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# **Deliverable D6.4** Dynamic simulation and control of the BioSFerA process

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# Contents









# <span id="page-4-0"></span>Executive Summary

This deliverable presents the dynamic simulation analysis of the BioSFerA main processes. A specific methodology for the development of the dynamic model was adopted and a strategy for the control of the key specification under various disturbances was built. For the plantwide control and the dynamic process simulation runs, the BioSFerA process was split into three parts, thermochemical, biological and thermocatalytic one and the analysis was performed separately.

For the thermochemical part, a Dual Fluidized Bed Gasification model that is exportable to Aspen Dynamics was developed in Aspen Plus. The model takes into account the hydrodynamics whereas the kinetic parameters of the main reactions were fine-tuned based in order to predict with good agreement the main products specifications as they have been reported in D6.2 for the large scale. The maintenance of the both reactors temperature and the syngas quality were the key conditions for the controllers setup.

The biological part of this deliverable presents a comprehensive dynamic simulation of syngas and acetate fermentation processes, designed for the continuous production of acetate and lipids, respectively. After fitting the model parameters using experimental and industrial-scale data, model validation for the industrial-scale data was conducted. Subsequently, the control systems for the validated model for both fermentation processes were integrated, using PID closed-loop controllers with feedback response mechanisms.

For the thermocatalytic part, an oil hydroprocessing model that is exportable to Aspen Dynamics was developed in Aspen Plus that includes both the fuel synthesis and recovery part. The model was verified against the respective data from the WP5 pilot activities. The control strategy was designed in such way that the reactor specifications (temperature, pressure and H2/oil ratio) are maintained at the desired levels and the bio-jet fraction is recovered with the appropriate properties, targeting to boiling point.





#### <span id="page-5-0"></span>1 **Introduction**

The present document encompasses the dynamic simulations of the three main sub-units (Dual Fluidized Bed Gasifier, Gas/Liquid Fermentation, TAGs hydroprocessing) of the BioSFerA concept. As initial point of reference, every investigated scenario was assumed to be in the steady state conditions presented in Deliverable D6.2 'Results of full-chain process simulations'. Dedicated unsteady conditions were applied in each sub-unit involving variations in feed streams flow rate and composition and changes in operating conditions. The dynamic simulations of the plant are carried out with the assistance of the commercial software Aspen Plus Dynamics™ (APD) and Matlab/Simulink.

The main scope of the present dynamic study is the primary evaluation of the plant adaptability in targeted operational fluctuations and the identification of the required measures to achieve system restore according to the desired conditions. The present dynamic study is focused on the investigation of the concept flexibility in selected unsteady conditions, rather than the optimization of the applied control scheme and optimal dynamic response of the system. The latter could be investigated within the framework of a more in-depth dynamic analysis dealing with difficult situations (i.e. emergency scenarios) and assessing their techno-economic impact would require further maturation of the technology.



#### <span id="page-6-0"></span> $\overline{2}$ Approach and methodology

The dynamic behavior of each main sub-unit (Dual Fluidized Bed Gasifier, Gas/Liquid Fermentation, TAGs hydroprocessing) is investigated as well as their transition over different operational modes via an appropriate control scheme that ensures a continuously safe and efficient performance. The adaptability of the BioSFerA concept is assessed under different possible conditions.

The scope of this dynamic study is primarily the evaluation of the plant adaptability in operational fluctuations and the identification of the required measures to achieve system restore according to the desired conditions. The level of detail for each model is strongly depended on the available data from the respective pilot activities and the scale up approach for the design of the basic components.

It should be clarified that the main objective of the present dynamic study is limited to the investigation of the concept flexibility and adaptability in selected unsteady conditions, rather than the optimization of the applied control scheme and optimal dynamic response of the system. The latter along with the involvement of equipment failure situations can be re-evaluated with the further maturation of the technology.





# <span id="page-7-0"></span>3 Thermochemical part

## <span id="page-7-1"></span>3.1 DFBG – Steady state model and export to Aspen Dynamics

A 0.5D DFBG process model that is exportable to Aspen (Plus) Dynamics (APD) has been developed and validated. The model takes into account the basic geometrical aspects as are reported in D6.1 [1] and verified in T6.3, the reaction kinetic models that were also applied in D6.1 the hydrodynamics along the risers, the pressure drop in the whole dual system and the basic heat and mass balance information as have been reported in D6.2, downscaled to 100M $W_{th}$  plant [2].

For the DFBG modeling, a different approach from that in D6.2 had to be adopted since the former model cannot be exported and executed in APD. Hence, a new DFBG model was developed in Aspen Plus™ that is able to address the following three key challenges:

1. Overcoming the inherent weakness of APD for not supporting solids and inclusion of all the involved solids in the process i.e. fuel, sand and ash.

2. Inclusion of Reactor basic dimensions and hydrodynamics as they play important role on the process performance

3. Development of a model that is consistent with the former DFBG model as it was presented in D6.2 in respect of the heat and mass balance.

In order to overcome the above mentioned bottlenecks, the following assumptions – simplifications are necessary to be taken into account:

- Hydrodynamics mechanisms and phenomena that take place in a circulating fluidized bed such as the internal solids recirculation and the core-annulus effect are not taken into account.

- The heat losses at the reactors are calculated as 1% of the fuel heat input on a LHV basis

- Physical properties of the user defined 'solids' such as density and heat capacity (*Cp*) are constant and independent of the temperature.

## <span id="page-7-3"></span><span id="page-7-2"></span>3.1.1 Physical properties of the user defined solids



*Table 1: Fuel stream composition (mol/mol)*

A special approach is introduced for the modelling of the streams that are normally represented from nonconventional and solids compounds. This approach is based on the construction of user defined components C, ash (represented by the compound of Ca), permanent at liquid phase maintaining the properties (molecular weight,





*Hcomb*, *ρ*, *Cp*, *ΔHform*, *ΔGform*) same to the corresponding real ones. As concerns the fuel stream, the feed stream that is considered is the devolatilized stream after the pyrolysis step. The composition of this stream is obtained from the model presented in [Table 1.](#page-7-3)

As already mentioned, ash is modelled as an inert compound like Ca. The sand is considered as 50% CaCO<sub>3</sub> and 50% SiO<sup>2</sup> on a mass basis. Moreover, as the enthalpy of the feed stream is not the same with the enthalpy of the original 'correct' fuel, the energy balance consistency is satisfied by adding an inlet heat stream with the value of that difference. This value is obtained from the devolatilization reactor (RYIELD) of the D6.2 process model [2].

Following the same approach with the former DFBG process model (developed within D6.2), the remaining char is modelled as C instead of CaHbOc.

For the definition of the user defined components (C, ash, CaCO<sub>3</sub>, SiO<sub>2</sub>), the main scope is to construct five components that will be permanently at the liquid phase (i.e. the extended Antoine vapor pressure to be equal to 0 and the normal boiling temperature to be very high) and will maintain all the physical properties of the real components [\(Table 2\)](#page-8-1).

<span id="page-8-1"></span>

#### *Table 2: User defined components physical properties.*

## <span id="page-8-0"></span>3.1.2 Model description

[Figure 1](#page-9-1) presents a schematic depiction of the DFBG model as it was developed in Aspen Plus in the framework of the dynamic process simulations. It should be mentioned that all the sub-blocks along the Gasifier riser were modeled as continuous stirred tank reactors (CSTR). Although the most suitable reactor block to model the freeboard zone is the plug flow reactor (RPLUG), this was not allowed by Aspen Plus due to the minimum requirement for solid void fraction (>0.01). At first, a separator block (SEP) is employed to separate the components that formulate the fluid parts (volatiles and moisture) with the real solid one i.e. the ash and the C that corresponds to the fixed carbon (char).





*Figure 1. Aspen flowsheet of DFBG unit*

## <span id="page-9-1"></span><span id="page-9-0"></span>3.1.3 Gasification reactions modeling

<span id="page-9-2"></span>For the impurities, tars CO/CO<sup>2</sup> formation, the reactions i[n Table 3](#page-9-2) are integrated in a RSTOIC:

	<b>Fractional conversion</b>	<b>Fractional conversion of component</b>	
$N_2 + 3H_2 \rightarrow 2NH_3$	0.8889	N <sub>2</sub>	R 1
$2C + N_2 + H_2 \rightarrow 2HCN$	0.0055	N <sub>2</sub>	R 2
$C+2H_2 \rightarrow CH_4$	0.1712		R 3
$2C+2H_2 \rightarrow C_2H_4$	0.1367		<b>R4</b>
$6C+3H_2 \rightarrow C_6H_6$	0.1021		R 5
10C+4H <sub>2</sub> $\rightarrow$ C <sub>10</sub> H <sub>8</sub>	0.0881		R 6
$S + H_2 \rightarrow H_2S$	0.95	S	<b>R7</b>
$2S + 2C + O2 \rightarrow 2COS$	0.05	S	R 8
$Cl2 + H2 \rightarrow 2HCl$		Cl <sub>2</sub>	R 9
$C+0.5O2\rightarrow CO$	0.7		R 10
$C+O2\rightarrow CO2$	1.0		R 11

*Table 3: Reactions specifications for impurities formation (reactions occur in series)*

The fractional conversion rates were calculated by means of Calculator block targeting to approach the respective mole fractions of the impurities in the raw syngas, as they had been calculated and presented in D2.5 [3]. Moreover, the remaining C that does not take part in the HCs, HCN and COS formation is converted into CO and CO<sub>2</sub> by a ratio of 0.7/0.3.

In the gasification step, it is assumed that only homogeneous reactions (water gas shift [R 12](#page-10-1) and gas oxidation reactions [R 13,](#page-10-2) [R 14\)](#page-10-3) take place in the bubble zone whereas in the emulsion zone both homogeneous and heterogeneous (water gas [R 15](#page-10-4) and Boudouard [R 16\)](#page-10-5) reactions are considered. [Table 4](#page-10-6) summarizes all the parameters for the gasification kinetic modeling. The homogeneous reactions were embedded straight at Aspen Plus as are first order Arrhenius whereas the kinetics for the heterogeneous were calculated via an external Fortran code.



<span id="page-10-6"></span><span id="page-10-4"></span><span id="page-10-3"></span><span id="page-10-2"></span><span id="page-10-1"></span>

#### *Table 4. Gasification reactions kinetic parameters*

<span id="page-10-5"></span>Molar concentrations are expressed in kmol/m

The oxidizer reactor is modelled in a simpler way assuming that all combustibles totally react with the oxygen producing combustion products in a stoichiometric reactor (RSTOIC). The ash and the hot sand are perfectly removed from the flue gas after reactor exit while the latter one are led to the gasifier.

## <span id="page-10-0"></span>3.1.4 Hydrodynamics

A 1D model for fast fluidization is set aiming to calculate the particle distribution along the two risers [6]. Like in the majority of detailed CFB gasification process models in the literature, Kunii-Levenspiel model for circulating fluidized bed was employed [7],[8]. Information such as fluidization agent characteristics (*U*, *μ*), gas/solids density (*ρg*, *ρp*), particles mean diameter (*dp*), reactor main dimensions (*D*, *Ht, At*), the gas distribution per nozzle (*Ao*) and total pressure drop (Δ*p*), are used as input in order to calculate the bed height *(Hbed*), dense zone/ freeboard volume area, solid volume fraction distribution as well as the bubble/emulsion phase volume ratio (*δ*). The gasifier unit main design parameters are given in [Table 5.](#page-10-7)



<span id="page-10-7"></span>





<sup>1</sup> the hydrodynamics calculations take place before the gasifier operation blocks and the current (syn)gas density is not yet available. To overcome that, it is considered that the gas density is the inlet gas density decreased by a certain correction factor

The superficial gas velocity *U<sup>0</sup>* is calculated as:

$$
U_0 = 1.05 \frac{F_g}{\rho_g A}
$$
 *Equation 1*

Where  $F_g$  is the gas inlet mass flow (in kg/s) and  $A=1/4 \cdot \pi \cdot D^2$  is the reactor cross flow area (in m<sup>2</sup>).

The minimum fluidization velocity is calculated as:

$$
U_{mf} = \frac{(\sqrt{c_1^2 + c_2 Ar} - c_1)\mu_g}{\rho_g d_p}, (C_1 = 27.2 \text{ and } C_2 = 0.0408)
$$
 *Equation 2*

where *Ar* is the Archimedes number:

$$
Ar = \frac{d_p^3 \rho_g (\rho_p - \rho_g)g}{\mu^2}
$$
 Equation 3

The saturated solids diameter  $d_p^*$ :

$$
d_p^* = d_p \left( \frac{g \cdot \rho_g \cdot (\rho_p - \rho_g)}{\mu^2} \right)^{1/3}
$$
 Equation 4

Whereas the saturated gas velocity is:

$$
U_g^* = U_0 \left( \frac{\mu_g \cdot g \cdot (\rho_p - \rho_g)}{\rho_g^2} \right)^{1/3}
$$
 Equation 5

The solids entrainment velocity *Up,se* is:

$$
U_{p,se} = 1.53 \sqrt{\frac{(\rho_p - \rho_g)d_p g}{\rho_g}}
$$
 Equation 6

And the solid flux in the riser  $G_s$  (kg/m<sup>2</sup>s) is:

$$
G_s = \frac{F_{cs}}{A}
$$
 Equation 7

Where  $F_{c,s}$  is the mass flow of the circulating solids.

The saturated terminal velocity is:

$$
U_t^* = \frac{1}{\frac{18}{d_p^*^2} + \frac{0.591}{d_p^{*0.5}}}
$$
 *Equation 8*

and the terminal velocity *U<sup>t</sup>* is calculated accordingly:

$$
U_t = U_t^* \left( \frac{\mu(\rho_p - \rho_g)g}{\rho_g^2} \right)^{1/3}
$$
 Equation 9



The saturated solids flux  $G_s^*$ :

$$
G_s^* = 23.7 \rho_p U_o exp\left(-5.5 \frac{U_t}{U_o}\right)
$$
 *Equation 10*

And the saturation capacity of the gas is estimated based on the following equation:

$$
\varepsilon_s^* = \frac{G_s^*}{(U_o - U_t)\rho_p}
$$
 *Equation 11*

For the hydrodynamics calculations at the dense zone, firstly it is estimated the fast fluidization input parameter alpha as

$$
a = \frac{5}{U_o}
$$
 Equation 12

It is also assumed that the solid void fraction at dense zone is constant at  $\varepsilon_{s,d}$  = 0.16. The solid void fraction at riser exit is estimated based on the following equation:

$$
\varepsilon_{s,e} = \varepsilon_s^* + (\varepsilon_{s,d} - \varepsilon_s^*) \cdot \exp(-a \cdot H_l) \tag{Equation 13}
$$

where  $H_l$  is the total height of lean zone calculated as:

$$
H_l = H_t - H_{bed}
$$

The solid inventory at dense zone is:

$$
W_{bed} = A \cdot \rho_p \cdot H_{bed} \cdot \varepsilon_{s,d}
$$

The mean solid void fraction at lean zone is:

$$
f = \varepsilon_s^* + \frac{\varepsilon_{s,d} - \varepsilon_{s,e}}{H_l \cdot a}
$$
 Equation 16

The solid inventory at lean zone is:

$$
W_l = A \cdot \rho_p \cdot H_l \cdot f
$$
 Equation 17

Based on the above equations a corrected calculation of the lean zone height can be expressed from the following equation:

$$
H_{l,new} = \frac{\frac{\varepsilon_{s,d} - \varepsilon_{s,e}}{a} + H_t \cdot \varepsilon_{s,d} - \frac{W}{A \cdot \rho_p}}{\varepsilon_{s,d} - \varepsilon_s^*}
$$
 Equation 18

The equations [Equation 14](#page-12-0) - [Equation 18](#page-12-1) are solved iteratively as soon the error *err* = 1- *Hl*/*Hl,new* becomes less than 0.0001.

In order to calculate the basic design characteristics in the dense zone (i.e. the volume of the emulsion and bubble phase zone) the following set of equations are used [9]. At first, the bubble diameter (*db*) is calculated from the following equation [10]:

<span id="page-12-1"></span><span id="page-12-0"></span>*Equation 15*





<span id="page-13-1"></span><span id="page-13-0"></span>*Equation 23*

$$
d_b = 0.54 \left( U_o - U_{mf} \right)^{0.4} \left( \frac{H_{bed}}{2} + 4\sqrt{A_o} \right)^{0.8} g^{-0.2}
$$
 *Equation 19*

The volume fraction of the bed consisting of bubbles (*δ*) is defined as:

$$
\delta = \frac{U_{vis}}{U_{vis} - U_{b\infty}}
$$
 *Equation 20*

Where the single bubble velocity ( $u_b$ ) is calculated as following [11]:

$$
U_{b\infty} = 0.71(gd_b)^{1/2}
$$
 Equation 21

And the visible bubble flow:

$$
U_{vis} = \psi \left( U_o - U_{mf} (1 - \delta) \right)
$$
  
Equation 22  

$$
\psi = \frac{0.26 + 0.70 \cdot \exp(-3300d_p)}{(0.15 + U_o - U_{mf})^{1/3}} \left( \frac{H_{bed}}{2} + 4\sqrt{A_o} \right)^{0.4}
$$
  
Equation 23  
Equation 23

The equations [Equation 20](#page-13-0) - [Equation 23](#page-13-1) are solved iteratively assuming an initial value for δ as soon the error *err2* = 1- *δ*/*δ,new* becomes less than 0.0001.

The freeboard height is divided into four consecutive sub-regions with the same height (and consequently the same volume as the bed diameter is constant). The mean solid fraction at each sub-region is calculated according to the following equation:

$$
\varepsilon_{s,l,i} = \varepsilon_s^* + (-\varepsilon_s^*) \cdot \exp(-a \cdot H_i) \tag{Equation 24}
$$

The pressure drop (in kPa) after both reactors at the dense zone is calculated as:

 $\frac{bea}{2} + 4\sqrt{A_o}$ 

$$
\Delta p_{d,i} = -\varepsilon_{s,d} H_d g \rho_p / 1000
$$

The pressure drop (in kPa) after each reactor at the lean zone is calculated as:

$$
\Delta p_{l,i} = -(1 - \varepsilon_{s,l,i})H_i g \rho_p / 1000
$$
 *Equation 26*

The pressure drop calculations for the rest part of the gasifier apart from the riser were based on the approaches of Karmakar and Datta [12] and Kaiser et al [13]. Firstly, the pressure drop at riser exit duct is calculated as [12]:

$$
\Delta p_{red} = G_{red} (2.84 + 0.0108 U_{red}^2)
$$
 *Equation 27*

where G<sub>red</sub> is solid flux (kg/m<sup>2</sup>-s) that based on the mass balance is:

$$
G_{red} = G_s \frac{A}{A_{red}} = G_s \frac{A}{0.25 \cdot \pi \cdot D_{red}^2}
$$
 *Equation 28*

and *Ured* the solids velocity (m/s):

$$
U_{red} = \frac{G_{red}}{\rho_p} \tag{Equation 29}
$$





The pressure drop at Cyclone is estimated as follows [13]:

$$
\Delta p_{cyc} = k_{cyc} \rho_g U_0^2
$$
 *Equation 30*

where *kcyc* is set equal to 30. For the downcomer, the voidage *εdc* is more than compact bed voidage, but less than voidage at minimum fluidization condition ( $ε<sub>mt</sub>$ ), so it is estimated as:

$$
\varepsilon_{dc} = \delta_b + (1 - \delta_b)\varepsilon_{mf} \tag{Equation 31}
$$

where the bubble fraction [13]:

$$
\delta_b = \frac{1}{1 + \frac{1.3(0.15 + U_{ls} - U_{mf})^{0.33}}{0.26 + 0.7e^{-3.3d_p}} (U_{ls} - U_{mf})^{-0.8}}
$$
 Equation 32

and the voidance at the minimum fluidization velocity is set at *εmf* = 0.5. The velocity at loop seal (*Uls*) is estimated according to the following relation:

$$
U_{ls} = X_{ls} U_{mf} \tag{Equation 33}
$$

The fluidization factor *Xls* is between 5 and 10 and in this study it is set at 6. The pressure drop is then calculated as:

$$
\Delta p_{dc} = -(1 - \varepsilon_{dc}) \rho_p g L_{dc}
$$
 Equation 34

At the loop seal, there is a pressure drop at the standpipe and the horizontal part[13]:

$$
\Delta p_{ls,st} = -(1 - \varepsilon_{dc})\rho_p g h_{st} \tag{Equation 35}
$$

$$
\Delta p_{ls,hor} = 3.5 \cdot \frac{U_{ls}}{2} \frac{A_{wall}}{A_{cross}} \rho_p (1 - \varepsilon_{mf})
$$
 *Equation 36*

where  $A_{wall} = 2 \cdot L_{ls} (h_{ls} + d_{ls})$  and  $A_{cross} = L_{ls} d_{ls}$ 

## <span id="page-14-0"></span>3.2 Model verification and sensitivity analysis

<span id="page-14-1"></span>The main operating conditions are presented in [Table 6.](#page-14-1)

*Table 6. Main operating conditions for the thermochemical part of the BioSFerA concept*







[Table 7](#page-15-0) compares some of the main simulation results from the process models developed in Task 6.2 and here. The small relative errors in all parameters indicate that both models are in good agreement and allow the use of the latter for the dynamic simulation analysis.

<span id="page-15-0"></span>

#### *Table 7. Dynamic DFBG model validation*

[Figure 2](#page-15-1) demonstrates how the voidance, reactor temperature, main syngas products composition and pressure evolves along the gasification riser. The last graph shows the pressure drop not only at the riser but also at the return system. The temperature is almost stable along the dilute zone whereas it is seen that the WGS reactor prevails at this region.



<span id="page-15-1"></span>*Figure 2. Steady state simulation results along the gasifier riser (the last graph depicts the pressure drop at the riser and return system as well)*



Several parametric investigations have been carried out in order to identify the most important and critical parameters, the change of which affect the performance and operation of the DFBG unit. Indicatively, [Table 8](#page-16-0) demonstrates how the exclusion of one of both combustibles gases ("Ext gas": the light gas coming from the fractionation unit after hydrotreatment, "Ferm gas": off gas after the gas fermenter) that are burnt in the oxidizer affects the overall process operation. As great variance in several parameters such syngas quality and composition is observed due to the necessary change in the flow of unconverted char, two of the examined scenarios will be to control the unit when the normal feed of these streams fail.

<span id="page-16-0"></span>

#### *Table 8. Influence of external combustible gas streams on process operation*

The following graphs show the impact of the operating pressure of the gasifier, as it is illustrated by the steam inlet pressure, on the syngas composition and quality:



*Figure 3. Impact of gasifier pressure on produced syngas quality and composition*

The increase in steam pressure inlet causes the lowering of the gasification temperature (autothermal conditions) and increase in H<sub>2</sub>/CO ratio that may have direct impact on the gas fermentation operation.

The Aspen Plus model has been successfully exported to Aspen Plus Dynamics and is able to run at steady state (for the time being) with no errors or problems (see Figure below).





*Figure 4. Temporal evolution of syngas main characteristics at stable conditions*

## <span id="page-17-0"></span>3.3 Control strategy and examined scenarios

The following table summarizes the scenarios and the respective control strategies that are going to be investigated at dynamic/unsteady conditions:



#### *Table 9. Examined scenarios and the control strategy (tentative)*

Before the run of the examined scenarios, the controllers are tuned in order to operate properly. The set point (SP), process variable (PV) and the operating parameter (OP) of each controller are seen in [Figure 5w](#page-18-2)hereas their respective specifications after tuning are summarized in [Table 10.](#page-18-3) All controllers are PI.





<span id="page-18-2"></span>*Figure 5. Aspen Plus Dynamics process flowsheet and controllers parameters*

<span id="page-18-3"></span>

#### *Table 10. DFBG model controllers specifications*

## <span id="page-18-0"></span>3.4 Dynamic simulations

## <span id="page-18-1"></span>3.4.1 Feedstock flowrate

In this scenario, the step change of feedstock flow through sudden decrease and increase is investigated. At first, the biomass flow rate drops by 20% and after 2 hours, it decreases again by 20% operating for 1.5h at 60% load. Then, the full load operation is restored (see [Figure 6](#page-19-0) blue line). The circulating solids flow [\(Figure 6](#page-19-0) grey line) is the process variable to keep the gasification temperature stable (set point) and its variation has strong similarities with that of syngas temperature [\(Figure 7](#page-19-1) blue line). It is observed that the syngas temperature restoration is accomplished with half an hour at all changes occurred. On the other hand, since the circulating solids flow is pretty large, no temperature change is observed.





*Figure 6. Evolution of DFBG operation parameters at load variations*

<span id="page-19-0"></span>

*Figure 7. Reactors temperature evolution*

<span id="page-19-1"></span>It takes around 30 min to restore the temperature level at the gasifier. On the contrary, the temperature at the oxidizer remains the same along the whole time period, as illustrated from the hot sand temperature. This is attribute to the large mass flow and the high heat capacity of the solids that does not favor the great temperature variations.

What is also observed and should be mentioned is that during the period that the DFBG operates at lower load the H2/CO increases, changing the syngas quality that exit the gasifier. In case that this variation has negative effect on gas fermentation performance and in case that the steam catalytic reforming that follows the gasification process is not at the position to restore the syngas composition at the initial desired level, an alternative control strategy should be adopted.



## <span id="page-20-0"></span>3.4.2 Steam flow rate

For this scenario, the temporal variation of inlet water/steam that is illustrated by the steam to biomass ratio is seen in red in [Figure 8](#page-20-1) (left). It was also observed that both the gasification and combustion process operate at the same temperatures without variation. To achieve that, the circulating solids flow and the unconverted char flow vary considerably. The control scheme operates in such a way that the H2/CO ratio almost remains constant at around 2.5. In other words, this control approach is appropriate to keep the syngas quality at a stable and desired level.



<span id="page-20-1"></span>







## <span id="page-21-0"></span>3.4.3 Gasifier temperature

*Figure 9. Effect of various DFBG process parameters on gasifier temperature variation*

In this section, the sudden change (increase) of gasifier temperature up to 800  $^{\circ}$ C in three steps is examined. Like in other cases, the oxidizer (or hot sand) temperature remains the same during the whole test run. To achieve that, the solids (hot sand) circulation flow should have been controlled properly. The char conversion at the gasifier increases as gasifier temperature gets higher, then the unconverted char drops and the circulating solids (control output) should increase in order to cover the elevated heat demands at the gasifier. The hydrogen and CO content in the syngas increases probably due to the higher conversions in water gas reaction [\(R 15\)](#page-10-4) and thus the H $_2$ /CO ratio does not vary considerably.

## <span id="page-21-1"></span>3.4.4 Fermentation off gas flow

[Figure 10](#page-22-0) shows how the main process parameters are affected from the change (sudden stop) of fermentation off gas flow rate. When this occurs, the main operation temperatures (i.e. gasification and hot sand) cannot maintain at their initial levels and drop. The unconverted char that increases is not sufficient enough to restore them. As a matter of that, the produced syngas decreases and the  $H_2$ /CO increases because the WGS reaction favors from the fact that more steam is available to react with CO as carbon water gas reaction rate decreases.



It should be mentioned that the control scheme in that case is set up in such way that the recirculating solids flow rate and steam to biomass ratio remain unstable. In case that the change in syngas composition does not favors the gas fermentation efficiency, an alternative control strategy should be established.



<span id="page-22-0"></span>*Figure 10. Effect of fermentation off gas flow on DFBG parameters* 





#### <span id="page-23-0"></span>Biological part  $\overline{4}$

The aim of this section is to conduct dynamic simulations for the biological part of the system and to design a plantwide control system to regulate syngas composition in accordance with strain specifications while minimizing byproducts. The biological part consists of two different fermentation processes: syngas fermentation and acetate fermentation. In the first one, which involves the gaseous substrate, syngas is converted into acetate under anaerobic conditions using advanced acetogenic bacteria. The acetyl-CoA pathway, also known as the Wood-Ljungdahl pathway, can utilize both CO and H<sub>2</sub> as electron donors and CO and CO<sub>2</sub> as carbon sources (Figure 11).



*Figure 11: Wood-Ljungdahl Pathway for acetate synthesis from CO/CO2/H<sup>2</sup> syngas*

The chemical reactions catalyzed by the cell are presented in [R 17](#page-23-1) and [R 18:](#page-23-2)

$$
4 CO + 2 H2O \rightarrow C2H4O2 + 2 CO2
$$
 R 17

<span id="page-23-2"></span><span id="page-23-1"></span>
$$
4 H_2 + 2 CO_2 \rightarrow C_2 H_4 O_2 + 2 H_2 O
$$
 R 18

In the second fermentation step, which involves the liquid substrate, the produced acetate is converted into targeted lipids, namely triacylglycerides (TAGs), under aerobic conditions, using oleaginous yeasts such as *Yarrowia lipolytica*. The metabolic model depicting this process is illustrated in Figure 12.





*Figure 12: Reaction mechanism for TAGs production through acetate fermentation*

There are two steps involved in this process, the growth phase and the lipid production phase. In the growth phase, biomass must be generated for the yeasts to accumulate lipids. The stoichiometric equation for this phase is expressed by [R 19.](#page-24-0)

 $CH_3COOH + 0.908 O_2 + 0.147 NH_3 \rightarrow 1.05 CH_{1.66}O_{0.54}N_{0.14} + 1.349 H_2O + 0.95 CO_2$  R 19 In the second phase, lipids are produced inside the cultivated yeasts. The stoichiometric equation for this phase is presented in [R 20.](#page-24-1)

<span id="page-24-1"></span><span id="page-24-0"></span>
$$
50.11 \text{ CH}_3\text{COOH} + 27.72 \text{ O}_2 \rightarrow C_{51}H_{98}O_6 + 51.22 \text{ H}_2\text{O} + 49.22 \text{ CO}_2
$$
 R 20

This section focuses on the dynamic simulation of the industrial-scale implementation of the two fermentation processes. The suggested double-stage fermentation scheme for the scale up is depicted in [Figure 13.](#page-24-2)



*Figure 13: Suggested double-stage fermentation scheme for the scale up*

<span id="page-24-2"></span>More particularly, in the gas fermentation process, syngas, that is the substrate, is distributed among 30 bioreactors



and enters the bioreactors in parallel. The liquid medium is fed in series from the first bioreactor stage to downstream stages [14]. The acetate concentration gradually increases in the bioreactors, and the final concentration must reach up to 30 g/L to prevent accumulation.

For the liquid fermentation process aimed at TAGs production, two phases are considered. The first phase involves a CSTR reactor serving as the biomass growth phase in a rich medium under continuous and aerobic conditions. The second phase is dedicated to TAGs production, operating under continuous and nitrogen-limited conditions. In this phase, the acetate previously produced is divided among 29 reactors and added in parallel. Additionally, air is added in parallel in both phases. Cells are harvested at the outlet of the final stage [15].

## <span id="page-25-0"></span>4.1 Dynamic process simulation

Based on the analyzed scale-up scheme, a mathematical model is developed using MATLAB and Simulink for dynamic process simulation. Firstly, the differential equations for mass balances are determined based on the biological reactions of each fermentation process. Two phases, gas and liquid, are considered, and the differential equations for components are derived according to their respective phases.

In the dynamic model, several kinetic parameters must be determined using MATLAB's optimization tool "fmincon". Firstly, for the syngas fermentation process, the kinetic parameters are fitted to the experimental data for cleaned syngas from D3.4 [16] and then to the experimental data from T4.3 when syngas is contaminated with impurities. For the acetate fermentation process, the kinetic parameters are fitted to the experimental data from D3.5 [17].Then, based on the industrial-scale data from D6.2 [2], the final fitted parameters for each case are found.

Once the kinetic parameters for the industrial-scale data from D6.2 [2] are fitted, the dynamic simulation is developed in Simulink, and validated by achieving the similar final product concentration with D6.2 [2]. Both fermentation processes are assumed to operate continuously in bioreactors. Feed is added while product is withdrawn, treating the bioreactors as simple continuous stirred tank reactors (CSTRs) where various biological reactions take place. Assumptions include perfect mixing within the reactor, constant pH, and constant flow rates for substrate feed and product output.

Subsequently, a plant-wide control system is developed for both syngas and acetate fermentation. The objective for syngas fermentation is to control acetate concentration by adjusting the agitation speed in response to changes in syngas composition. For acetate fermentation, the substrate flowrate is the varying parameter that affects the final TAGs concentration and acetate residual. By manipulating the biomass concentration produced in growth phase -by adjusting the C/N ratio, oxygen flowrate and agitation speed- the final acetate concentration is controlled to the desired value that must be close to zero.

The following sections, 4.1.1 and 4.1.2, describe the dynamic simulation of the two fermentation processes in detail.

## <span id="page-25-1"></span>4.1.1 Syngas Fermentation

### 4.1.1.1 Model description and parameter fitting

#### **Mass balance equations:**

As previously described, the dynamic model for syngas fermentation occurs in a continuous stirred tank reactor (CSTR). The components are divided into non-condensable and condensable and they exist either in liquid or gas phase. The differential equations, [Equation 37](#page-26-0)[-Equation 41,](#page-26-1) assume isothermal and isobaric operation, as well as homogeneity and constant liquid and gas volumes in the reactor [18].

For non-condensable components, j, that are CO, H<sub>2</sub>, CO<sub>2</sub>, and the impurities NH<sub>3</sub>, H<sub>2</sub>S, HCN, COS, Benzene:





In the gas phase:

<span id="page-26-0"></span>
$$
\frac{dC_{G,j}}{dt} = \left(\frac{1}{V_G}\right) \left(Q_{G,in} \cdot C_{G,j,in} - Q_{G,out} \cdot C_{G,j}\right) - k_L a_j \left(\frac{C_{G,j}}{m_{j,NC}} - C_{L,j}\right) \left(\frac{V_L}{V_G}\right)
$$
   
Equation 37

In the liquid phase:

$$
For CO, H_2, CO_2: \frac{dC_{L,j}}{dt} = \left(\frac{Q_L}{V_L}\right) \left(C_{L,j,in} - C_{L,j}\right) + k_L a_j \left(\frac{C_{G,j}}{m_{j,NC}} - C_{L,j}\right) + v_j \cdot C_X
$$
  
For impurities: 
$$
\frac{dC_{L,j}}{dt} = \left(\frac{Q_L}{V_L}\right) \left(C_{L,j,in} - C_{L,j}\right) + k_L a_j \left(\frac{C_{G,j}}{m_{j,NC}} - C_{L,j}\right)
$$

For condensable components,  $i$ , that are acetate,  $H_2O$ :

In the gas phase:

$$
\frac{dC_{G,j}}{dt} = \left(\frac{1}{V_G}\right) \cdot \left(Q_{G,in} \cdot C_{G,j,in} - Q_{G,out} \cdot C_{G,j}\right) + k_L a_j \left(\frac{C_{L,j}}{m_{j,C}} - C_{G,j}\right) \left(\frac{V_L}{V_G}\right)
$$
\nEquation 39

In the liquid phase:

$$
\frac{dC_{L,j}}{dt} = \left(\frac{Q_L}{V_L}\right) \cdot \left(C_{L,j,in} - C_{L,j}\right) - k_L a_j \left(\frac{C_{L,j}}{m_{j,C}} - C_{G,j}\right) + v_j \cdot C_X
$$
 Equation 40

For the biomass concentration,  $C_X$ , in the liquid phase:

<span id="page-26-1"></span>
$$
\frac{dC_X}{dt} = \left(\frac{Q_L}{V_L}\right)(-C_X \cdot XP) + \mu \cdot C_X - r_d
$$
 \tEquation 41

Where  $m_{j,NC}, m_{j,C}$  parameters are the gas-liquid equilibrium factors for non-condensable and condensable components, listed in [Annexes.](#page-59-0)

The industrial-scale fermentation consists of 30 CSTR, where  $V_G, V_L$ , are the volumes of gas and liquid inside each reactor and equal to 50000 L and 250000 L, respectively. The volumetric flow rate in,  $Q_{G,in}$ , and out,  $Q_{G,out}$ , of the reactor are both at 656043  $\frac{L}{hr}$ , and  $XP=0$ , meaning that all cells are recycled to the reactor. The above values are obtained from D6.2 [2].

#### **Reaction rates:**

The specific consumption/production rates of species CO and H<sub>2</sub>,  $v_{CO, H2}$   $\left(\frac{mol}{ab} \right)$  $\frac{m_{\rm \scriptscriptstyle HI}}{g\cdot h r}$  are estimated i[n Equation 42](#page-26-2) [18]:

<span id="page-26-2"></span>
$$
v_{CO, H2} = -\frac{v_{max,j}C_{L,j}}{K_{S,j} + C_{L,j}} I_i I_{CO}
$$
   
Equation 42

Where i are the Acetic acid, NH<sub>3</sub>, H<sub>2</sub>S, HCN, COS, Benzene components, and  $I_i = \frac{1}{\sqrt{2}}$  $\frac{C_{L,i}}{1+\frac{C_{L,i}}{V}}$  $K_{l,i}$  $I_{CO,j=H_2}$  = 1  $1+\frac{C_{L,CO}}{V}$ ,  $I_{CO,j=CO} = 1$ .

The specific biomass growth rate  $\mu$   $(hr^{-1})$  is calculated from these rates and yield coefficients  $Y_{X,CO}$  and  $Y_{X,H2}$ , [Equation 43:](#page-26-3)

<span id="page-26-3"></span>
$$
\mu = -v_{CO}Y_{X,CO} - v_{H2}Y_{X,H2}
$$
 *Equation 43*

The death rate  $r_d$  is a function of cell concentration, as shown in [Equation 44:](#page-27-0)



 $K_{l,CO}$ 



<span id="page-27-0"></span>
$$
r_d = k_d C_X
$$
 Equation 44

where  $k_d$  is the death constant.

The reaction rates from Reactions 1 and 2 are calculated from [Equation 45](#page-27-1) an[d Equation 46:](#page-27-2)

<span id="page-27-3"></span><span id="page-27-2"></span><span id="page-27-1"></span>
$$
v_{1R} = -\frac{v_{CO}}{4}
$$
 Equation 45  

$$
v_{2R} = -\frac{v_{H2}}{4}
$$
 Equation 46

4 The total consumption/production rates of other components are calculated fro[m Equation 47-](#page-27-3) [Equation 49:](#page-27-4)

<span id="page-27-4"></span>
$$
v_{CO2} = 2v_{1R} - 2v_{2R}
$$
 *Equation 47*

$$
v_{HAC} = v_{1R} Y_{ACCO} + v_{2R} Y_{ACH2}
$$
 *Equation 48*

<span id="page-27-5"></span>
$$
v_{H20} = -2v_{1R} + 2v_{2R}
$$
 *Equation 49*

So, the unknown kinetic parameters that are estimated are the  $k_d$ , the yield coefficients used in these calculations,  $v_{max,CO}$ ,  $v_{max,H2}$ ,  $K_{S,j}$ ,  $K_{l,j}$ .

#### **Mass transfer coefficients:**

To calculate the mass transfer coefficients  $k<sub>L</sub>a<sub>i</sub>$  several equations must be determined based on literature [18]. Firstly, the mass transfer coefficient  $k<sub>L</sub>a$  for air in water at 20 °C is described by [Equation 50:](#page-27-5)

$$
k_L a^{(20)}(hr^{-1}) = f_0 k_L a_0^{(20)} + (1 - f_0) k_L a_1^{(20)}
$$
 Equation 50

Where  $k_L a_0^{(20)}$  and  $k_L a_1^{(20)}$  are the nonalescing and coalescing broth according to the correlations proposed by van't Riet (1979) for air in water and are described by [Equation 51](#page-27-6) and [Equation 52,](#page-27-7) respectively:

$$
k_L a_0^{(20)}(hr^{-1}) = 3600(0.002 \left(\frac{P_g}{V_L}\right)^{0.7} (u_s)^{0.2}
$$
   
Equation 51

$$
k_L a_1^{(20)}(hr^{-1}) = 3600(0.026 \left(\frac{P_g}{V_L}\right)^{0.4} (u_s)^{0.5}
$$
 Equation 52

Where  $u_s$ , is the superficial gas velocity and  $\frac{p_g}{v_L}$  is the impeller power per unit volume, which is estimated from the impeller ungassed power  $P_{ug}$  [\(Equation 53\)](#page-27-8) and the correlation for the  $P_g$  i[n Equation 54.](#page-27-9)

<span id="page-27-10"></span><span id="page-27-9"></span><span id="page-27-8"></span><span id="page-27-7"></span><span id="page-27-6"></span>
$$
P_{ug} = N_p \rho_L N^3 d_i^5
$$
 *Equation 53*

$$
\frac{Q_{G,in}N^{0.25}}{d_i^2} \le 0.055 \to P_g = -(9.9 \left(\frac{Q_{g,in}N^{0.25}}{d_i^2}\right) P_{ug} - P_{ug})
$$
  
\n
$$
\frac{Q_{G,in}N^{0.25}}{d_i^2} \ge 0.055 \to P_g = -(0.52 + 0.62 \left(\frac{Q_{g,in}N^{0.25}}{d_i^2}\right) P_{ug} - P_{ug})
$$
  
\nEquation 54

The ungassed power number,  $N_p$ , is described by [19] for 1 impeller[, Equation 55:](#page-27-10)

$$
N_p = Agspeed^3(39.3701 \cdot d)^5(4.5 \cdot 10^{-13})
$$
 *Equation 55*

Where  $Agspeed$  is the agitation speed in rpm, and  $N(s^{-1}) = 0.01667 \cdot Agspeed$ 

In all cases, it is assumed that the reactor has a Height/Diameter ratio of 3 and an impeller diameter,  $d$ , of 40% the reactor diameter, D.

The mass transfer coefficient at different temperature is expressed by [Equation 56:](#page-28-0)





$$
k_L a^{(T)}(hr^{-1}) = \frac{k_L a^{(20)}}{1.024^{(20-T)}}
$$
 Equation 56

The individual  $k_L a_j$  for each component is calculated by [Equation 57,](#page-28-1) by applying the penetration theory:

<span id="page-28-1"></span><span id="page-28-0"></span>
$$
k_L a_j = k_L a^{(T)} \left(\frac{D_{f,j}}{D_{f,air}}\right)^{\frac{1}{2}}
$$
 Equation 57

Where  $D_{f,j}$  is the mass diffusivity of species j in water and are expressed in [Annexes.](#page-59-0) Based on the above, the unknown parameters that are estimated are the weighting factor,  $f_0$  and the agitation speed.

#### **Estimation of model parameters:**

The unknown parameters expressed above, that are the  $k_d$ , the yield coefficients,  $v_{max,CO}$ ,  $v_{max,H2}$ ,  $K_{S,j}$ ,  $K_{l,j}$ ,  $f_0$ , AgSpeed are estimated using the maximum likelihood principle (MLP), [18], with data retrieved from experimental data from D3.4 [16] for cleaned syngas, then from experimental data from T4.3 for syngas with impurities and finally from industrial-scale data from D6.2 [2], that are these being used in the dynamic simulation. The fitted parameters and the dynamic profile of each case are illustrated in [Annexes.](#page-59-0) Given the high nonlinearity of the MLP method, the objective function is minimized using the "fmincon" function in MATLAB. This approach ensures a good initial estimation of parameters, enhancing the accuracy of the MLP results.

#### **Model simulation:**

<span id="page-28-2"></span>After the kinetic parameters are fitted to the industrial scale data, the syngas fermentation process is simulated in Simulink and its results are compared against those from D6.2 [2], to validate the model. The dynamic syngas fermentation model is a nonlinear algebraic-differential system, demanding numerical solvers suitable for stiff problems. Therefore, the ode15s method from MATLAB is used for time integration from the initial conditions based on D6.2 [2]. The initial feedstock and operating conditions are outlined in [Table 11.](#page-28-2)





The Simulink model, depicting two of the thirty reactors for syngas fermentation, is illustrated in [Figure 14](#page-29-0) and [Figure 15.](#page-29-1) These figures provide a visual representation of the setup and operation of the fermentation process within the reactors.





*Figure 14: Syngas fermentation model in Simulink*

<span id="page-29-0"></span>

*Figure 15: Subsystem depicting the reactor of syngas fermentation*

<span id="page-29-1"></span>The Simulink model, shown in [Figure 14,](#page-29-0) comprises inputs including the syngas flow rate and composition, operating conditions that are agitation speed and temperature, and the medium flow rate. These inputs are connected to a subsystem representing the first reactor. The outputs from this subsystem are the gas and liquid concentrations exiting the reactor. The syngas fermentation process, as previously explained, consists of a total of 30 reactors, where the liquid medium enters each reactor in series, while syngas enters each reactor in parallel.

For the second reactor, the liquid inlet is the liquid output from the first reactor, and the syngas flow rate and composition remain the same as in the first reactor. Acetate is transferred from one reactor to the next until it exits



from the final reactor. It is crucial to note that the final acetate concentration should not exceed 30 g/L to avoid accumulation.

In [Figure 15,](#page-29-1) the reaction subsystem is depicted, which consists of two MATLAB functions. One function is for the gas phase and the other for the liquid phase. Each function solves the mass balances as previously described.

### 4.1.1.2 Validation of the dynamic simulation

The validation of the dynamic simulation of syngas fermentation involves comparing the concentration of the produced components with the results from D6.2 [2]. In the dynamic model, the produced acetate concentration from the first and the second reactor is the same, equal to 0.0164 mol/L. Therefore, it can be assumed that all the reactors will exhibit the same behavior, and the total concentration of the products is the concentration after the first reactor multiplied by the number of reactors, which is 30. The final acetate concentration after the 30<sup>th</sup> reactors equals to 29.5 g/L. The results from D6.2 [2] and from the dynamic simulation are summarized in [Table 12](#page-30-0) for comparison.

*Table 12: Results from D6.2 and dynamic simulation*

Final concentration (mol/L)	<b>Dynamic simulation</b>	$D6.2$ [2]
Acetate	29.5	30

<span id="page-30-0"></span>[Table 12](#page-30-0) indicates that the final acetate concentration from the dynamic simulation of syngas fermentation closely match that of D6.2 [2]. This validation confirms that the model accurately represents the behavior of the syngas fermentation process[. Figure 16](#page-30-1) illustrates the concentration profiles of CO, H<sub>2</sub>, CO<sub>2</sub> and the produced acetate from the first reactor.



*Figure 16: Concentration profiles of CO, H2, CO<sup>2</sup> and acetate*

<span id="page-30-1"></span>Based o[n Figure 16,](#page-30-1) it can be observed that syngas is consumed in the first hour, while acetate is produced. The conversion rates  $\left(equal\ to\ \frac{final-initial\ concentration}{initial\ concentration}\right)$  of CO, H<sub>2</sub> and CO<sub>2</sub> are then 67.5%, 70% and 69%, respectively.



## <span id="page-31-0"></span>4.1.2 Acetate fermentation

### 4.1.2.1 Model description and parameter fitting

#### **Mass balance equations:**

As for acetate fermentation, the dynamic model of the growth and lipid production phases is developed based on the mass balance equations of the process, assuming that the bioreactors operate continuously [20]. Hence, the liquid volume, V<sub>l</sub>, variation is stable and equal to its initial value. In both stages, pH regulation is crucial and can be controlled at 7 by adding a pH regulatory solution. During continuous culture, pH is only controlled by KOH addition, considered negligible compared to the feed medium addition [20]. As for the feed medium flow rate, F, in both phases, it is set at 30623 L/h. In growth phase, it consists of acetate and ammonium as carbon and nitrogen sources with compositions of 30.5 g/L and 0.18 g/L, respectively. The C/N ratio (kg/kg) of this mixture is 45. In the lipid production phase, the composition of the feed is the one produced from the syngas fermentation, with 28.3 g/L acetate and 0.057 g/L NH3. Hence, the C/N ratio at this stage is 130. The acetate fermentation operates in aerobic conditions, so air enters with flow rate at 764245 L/h.

The model can then be described by the cell mass concentration, X, changes in the reactor, as shown in [Equation](#page-31-1)  [58,](#page-31-1) and the concentration variation in the reactor for liquid solute or a dissolved gas compound i, [Equation 59.](#page-31-2) The components, i, are the acetate, TAGs,  $NH_3$ , and  $O_2$  and  $CO_2$  in the liquid phase.

<span id="page-31-3"></span><span id="page-31-2"></span><span id="page-31-1"></span>
$$
\frac{dX}{dt} = \frac{1}{V_l}(R_X - X \cdot V_l - F \cdot X)
$$
\nEquation 58\n
$$
\frac{dS_i}{dt} = \frac{1}{V_l}(R_X - X \cdot V_l - F \cdot X)
$$
\nEquation 59\nEquation 59

$$
\frac{dS_i}{dt} = \frac{1}{V_l} \Big( R_{Si} - S_i \cdot V_l + F(S_{ifeed} - S_i) \Big) + k_l a (S_i^* - S_i)
$$

The mass balance of the  $O_2$  and  $CO_2$  in the gas phase, j, is described by [Equation 60.](#page-31-3)

$$
\frac{d c_{G,j}}{dt} = \left(\frac{1}{v_G}\right) \left(Q_{G,in} \cdot C_{G,j,in} - Q_{G,out} \cdot C_{G,j}\right) - k_L a_j \left(\frac{c_{G,j}}{m_{j}} - C_{L,j}\right) \left(\frac{v_L}{v_G}\right)
$$
 Equation 60

In the growth phase, the differential equation for TAGs production is set to 0, as TAGs are not produced during this phase. In the lipid production phase, the differential equation for cell mass concentration, X, equals to 0, as TAGs are produced within the cells, resulting in a constant amount. The equations above involve the mass transfer coefficient,  $k<sub>L</sub>a$ , calculated as in syngas fermentation and the dissolved gas concentration in saturation conditions,  $S_i^*$ . The parameters  $m_j$  are the gas-liquid equilibrium factors, listed in [Annexes](#page-59-0) along with the diffusion coefficients at infinite dilution in water, Df.

#### **Reaction rates:**

The production rates of biomass,  $R_X$ , and the compounds involved in the reaction,  $R_{Si}$ , are calculated using algebraic equations [20]. The biomass production rate,  $R_X$ , is associated to the Monod kinetic law, as given b[y Equation 61,](#page-31-4) [20].

$$
R_X = \mu_{max} \cdot \frac{S_{ac}}{K_S + S_{ac}} \cdot \frac{S_N}{K_{SN} + S_N} \cdot \frac{S_{O2}}{K_{SO2} + S_{O2}} \cdot X
$$
 \tEquation 61

Where  $\mu_{ma}$  represents the maximal growth rate, and  $K_s,K_{SN},K_{SO2},$  are the "half velocity" constants for the respective concentrations.

The production rate of the compounds involved in reactions,  $R_{si}$  are calculated as in [Equation 62,](#page-31-5) [20]:

<span id="page-31-5"></span><span id="page-31-4"></span>
$$
R_{Si} = \frac{1}{Y_{X/i}} \cdot R_X
$$
 Equation 62





With  $Y_{X/i}$  the mass yield of each compound i calculated from the stoichiometric equations.

The lipid production rate,  $R_L$ , is expressed as shown [iEquation 63,](#page-32-0) [20].

$$
R_L = B \cdot X \cdot \frac{S_{ac}}{K_S + S_{ac}} \cdot \frac{S_N}{K_{SN} + S_N} \cdot Shift_{RL} \cdot lim_{lip}
$$
 \tEquation 63

e B, is the lipid formation rate.

ShiftRL, is the metabolic shift toward lipid production calculated from [Equation 64:](#page-32-1)

<span id="page-32-1"></span><span id="page-32-0"></span>
$$
Shift_{RL} = 1 - \frac{1 + e^{-100\frac{1 - 11}{11}}}{1 + e^{-100\frac{lim N - 11}{11}}}
$$
 *Equation 64*

Up to a certain lipid content, lipid production stopped and this is expressed by  $lim_{lip}$ , as [Equation 65:](#page-32-2)

<span id="page-32-2"></span>
$$
lim_{lip} = 1 - \frac{1 + e^{-100 \frac{1 - l2}{l2}}}{1 + e^{-100 \frac{\% lip - l2}{l2}}} \tag{Equation 65}
$$

Where  $\%lip = \frac{S_{lip}}{DGM}$  $\frac{\partial u_p}{\partial C}$ , with DCW being the total dry cell weight.

The maintenance is described by the acetic acid oxidation without cell mass or storage compounds synthesis, described b[y R 21,](#page-32-3) [20]:

<span id="page-32-4"></span><span id="page-32-3"></span>
$$
Acetate + 2 O_2 \rightarrow 2 CO_2 + H_2O
$$
 R 21

The associated kinetic law could be written as i[n Equation 66,](#page-32-4) [20]:

$$
R_m = mX \frac{S_{ac}}{Ks + S_{ac}} \frac{S_{02}}{K_{02} + S_{02}} \left(1 - \frac{R_X}{\mu_{max} X}\right) \left(1 - \frac{R_L}{BX}\right)
$$
 Equation 66

Where m is the maintenance term.

<span id="page-32-5"></span>The  $R_{Si}$  equation of each compound is finally illustrated in [Table 13:](#page-32-5)

*Table 13: Production rate equations of the compounds involved in reactions*



#### **Estimation of model parameters:**

The unknown parameters expressed above, that are the  $\mu_{max}$ ,  $K_S$ ,  $K_{SN}$ ,  $K_{SO2}$ ,  $B$ ,  $I_1$ ,  $I_2$ ,  $Y_{Xac}$ ,  $Y_{XO2}$ ,  $Y_{XN}$ ,  $m$ ,  $f_0$ ,  $Y_{lac}$ ,  $Y_{XCO2}$ and Agitation Speed are estimated using the maximum likelihood principle (MLP), [18], with data retrieved from experimental data from D3.5 [17] and then from industrial-scale data from D6.2 [2], that are these being used in the dynamic simulation. The fitted parameters and the dynamic profile of each case are illustrated in [Annexes,](#page-59-0) assuming that the acetate fermentation operates at 28 °C and 1 bar. Given the high nonlinearity of the MLP method, the objective function is minimized using the "fmincon" function in MATLAB. This approach ensures a good initial estimation of parameters, enhancing the accuracy of the MLP results.





#### **Model simulation:**

After the kinetic parameters are fitted to the industrial-scale data, the acetate fermentation process is simulated in Simulink and its results are compared against those from D6.2 [2] to validate the model. The dynamic model of acetate fermentation is a nonlinear algebraic-differential system, demanding numerical solvers suitable for stiff problems. Therefore, as syngas fermentation, the ode15s method from MATLAB is used for time integration setting as feedstock the final concentration of syngas fermentation. The Simulink models depicting the growth reactor are illustrated i[n Figure 17,](#page-33-0) and three of the 29 reactors for the lipid production phase are shown in [Figure 18](#page-33-1) and [Figure](#page-34-0)  [19.](#page-34-0) These figures provide a visual representation of the setup and operation of the fermentation process within the reactors.



*Figure 17: Growth production phase model in Simulink and subsystem of the reactor*

<span id="page-33-0"></span>

<span id="page-33-1"></span>*Figure 18: Lipid production phase model in Simulink*





*Figure 19: Subsystem depicting the reactor for lipid production*

<span id="page-34-0"></span>In the growth phase, [Figure 17,](#page-33-0) the main input values are the medium and air, with the biomass concentration exiting the reactor. This biomass, along with the medium flow rate and composition, operating conditions such as agitation speed and temperature, and the oxygen concentration and flow rate are the inputs for the lipid production phase, [Figure 18.](#page-33-1) These inputs are connected to a subsystem representing the first reactor. The outputs from this subsystem are the gas and liquid concentrations exiting the reactor. The acetate fermentation process, as previously explained, consists of 30 reactors: one for the growth phase and the others for lipid production. In the lipid production reactors, the biomass enters each reactor in series, while acetate produced in syngas fermentation and air enter each reactor in parallel. [Figure 19](#page-34-0) depicts the reaction subsystem, which consists of two MATLAB functions. One function is for the gas phase and the other for the liquid phase. Each function solves the mass balances as previously described.

### 4.1.2.2 Validation of the dynamic simulation

The validation of the dynamic simulation of acetate fermentation involves a comparison of the concentration of the produced components with the results from D6.2 [2]. During the growth phase, an increase in the C/N ratio is expected as the biomass is cultured. According to the dynamic model, the final C/N ratio after biomass production is expected to reach 230.

[Figure 20](#page-35-0) illustrates the concentration profiles of biomass, acetate, TAGs and NH<sup>3</sup> as well as the C/N ratio profile of the growth phase.





*Figure 20: Concentration profiles of components and C/N ratio in the growth phase*

<span id="page-35-0"></span>Analysis of [Figure 20](#page-35-0) reveals that during the growth phase, acetate and NH<sub>3</sub> are consumed, resulting in the production of biomass. Hence, as expected, the C/N ratio increases. It is noteworthy that TAGs are not produced during this growth phase.

<span id="page-35-1"></span>As for the liquid production phase, based on the dynamic model, the produced lipid concentration from the first and the second reactor is the same, equal to 0.24 g/L. Therefore, it can be assumed that all the reactors will exhibit the same behavior, and the total concentration of the products is the concentration after the first reactor multiplied by the number of reactors, which is 29. So, a final TAGs concentration after the 29<sup>th</sup> reactor of 6.96 g/L is achieved. The results from D6.2 [2] and from the dynamic simulation are summarized in [Table 14](#page-35-1) for comparison.





[Table 14](#page-35-1) indicates that the final concentrations from the dynamic simulation of acetate fermentation closely match those from D6.2 [2]. This validation confirms that the model accurately represents the behavior of the acetate fermentation process.

[Figure 21](#page-36-2) illustrates the concentration profiles of biomass, acetate, TAGs and NH<sub>3</sub> of the first reactor of lipid production phase. It is observed that acetate is consumed in the first hour, while TAGs are produced in the biomass cells, which remain constant.





*Figure 21: Concentration profiles of components for lipid production phase*

## <span id="page-36-2"></span><span id="page-36-0"></span>4.2 Control system

To develop a plant-wide control design system, it is crucial to carefully select the controlled variables. The plantwide control design refers to the structure of the controller rather than the control algorithm or control law itself. It is crucial to ensure that each controlled variable is sensitive to its corresponding manipulated variable.

In the syngas fermentation process, the controlled variable is the acetate concentration which can be manipulated by adjusting the agitation speed. In acetate fermentation, the lipid and acetate concentrations are controlled by adjusting the biomass concentration via operational parameters of the growth phase such as agitation speed, C/N ratio and oxygen flowrate. The control system for the growth phase is not executed in this deliverable.

For each fermentation process, a PID controller with closed-loop feedback control is created and tuned. In industrial applications, PID controller tuning is often done empirically [21]. The model is then utilized to analyze the impacts of varying syngas composition (molar concentrations of CO,  $H_2$  and CO<sub>2</sub>) in syngas fermentation and liquid flow rate and acetate concentration in acetate fermentation, on the manipulated variable.

Following, in paragraphs 4.2.1 and 4.2.2, the control system for the two fermentation processes is described in detail.

## <span id="page-36-1"></span>4.2.1 Syngas Fermentation

### 4.2.1.1 Description and validation

To develop the control system for the syngas fermentation process, it must be proved that the controlled variable, acetate concentration, is influenced by the manipulated variable, agitation speed. To accomplish this, several agitation speeds are set in the model, and the acetate concentration from first reactor is retrieved. [Figure 22](#page-37-0) depicts the sensitivity of acetate concentration with time to agitation speed.





*Figure 22: Diagram of acetate concentration profile for different agitation speeds*

<span id="page-37-0"></span>Subsequently, the control system is developed by incorporating a PID (Proportional-Integral-Derivative) controller with feedback control into the Simulink model, as illustrated in [Figure 23.](#page-37-1)



*Figure 23: Control system for syngas fermentation*

<span id="page-37-1"></span>The PID controller operates by calculating an error signal, which is the difference between the desired set point of acetate concentration (0.0164 mol/L) and the actual output of the system. This error signal is then used to adjust the agitation speed to the system in order to minimize the error and keep the system operating at the desired set





point. The PID controller operates in discrete time, which means that the control signal is updated only at these discrete points in time.

To tune the PID parameters  $(K_p, T_i, T_d)$ , a trial-and-error method is used in each case until the controller achieves a steady state. To validate the PID controller for the base case previously described, the agitation speed is adjusted to 25 rpm to achieve the desired acetate concentration, as illustrated in [Figure 24.](#page-38-0)





<span id="page-38-0"></span>[Figure 24](#page-38-0) shows that after 5 hours, the acetate concentration is fixed to the set point, as the agitation speed remains constant at 25 rpm. Therefore, the control system is deemed valid.

### 4.2.1.2 Results for different cases

<span id="page-38-1"></span>The model is then utilized to analyze the impacts of varying syngas composition (molar concentrations of CO,  $H_2$ and CO2) on the agitation speed. The studied cases are described in [Table 15.](#page-38-1)





Analyzing [Table 15,](#page-38-1) case A represents the initial syngas composition. In cases B and C, the CO molar concentration increases and decreases by 50%, respectively, while the  $H_2$  and  $CO_2$  molar concentrations remain constants. In





cases D and E, H<sub>2</sub> molar concentration increases and decreases by 50% and 40%, while the CO and CO<sub>2</sub> molar concentrations remain constants. In F case, CO<sub>2</sub> molar concentration increase by 50%. Finally, in case H, all molar concentrations increase by 50%.

By inserting each variation through a step input in the controlled model, the agitation speed is adjusted to maintain the acetate concentration at the desired point of 0.0164 mol/L, after the first reactor. The results of the agitation speeds for all cases and the  $K_p$ , T<sub>I</sub>, T<sub>D</sub> coefficients of each PID controller are shown in [Table 16.](#page-39-0)

<span id="page-39-0"></span>

#### *Table 16: PID coefficients and adjusted agitation speed in different cases*

From [Table 16,](#page-39-0) it can be observed that varying the CO and H<sup>2</sup> molar concentrations (Cases B-E) results in changes in agitation speed, indicating that CO and H<sub>2</sub> have an influence on the final concentration of acetate. Conversely, CO<sub>2</sub> (Case F) does not significantly impact the final acetate concentration, as the agitation speed of the controlled system remains constant. Upon analyzing the results of agitation speed from the PID controller for cases B and C, it is noted that agitation speed exhibits a small change in response to CO variations. However, when varying the  $H_2$ molar concentration (cases D and E), the agitation speed changes considerably, highlighting its importance.

The dynamic responses of reactants and acetate, as well as the responses of the PID controller for cases B and D, are illustrated in [Figure 25](#page-39-1) and [Figure 26,](#page-40-0) respectively.



<span id="page-39-1"></span>*Figure 25: Diagram of agitation speed and concentration of components to time for controlled system of case B*





<span id="page-40-0"></span>*Figure 26: Diagram of agitation speed and concentration of components to time for controlled system of case D*

[Figure 25](#page-39-1) and [Figure 26](#page-40-0) illustrate the dynamic responses of the concentrations of the components by varying the syngas composition in the first hour. Additionally, the figures display the agitation speed at every discrete time for 10 hours. In both cases, the controller reaches the set point after 7 hours.

As previously mentioned, cases D and E are the two with the more obvious variation. After implementing the PID controller in the dynamic simulation, the comparison between the dynamic response of the system without the agitation speed adjustment (uncontrolled) and the controlled system is illustrated in [Figure 27](#page-40-1) and [Figure 28](#page-41-1) for cases D and E, respectively.



<span id="page-40-1"></span>*Figure 27: Dynamic response of acetate concentration for case D, comparing the controlled and uncontrolled system*





<span id="page-41-1"></span>*Figure 28: Dynamic response of acetate concentration for case E, comparing the controlled and uncontrolled system*

These visualizations further emphasize the significant impact of varying the H<sub>2</sub> molar concentration on the acetate concentration and highlight the importance of the control system in maintaining the desired acetate concentration by adjusting the agitation speed to avoid accumulation. The delay of the controlled system in achieving a steady state condition is due to the controller and it could be decreased by changing its coefficients.

## <span id="page-41-0"></span>4.2.2 Acetate fermentation

### 4.2.2.1 Description and validation

As it was previously described, the control system is developed to adjust the biomass concentration -through manipulating the operational parameters of biomass growth phase- due to variations in liquid flow rate or acetate concentration. To develop the control system for the lipid production phase of acetate fermentation process, it must be proved that the controlled variable, the final acetate concentration, is influenced by the manipulated variable, biomass concentration. To accomplish this, several initial biomass concentrations are set in the model, and the final acetate concentration from first reactor is retrieved. The different studied cases are the biomass concentration for the fitted model (Case 1), 30% over Case 1 (Case 2), and 30% less than Case 1 (Case 3). [Table 17](#page-41-2) depicts the sensitivity of acetate concentration when varying the biomass concentration.



<span id="page-41-2"></span>

These results show the final acetate concentrations achieved for the different cases. It is also noteworthy that in all cases, the TAGs concentration remains constant at 0.24 g/L. Subsequently, the control system is developed by incorporating a PID (Proportional-Integral-Derivative) controller with feedback control into the Simulink model, as illustrated in [Figure 29.](#page-42-0)







*Figure 29: Control system for lipid production phase of lipid fermentation*

<span id="page-42-0"></span>The PID controller operates by calculating an error signal, which is the difference between the desired set point of acetate concentration at 0.06 g/L and the actual output of the system. This error signal is then used to adjust biomass concentration via biomass growth operational parameters to the system in order to minimize the error and keep the system operating at the desired set point. The PID controller operates in discrete time, which means that the control signal is updated only at these discrete points in time.

To tune the PID parameters  $(K_p, K_l, K_d)$ , a trial-and-error method is used until the controller achieves a steady state. To validate the PID controller for the base case previously described, the biomass concentration is adjusted to 1.4 g/L to achieve the desired acetate concentration, as illustrated in [Figure 30.](#page-43-0)





*Figure 30: Diagram of biomass concentration to time*

<span id="page-43-0"></span>

*Figure 31: Diagram of final acetate and TAGs concentration to time*

<span id="page-43-1"></span>[Figure 30](#page-43-0) and [Figure 31](#page-43-1) show that after an hour, the final acetate concentration is fixed to the set point, as the biomass concentration remains constant at 1.4 g/L. Therefore, the control system is deemed valid.

### 4.2.2.2 Results for different cases

The model is then utilized to analyze the impacts of varying liquid flow rates and acetate concentration on the final acetate concentration. The second variation would occur in case of difficulty in controlling syngas fermentation. The studied cases are described in [Table 18.](#page-44-0)

Analyzing [Table 18,](#page-44-0) Case A represents the initial liquid flow rate and acetate concentration. In cases B and C, the liquid flow rate decreases and increases by 20%, respectively. In cases D and E, the acetate concentration decreases and increases by 20%, respectively.





<span id="page-44-0"></span>

#### *Table 18: Different cases of liquid flow rate*

By inserting each variation through a step input in the controlled model, the biomass concentration is adjusted to maintain the final acetate concentration at the desired point of 0.06 g/L, after the first reactor. The results of the biomass concentration for all cases and the  $K_p$ ,  $T_l$ ,  $T_D$  coefficients of each PID controller are shown in [Table 19.](#page-44-1)

*Table 19: Adjusted biomass concentration in different cases*

<span id="page-44-1"></span>

Based on the data presented in [Table 19,](#page-44-1) it is evident that an increase in the liquid flow rate or in acetate concentration corresponds to a higher biomass concentration. Conversely, a decrease in the liquid flow rate is associated with a lower biomass concentration.



<span id="page-44-2"></span>*Figure 32: Diagram of biomass concentration and TAGs and acetate concentration to time for controlled system of case C*





<span id="page-45-0"></span>*Figure 33: Diagram of biomass concentration and TAGs and acetate concentration to time for controlled system of case E*

[Figure 32](#page-44-2) and [Figure 33](#page-45-0) illustrate the dynamic response of the acetate and TAGs concentrations by varying the liquid flow rate or the acetate concentration in the first hour. Additionally, the figure displays the biomass at every discrete time for 10 hours. In both cases, the PID controllers achieve the desired set point in an hour.

After implementing the PID controller in the dynamic simulation, the comparison between the dynamic response of the system without the biomass adjustment (uncontrolled) and the controlled system is illustrated i[n Figure 34](#page-45-1) and [Figure 35](#page-46-0) for cases C and E, respectively.



<span id="page-45-1"></span>*Figure 34: Dynamic response of acetate concentration for case C, comparing the controlled and uncontrolled system*





<span id="page-46-0"></span>*Figure 35: Dynamic response of acetate concentration for case E, comparing the controlled and uncontrolled system*

These visualizations further emphasize the significant impact of different liquid flow rates or acetate concentrations on the biomass concentrations and highlight the importance of the control system in maintaining the desired final acetate concentration.





# <span id="page-47-0"></span>5 Thermocatalytic part

## <span id="page-47-1"></span>5.1 Steady state model and export to Aspen Dynamics

An new Aspen model had to be developed because the model that was built and used for the simulation runs in Task 2.5 and Task 6.2 could not be exported to Aspen Dynamics and be the basis for the dynamic simulation analysis.

## <span id="page-47-2"></span>5.1.1 Model description

[Figure 36](#page-49-0) shows the Aspen model for the thermocatalytic part. As in the respective pilot unit at Task 5.4, the hydroprocessing is carried out in two reactors. In the first one, the conversion reactions of triglycerides into paraffins take place whereas in the second reactor, the isomerization and hydrocracking reactions occur. Since no CO or CO<sup>2</sup> were detected in gas analysis at the pilot campaign (D5.4 [22] see [Table 25\)](#page-50-1) the decarboxylation and decarbonylation reactions are discarded assuming that they do not take place in the reactor.

<span id="page-47-3"></span>

#### <span id="page-47-5"></span><span id="page-47-4"></span>*Table 20. Initial reactions at 1st reactor*

The modeling of the first reactor is split into two parts. In the first one (HDR-01), the initial reactions of triglycerides decomposition (depropanation) are considered assuming total conversion into their respective free fatty acids, using a stoichiometric reactor (RSTOIC). Moreover, the linoleic acid ( $C_{18}H_{32}O_2$ ) is converted into oleic ( $C_{18}H_{34}O_2$ ) and stearic acid (C<sub>18</sub>H<sub>36</sub>O<sub>2</sub>) through hydrogenation reaction (see [Table 20,](#page-47-3) [R 22](#page-47-4) - [R 27\)](#page-47-5). In the second part (HDR-02), the hydrodeoxygenation (HDO) reactions of the fatty acids are considered. This reactor is modelled as a plug flow reactor (RPLUG). The kinetic models for stearic acid [\(R 28](#page-48-0)[-R 30\)](#page-48-1), oleic acid [\(R 31,](#page-48-2) [R 32\)](#page-48-3) and palmitic acid [\(R 33\)](#page-48-4) HDO reactions were obtained from [23], [24] and [25] respectively but the kinetic constants were fine-tuned where needed in order the product yields to reach an agreement with the respective experimental data.





#### <span id="page-48-4"></span><span id="page-48-3"></span><span id="page-48-2"></span><span id="page-48-1"></span><span id="page-48-0"></span>*Table 21. HDO reactions kinetics parameters*

 $1$  Molar concentrations are expressed in kmol/m $3$  and partial pressures in kPa

For the hydrocracker reactor modeling (HCR) a stoichiometric reactor is considered as there are no available kinetic data from the lab/pilot experiments or from the literature. The fractional conversion of each reaction was fine-tuned accordingly in order the product yields to reach an agreement with the respective measured fuels yields (see [Table](#page-48-5)  [22\)](#page-48-5). Moreover, as there was no relevant information about the produced isomers and their yields, no isomerization reaction is considered.



<span id="page-48-5"></span>





*Figure 36. Aspen Plus flowsheet for TAGs hydroprocessing and refining*

<span id="page-49-0"></span>The fuels products separation and refining unit was inspired by [26] considering two distillation columns. Before them, the water is separated from the bio-crude in a decanter. At the first column, the light gas mainly consisting in propane is retrieved from the top. The naphtha, jet fuel and diesel fractions are separated in the second column. The lighter fractions naphtha and kerosene are obtained as gaseous and liquid distillates at the condenser, respectively, setting a condenser temperature of 180°C.

The Peng-Robinson property method was selected for the hydroprocessing reactor part, the gas recycling and gas/liquid products separation, while the Non-random two-liquid model for liquid activity coefficients calculations combined with Redlich-Kwong Soave Equation of state with Henry's law (NRTL-RK) was used for the refining part at the two distillation columns. Especially for the first column, the convergence option of Petroleum/Wide-boiling was selected as it was observed that better results were achieved.

<span id="page-49-1"></span>

#### *Table 23. TAGs stream specifications*

The characteristics of the microbial oil feed stream were obtained from D6.2 [2] and are presented in [Table 23](#page-49-1) while the basic specifications for the overall unit are summarized in [Table 24.](#page-50-2) The reactors operating parameters were obtained from D5.4 [22]. For the separate retrieve of medium and heavy kero, the heavy fraction is obtained from stage 20 in liquid form with a flow ratio of 0.24 (mass basis). The reactor dimensions and designs that are taken into consideration at the HDR-02 model are L=0.954m, D=27.94mm and 0.326 kg catalyst loading (2300 kg/m<sup>3</sup>) density).





<span id="page-50-2"></span>

#### *Table 24. Thermocatalytic part process specifications (main case)*

## <span id="page-50-0"></span>5.1.2 Model validation

At first, the model validation is assessed against the pilot campaign data. The model was set according to the TRL5 reactor size and the inlet gas streams and the simulation results were compared against the respective pilot results. Graphs in [Figure 37](#page-51-1) shows the comparison of these results.



<span id="page-50-1"></span>

As seen in [Figure 37,](#page-51-1) the model is able to predict in good agreement not only the fractions of the main product streams but also the particular liquid fuel fractions after refining. Regarding the gas streams, there is an overprediction in propane and the model foresees steam at the gas stream after the flash separation, whereas it is not







detected in gas measurements. Overall, the model enable us to proceed with the scale up and the dynamic simulations.

*Figure 37. Comparison of measured and model predicted product results for pilot scale*

<span id="page-51-1"></span>For the model upscaling at full scale, given the input data of initial oil feed stream flow rate, a scale factor of 18000 is considered and applied for the increase of reactor 2 dimensions and catalyst loading.

## <span id="page-51-0"></span>5.2 Control strategy and investigated scenarios

The focus of the control strategy of this unit is given to the maintenance of the desired specifications at the reactors (H2/oil ratio at the inlet, temperature), separation vessels (flash, decanter and columns) temperatures that will secure stable and consistent fuels production and retrieval despite 'disturbances' (e.g. oil flow ramp up/down).

Dedicated PI controllers have been applied for: [\(Figure 38\)](#page-52-0)

- Maintenance of hydrogen concentration at reactor inlet (0.9728) via regulating the makeup hydrogen flow
- Maintenance of kerosene main specifications like boiling temperature (180 °C) via column 2 condenser duty regulation
- Maintenance of cooling steam temperature outlet from the reactors (400  $^{\circ}$ C) via flow control





*Figure 38. APD oil hydroprocessing and main controllers setup*

<span id="page-52-0"></span>The following table summarizes the scenarios and the respective control strategies that are going to be investigated at dynamic/unsteady conditions:





The fist scenario is associated with the disturbance that plant operation load drops below its design point and the reduced amount of microbial oil enters the hydrotreatment unit. In the second one, the ability of the unit to operate flexibly when the operator wish to modify (increase) the diesel/jet fuel ratio, by reducing the amount of hydrocarbons that undergo hydrocracking and hydroisomerization.



## <span id="page-53-0"></span>5.3 Dynamic process simulations

## <span id="page-53-1"></span>5.3.1 TAGs flow rate variation

As seen i[n Figure 39a](#page-53-2), the crude hydrotreated oil follows the trend of oil flow change, and the stability in the updated flow rate is restored in a very short time (a few minutes) after each step change. This fact verifies that the reactors system and the gas/liquid separation unit (including water removal) have a very robust control system that restores very quickly at the desired conditions according to the set points. Interestingly, the hydrogen supply operate in an effective way and the total hydrogen flow rate at the reactor inlet has similar behavior with the produced bio-crude [\(Figure 39b](#page-53-2)). The reactors temperatures are not perfectly closed to the set point temperatures (340  $\degree$ C) as depicted in [Figure 39c](#page-53-2) but no considerable deviation is detected. On the other hand, the control system in Column 2 fail to keep it at a desired operation framework and the jet fuel production is eliminated after the first step change [\(Figure](#page-53-2)  [39a](#page-53-2)). The jet fuel steam is appeared again at t=4.5 h but could not reach a stable condition again as the initial biocrude feed drops by 20%. What is inferred is that that column can operate normally only at the design point under the given control strategy. Moreover, the temperature and pressure outlet of the jet fuel stream did not remain stable during the whole process but the jet fuel yield starts varying even when temperature and pressure were close to the respective set points.



<span id="page-53-2"></span>*Figure 39. Main process outputs behavior at sudden drops in oil flow rate*



## <span id="page-54-0"></span>5.3.2 Bypass at Hydrocracker reactor

When the bypass stream flow increases and the inlet flow at Reactor 2 drops [\(Figure 40a](#page-54-1)), the diesel yield linearly increases as less long chain hydrocarbons are broken into smaller one [\(Figure 40b](#page-54-1)). On the other hand, naphtha flow remains almost stable and the restricted hydrocracking reactions have impact merely on the jet fuel fraction. Contrary to the previous case, column 2 operates at the desired set point as concerns the jet fuel specified temperature and pressure [\(Figure 40c](#page-54-1)). It is easily concluded that the present control at the second distillation column works effectively at variable conditions provided that the inlet flow rate remains stable and at the design point. Finally, the behavior at the make-up hydrogen flow and the specified H<sub>2</sub> concentration at the reactor inlet is quite expectable and according to the initial control strategy [\(Figure 40d](#page-54-1)).



<span id="page-54-1"></span>*Figure 40. Main process outputs behavior when decreasing the inlet stream at hydrocracker (reactor 2)*



# <span id="page-55-0"></span>**6 Conclusions**

The key conclusions that are derived from the Task 6.4 activities are summarized below.

**Thermochemical part**: While the reactors temperatures maintained at the desired levels, the H<sub>2</sub>/CO increased when the feedstock flow rate decreased. Changes in the gasifier temperature can be handled and the process performance can be restored and maintained at the initial specifications through the appropriate control of the circulating solids flow rate. Variations in the water/steam flow and external gas that burnt in the oxidizer affect the stability of the syngas composition and a difference control strategy approach should be adopted in case that the different H2/CO ratio has negative impact on the gas fermentation performance and cannot be restored at the desired levels after the ATR.

**Biological Part**: In the syngas fermentation process, variations in syngas composition were considered, prompting the testing of various scenarios involving CO,  $CO<sub>2</sub>$  and  $H<sub>2</sub>$  to control the acetate production by fine-tuning agitation speed. Hydrogen variations found to notably influence agitation speed, while the impact of CO was comparatively minor. In the acetate fermentation process, fluctuations in acetate concentration and liquid flow rate are exhibited, often attributed to potential malfunctions in the syngas fermentation controllers. To counteract this, the controlled system primarily focuses on optimizing the lipid production phase. By maintaining acetate residuals close to zero, and ensuring satisfactory TAGs production, biomass concentration was adjusted accordingly. Adjusting biomass concentration produced in the growth phase is crucial for achieving desired final product outcomes and residuals. This adjustment involves optimizing operational parameters during the growth phase, such as agitation speed, oxygen flow and C/N ratio. While this task is not executed in this deliverable, it was studied that other operational parameters in the lipid production phase, such as agitation speed and oxygen flow rate exert a smaller influence on final acetate concentration compared to biomass concentration.

**Thermocatalytic part**: The fuel production and bio-crude recovery part that consists the reactors and the gas/liquid separation step has a good behaviour and stability at the process specifications when the operation load (oil flow rate) or the diesel/jet fuel ratio set-point change. The proper control of the make-up hydrogen and the temperature at both reactors has been accomplished, achieving thus a smooth operation of the reactors that led to the expected oil to hydrotreated TAGs conversion. On the other hand, the second column operated improperly at lower loads, failing to separate the bio-jet fraction and under the specified condenser temperature. More effort should be paid in order to setup a robust control of the distillation column to secure its proper operation when lower smaller bio-crude flow rate is fed. When the by-pass fraction variation at the second reactor was examined, no similar issues were observed although the feed composition varied.

# <span id="page-55-1"></span>Future work and outlook

There is still some room for improvement of the DFBG dynamic model, especially focusing on the better prediction of solids distribution along the riser and the inclusions of hydrodynamics and reactions at the oxidizer. Moreover, other parts like the loop seal the connection duct between the two reactors for an improved illustration of the hydrodynamics and solids in the system could be considered. Last but not least, the inclusion of the tars reforming at the autothermal reformer should be performed in a follow-up study to verify whether the syngas composition can be regulated there.

Integration of energy balances in both syngas and acetate fermentation processes could be executed in the models, focusing on examining the influence of cooling water supply in maintaining the reaction temperature and its impact on variations in the syngas or acetate composition. Further efforts will be directed towards refining the control of biomass concentration during the growth phase through adjustments in operational parameters, as previously



analyzed. Additionally, ongoing improvements in PID coefficients for the syngas fermentation process are anticipated, with the goal of achieving a faster steady-state condition to enhance process efficiency and stability. This future work will contribute to advancing the understanding and control of these fermentation processes for improved acetate and lipid production.

Finally, at the thermocatalytic section, the control strategy at the second distillation column should be improved when the feed stream flow rate drops. Moreover, a more detailed reactor modeling especially on the reactions associated to the hydrocracking and hydroisomerization, depending on what data are available from the respective real tests, would lead to fuel products with advanced properties. In that case, a better evaluation on their specifications would be carried out. Furthermore, an analytical modeling on the hydrogen recovery before its recirculation is scheduled as a future work.





# <span id="page-57-0"></span>**8 References**

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# <span id="page-59-0"></span>Annexes

<span id="page-59-1"></span>The gas-liquid equilibrium factors for non-condensable and condensable for components in syngas fermentation are listed i[n Table 27.](#page-59-1)



*Table 27: Gas-liquid equilibrium factors for syngas fermentation*

<span id="page-59-2"></span>The mass diffusivity of species j in water,  $D_{f,j}$  in syngas fermentation process are listed in [Table 28.](#page-59-2)



*Table 28: Mass diffusivity of components in water for syngas fermentation*

The gas-liquid equilibrium factors for components in acetate fermentation are listed i[n Table 29:](#page-60-0)



<span id="page-60-0"></span>

#### *Table 29: Gas-liquid equilibrium factors for acetate fermentation*

<span id="page-60-1"></span>The mass diffusivity of species j in water,  $D_{f,j}$  in acetate fermentation process are listed in [Table 30.](#page-60-1)

*Table 30: Mass diffusivity of components in water for acetate fermentation*



<span id="page-60-2"></span>The fitted parameters for syngas fermentation for the studied cases are shown i[n Table 31](#page-60-2) .

#### *Table 31: Fitted parameters for syngas fermentation*







The dynamic profile of biomass, acetate, CO,  $H_2$ , CO<sub>2</sub> concentrations after parameter fitting during syngas fermentation are illustrated in [Figure 41,](#page-61-0) [Figure 42](#page-62-0) and [Figure 43](#page-62-1) for the experimental data from D3.4 [16] (cleaned syngas) and T4.3 (syngas with impurities).



<span id="page-61-0"></span>*Figure 41: Dynamic profile of biomass for experimental data from D3.4* [16] *and T4.3*





<span id="page-62-0"></span>*Figure 42: Dynamic profile of acetate production for experimental data from D3.4* [16] *and T4.3*



<span id="page-62-1"></span>*Figure 43: Dynamic profile of CO, H2, CO<sup>2</sup> consumption after parameter fitting in experimental D3.4* [16] *data*

Upon analysis of [Figure 41.](#page-61-0) [Figure 42](#page-62-0) and [Figure 43](#page-62-1) it is evident that the biomass and acetate production rates vary depending on the inclusion of impurities in the initial feedstock. As for the acetate fermentation, the fitted parameters for the studied cases are shown in [Table 32.](#page-63-0)



<span id="page-63-0"></span>

#### *Table 32: Fitted parameters for acetate fermentation*

The  $K_{s1},K_{s2},K_{s3},\mu_{max1}$  parameters are the equivalent  $K_s,K_{SN},K_{SO2},\mu_{max}$  parameters used in the  $R_X\,$  equation during the lipid production phase.

The dynamic response of acetate, CDW (Cell Dry Weight) and TAGs concentrations during acetate fermentation after parameter fitting with the experimental data from D3.5 [17] is illustrated i[n Figure 44.](#page-64-0)





<span id="page-64-0"></span>*Figure 44: Predicted dynamic profiles of acetate, TAGs and CDW and experimental points for D3.5* [17]

Upon analysis of [Figure 44](#page-64-0) it is evident that the model parameters accurately describe the acetate, TAGs and CDW concentration profiles.

