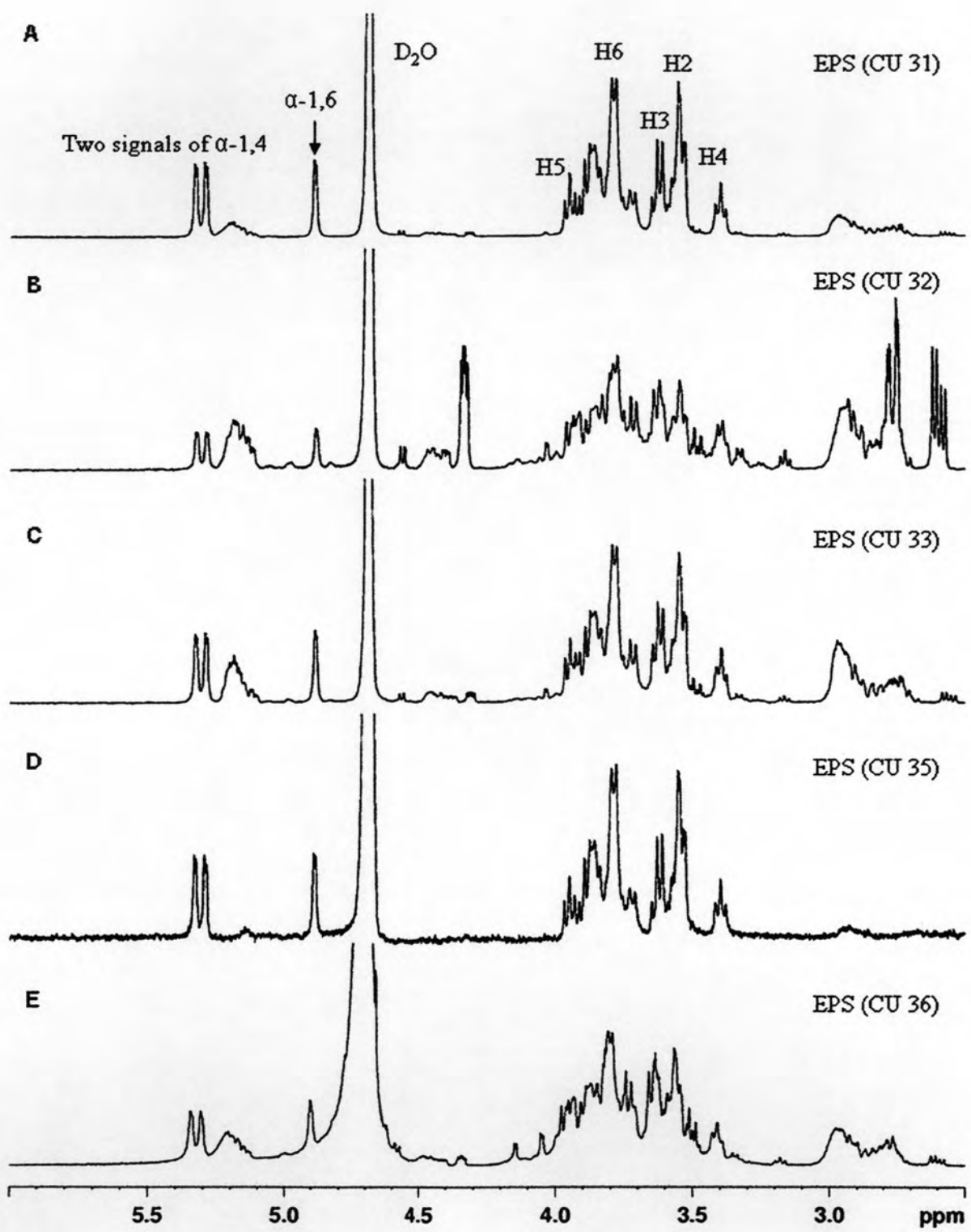
n.d. = not determined.

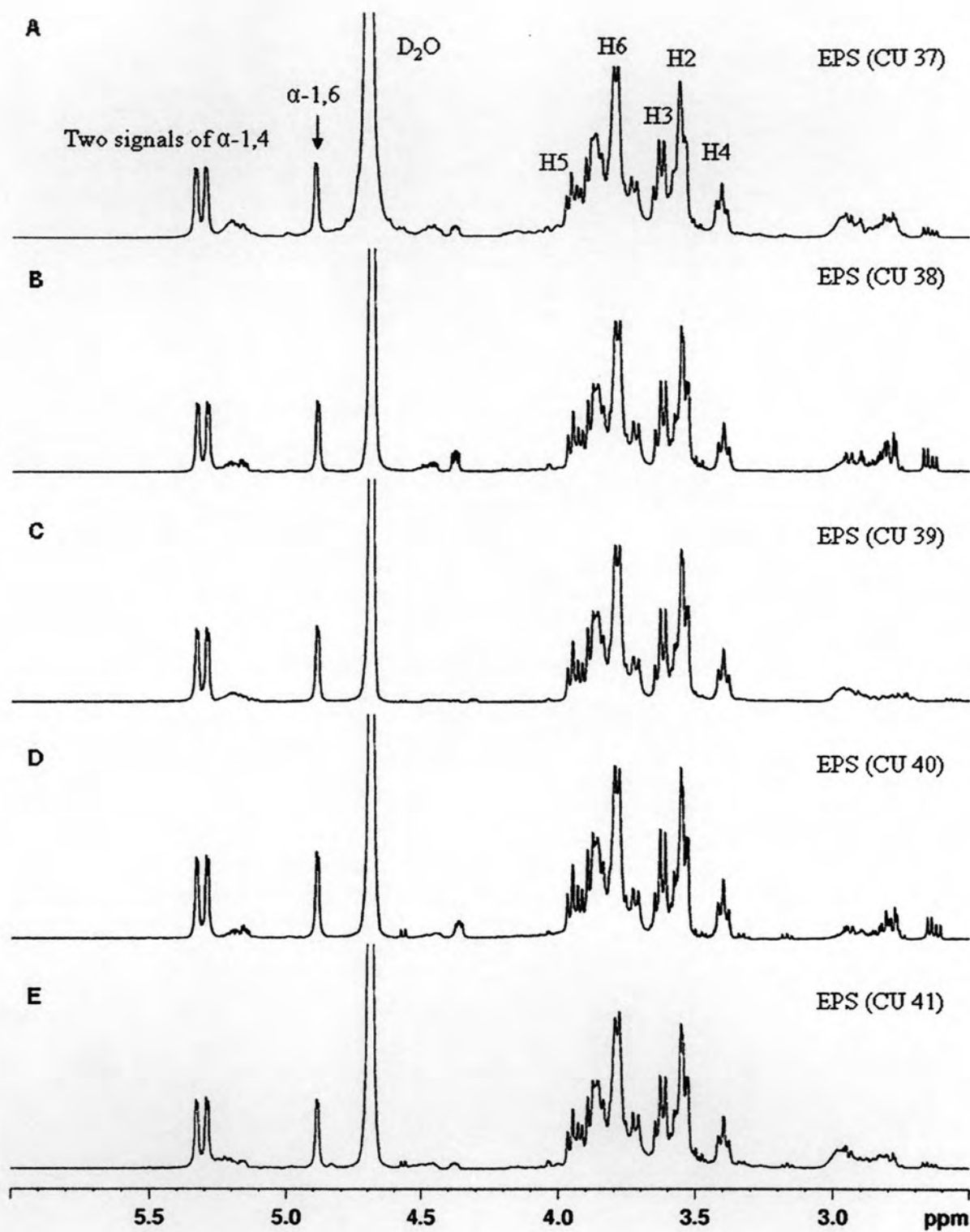


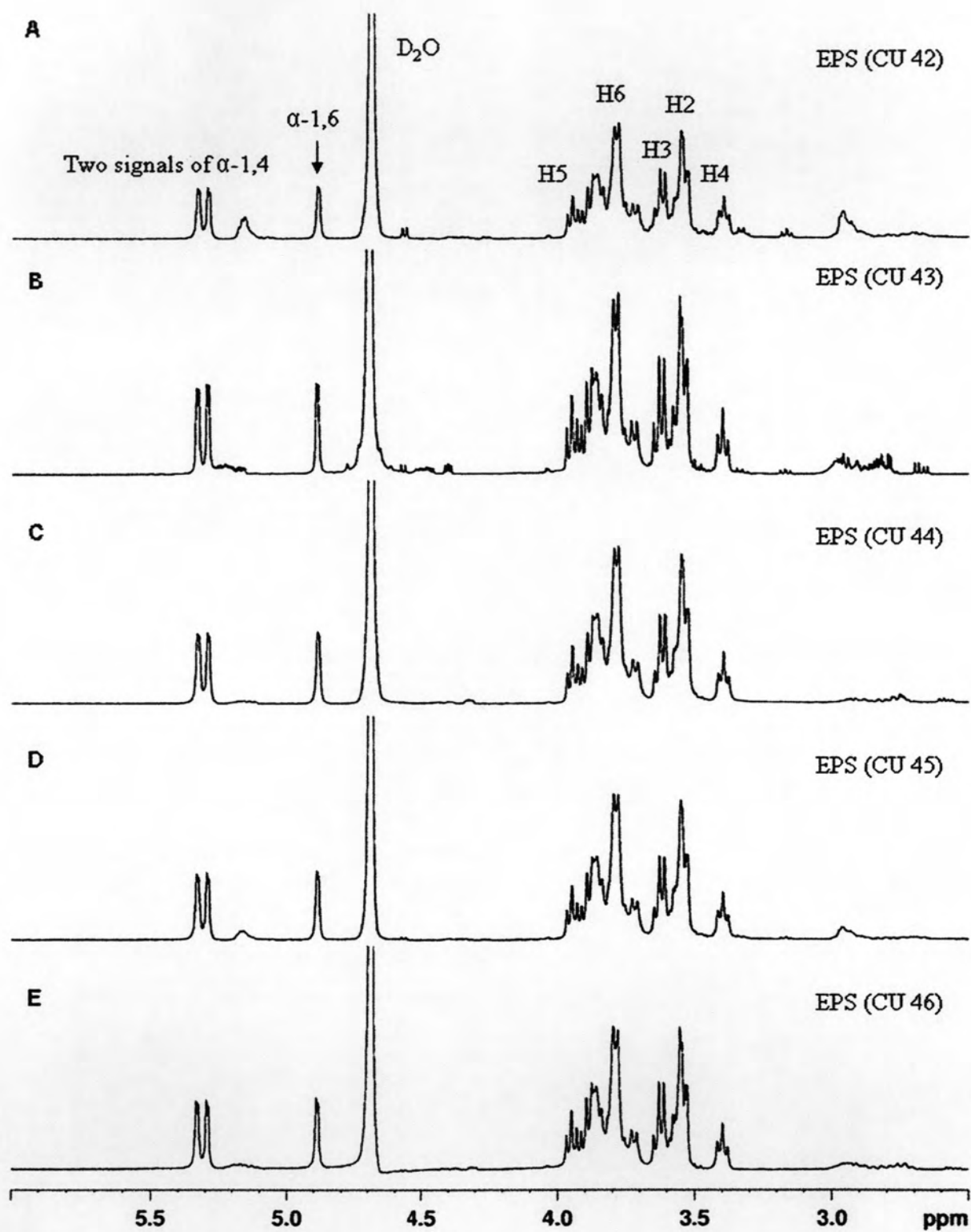


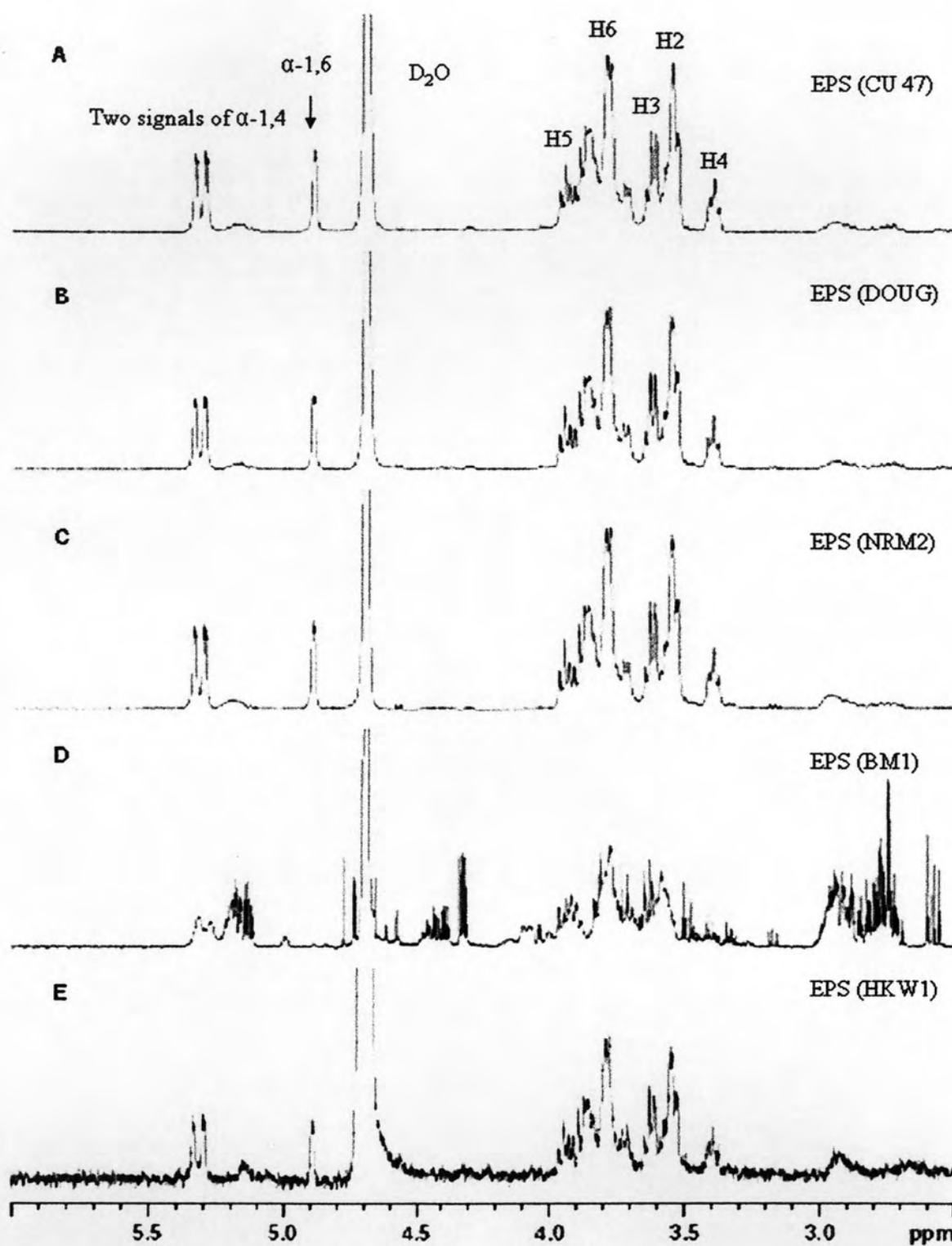




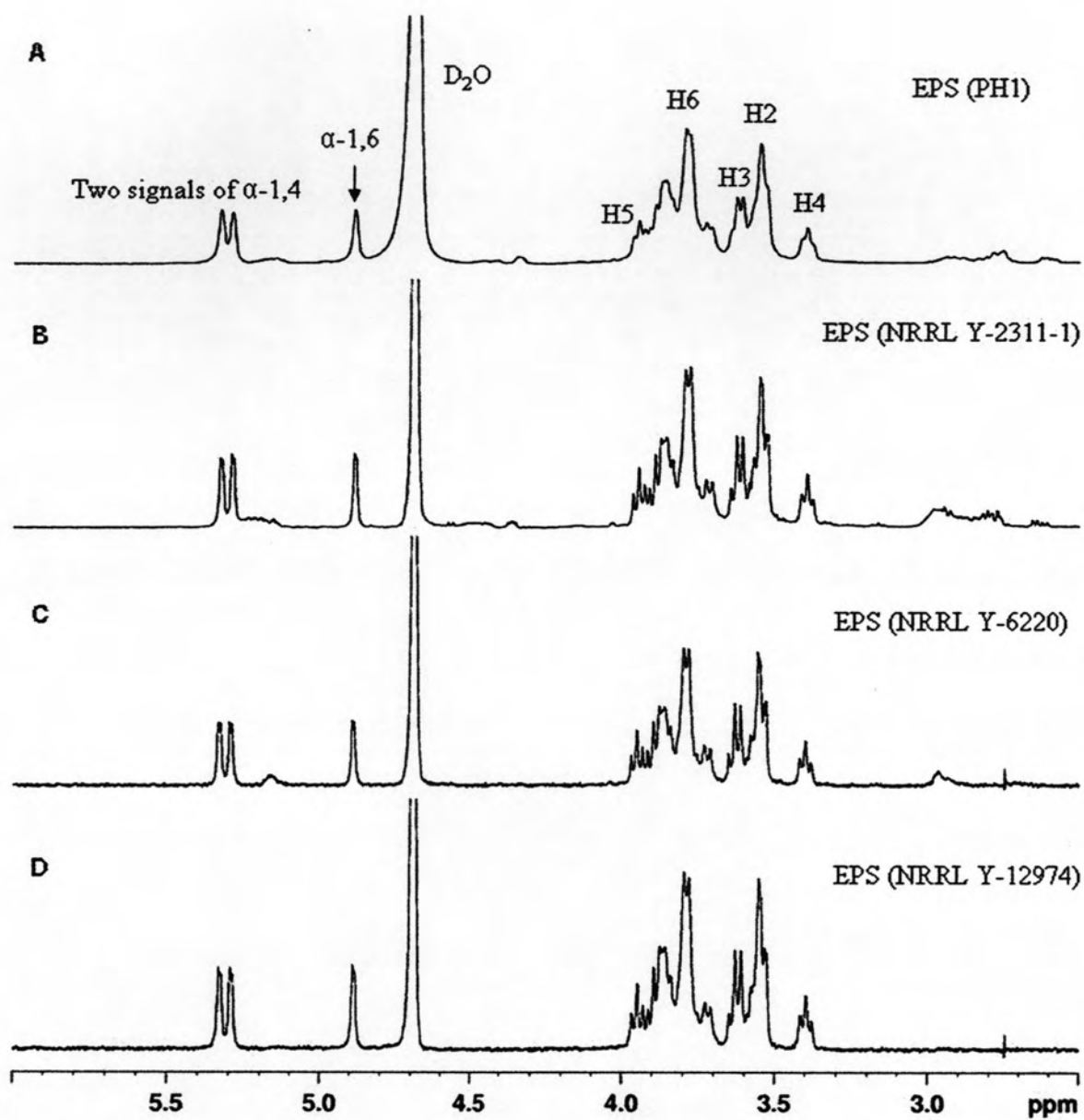


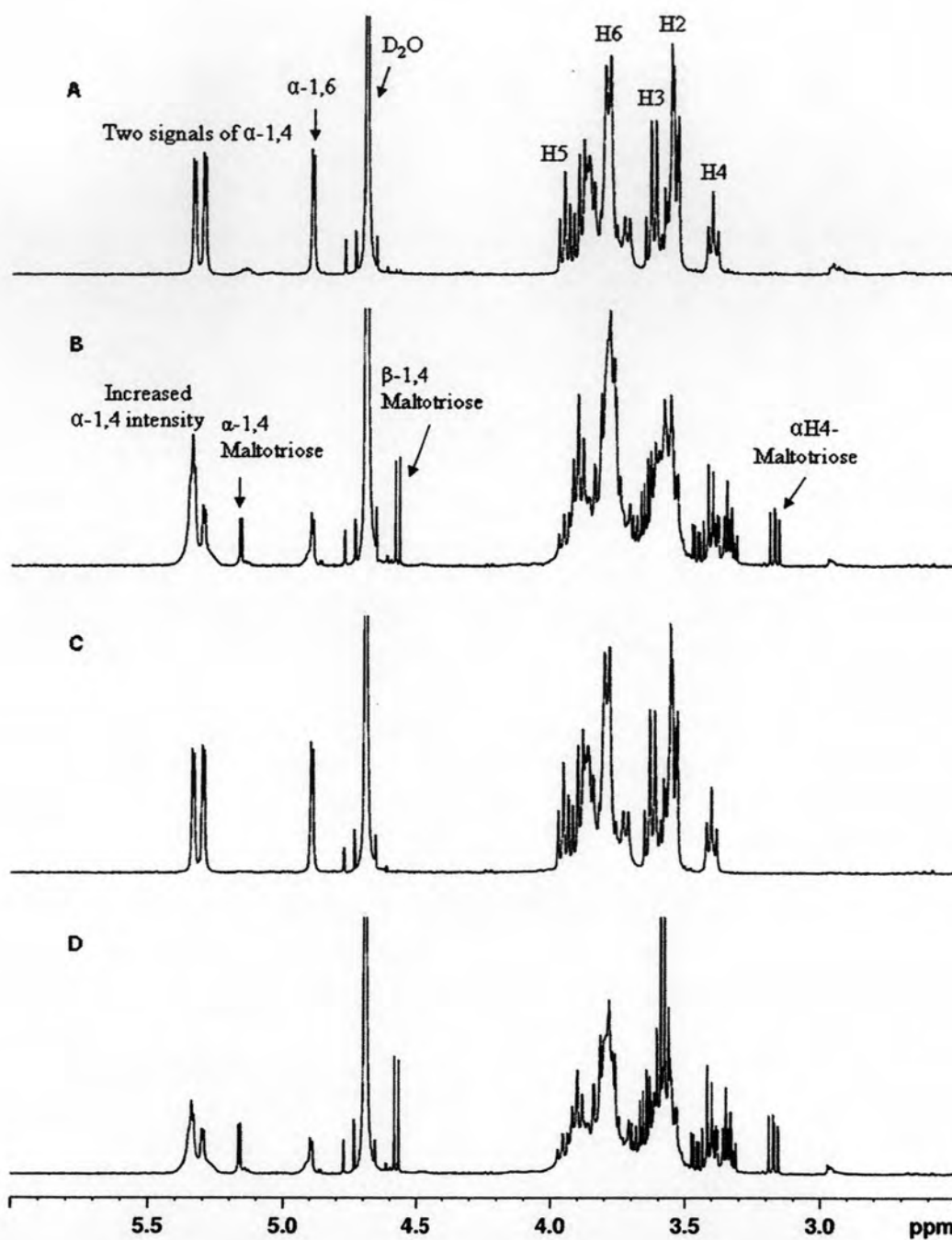






Note that: Since the EPS of BM1 was hardly dissolved, It contained dark pigment and other insoluble polymers, therefore the NMR spectra shows many unknown peaks and no peak of α -1,6 EPS.





$^1\text{H-NMR}$ spectra of exopolysaccharides (EPS) of *Aureobasidium pullulans* NRRL Y-12974 precipitated from pullulan production medium containing (A) 5.0% sucrose, 0.1% N-sources (B) 5.0% starch, 0.1% N-sources (C) 5.0% sucrose, 0.3% N-sources (D) 5.0% starch, 0.3% N-sources.

Appendix G

Morphological studies of each isolates

CU 1



CU 2



CU 3



CU 4



CU 5



CU 6



MEA
(day 7)

YMA
(Day 7)

YMA
(after 2 months)

PM
(Day 7)

Microscopy
(YMB, day 1-3)
(bar = 10 µm)

CU 7



CU 8



CU 9



CU 10



CU 11



CU 12



MEA (day 7) YMA (Day 7) YMA (after 2 months) PM (Day 7) Microscopy (YMB, day 1-3) (bar = 10 μm)

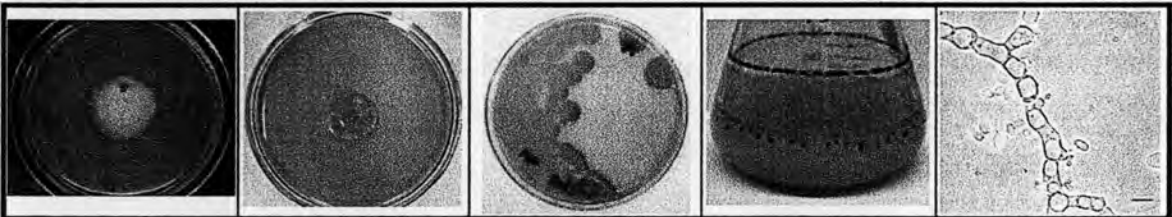
CU 13



CU 14



CU 15



CU 16



CU 17



CU 18



MEA
(day 7)

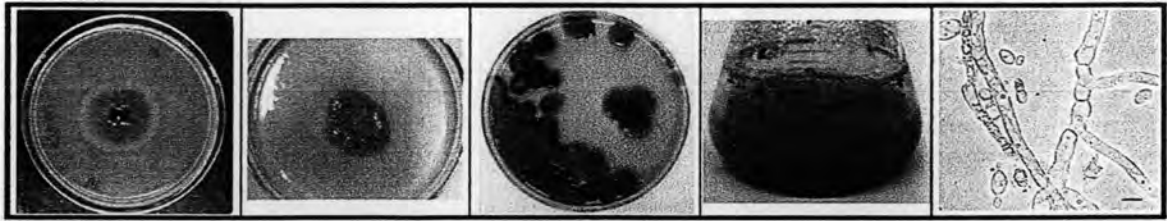
YMA
(Day 7)

YMA
(after 2 months)

PM
(Day 7)

Microscopy
(YMB, day 1-3)
(bar = 10 μ m)

CU 19



CU 20



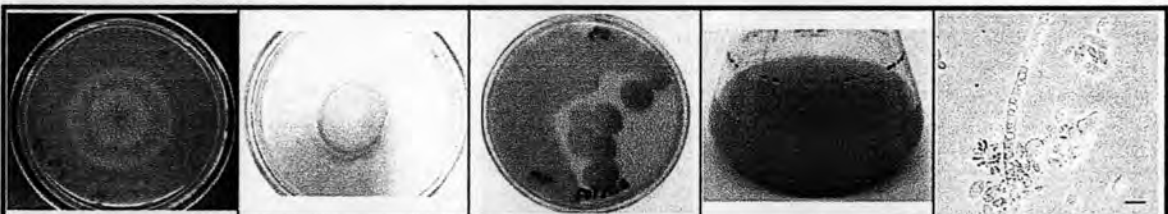
CU 21



CU 22



CU 23



CU 24

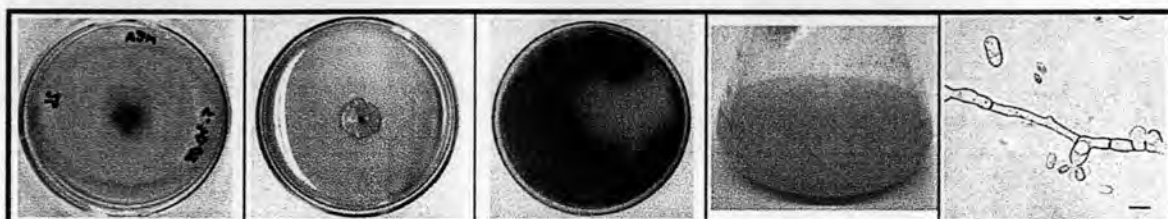


MEA	YMA	YMA	PM	Microscopy
(day 7)	(Day 7)	(after 2 months)	(Day 7)	(YMB, day 1-3)
				(bar = 10 μ m)

CU 25



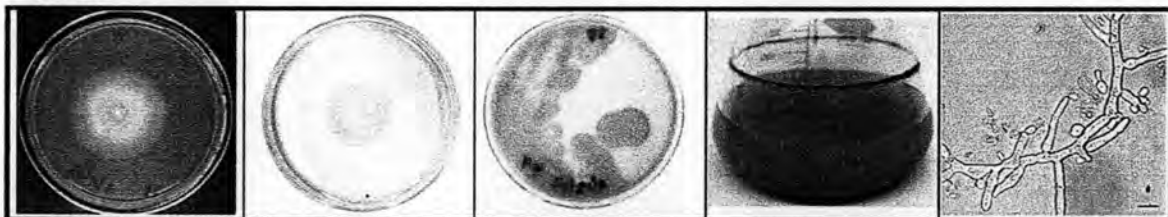
CU 26



CU 27



CU 28



CU 29



CU 30



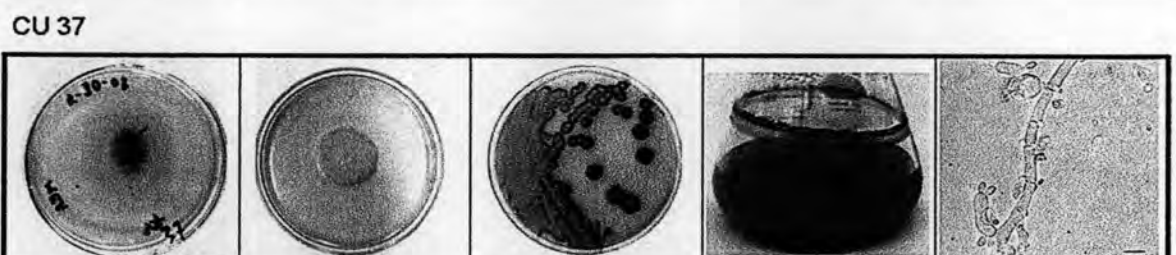
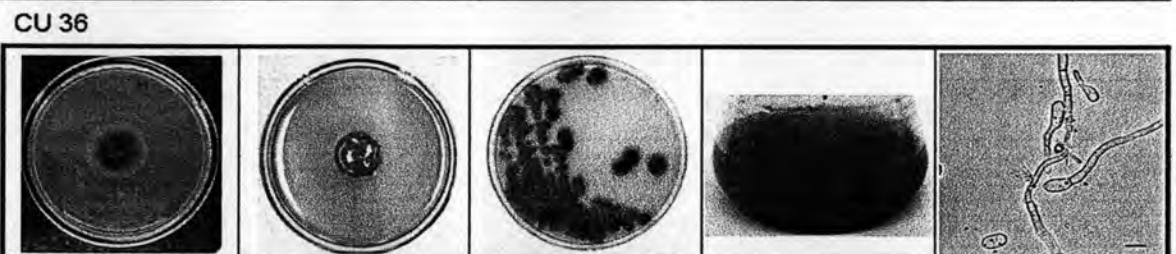
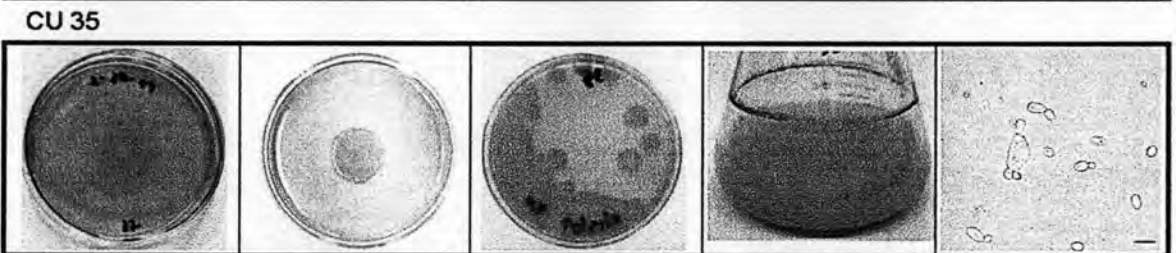
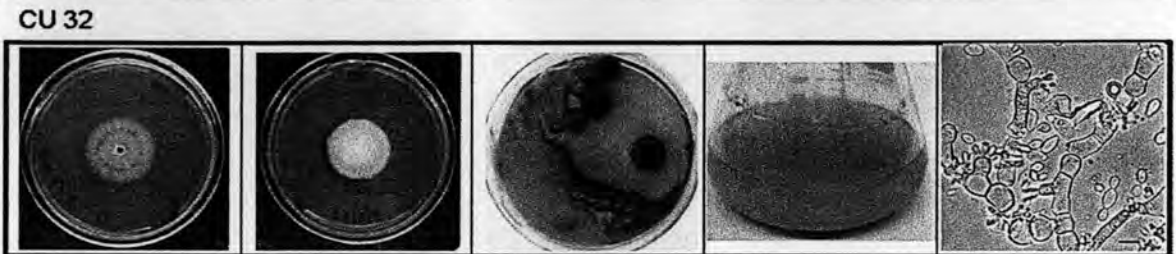
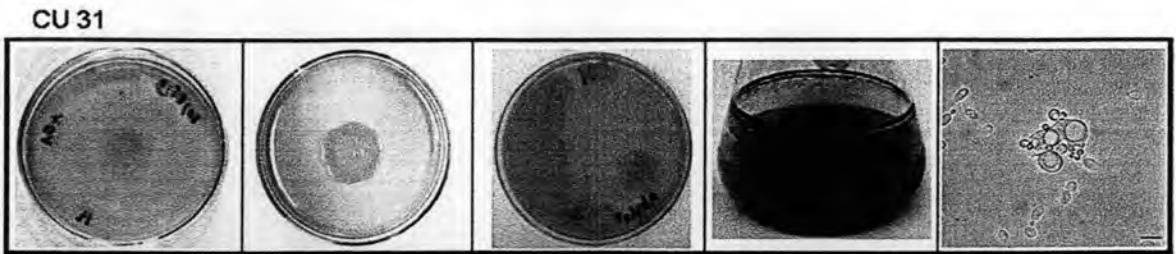
MEA
(day 7)

YMA
(Day 7)

YMA
(after 2 months)

PM
(Day 7)

Microscopy
(YMB, day 1-3)
(bar = 10 μ m)

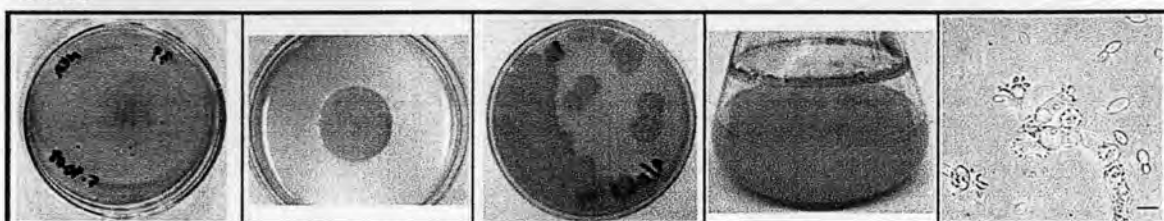


MEA	YMA	YMA	PM	Microscopy
(day 7)	(Day 7)	(after 2 months)	(Day 7)	(YMB, day 1-3)
				(bar = 10 μm)

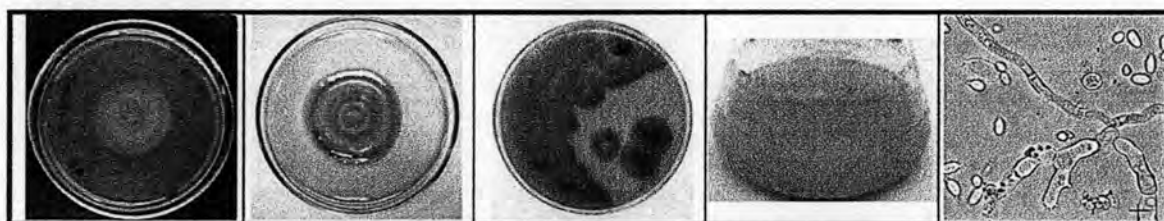
CU 38



CU 39



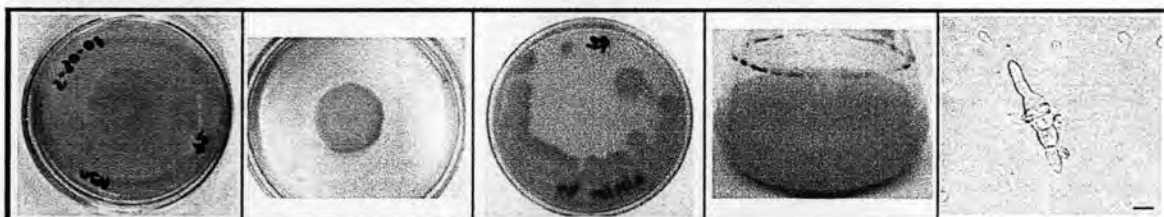
CU 40



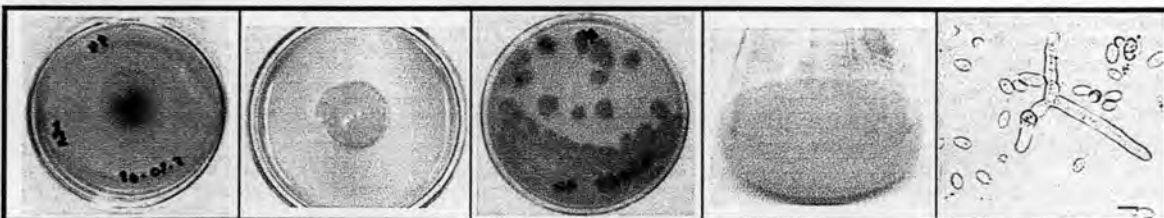
CU 41



CU 42



CU 44



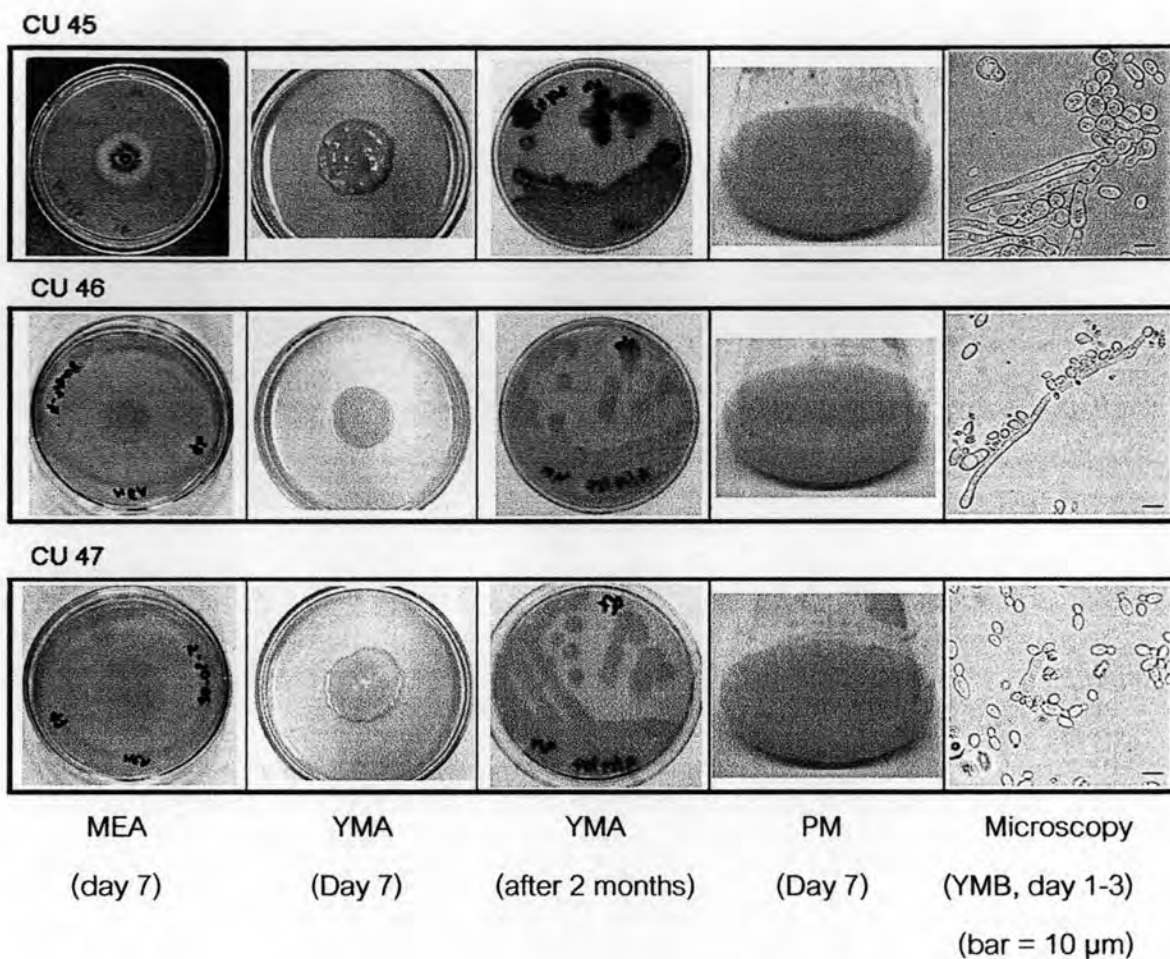
MEA
(day 7)

YMA
(Day 7)

YMA
(after 2 months)

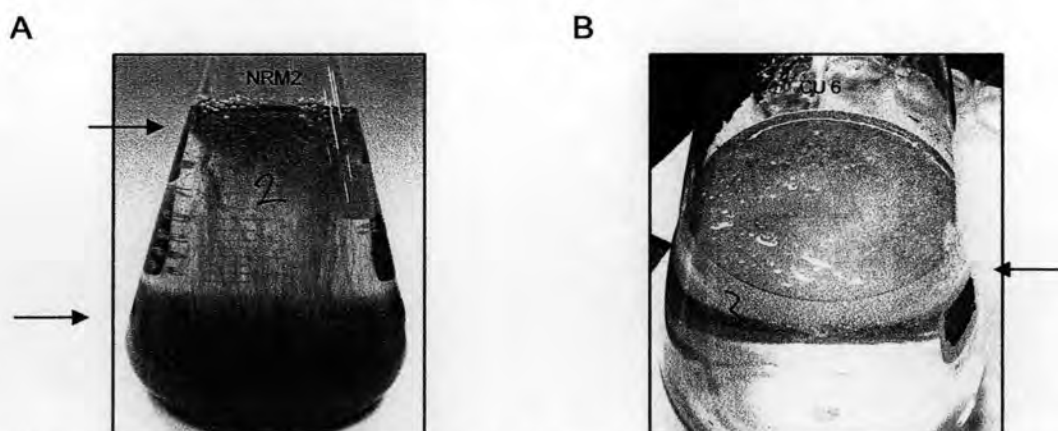
PM
(Day 7)

Microscopy
(YMB, day 1-3)
(bar = 10 μ m)



Appendix H

Two fractions of EPS obtained after precipitation by Ethyl alcohol



(A) Two distinct fractions obtained after ethanol precipitation of NRM2 culture supernatant (B) Single fraction of ethanol precipitate from CU 6. Both were cultured in standard PM and harvested at day 7.

Appendix I

Accession number of isolates in this study, deposited in Fungal Section,
Professor Kasin Suvatabhandhu Herbarium, Department of Botany, Faculty of Science,
Chulalongkorn University, Bangkok, Thailand.

Isolate	Accession number	Isolate	Accession number
CU 1	Pennapa 1 (BCU)	CU 26	Pennapa 26 (BCU)
CU 2	Pennapa 2 (BCU)	CU 27	Pennapa 27 (BCU)
CU 3	Pennapa 3 (BCU)	CU 28	Pennapa 28 (BCU)
CU 4	Pennapa 4 (BCU)	CU 29	Pennapa 29 (BCU)
CU 5	Pennapa 5 (BCU)	CU 30	Pennapa 30 (BCU)
CU 6	Pennapa 6 (BCU)	CU 31	Pennapa 31 (BCU)
CU 7	Pennapa 7 (BCU)	CU 32	Pennapa 32 (BCU)
CU 8	Pennapa 8 (BCU)	CU 33	Pennapa 33 (BCU)
CU 9	Pennapa 9 (BCU)	CU 35	Pennapa 34 (BCU)
CU 10	Pennapa 10 (BCU)	CU 36	Pennapa 35 (BCU)
CU 11	Pennapa 11 (BCU)	CU 37	Pennapa 36 (BCU)
CU 12	Pennapa 12 (BCU)	CU 38	Pennapa 37 (BCU)
CU 13	Pennapa 13 (BCU)	CU 39	Pennapa 38 (BCU)
CU 14	Pennapa 14 (BCU)	CU 40	Pennapa 39 (BCU)
CU 15	Pennapa 15 (BCU)	CU 41	Pennapa 40 (BCU)
CU 16	Pennapa 16 (BCU)	CU 42	Pennapa 41 (BCU)
CU 17	Pennapa 17 (BCU)	CU 44	Pennapa 43 (BCU)
CU 18	Pennapa 18 (BCU)	CU 45	Pennapa 44 (BCU)
CU 19	Pennapa 19 (BCU)	CU 46	Pennapa 45 (BCU)
CU 20	Pennapa 20 (BCU)	CU 47	Pennapa 46 (BCU)
CU 21	Pennapa 21 (BCU)	DOUG	Eveleigh 1 (BCU)
CU 22	Pennapa 22 (BCU)	NRM2	Sehanat 1 (BCU)
CU 23	Pennapa 23 (BCU)	BM1	Patcharawan 1 (BCU)
CU 24	Pennapa 24 (BCU)	HKW1	Patcharawan 2 (BCU)
CU 25	Pennapa 25 (BCU)	PH1	Patcharawan 3 (BCU)

VITA

Miss Pennapa Manitchotpsit was born on May 28th, 1972 in Bangkok, Thailand. She graduated from Satrividhaya School in March 1990 and received the Bachelor of Science degree with a major in Genetics from Chulalongkorn University in April 1995. She continued to study for the Master degree of Science in Molecular Genetics and Genetic Engineering, at Institute of Molecular Biology and Genetics, Salaya, Mahidol University and graduated in October 1999. After that, she worked as a research assistant at Institute of Molecular Biology and Genetics, Salaya, Mahidol University until October 2000 and then she had become the faculty member of Department of Medical Science, Faculty of Science, Rangsit University. Since November 2004, she has studied the degree of Philosophy of Science in Biological Science s Program, at the Faculty of Science, Chulalongkorn University.

