



The 2nd International
Resource Recovery
Conference



VIA-NITRITE SELECTION OF PHA STORING BIOMASS THROUGH FERMENTATION LIQUID OF CELLULOSIC PRIMARY SLUDGE

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Nutrients

Water

Energy

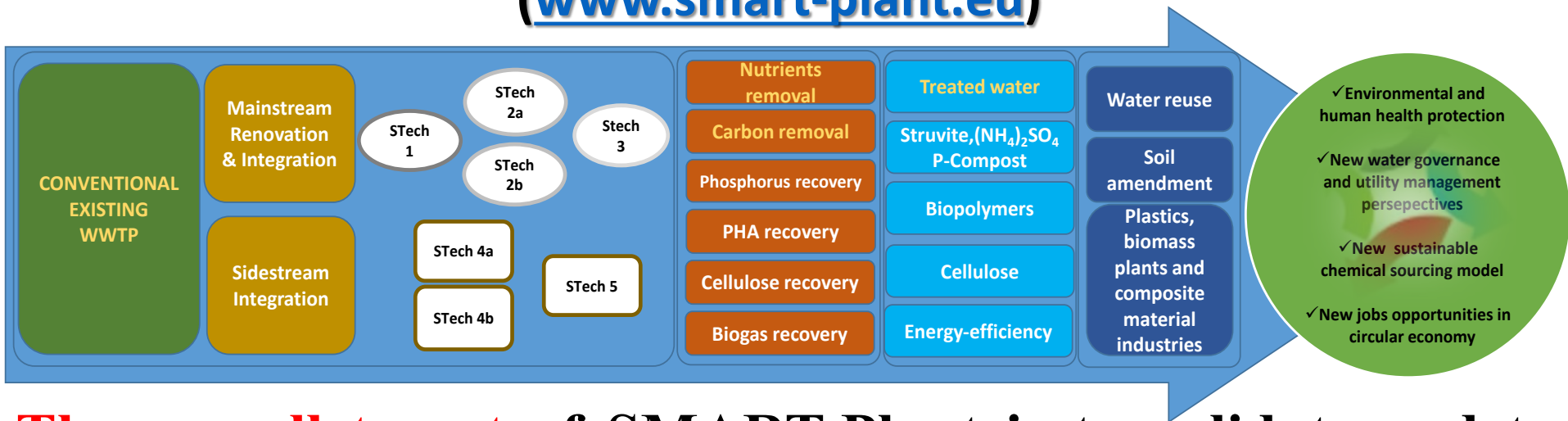
Outline

- The Horizon2020 «SMART-Plant» Innovation Action
- Cellulose in wastewater and recovery/valorization routes
- Controlled fermentation to recover propionate-rich volatile fatty acids
- Bio(co)polymers (PHB-coPHV) production in the sidestream SCEPPHAR system
- What's next: scale-up to demo in real environment at Carbonera WRRF

The Horizon2020 “SMART-Plant” Innovation Action

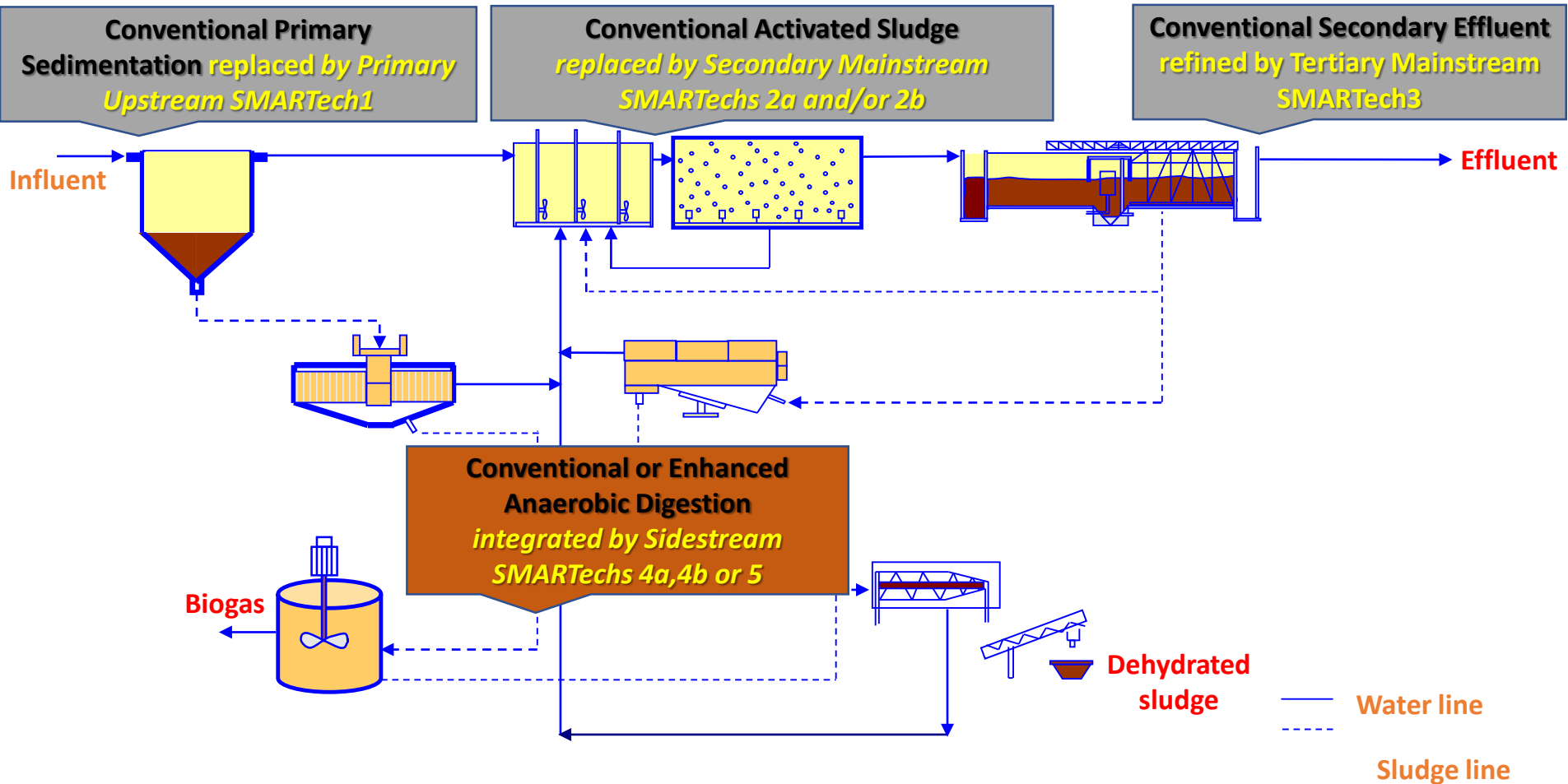
Scale-up of low-carbon footprint **MA**terial **R**ecovery **T**echniques for upgrading existing wastewater treatment **Plants**

(www.smart-plant.eu)



The overall target of SMART-Plant is to validate and to address to the market a portfolio of SMARTechnologies that, singularly or combined, can **renovate and upgrade existing wastewater treatment plants** and give the added value of instigating the **paradigm change towards efficient wastewater-based bio-refineries.**

The **SMART**Technologies to integrate and renovate existing WWTPs

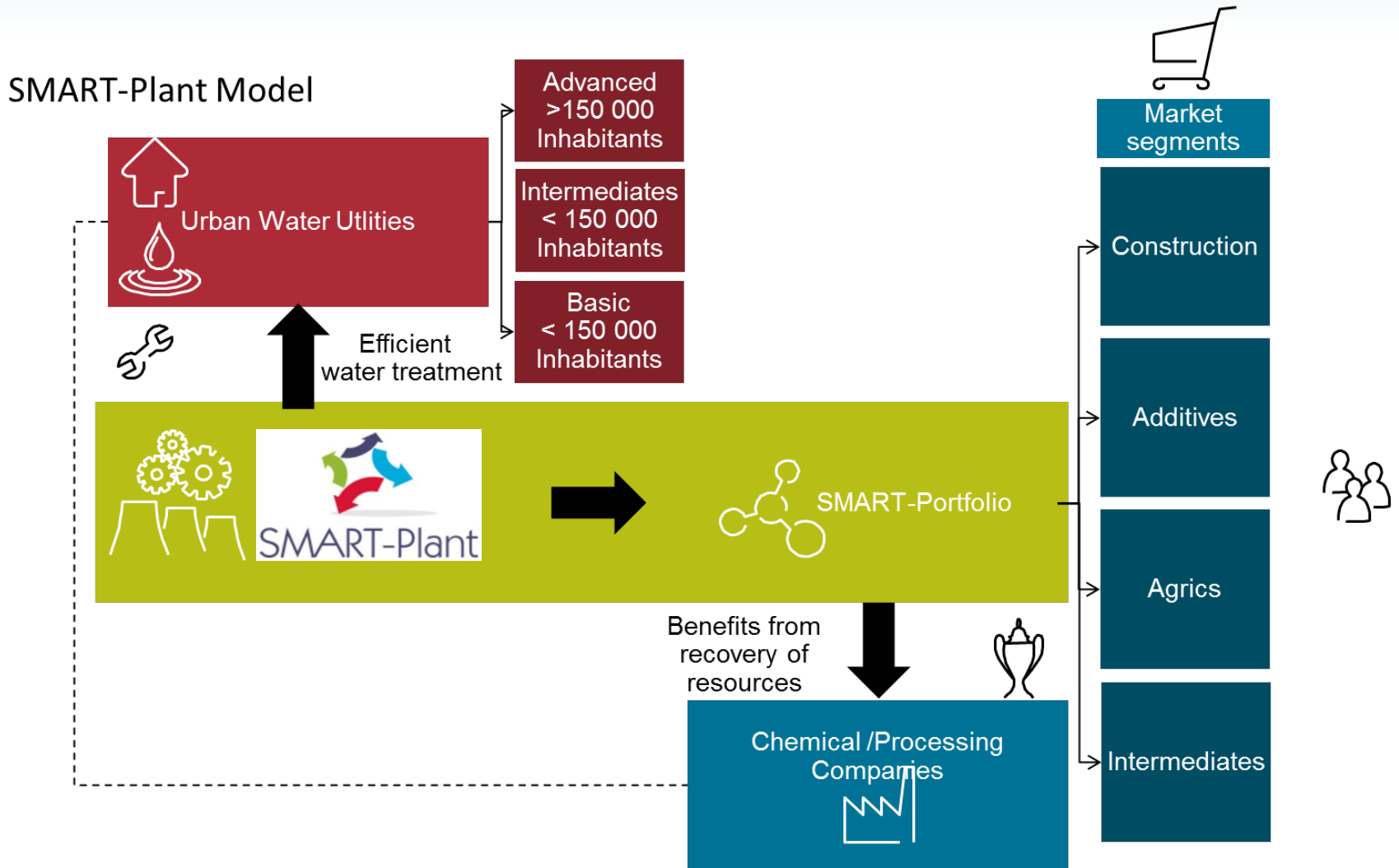


Key Enabling Strategy: upstream solid concentration, integration and innovation of the sewage sludge treatment, energy efficient mainstream treatment for water reuse

The WWTPs integrated by SMART-Plant to WRRFs

SMARTech n.	Integrated municipal WRRF	Key enabling process(es)	SMART-product(s)
1	Geestmerabacht (Netherlands)	Upstream dynamic fine-screen and post-processing of cellulosic sludge	Cellulosic sludge, refined clean cellulose
2a	Karmiel (Israel)	Mainstream polyurethane-based anaerobic biofilter	Biogas, Energy-efficient water reuse
2b	Manresa (Spain)	Mainstream SCEPPHAR	P-rich sludge, PHA
3	Cranfield (UK)	Mainstream tertiary hybrid ion exchange	Nutrients
4a	Carbonera (Italy)	Sidestream SCENA+conventional AD	P-rich sludge, VFA
4b	Psytalia (Greece)	Sidestream SCENA+enhanced AD	P-rich sludge
5	Carbonera (Italy)	Sidestream SCEPPHAR	PHA, struvite, VFA

SMART-Plant value chains, business plan and market deployment strategy

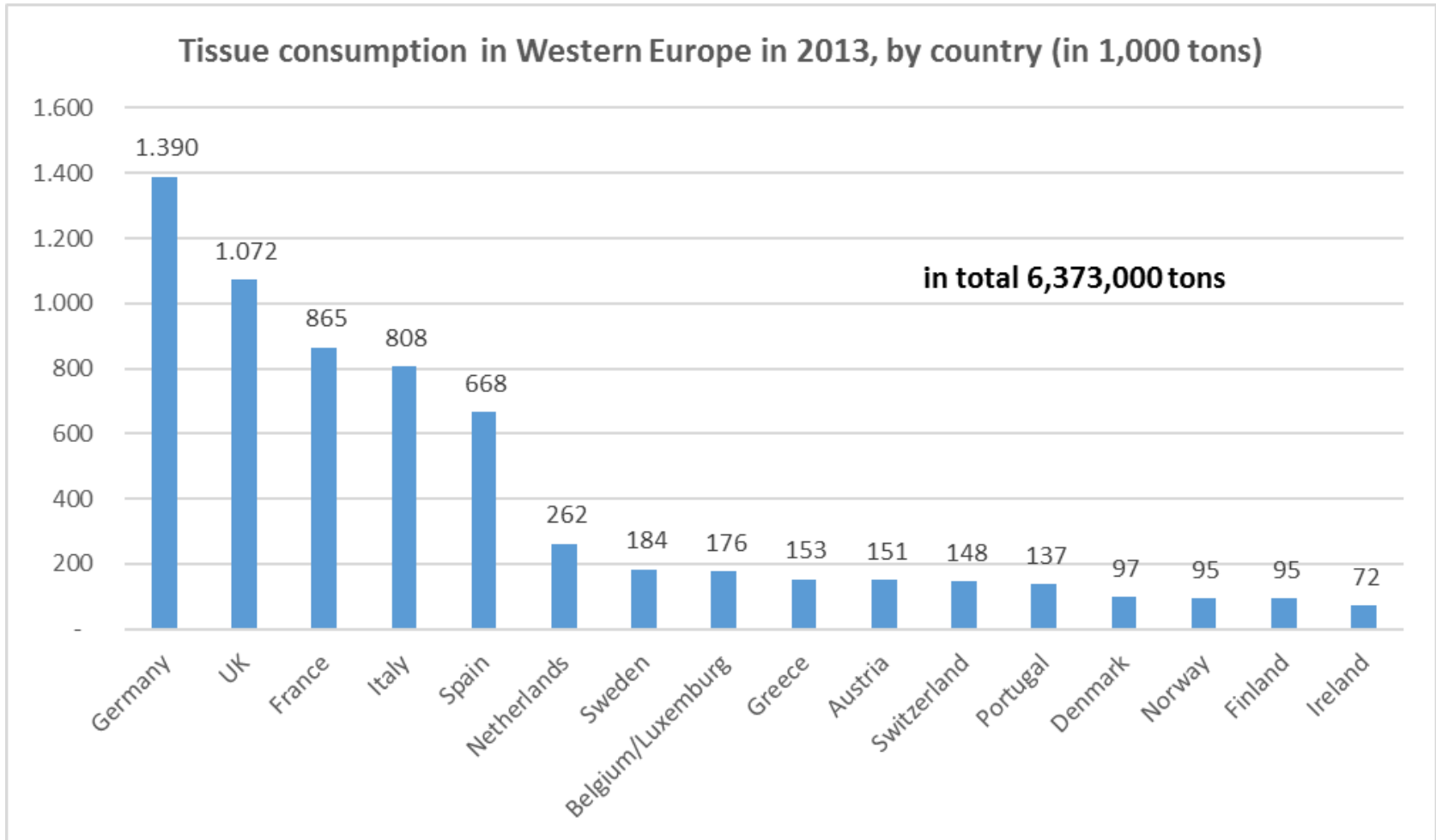


Resource recovery optimization? What about toilet paper?



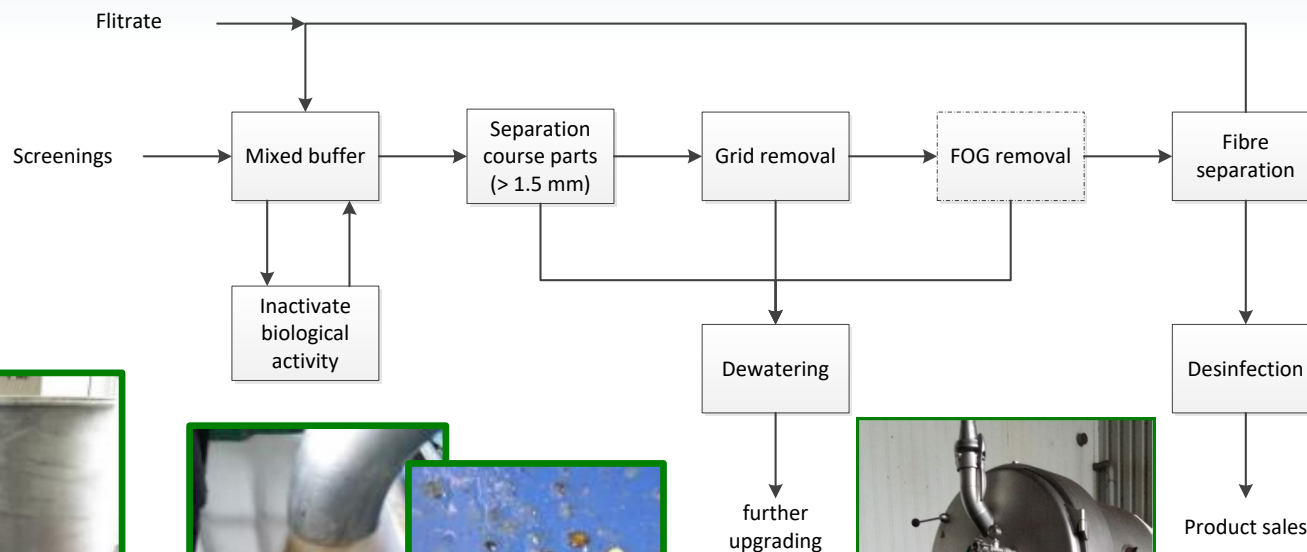
- ✓ 12 to 18 kg per person per year
- ✓ 8.5 pieces of paper per visit to the restroom
- ✓ On average, a person spends 43 hours a year on the toilet
- ✓ 70% folds the sheets before using them, 29% make a proper use

Facts and figures



Source: www.statista.com

SMARTech1: Primary (upstream) dynamic sieving and clean cellulose recovery



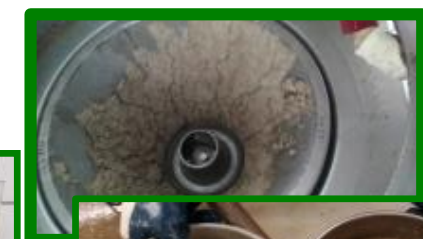
Inactivation biological activity



Separation course parts



Sand-/grid removal



Fibre separation

SMARTech1: Primary (upstream) dynamic sieving and clean cellulose recovery

- 79% cellulose fiber,
- 5 % other organics,
- 6% inorganic (ash),
- 10% other contaminants (average in The Netherlands).

Potentially marketable product, but the economic feasibility depends mainly on savings at the WWTP

Market development

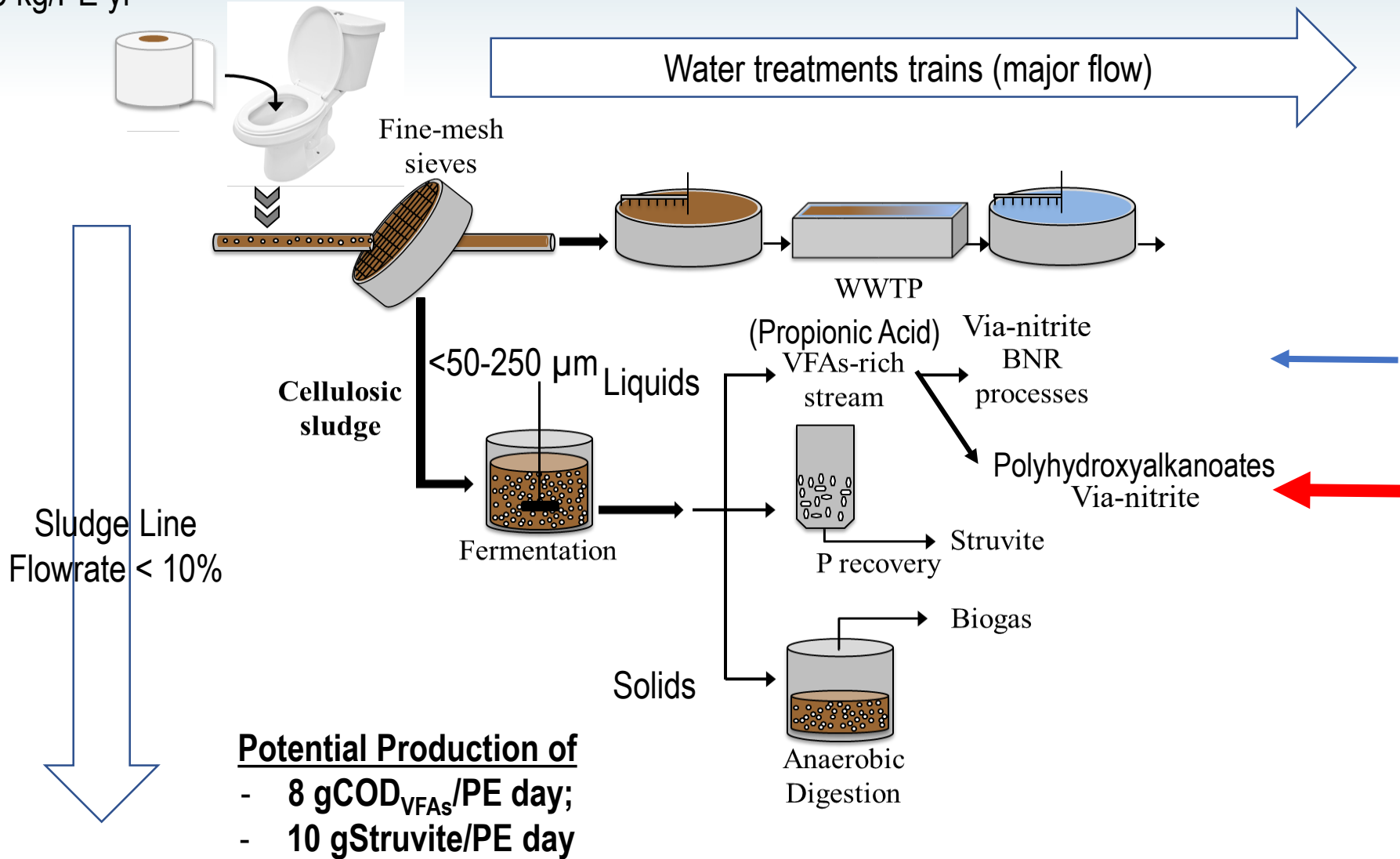
Marketing and valorization of recovered cellulose

- ✓ Reuse in asphalt
- ✓ Raw material for composite
- ✓ Insulation materials (In development, not sure yet)



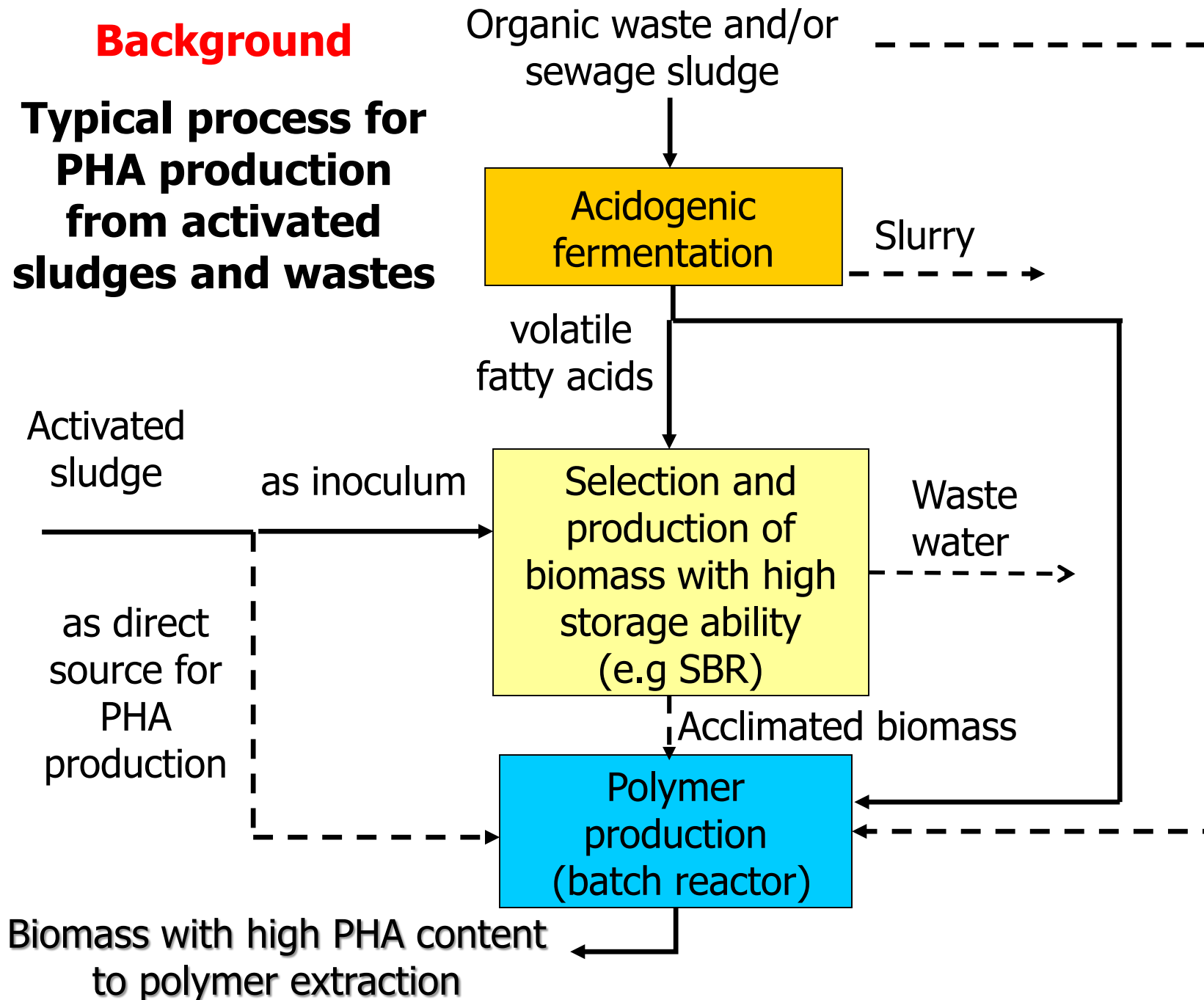
Novel SMART-Plant demos for valorization of cellulosic sludge

5-15 kg/PE yr



Background

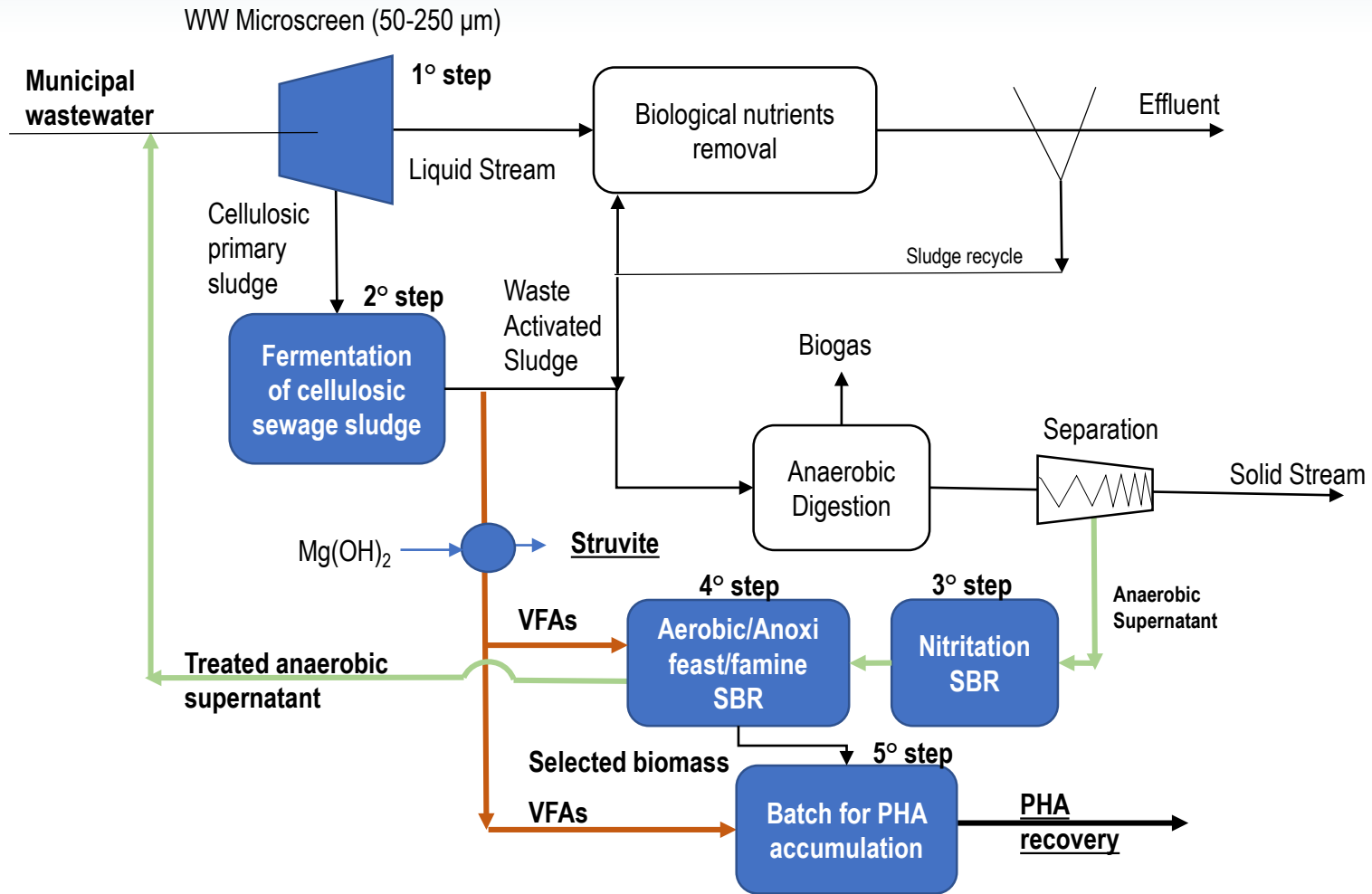
Typical process for PHA production from activated sludges and wastes



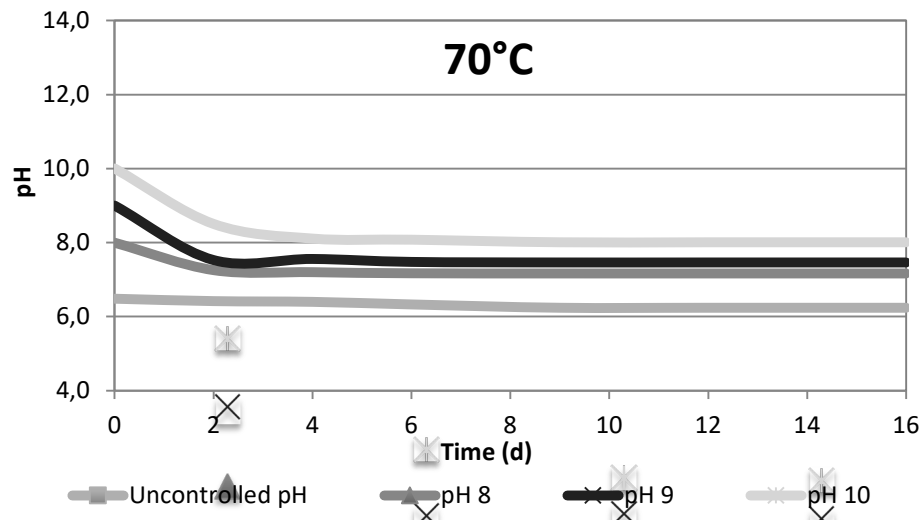
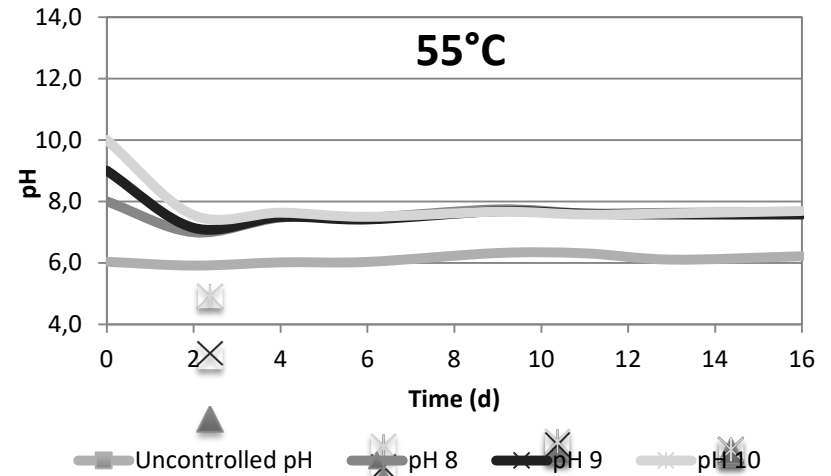
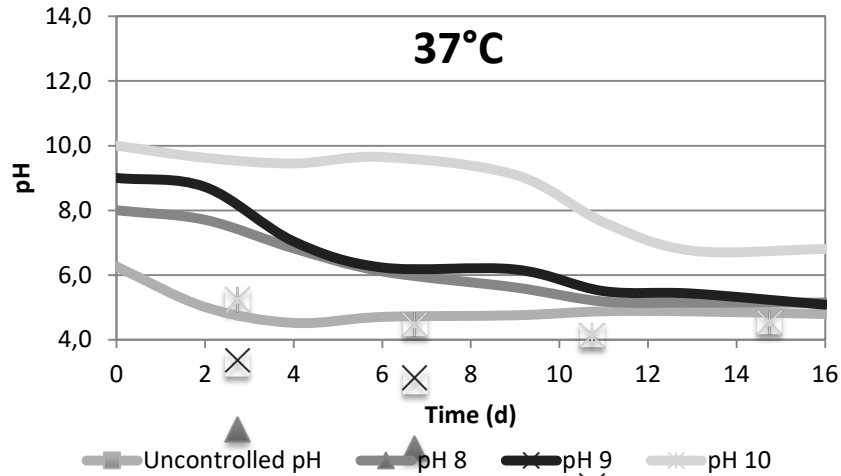
The SMART-Plant innovation:

- To integrate the energy-efficient via-nitrite nitrogen removal with the PHA recovery → major interest of the water utility
- To adopt anoxic (via-nitrite) famine conditions to optimize energy consumptions
- Phosphorus (struvite) recovery even to support the balance of nitrogen and phosphorus to the PHA recovery system
- **To optimize the system using primary cellulosic sludge → IRRC2017**

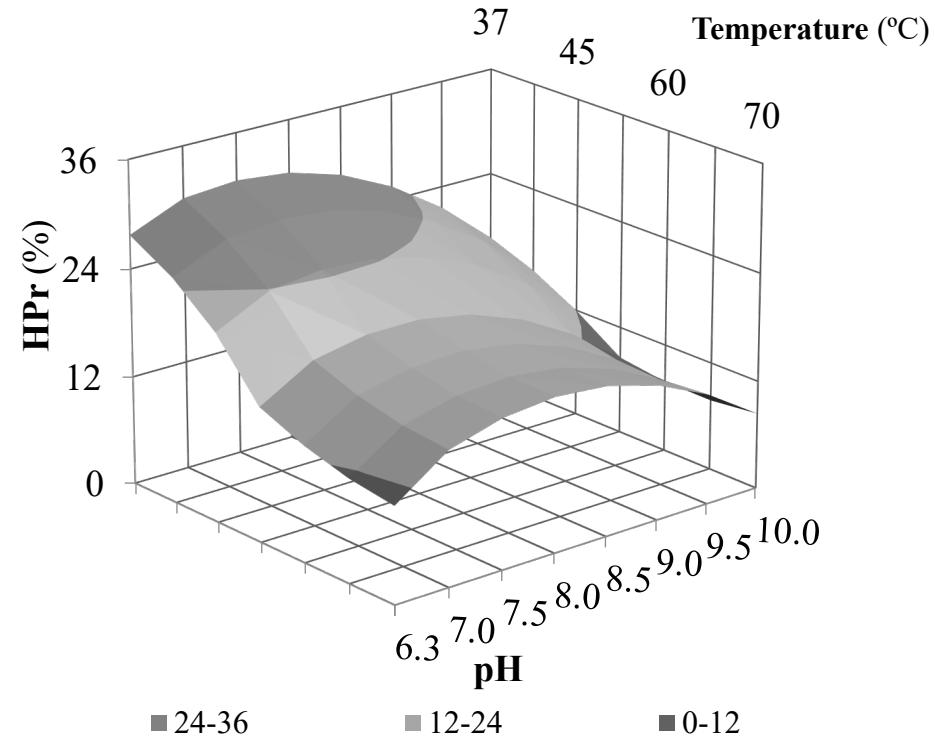
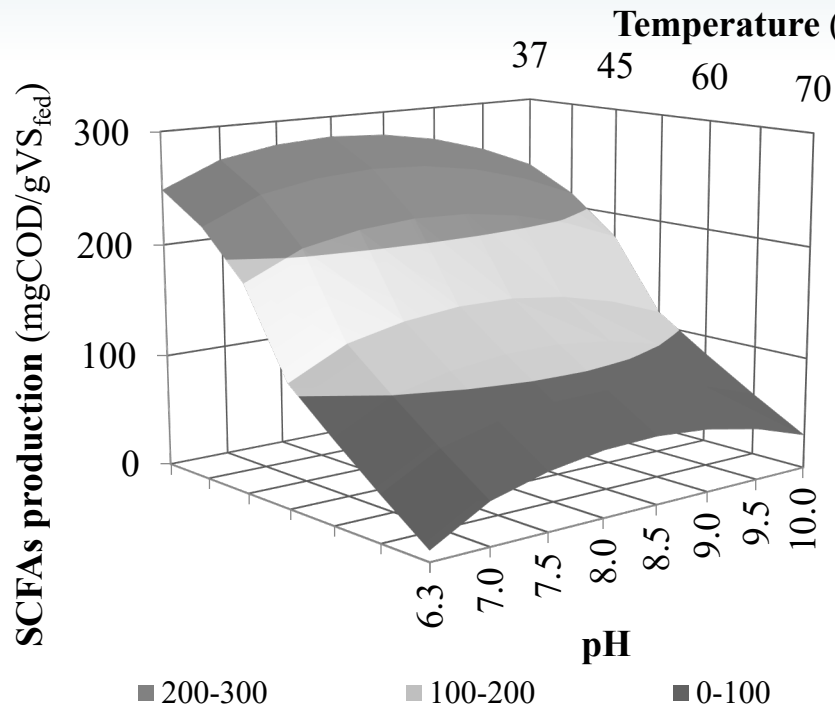
Sidestream S.C.E.P.P.H.A.R.: Short-Cut Enhanced Phosphorus and PHA recovery



Outline of the batch Cellulosic Fermentation experiments

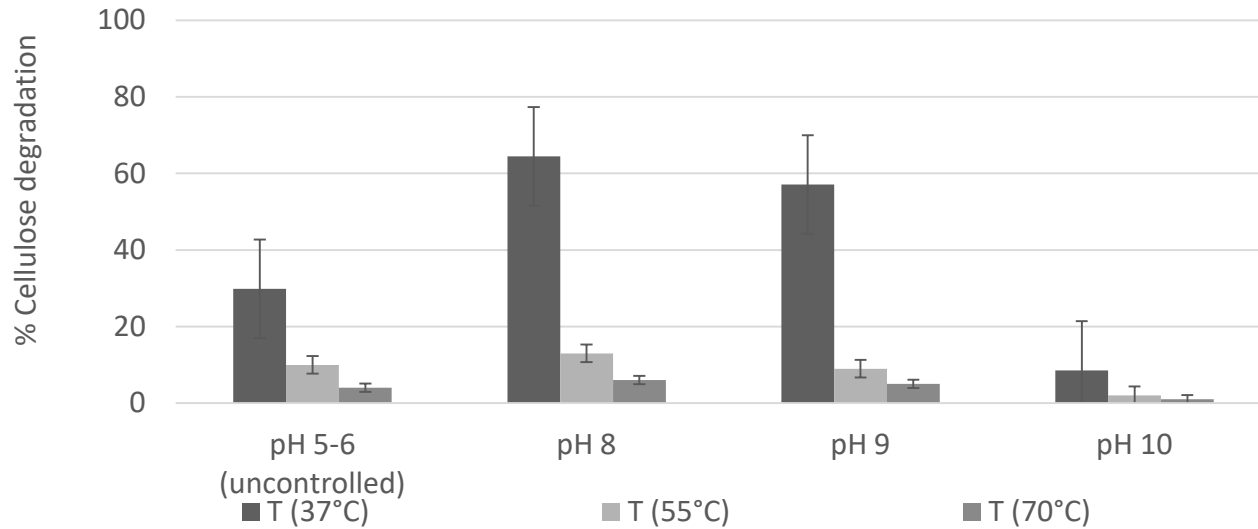


Optimal parameters for the recovery of Propionic Acid



- Initial pH 7.5-8
- Temperature 37°C

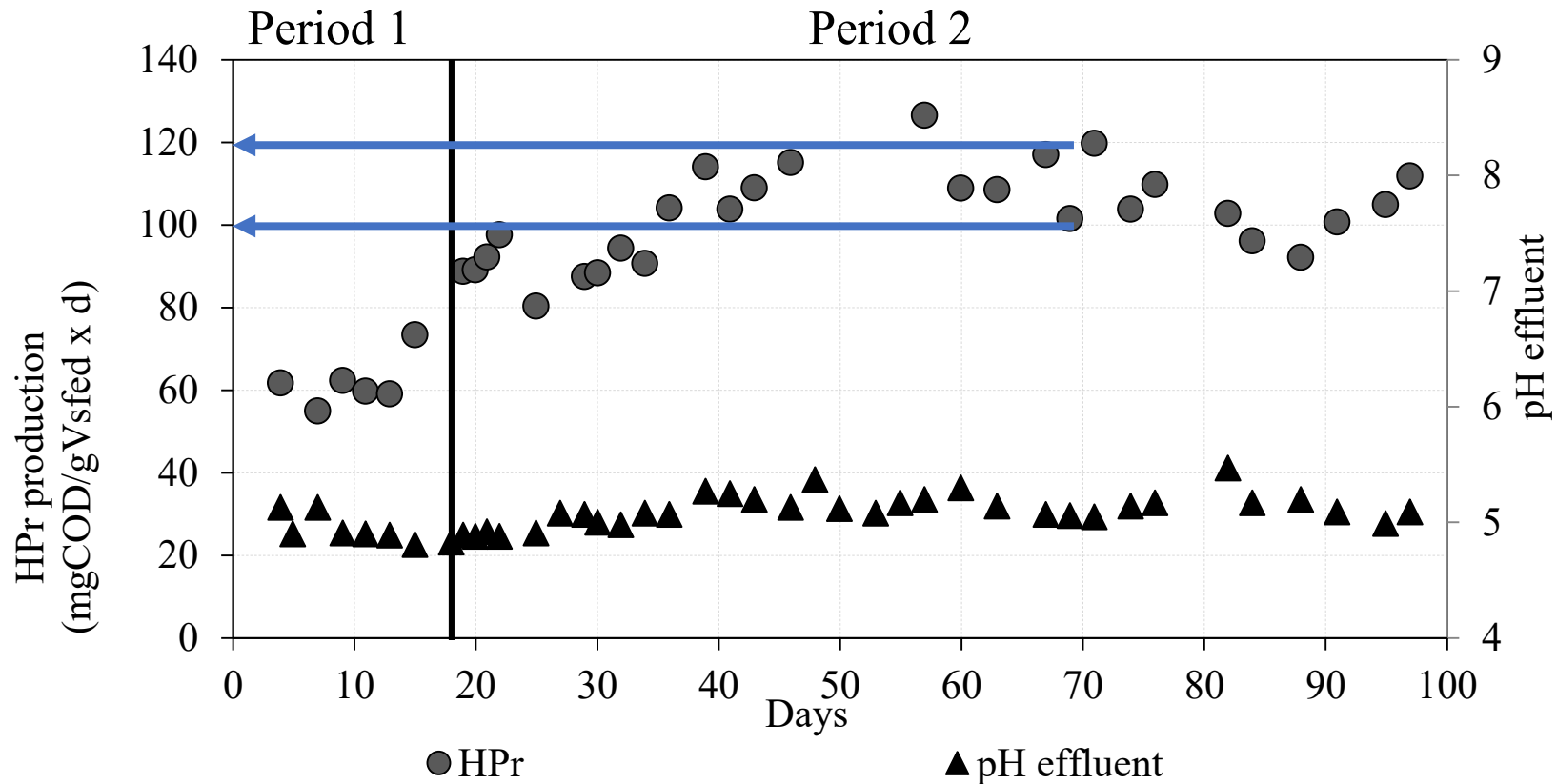
Cellulose biodegradation



Residual sieved material
(250 μ m) after fermentation

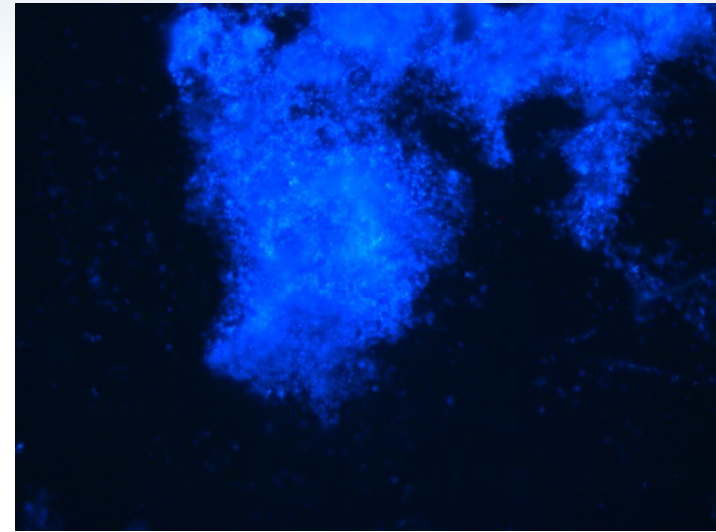
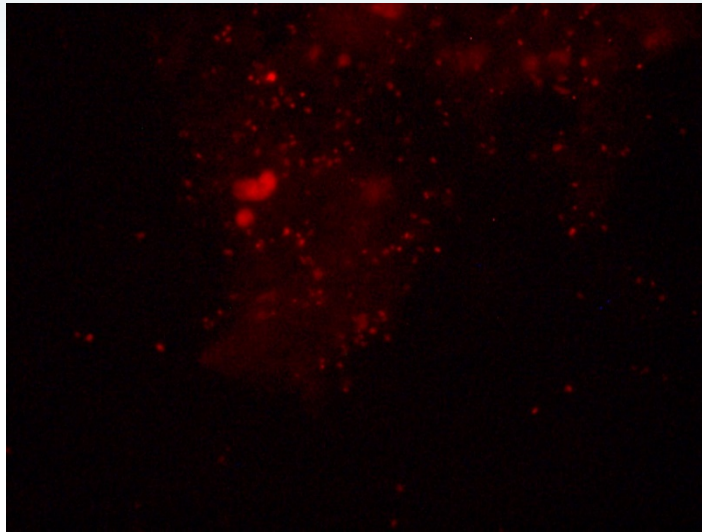


Sequencing Batch Fermentation Reactor



- Period 1: uncontrolled pH; **Period 2: influent pH 8**
- Stable production of propionic acid during period 2
- Propionic acid: 30-35 % SCFA

FISH Analyses on *Propionibacterium acidopropionici*

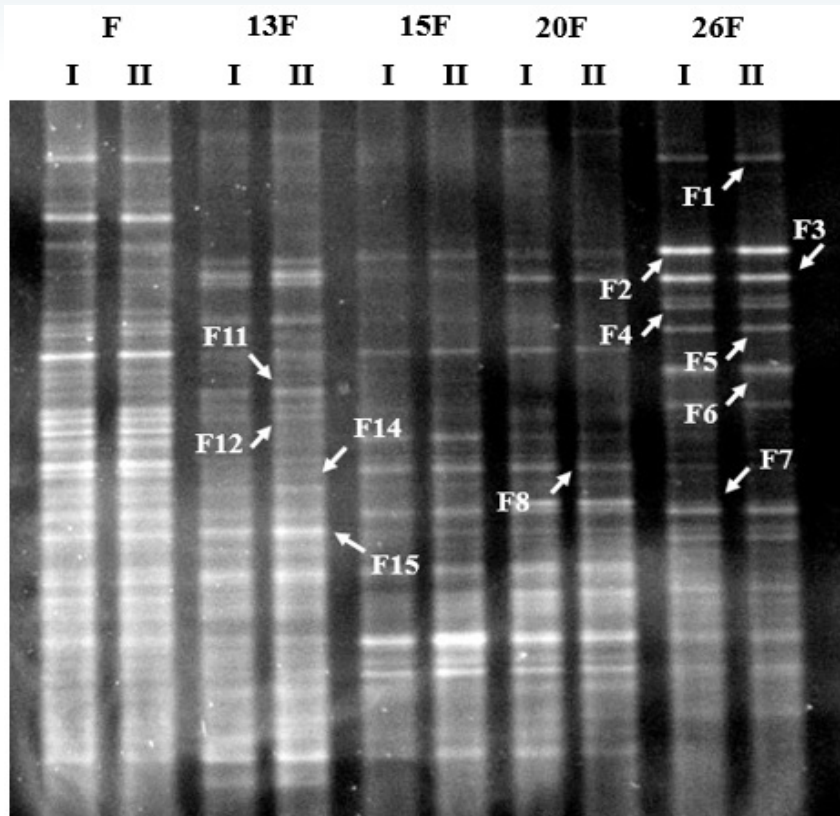


Temperature (°C)	37	55
	Aver. ± St.Dev	Aver. ± St.Dev.
Bacteria detected with probe Apr820 (%)	34±15.7	24±28
Propionate Production (mgCOD/gVSS)	100-120	20-40

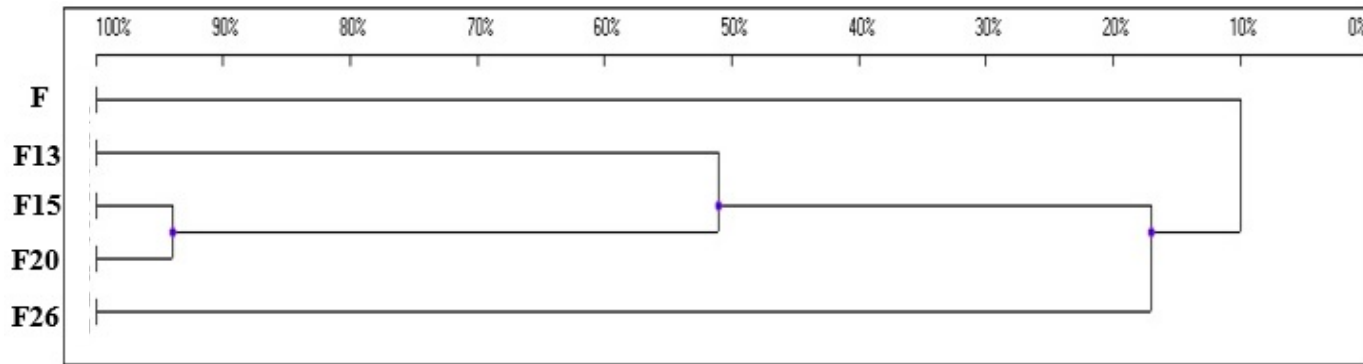
- *Propionibacterium acidopropionici* are more present at a temperature of 37 rather than 55° C.
- % of bacteria is correlated with the amount of propionate and they both decrease as the temperature increases

Microbial Community Analyses of the Fermentation Unit

DGGE Analyses



The PCR-DGGE results showed heterogeneous bacterial population in the fermentation unit



Phylogenetic analyses

Band	Closest bacterial strain	Accession number	Percentage of identity (%)	Class	Phylum
F1	Uncultured <i>Rhodopila</i> sp. CJ	FJ495222	99	Alphaproteobacteria	Proteobacteria
	Uncultured <i>Parvibaculum</i> sp. CH	FJ495220	99	Alphaproteobacteria	Proteobacteria
F2	Uncultured bacterium	AF390917	100	Betaproteobacteria	Proteobacteria
	Uncultured beta proteobacterium	HE856465	99	Betaproteobacteria	Proteobacteria
	<i>Comamonas</i> sp. RPWA5.3	KC584758	98	Betaproteobacteria	Proteobacteria
	<i>Curvibacter delicatus</i>	NR_028713	98	Betaproteobacteria	Proteobacteria
	<i>Acidovorax</i> sp. DQS-01	KP126996	98	Betaproteobacteria	Proteobacteria
F3	Uncultured alpha proteobacterium	JQ919505	99	Alphaproteobacteria	Proteobacteria
	Uncultured <i>Rhodopila</i> sp. CJ	FJ495222	99	Alphaproteobacteria	Proteobacteria
	Uncultured <i>Parvibaculum</i> sp. CH	FI495220	99	Alphaproteobacteria	Proteobacteria
F4	Uncultured <i>Proteiniphilum</i> sp. partial 16S rRNA gene, isolate OTU 224	LT625110	98	Cytophagia	Bacteroidetes
	Uncultured <i>Bacteroidetes</i> bacterium	FJ754858	98		Bacteroidetes
	<i>Bacteroidales</i> bacterium	GU129048	98		Bacteroidetes
F5	Uncultured alpha proteobacterium	JQ919505	99	Alphaproteobacteria	Proteobacteria
	Uncultured <i>Rhodopila</i> sp. CJ	FJ495222	98	Alphaproteobacteria	Proteobacteria
	Uncultured <i>Parvibaculum</i> sp. CH	FI495220	98	Alphaproteobacteria	Proteobacteria
F6	Uncultured <i>Bacteroidetes</i> bacterium	CU917950	100		Bacteroidetes
	<i>Bacteroidales</i> bacterium	GU129081	99		Bacteroidetes
	<i>Proteiniphilum</i> sp. I28	KP262056	99	Cytophagia	Bacteroidetes
	<i>Porphyromonadaceae</i> bacterium enrichment culture	HQ133064	99	Bacteroidales	Bacteroidetes
F7	Uncultured alpha proteobacterium	JQ919505	99	Alphaproteobacteria	Proteobacteria
	Uncultured <i>Rhodopila</i> sp. CJ	FJ495222	99	Alphaproteobacteria	Proteobacteria
	Uncultured <i>Parvibaculum</i> sp. CH	FI495220	99	Alphaproteobacteria	Proteobacteria
F8	Uncultured bacterium clone	KP279931	100		
	<i>Flavobacterium</i> sp. enrichment culture clone SA NR2 1	GU726988	99	Flavobacteria	Bacteroidetes
	Uncultured <i>Leadbetterella</i> sp. SSP-AOLR-2-1	KR705991	99	Cytophagia	Bacteroidetes
	<i>Cytophaga</i> sp. INT-18	AB046729	98	Cytophagia	Bacteroidetes
F11	Uncultured bacterium gene	AB746716	96		
F12	Uncultured alpha proteobacterium	JQ919505	99	Alphaproteobacteria	Proteobacteria
	Uncultured <i>Rhodopila</i> sp. CJ	FJ495222	98	Alphaproteobacteria	Proteobacteria
	Uncultured <i>Parvibaculum</i> sp. CH	FJ495220	98	Alphaproteobacteria	Proteobacteria
F14	Uncultured <i>Saprospiraceae</i> bacterium	KP717501	100	Sphingobacteria	Bacteroidetes
	Uncultured <i>Halicomenobacter</i> sp. C-115	JX415430	99	Sphingobacteria	Bacteroidetes
F15	Uncultured <i>Clostridium</i> sp. D7	KM494503	96	Clostridia	Firmicutes

Focus on cellulolytic activity

Firmicutes

Firmicutes, (e.g. like Clostridium and Bacillus) are able to produce cellulases such as carboxymethyl cellulase (CMCase). It plays an important role to hydrolyse cellobiose to fermentable glucose

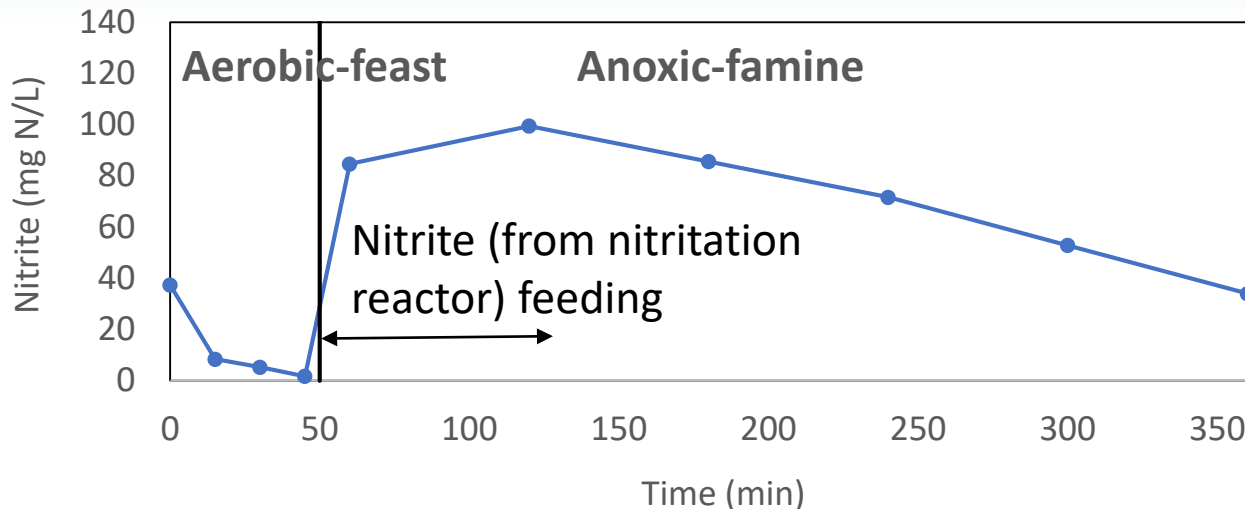
Bacteroidetes

Bacteroidetes are able to decompose long chained polymers such as cellulose and so contribute to the hydrolysis step in fermentation.

Operating conditions for the PHA-storing community selection

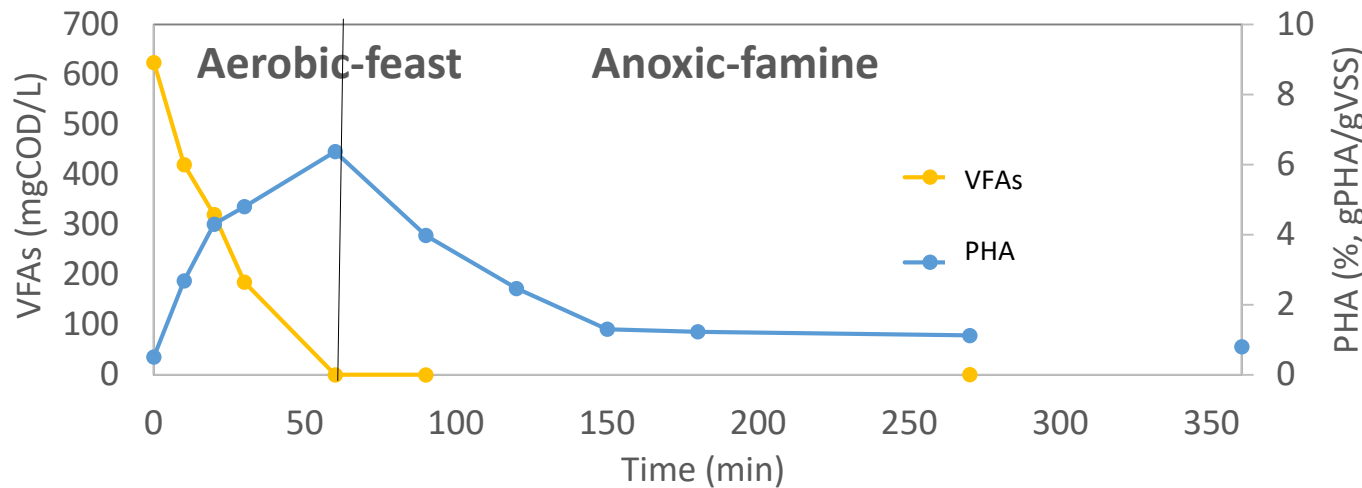
Parameter	Period I (0-39 days)	Period II.1 (40-106 days)	Period II.2 (107-145 days)
vNLR (kgN/m ³ d)	0,610	0,480	0,56
vOLR(kgCOD/m ³ d)	1,3	1,32	1,2
HRT (d)	1,5	1,1	1,2
Aerobic/Anoxic	1	0,44	0,10-0,20
SRT (d)	5	7-10	7-10
Carbon source	VFAs from synthetic acetic and propionic acid	VFAs from synthetic acetic and propionic acid	VFAs from real cellulosic primary sludge (C.P.S)
Nitrified anaerobic supernatant	NO ₂ and NH ₄ ⁺ from synthetic anaerobic supernatant	NO ₂ and NH ₄ ⁺ from synthetic anaerobic supernatant	NO₂ and NH₄⁺ from real anaerobic supernatant

Selection of PHA-storing microorganisms: S.C.E.P.P.H.A.R. cycle



Nitrite denitritation
driven by PHA

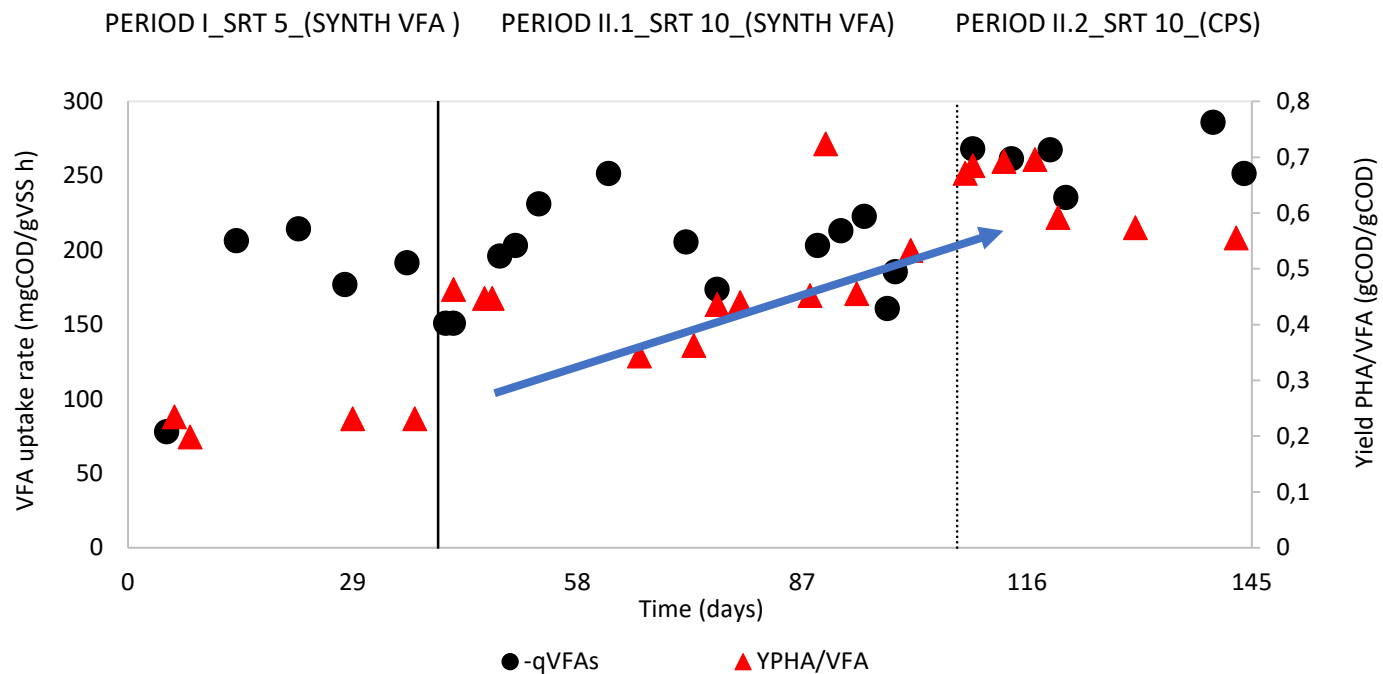
$K_{den}(20^{\circ}\text{C}) =$
6-8 $\text{mgNO}_2\text{-N/gVSS h}$



Feast phase 16%
of the total cycle

Performance of the selection of PHA-storing microorganisms

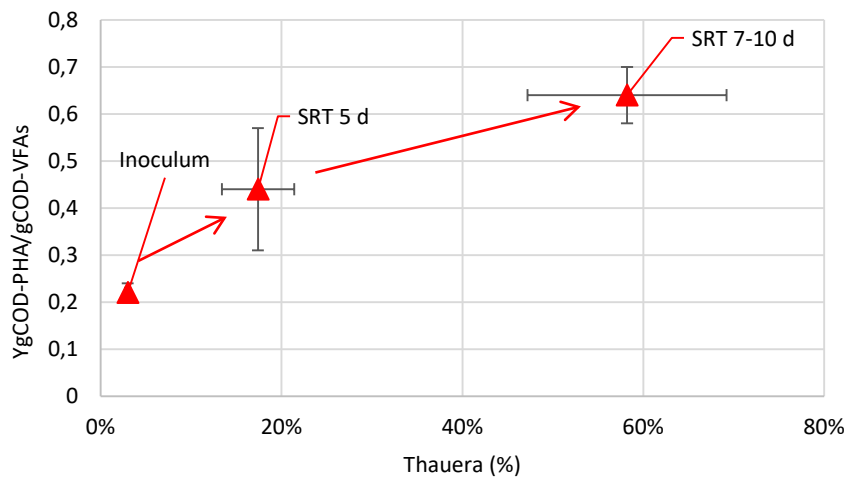
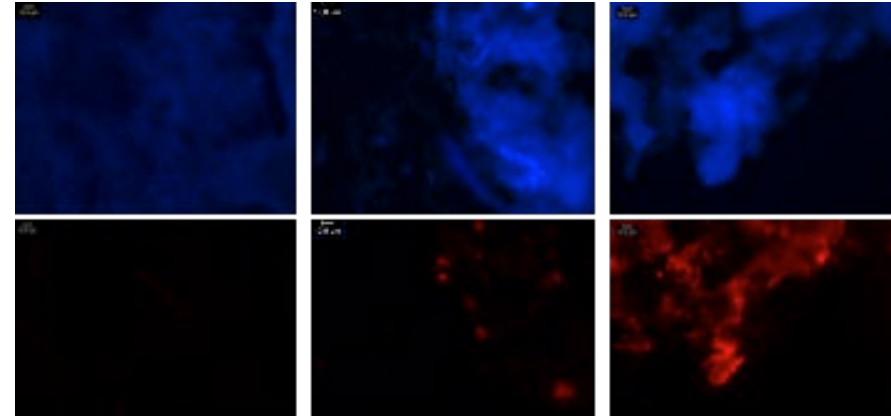
From the day 40, the $Y_{\text{PHA/VFA}}$ gradually increased (from 0.22 to 0.51 $\text{gCOD}_{\text{PHA}}/\text{gCOD}_{\text{VFA}}$), reaching the better results when the CPS was used as C-source in the period II.2 (up to $0.65 \text{ gCOD}_{\text{PHA}}/\text{gCOD}_{\text{VFA}}$).



Analyses of the microbial community

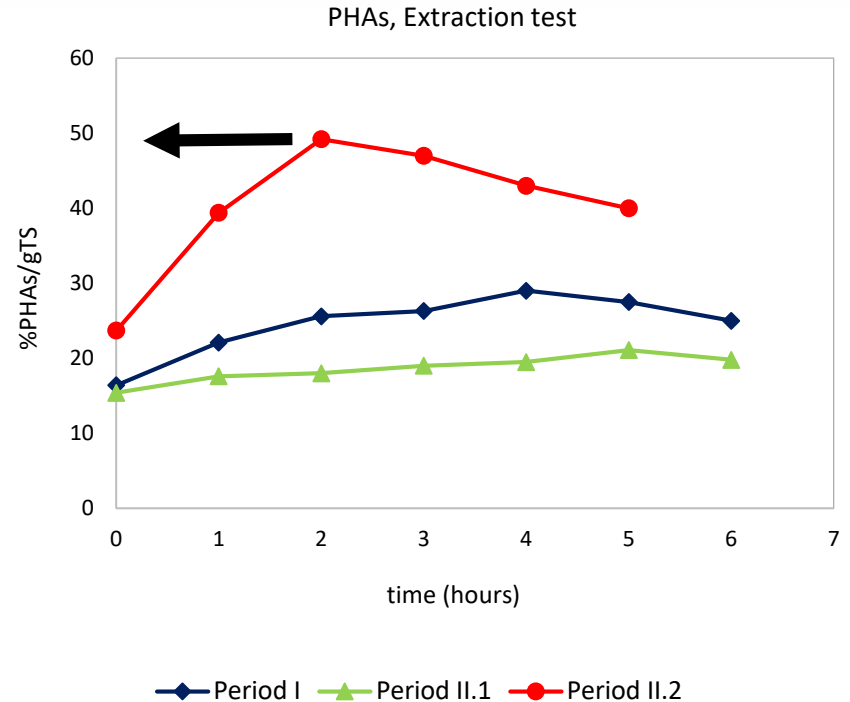
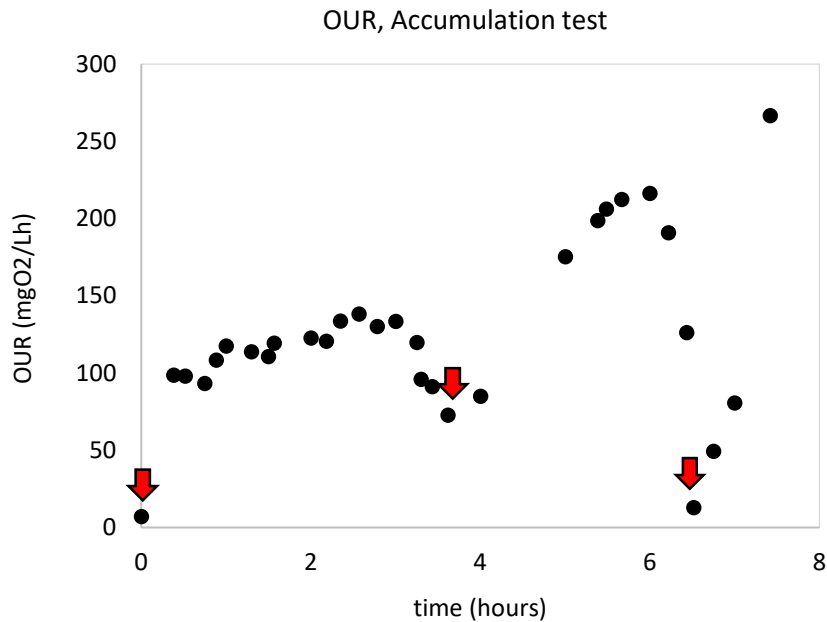
Link between process performance characteristics and microbial population

Significant increase of **Thauera** concentration from 3 ± 0 (Period I) to $58 \pm 11\%$ (Period II.2), according with the increase of the PHA storage yields at SRT of 7-10 days.



The increase of the $Y_{PHA/VFA}$ from 0.42 (with SRT 5 days) to 0.64 (with SRT 7-10) $gCOD_{PHA}/gCOD_{VFAS}$ could be attributed to the presence of other type of organisms such as Paracoccus and Azoarcus.

Accumulation and PHAs Extraction



- The presence of carbon substrate was monitored via the Oxygen Uptake Rate (O.U.R) profiles;
- No significant differences between the period I and II.1 in terms of %PHAs/gTS (25% PHA/gTS);
- The carbon source from cellulosic sludge maximized the PHAs content up to 50 %PHA/gTS.

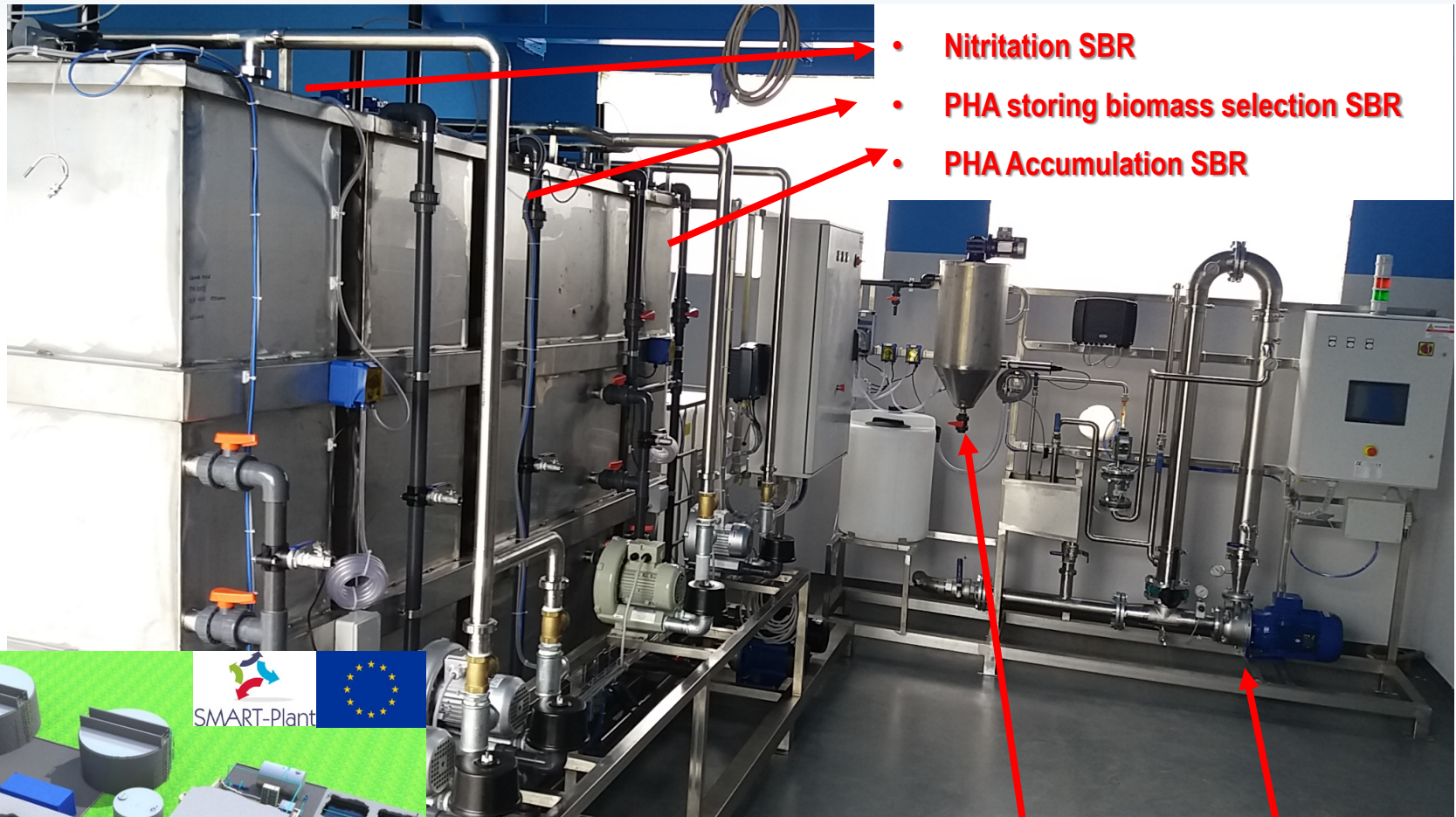
From SCEPPHAR 1.0 to 2.0 (using cellulosic sludge)

Parameter	Original configuration SRT 15 days (Frison et al., EST, 2015)	Period II.2 (107-145) SRT 10 days (current study)
Carbon Source	VFAs from mix sludge fermentation	VFAs from real cellulosic primary sludge
Denitrification efficiency (%)	89,4 ± 4	86,21 ± 4,85
F/M (kgCOD/KgMVLSS) feast/famine	0,37 ± 0,07	0,63 ± 0,09
-qVFA (mgCOD/gVSS h)	0,13	0,10 ± 0,01
<u>Y_{PHA/VFAs}</u> (gCOD-PHA/gCOD-VFA)	239 ± 7	260,24 ± 18,77
	<u>0,42 ± 0,03</u>	<u>0,64 ± 0,06</u>

Conclusions: via-nitrite PHA production

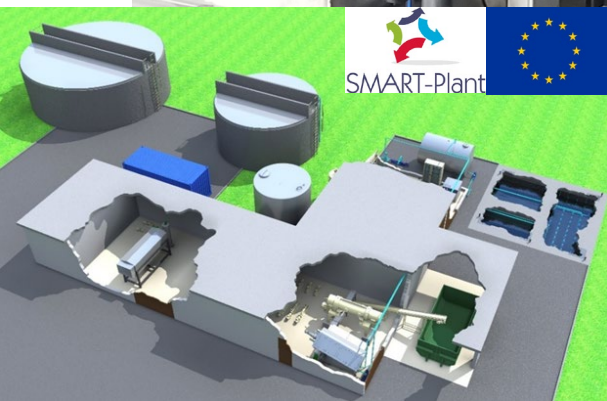
- The fermentation of cellulosic primary sludge from sieved wastewater provides a suitable source of propionate-rich VFAs for the PHA production;
- Aerobic/Feast with Anoxic/Famine regime was coupled with the via-nitrite route to treat high nitrogenous anaerobic effluent;
- After 2 hours of accumulation, the PHA content was as much as 50% (gPHA/gTVS).
- Struvite recovery from cellulosic primary sludge can be a strategy to promote the PHA storage during the accumulation stage.
- The Sidestream S.C.E.P.P.H.A.R. is the **SMARTech5** of the **Horizon2020 Smart-Plant** which will be scaled up at pilot scale (potential 0.5-0.8 kgPHA/d) within the WWTP of Carbonera (TV);

SCEPPHAR pilot scale at the Carbonera WWTP (Smartech 5)



- Nitritation SBR
- PHA storing biomass selection SBR
- PHA Accumulation SBR

- Struvite Crystallizer
- Solid liquid separation



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