

## Supplementary material

### Study of the biomass, sand, and biochar fluidization in a typical bed of fast pyrolysis - statistical analysis

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#### *S1.0 Segregation behavior*

To compare the different measures of segregation percentage, we used the analysis of variance (ANOVA) (Palkar and Shilapuram, 2017; Yusup et al., 2014). ANOVA allows the significance of all main factors and their interactions to be evaluated by comparing the mean squares against an estimate of the experimental error at a specified confidence level. Table S1 shows the degrees of freedom, the sum-of-squares, the contribution of each parameter in the prediction model, the modified sum of the squares, the modified average sum of the squares, and the P-value for each studied fluidization parameter. Alpha was set at 0.1 (confidence level of 90%). Curvature was found nonsignificant, and therefore the linear model gave a better fit for the segregation percentage.

**Table S1.** ANOVA results of segregation percentage for the linear model.

	All variables				Significant variables			
	(SS)	(df)	(MS)	P	(SS)	(df)	(MS)	P
Curvature	41.621	1	41.621	0.142115				
d <sub>sand</sub>	367.353	1	367.353	0.019724	367.353	1	367.353	0.019724
d <sub>bio</sub>	210.666	1	210.666	0.033660	210.666	1	210.666	0.033660
w <sub>bio</sub>	521.508	1	521.508	0.014016	521.508	1	521.508	0.014016
w <sub>biochar</sub>	3.765	1	3.765	0.551259				
d <sub>sand</sub> -d <sub>bio</sub>	23.641	1	23.641	0.217129				
d <sub>sand</sub> -w <sub>bio</sub>	40.185	1	40.185	0.146134				
d <sub>sand</sub> -w <sub>biochar</sub>	250.100	1	250.100	0.028578	250.100	1	250.100	0.028578
d <sub>bio</sub> -w <sub>bio</sub>	343.533	1	343.533	0.021049	343.533	1	343.533	0.021049
d <sub>bio</sub> -w <sub>biochar</sub>	45.777	1	45.777	0.131645				
w <sub>bio</sub> -w <sub>biochar</sub>	22.709	1	22.709	0.223274				
d <sub>sand</sub> -d <sub>bio</sub> -w <sub>bio</sub>	3.520	1	3.520	0.563235				
d <sub>sand</sub> -d <sub>bio</sub> -w <sub>biochar</sub>	596.207	1	596.207	0.012292	596.207	1	596.207	0.012292
d <sub>sand</sub> -w <sub>bio</sub> -w <sub>biochar</sub>	382.142	1	382.142	0.018983	382.142	1	382.142	0.018983
d <sub>bio</sub> -w <sub>bio</sub> -w <sub>biochar</sub>	5.313	1	5.313	0.487704				
Lack of Fit	208.755	1	208.755	0.33953	395.285	9	43.9205	0.153678

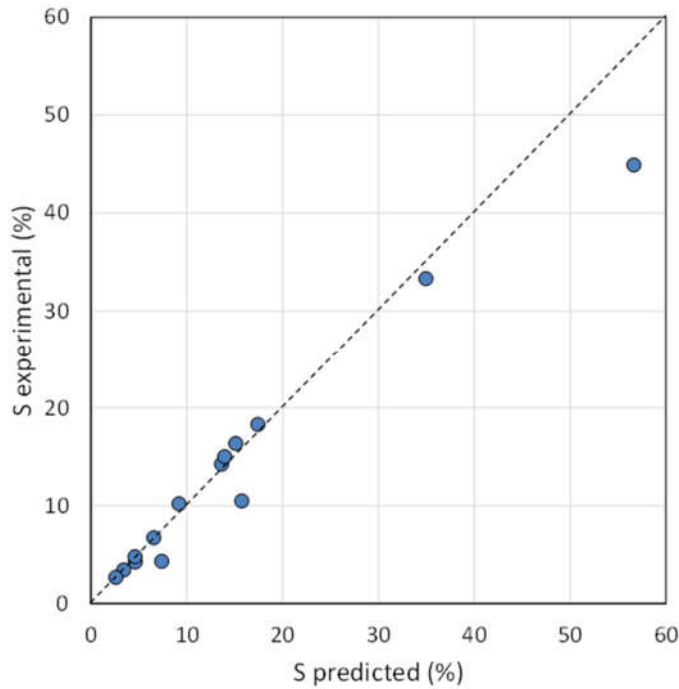
Pure Error	14.932	2	7.4659	14.932	2	7.4659
Total SS	3081.726	18		3081.726	18	
R <sup>2</sup>	0.93			0.87		
R <sup>2</sup> adj	0.56			0.78		

The main variable ( $w_{bchar}$ ), the first-order interactions ( $d_{sand-d_{bio}}$ ,  $d_{sand-w_{bio}}$ ,  $d_{bio-w_{bchar}}$ ,  $w_{bio-w_{bchar}}$ ), and second-order interaction ( $d_{sand-d_{bio-w_{bchar}}}$ ,  $d_{bio-w_{bio-w_{bchar}}}$ ) were found to be nonsignificant, and were removed from the calculation. The other response variables had P-values below 0.1. The coefficient of determination ( $R^2$ ) shows how well a change in response was explained by the model. The model better predicted the segregation percentage when  $R^2$  was close to unity, as shown in Table 5 (0.93).  $R^2$  was adjusted ( $R^2$  adj) to take account of the number of predictors in the model, allowing models with different numbers of predictors to be compared. Table S1 shows that  $R^2$  adj = 0.56 when all test variables were considered. However, the adjusted  $R^2$  remained closer to  $R^2$  when the nonsignificant variables were removed from the model ( $R^2$  adj = 0.78), suggesting that it was a better predictor of segregation percentage. Pure-error and lack-of-fit are taken account of in the residual sums of squares when testing a hypothesis. The pure-error cannot be predicted by including additional terms, whereas lack-of-fit can be predicted by including additional terms for the predictor variables. Table S1 shows a non-significant P-value ( $> 0.1$ ) for lack-of-fit. A test of lack-of-fit for the model without the additional terms was performed using the mean square pure error as the error term. We determined a mean square pure error of 7.4659, which suggests that the calculated model values had medium variability around the mean.

Measures of adjustment quality were used to assess the divergence between the observed values and the values predicted by the model. According to the calculated statistical parameters, the sensitivity analysis is sufficiently accurate. The final model used to assess the segregation percentage was the first-order polynomial regression model (Eq. (S1)):

$$S = 10.203 + 5.792 d_{sand} - 3.629 d_{bio} + 5.709 w_{bio} - 3.954 d_{sand} w_{bchar} - 4.634 d_{bio} w_{bio} + 6.001 d_{sand} d_{bio} w_{bchar} - 4.987 d_{sand} w_{bio} w_{bchar} \quad (S1)$$

The model was verified using experimental data. Fig. S1 compares the measured segregation percentage values with the values calculated using Eq. (S1). Table 2 shows the experimental segregation percentage data used for model verification, as well as the independent variable coded data used to predict the segregation percentage. The results were sufficiently close to confirm the predictive accuracy of the model.



**Fig. S1.** Experimental end predicted segregation percentage.

From Eq. (S1) the terms that most influenced the S value were  $d_{sand}$ ,  $w_{bio}$ , and the interaction between  $d_{sand}$ ,  $d_{bio}$ , and  $w_{bchar}$ . These three terms caused increases in S, though the term  $d_{bio}$  and interactions  $d_{sand}-w_{bchar}$ ,  $d_{bio}-w_{bio}$  and  $d_{sand}-w_{bio}-w_{bchar}$  have antagonistic effects on S. The effects of the main variables  $d_{sand}$ ,  $d_{bio}$  and  $w_{bio}$  are well recognized in the literature (Cluet et al., 2015; Shao et al., 2016). However, the variation of S caused by the synergy between  $d_{sand}$ ,  $d_{bio}$ ,  $w_{bio}$ , and  $w_{bchar}$  is a novel aspect that arose in this work.

### **S2.0 $U_{ff}$ behavior**

To evaluate the different measures of segregation percentage between them, the analysis of variance (ANOVA) was also used (Table S2). The main variable ( $w_{bchar}$ ) and the first-order interactions, ( $d_{sand}-d_{bio}$ ,  $d_{sand}-w_{bio}$ ,  $w_{bio}-w_{bchar}$ ) were nonsignificant, and were removed from the calculation. The other response variables had P-values below 0.1. In order to obtain a model that better predicted the final fluidization velocity,  $R^2$  should be close to unity. From Table S2, it can be observed that when all the variables are considered,  $R^2$  is 0.90, but the adjusted  $R^2$  is 0.77. However, when the nonsignificant variables were removed from the model, the adjusted  $R^2$  (0.81) gets closer to  $R^2$  (0.89), suggesting that it was a better

predictor of final fluidization velocity. Table S2 shows a non-significant P-value ( $> 0.1$ ) for lack-of-fit. Mean square pure error of 0.000867 suggests that the calculated model values had low variability around the mean.

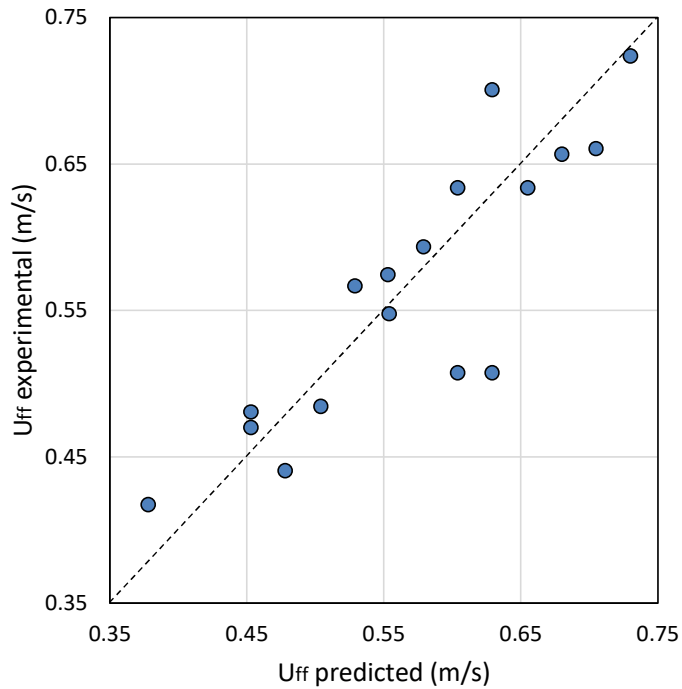
**Table S2.** ANOVA results of final fluidization velocity for the quadratic model.

	All variables				Significant variables			
	(SS)	(df)	(MS)	P	(SS)	(df)	(MS)	P
$d_{\text{sand}}$	0.088573	1	0.088573	0.009647	0.088573	1	0.088573	0.009647
$d_{\text{bio}}$	0.017822	1	0.017822	0.045364	0.017822	1	0.017822	0.045364
$w_{\text{bio}}$	0.026934	1	0.026934	0.030714	0.026934	1	0.026934	0.030714
$w_{\text{bchar}}$	0.000938	1	0.000938	0.407610				
$d_{\text{sand}}^2$	0.013445	1	0.013445	0.058851	0.013445	1	0.013445	0.058851
$d_{\text{bio}}^2$	0.030368	1	0.030368	0.027383	0.030368	1	0.030368	0.027383
$w_{\text{bio}}^2$	0.007584	1	0.007584	0.097836	0.007584	1	0.007584	0.097836
$w_{\text{bchar}}^2$	0.007584	1	0.007584	0.097836	0.007584	1	0.007584	0.097836
$d_{\text{sand}}$ by $d_{\text{bio}}$	0.000600	1	0.000600	0.492901				
$d_{\text{sand}}$ by $w_{\text{bio}}$	0.007832	1	0.007832	0.095159				
$d_{\text{sand}}$ by $w_{\text{bchar}}$	0.010201	1	0.010201	0.075493	0.010201	1	0.010201	0.075493
$d_{\text{bio}}$ by $w_{\text{bio}}$	0.003969	1	0.003969	0.165764	0.003969	1	0.003969	0.165764
$d_{\text{bio}}$ by $w_{\text{bchar}}$	0.000600	1	0.000600	0.492901	0.007832	1	0.007832	0.095159
$w_{\text{bio}}$ by $w_{\text{bchar}}$	0.000156	1	0.000156	0.712492				
Lack of Fit	0.020568	10	0.002057	0.332822	0.022862	14	0.001633	0.400552
Pure Error	0.001734	2	0.000867		0.001734	2	0.000867	
Total SS	0.214484	26			0.214484	26		
$R^2$	0.90				0.89			
$R^2$ adj	0.77				0.81			

Measures of adjustment quality were also used to assess the divergence between the observed values and the values predicted by the model. The final model used to assess the final fluidization velocity was the second-order polynomial regression model (Eq. (S2)):

$$U_{\text{ff}} = 0.470 + 0.061 d_{\text{sand}} + 0.027 d_{\text{bio}} + 0.033 w_{\text{bio}} - 0.025 d_{\text{sand}} w_{\text{bchar}} + 0.016 d_{\text{bio}} w_{\text{bio}} - 0.022 d_{\text{sand}} w_{\text{bio}} + 0.025 d_{\text{sand}}^2 + 0.038 d_{\text{bio}}^2 + 0.019 w_{\text{bio}}^2 + 0.019 w_{\text{bchar}}^2 \quad (\text{S2})$$

Fig. S2 compares the measured final fluidization velocity values with the values calculated by Eq. (S2). Table 2 shows the experimental final fluidization velocity data used for model verification, as well as the independent variable coded data used to predict the final fluidization velocity. The results were sufficiently close to confirm the predictive accuracy of the model.



**Fig. S2.** Experimental end predicted segregation percentage.

From Eq. (S2), the term  $d_{\text{sand}}$  and  $d_{\text{bio}}^2$  were the ones that most influenced the final fluidization velocity, while  $d_{\text{bio}}$  was the least sensitive. Most of the terms increased the final fluidization velocity. However, the interactions  $d_{\text{sand}}-w_{\text{bchar}}$  and  $d_{\text{sand}}-w_{\text{bio}}$  have the opposite effect. The effects of the main variables  $d_{\text{sand}}$ ,  $d_{\text{bio}}$  and  $w_{\text{bio}}$  are well recognized in the literature (Vasconcelos et al., 2018). Nevertheless, the variation of final fluidization velocity caused by the synergy between  $d_{\text{sand}}$ ,  $d_{\text{bio}}$ ,  $w_{\text{bio}}$ , and  $w_{\text{bchar}}$  is a novel aspect that arose in this work.

## References

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