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RESEARCH ARTICLE

Corncob as an effective, eco-friendly, and economic biosorbent for removing the azo dye Direct Yellow 27 from aqueous solutions

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Abstract

The corncob is an agricultural waste generated in huge quantities during corn processing. In this paper, we tested the capacity of corncob particles for water purification by removing the azo dye Direct Yellow 27 (DY27) via biosorption. The biosorption process was investigated in terms of the kinetics, equilibria, and thermodynamics. Batch biosorption studies showed that the biosorption performance has strong inverse correlations to the solution pH and the corncob particle size, and it increases quickly with increasing contact time and initial dye concentration. The pseudo-second-order kinetic model provides the best fit to the experimental data, whereas the Redlich-Peterson isotherm model is most suitable for describing the observed equilibrium biosorption. The biosorption process is exothermic, spontaneous, and physisorption in character. Fourier transform infrared (FTIR) spectroscopy and confocal scanning laser microscopy (CSLM) studies suggest that lignocellulose and proteins play key roles in the biosorption of DY27 from aqueous solutions by corncob. Furthermore, after biosorption onto the corncob, the dye can be effectively desorbed using 0.1 M NaOH solution. Therefore, the corncob can be used as a promising biosorbent to remediate DY27-contaminated water and wastewater.

Introduction

Synthetic organic dyes are extensively used in many technologies, such as textiles, printing, paper making, leather tanning, rubber, food processing, plastics, cosmetics, hair coloring, pharmaceuticals, photography, agricultural research, light-harvesting arrays, and photoelectrochemical cells [1–3]. Currently, there are estimated to be more than 100,000 commercially available dyes with a total production rate of more than $0.7-1 \times 10^6$ tons per year [1].

Approximately 12% of the synthetic dyes produced annually in the world are believed to be lost during manufacturing and processing operations, and about 20% of the lost dyes enter

industrial wastewaters [4]. Such wastewaters are often discharged directly into natural water bodies without treatment. This has caused considerable adverse impacts on the environment and human health [2, 5]. For example, even a very small amount of dye (less than 1 mg L⁻¹) can markedly color the water, making it unfit for human consumption [6, 7]. Such colored water also has reduced sunlight penetration, which lowers photosynthetic activities in the water and increases the heterotrophic activity that depletes dissolved oxygen [8]. Furthermore, dyes may also cause eutrophication; prevent re-oxygenation of water bodies; pollute ground water; and be toxic to aquatic plants, microorganisms, fish, and mammals [9]. Especially, the impurities and (bio)transformation products of certain synthetic dyes directly affect human health due to their acute and/or chronic toxicity; allergenic, mutagenic, carcinogenic, genotoxic, cytotoxic, estrogenic, and anti-estrogenic activities; and immune suppression effects [9– 12].

Many chemical, physical, and biological technologies have been developed to remove dyes from aqueous solutions. They include: coagulation-flocculation, precipitation-flocculation, electrokinetic coagulation, electrocoagulation-electroflotation, photocatalysis, oxidation, advanced oxidation, electrochemical treatment, adsorption on activated carbon, membrane separation, ion exchange, irradiation, incineration, biodegradation, and decolorization by bacterial, white-rot fungal, or mixed cultures [1, 7, 13]. However, each of these technologies has drawbacks, such as operational problems, high cost, and the generation of secondary wastes that could be even more toxic and/or difficult and expensive to remove/degrade [14–16]. In comparison, biosorption has been considered a promising technology for the removal of dyes from aqueous solutions, owing to its higher treatment efficiency and efficacy, faster removal rate, high flexibility, and low capital investment and operating costs. This method is also easy to design, implement, and operate. Furthermore, being an environmentally friendly method, it minimizes secondary waste production and allows possible dye recovery and biosorbent regeneration afterwards [15, 17].

Agricultural wastes have attracted significant attention for biosorption of dyes from aqueous solutions. Many of these materials display excellent performance, low or zero cost, natural abundance, high availability, eco-friendliness, and renewability. The functional groups on their surface could bind to many classes of dyes with selectivity [1, 6, 18]. Corn is one of the most widely planted crops in the world. During the processing of corn, a large volume of corncobs is generated as agricultural waste [19, 20]. The amount of corncobs generated worldwide is approximately 144 million tons per year [21], and most of it is discarded or burnt, causing serious environmental pollution and no benefit [22, 23]. Considering this, corncobs may be used as a value-added biosorbent for the removal of toxic organic and inorganic pollutants from wastewaters [22, 24].

In the present work, we assess the potential of corncobs to remove the Direct Yellow 27 (DY27), which is a monoazo direct dye with good light fastness [25]. DY27 is extensively used for the dyeing and printing of paper, leather, as well as of cotton, nylon, rayon, and viscose fabrics [3]. Consequently, DY27 has been found in many industrial wastes in aqueous environments [26]. The biosorbent made from corncobs was characterized by its chemical composition, pH at the point of zero charge (pH_{pzc}), Fourier transform infrared (FTIR) spectroscopy, scanning electron microscopy (SEM), and confocal scanning laser microscopy (CSLM). The influence of different process parameters (such as solution pH, corncob particle size, shaking contact time, and initial DY27 concentration) on the kinetics of DY27 biosorption from aqueous solutions by corncob was studied. Additionally, the isotherms and thermodynamics of DY27 biosorption were examined. Finally, the optimal elute for DY27 desorption from the used biosorbent and the associated kinetics were also discussed. It is worth it to mention that no previous works have addressed the biosorption of DY27 from aqueous solution.

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Materials and methods

Ethics statement

No specific permits were required for the corncob sample collection for this study. We used corncob material that came from the private lands from one of the authors (NKBV). No material was taken from protected land or National Parks. The studies conducted in this work did not involve any endangered or protected species.

Chemicals

DY27 dye (molecular formula: $C_{25}H_{20}N_4Na_2O_9S_3$, molecular weight: 662.62 g mol⁻¹, Fig.1) [27] was purchased from Sigma-Aldrich Chemicals. All other reagents were of analytical grade. A DY27 stock solution of 2 g L⁻¹ was prepared by dissolving an accurately weighed amount of DY27 in distilled deionized water. Test solutions of DY27 were prepared by diluting the stock solution with distilled deionized water. In this work, the initial DY27 concentration (C_o) varied from 7 to 293 mg L⁻¹, and the pH of each solution was adjusted to the desired value using 0.1 M HCl or NaOH solutions.

Biosorbent preparation

Corncobs (*Zea mays* L.) were collected in the municipality of Villa del Carbón (Latitude: 19° 43' 41"N; Longitude: 99° 28' 28"W), State of Mexico, Mexico. The collected corncobs were first washed thoroughly with distilled deionized water, and then oven-dried at 60°C until the weight was constant. The dried samples were subsequently ground in a hammer mill (Glen Creston Ltd.), and the resulting particles were screened using ASTM standard sieves to obtain fractions with different particle sizes. The sieved fractions were stored in airtight plastic containers until use.

Proximate chemical composition of corncob

The biosorbent's proximate chemical composition was analyzed in triplicate, following methods described by the Association of Official Analytical Chemists [28]. The total protein (*TP*) was analyzed by the Kjeldahl method, using 6.25 as the conversion factor from total nitrogen to total protein. Ash (*A*) was determined by incineration of samples in a muffle furnace at 600°C until the weight was constant. Ether extract (*EE*) was determined by the Soxhlet method. Crude fiber (*CF*) was measured as the loss on ignition of dried lipid-free residues



Fig 1. Chemical structure of Direct Yellow 27.

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after digestion with H_2SO_4 and NaOH standard solutions. Nitrogen-free extract (*NFE*) was calculated as the difference between dry matter and the sum of total protein, ether extract, ash, and crude fiber (Eq 1):

$$\% NFE = 100 - \% A - \% TP - \% EE - \% CF$$
(1)

Determination of point of zero charge

The biosorbent's point of zero charge (pH_{pzc}) is defined as the pH at which the amounts of positive and negative charges on its biosorbent surface are equal, leading to a neutral net charge [29]. In the present work, pH_{pzc} was determined using a batch equilibrium method, following the procedures described by Lim et al. [30]. Briefly, corncob samples (1 g L⁻¹) were mixed with 0.1 M KNO₃ solutions which were previously adjusted to initial pH (pH_i) values of 2, 4, 6, 8, 10, and 12 by the addition of 0.1 M HNO₃ and/or 0.1 M NaOH. The suspensions were agitated continuously at 150 rpm at $18 \pm 1^{\circ}$ C for 48 h to ensure that an equilibrium pH was reached in each case. Subsequently, the suspensions were filtered through filter paper (Whatman, grade 42), and the final pH (pH_f) of each filtrate was recorded. A graph of the pH change in the KNO₃ solution (Δ pH = pH_i-pH_f) against pH_i was then constructed to determine the pH_{pzc} of corncob.

Batch kinetic experiments for DY27 biosorption

Batch kinetic biosorption studies were conducted to evaluate the effect of relevant operational parameters, such as solution pH, corncob particle size, shaking contact time, and C_o on the biosorption of DY27 by corncob. All experiments were performed in 500-mL Erlenmeyer flasks, using 100 mL of DY27 solution of known pH and dye concentration and 1 g (dry weight) L⁻¹ of biosorbent. During each experiment, the pH of the solution was maintained at a constant value (± 0.1 pH unit) by periodic checks and adjustments with 0.1 M HCl or NaOH solution. The flasks were placed in a shaker bath (Cole Parmer®) at a constant shaking speed of 150 rpm for 72 h.

The influence of solution pH was examined at $C_o = 50 \text{ mg L}^{-1}$ at different pH values (1.5, 2.0, 2.5, 3.0, 4.0, 5.0, and 6.0) at 18 ± 1 °C. The effect of particle size was explored in DY27 solution at $C_o = 50 \text{ mg L}^{-1}$, pH 1.5, and with different corncob particle sizes (0.297–0.50, 0.50–0.80, 0.8–1.0, 1.0–1.18, 1.18–1.41, 1.41–1.70, and 1.70–2.0 mm) at 18 ± 1 °C. The influence of shaking contact time was investigated for 0, 0.1, 0.25, 0.5, 0.75, 1.0, 1.5, 2.0, 2.5, 3.0, 4.0, 5.0, 6.0, 7.0, 24, 48, and 72 h in solutions with different initial dye concentrations (7, 13, 28, 50, 74, 85, 100, 145, and 293 mg L⁻¹) at pH 1.5 and 18 ± 1 °C. These C_o values were also used to examine the kinetic biosorption profiles at pH 1.5 and 18 ± 1 °C.

Corncob-free controls were used to evaluate the loss of DY27 dye arising from precipitation and/or adsorption onto glass wall.

For the kinetic studies, after certain contact times, samples were taken from the mixture and centrifuged at 5000 rpm for 10 min. The DY27 concentration in the supernatants was analyzed by spectrophotometry (Thermo Scientific^{**} Evolution 201) at 398 nm. The time-dependent DY27 biosorption capacity (q_p , mg g⁻¹) was estimated using the following mass balance relationship:

$$q_t = \frac{(C_0 - C_t)}{M} \tag{2}$$

Here, C_o and C_t are the initial and residual DY27 concentrations (mg L⁻¹) at time t = 0 and t (h), respectively, and M is the corncob concentration (g L⁻¹).

Isotherm studies of DY27 biosorption

For DY27 equilibrium biosorption, corncob biomass was added at 1 g (dry weight) L^{-1} to DY27 solutions at pH = 1.5 at $C_o = 7-293$ mg L^{-1} and different temperatures (18, 35, 50, and 60°C), with a constant agitation of 150 rpm for 72 h to ensure that biosorption equilibrium was reached. Afterwards, samples were taken, centrifuged at 5000 rpm for 10 min, and then the dye concentrations in the supernatants were measured. The DY27 equilibrium biosorption capacity ($q_{e exp}$) was calculated using Eq (2), by replacing C_t with C_e (the equilibrium concentration of DY27 in the aqueous solution in mg L^{-1}).

Kinetic modeling of DY27 biosorption

Modeling the kinetics of target pollutant removal is crucial for understanding relevant aspects of the biosorption process, such as the biosorption rate and its influencing factors, designing effective and efficient biosorption systems, selecting operation conditions for full-scale batch process, and establishing the time dependence of biosorption systems under different process conditions [31, 32]. In the present work, the dynamics of DY27 biosorption onto corncob biomass was examined at different solution pH levels, corncob particle sizes, and C_o . The data were analyzed using five different kinetic models: pseudo-first-order, pseudo-second-order, Elovich, intraparticle diffusion, and fractional power (Table 1).

Modeling of DY27 biosorption equilibrium isotherms

Equilibrium biosorption isotherm models are useful for describing the interactions between the adsorbate and biosorbent, and therefore they are essential for the design of biosorption facilities on a large scale [32]. Various two-parameter (Langmuir, Freundlich, Temkin, Halsey, and Dubinin-Radushkevich) and three-parameter (Sips, Toth, Radke-Prausnitz, and Redlich-Peterson) isotherm models (Table 1) were applied in the present work to fit the experimental equilibrium data of DY27 biosorption.

Thermodynamic analysis

The Gibbs free energy change (ΔG^o , J mol⁻¹), enthalpy change (ΔH^o , J mol⁻¹), and entropy change (ΔS^o , J mol⁻¹ K⁻¹) were calculated in order to describe the thermodynamic behavior of DY27 biosorption onto corncob biomass (Table 1).

Desorption studies

Corncob samples were mixed with a 293 mg L^{-1} DY27 solution (100 mL) at a ratio of 1 g L^{-1} , pH 1.5, and 18 ± 1°C with constant agitation at 150 rpm for 72 h to ensure biosorption equilibrium. Afterwards, the DY27-loaded sample was separated from the dye solution by centrifugation at 5000 rpm for 10 min, and the concentration of residual DY27 in the supernatant was measured to calculate the amount of dye biosorption. The DY27-loaded precipitate was gently washed with distilled deionized water to remove any unbiosorbed dye. After another centrifugal separation, the dye-loaded corncob was oven-dried at 60°C until a constant dry weight was obtained, and stored in an airtight plastic container until use.

All batch desorption experiments were conducted in 500-mL Erlenmeyer flasks containing 100 mL of given desorption solution and 1 g (dry weight) L^{-1} of DY27-loaded corncob. The flasks were agitated in a shaker bath (Cole-Parmer®) at 150 rpm and $18 \pm 1^{\circ}$ C.

To select the best desorption solutions, different acids (0.1 M HCl, H_2SO_4 , HNO₃, NH₄Cl, and (NH₄)₂SO₄), alkali (0.1 M NaOH, KOH, and Na₂CO₃), neutral salts (0.1 M NaCl and NaNO₃), and organic compounds (0.1 M acetone and chloroform) were mixed with

Table 1. Different kinetic, isotherm, and thermodynamic models.

		Kinetic models	
	Equation	Nomenclature	Reference
Pseudo-first-order	$\log (q_{e1} - q_t) = \log q_{e1} - \frac{k_1}{2.303} t$	q_{e1} and q_t : biosorption capacity [mg g ⁻¹] at equilibrium and at any time <i>t</i> [h], respectively; k_1 : rate constant of the model [h ⁻¹]	[33]
Pseudo-second- order	$\frac{t}{q_t} = \frac{1}{k_2 q_{e2}^2} + \frac{1}{q_{e2}} t$	q_{e2} and q_t : biosorption capacity [mg g ⁻¹] at equilibrium and at any time <i>t</i> [h], respectively; k_2 : rate constant of the model [h ⁻¹]	[34]
Intraparticle diffusion	$q_t = k_{id}t^{0.5} + c$	q_i : biosorption capacity [mg g ⁻¹] at any time t [h]; k_{id} : intraparticle diffusion rate constant [mg g ⁻¹ h ^{-0.5}]; c : model intercept	[35]
Elovich	$q_t = \frac{1}{B_E} \ln(A_E B_E) + \frac{1}{B_E} \ln t$	A_E : initial biosorption rate [mg g ⁻¹ h ⁻¹]; B_E : desorption constant of the model [g mg ⁻¹]	[16]
Fractional power	$q_t = k_{fp} t^{\nu}$	<i>v</i> : fractional power rate constant $[h^{-1}]$; k_{fp} : fractional power model constant $[mg g^{-1}]$	[16]
		Isotherm models	
Two-parameter mod	lels		
Langmuir	$q_e = \frac{q_m K_L C_e}{1 + K_L C_e}$	q_e and q_m : biosorption capacity at equilibrium and maximum biosorption capacity [mg g ⁻¹], respectively; C_e : liquid phase concentration of dye at equilibrium [mg L ⁻¹]; K_L : Langmuir constant [L mg ⁻¹]	[<u>30</u>]
Freundlich	$q_e = K_F C_e^{\frac{1}{n_F}}$	K_{F} : constant of the Freundlich model [(mg g ⁻¹)(mg L ⁻¹) ^{-1/nF}]; n_{F} : heterogeneity factor.	[16]
Temkin	$q_e = \frac{_{B_T}}{_{B_T}} \ln(A_T C_E)$	<i>T</i> : absolute temperature [K]; <i>R</i> : ideal gas constant [8.314 J mol ⁻¹ K ⁻¹]; <i>B</i> _T : constant related to heat of biosorption [J mol ⁻¹]; <i>A</i> _T : Temkin isotherm constant [L mg ⁻¹]	[35]
Halsey	$q_e = \left(K_H/C_e\right)^{1/n_H}$	K_{H} : Halsey isotherm model constant [L g ⁻¹] ^{-1/nH} ; n_{H} : Halsey model exponent	[16]
Dubinin- Radushkevich	$q_e = q_m \exp(-B_{DR} E_{DR}^2)$	B_{DR} : biosorption energy constant [mol ² J ⁻²]; E_{DR} : Polanyi potential [kJ mol ⁻¹]	[36]
Three-parameter mo	odels		
Sips	$q_e = q_m rac{(K_S C_e)^{1/n_s}}{1 + (K_S C_e)^{1/n_s}}$	K_s : Sips constant; $1/n_s$: Sips model exponent	[30]
Toth	$q_e = rac{q_m b_T Ce}{(1 + (b_T Ce)^{n_T - 1})^{n_T}}$	b_T : Toth model constant [L mg ⁻¹] ^{-1/nT} ; n_T : Toth model exponent	[36]
Redlich-Peterson	$q_e = rac{K_{RP}C_e}{1+A_{RP}C_e^{B_{RP}}}$	K_{RP} : Redlich-Peterson model isotherm constant [L g ⁻¹]; A_{RP} : Redlich-Peterson model constant [L mg ⁻¹] ^{BRP} ; B_{RP} : Redlich-Peterson model exponent	[<u>36</u>]
Radke-Prausnitz	$q_e = rac{A_R R_R C_e^{B_R}}{A_R + R_R C_e^{B_{R-1}}}$	A_R : [L g ⁻¹] and R_R [L mg ⁻¹]: Radke-Prausnitz model constants; B_R : Radke-Prausnitz model exponent	[16]
		Thermodynamic models	
Gibbs free energy change	$\Delta \mathbf{G}^{\circ} = -R T \ln K_{c}$ $K_{c} = \frac{q_{c}}{C_{e}}$	<i>T</i> : absolute temperature [K]; <i>R</i> : ideal gas constant [8.314 J mol ⁻¹ K ⁻¹]; K_c : equilibrium constant [L g ⁻¹]; $\Delta \mathbf{G}^\circ$: Gibbs free energy change [J mol ⁻¹]	[37]
Entropy change	$\partial \Delta \mathbf{G}^{\circ} = \partial T \partial \Delta \mathbf{S}^{\circ}$	<i>T</i> : absolute temperature [K]; ΔS° : entropy change [J mol ⁻¹ K ⁻¹]	[37]
Enthalpy change	$\Delta \mathbf{H}^{\circ} = \Delta \mathbf{G}^{\circ} - T \Delta \mathbf{S}^{\circ}$	ΔH° : enthalpy change [J mol ⁻¹]	[16]

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DY27-loaded corncob over a 48 h period. The best agent for DY27 desorption was used for batch desorption kinetic studies. The amount of desorbed DY27 was determined spectrophotometrically at a wavelength of 398 nm. The desorption efficiency (E_{Db} %) was estimated according to the following equation:

$$E_{Dt} = \frac{C_{Dt}}{q_e M} \times 100 \tag{3}$$

Here, C_{Dt} is the residual DY27 concentration (mg L⁻¹) in the desorption solution after time t_D , and q_e is the initial ($t_D = 0$ h) dye concentration in the corncob (mg g⁻¹), i.e., the equilibrium biosorption capacity.

Statistical data analysis

All biosorption and desorption experiments were repeated three times, and the values reported herein represent the average values. The biosorption and desorption data were statistically

analyzed by analysis of variance (ANOVA) with Tukey's test (overall confidence level = 0.05) using GraphPad Prism software (version 7.0; GraphPad Software, Inc.). The same software was also used to estimate all the biosorption model parameters by nonlinear regression analysis. The different biosorption models used for fitting were assessed using the coefficient of determination (r^2), standard deviation of residuals (*Sy.x*), absolute sum of squares (*ASS*), and Aikake information criterion (*AIC*). A high value of r^2 (close to 1.0) and small *Sy.x*, *ASS*, and *AIC* values correspond to better representation of experimental data by a particular model.

SEM analysis

To analyze the morphological features and surface characteristics of native (i.e. not loaded with DY27) and DY27-loaded corncob samples, micrographs were obtained using a high-resolution scanning electron microscope (JSM-7800F, JEOL) at an accelerating voltage of 5 kV after gold coating.

CSLM analysis

DY27-loaded corncob samples were stained with 1% fluorescein or calcofluor white M2R (Sigma-Aldrich) in order to detect proteins and cellulose, respectively. Excess dye was removed with deionized water, and the samples were dried at 60° C for 24 h. Micrographs were obtained using a multiphoton laser scanning confocal microscope (LSM710 NLO, Carl Zeiss) equipped with an argon/krypton laser with four excitation lines (405, 488, 561, and 633 nm) and powers of 100, 100, 2, and 2%. All images were acquired at 10× magnification without zooming, and saved in TIFF format at 512 × 512 pixels.

FTIR analysis

FTIR spectroscopic analysis of native and DY27-loaded corncob samples was used to identify the main functional groups on the corncob surface that may be involved in the biosorption of DY27 from aqueous solutions. For this purpose, finely ground native or DY27-loaded corncob were thoroughly mixed with dried spectroscopic KBr in a 1:3 ratio, and immediately analyzed using a Thermo Scientific Nicolet iS10 FTIR spectrometer equipped with an EasiDiff diffuse reflectance hemisphere accessory (Pike Technologies). FTIR spectra were obtained using 64 scans over the range of 4000–400 cm⁻¹ with a resolution of 1 cm⁻¹. The Kubelka-Munk function was calculated from the diffuse reflectance data using the OMNIC software (Thermo Fischer Scientific). The FTIR spectra in the present work are presented in Kubelka-Munk function vs. wavenumber.

Results and discussion

Chemical composition of corncobs

The proximate chemical composition of corncobs (dry matter basis) was found to be: total protein, $4.28 \pm 0.63\%$; total ash, $3.44 \pm 0.08\%$; ether extract, $2.03 \pm 0.22\%$; crude fiber, $26.29 \pm 0.52\%$; and nitrogen-free extract, $63.96 \pm 1.5\%$. Therefore, the corncob has low fat, protein, and mineral contents, and rich in carbohydrate and lignin contents. This is consistent with the knowledge that the main components of corncobs are cellulose, hemicellulose, and lignin, which together comprise approximately 90% of the dry matter [20, 21]. The polymeric components in corncob possess many functional groups that can be active sites for the biosorption of dyes [20].

pH_{pzc} of corncob

The solution pH strongly affects the surface charge of a biosorbent and its biosorption capacity. The surface charge can be qualitatively assessed by the point of zero charge (pH_{pzc}) or isoelectric point [2, 23]. The pH_{pzc} of corncob was determined to be 6.83 (Fig 2). Therefore, for solution pH below this value, the corncob particles carry positive charges, and vice versa. The value of 6.83 is close to that reported by Leyva-Ramos et al. [38] (pH_{pzc} = 6.2), who attributed the weak acidity of natural corncob to the slightly higher concentration of acid sites than that of the basic ones.

SEM analysis

SEM has been widely used to directly observe the surface structure and morphology of different biosorbents [18]. Fig 3 shows the SEM micrographs of DY27-unloaded (native) and DY27-loaded corncob. SEM micrographs revealed that native corncob has a rough and porous surface (Fig 3A), with pores of different shapes and with sizes exceeding 50 nm, thus indicating that corncob has a macroporous structure. Likewise, corncob has an irregular structure, which makes biosorption of DY27 dye on different parts of the biosorbent possible. From the SEM





Fig 3. SEM micrographs of native (A) and DY27-loaded (B) corncob at 1200× magnification. https://doi.org/10.1371/journal.pone.0196428.g003

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images of the native (Fig 3A) and DY27-loaded corncob (Fig 3B), there is no evidence of changes in the surface morphology of the biosorbents. Contrastingly, some studies have mentioned that the surface morphology of biosorbents may change once dyes have been loaded [18, 39].

Effect of solution pH on DY27 biosorption kinetics

No measurable changes in the DY27 dye concentration were detected in any of the assayed corncob-free controls, thus indicating that any evident DY27 removal was solely due to the biosorbent.

As mentioned earlier, the pH can affect the biosorption of dyes from aqueous solutions via changing the surface charge of biosorbents. The solution pH may also affect the kinetics of biosorption through the ionization state and solubility of dyes [2].

Fig 4 depicts the kinetic profiles of DY27 biosorption by corncob at different solution pH (1.5–6.0). Solutions with pH < 1.5 could not be assayed because of dye precipitation. Corncob was found to take up DY27 from aqueous solution at all other pH levels, although the rate and extent of biosorption varied greatly. The capacity of DY27 biosorption increased as the solution pH decreased, reaching its highest level at pH 1.5. This trend can be explained with the aid of pH_{pzc}. All the DY27 solutions assayed had pH < pH_{pzc}. Under these conditions, the surface charge of corncob is positive because the active biosorption binding sites are protonated. Such protonation favors the biosorption of anionic DY27 molecules as a result of electrostatic attraction. The lower the solution pH, the stronger the corncob surface is protonated and the greater the biosorption of DY27.

The present results clearly show that the optimal solution pH for the biosorption of DY27 onto corncob is 1.5, and this pH was therefore used in all further studies. To the best of our knowledge, no previous works have addressed the biosorption of DY27 from aqueous solution. Only a single study reported the removal of DY27 from aqueous solution using an inorganic adsorbent (crystalline hydroxyapatite), for which the optimal pH for adsorption was found to be 3.5 (the lowest pH value assayed by those authors) [26].

Influence of corncob particle size on DY27 biosorption

Fig 5 displays the effects of corncob particle size on DY27 biosorption at pH 1.5. It is apparent that the rate and capacity of biosorption increased for finer corncob particles. This trend is





attributed to the enlarged specific surface area, as well as a lower intraparticle diffusion resistance that improves accessibility of internal sites on the corncob [1, 40, 41]. Based on this result, the smallest particle size (0.297–0.50 mm) was used in further studies.

Effect of contact time and initial DY27 concentration on biosorption

From the results shown in Fig 6, it is evident that the contact time between biosorbent and DY27 solution has a significant effect on the biosorption. At all assayed C_o values, the biosorption capacity gradually increased with the experiment time, eventually reaching a maximum constant value that corresponded to the equilibrium biosorption capacity ($q_{e exp}$). After this point, the DY27 removal ceased. Furthermore, irrespective of C_o , the initial biosorption rate was quite fast because (1) there were a large number of positively charged binding sites available for biosorption and (2) the driving force to transfer DY27 molecules from the aqueous solution to the surface of corncob was high [2, 30]. Afterwards, both the number of vacant binding sites and the mass transfer driving force decreased gradually, therefore the biosorption rate decreased and eventually reached dynamic equilibrium [1, 42].

Fig 6 also shows that the DY27 equilibrium biosorption capacity increased from 6.5 to 73.00 mg g⁻¹ as C_o was increased from 7 to 145 mg L⁻¹. The higher availability of DY27 molecules in the solution increased their interactions with the corncob and the equilibrium biosorption capacity. A higher C_o also increases the DY27 concentration gradient near the

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Fig 5. Effect of corncob particle size on DY27 biosorption capacity of corncob.

particle surface, which is the thermodynamic driving force for the mass transfer of DY27 from the solution to the corncob surface. Consequently, the probability of interaction between the dye molecules and the active biosorption sites increases, and so does the DY27 biosorption capacity [16]. However, the equilibrium biosorption capacities were very similar at $C_o = 145$ and 293 mg L⁻¹, probably due to saturation of the biosorption sites.

The above results clearly indicate that there is a direct relation between the dye concentration and the available biosorption sites on the corncob surface, the latter is the limiting factor for DY27 biosorption, and that the saturation of biosorbent depends strongly on C_o [1, 16, 18].

Modeling of biosorption kinetics

The kinetics of DY27 biosorption onto corncob was analyzed using five different models at pH = 1.5–6.0, corncob particle size from 0.297–0.5 to 1.7–2.0 mm, and $C_o = 7–293$ mg L⁻¹. Tables 2–4 present the experimental equilibrium biosorption capacity ($q_{e exp}$), the kinetic parameter values (pseudo-first-order model: q_{e1} and k_1 , pseudo-second-order model: q_{e2} and k_2 , Elovich model: A_E and B_E , intraparticle diffusion model: k_{id} and c, and fractional power model: k_{fp} and v) under the different solution pH levels, corncob particle sizes, and C_o values assayed, together with the corresponding r^2 , ASS, Sy.x, and AIC values.

The r^2 values were higher with the pseudo-second-order kinetic model; and the corresponding *ASS*, *Sy.x*, and *AIC* values were lower than those obtained with the other four kinetic

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models. Therefore, the pseudo-second-order model is the most suitable for describing the experimental data obtained in this study, as shown by the solid lines in Figs 4–6. In addition, the equilibrium biosorption capacity calculated from this model (q_{e2}) matched the experimental value ($q_{e exp}$) closely.

The pseudo-second-order model has also been successfully used to describe the biosorption kinetics of other dye-biosorbent systems. Examples include red 27 onto the leaves of *Eichhor-nia crassipes* [16]; direct red-31 and direct orange-26 onto rice husks [18]; methylene blue onto tamarind fruit shell [39]; crystal violet onto *Artocarpus altilis* (breadfruit) skin [30]; acid orange 7, basic red 46, and basic blue 3 onto *Spirogyra* sp. [43]; and reactive black 5 onto *Laminaria* sp. [44].

Tables 2–4 show the variation in the pseudo-second-order model rate constant (k_2) according to the solution pH, corncob particle size, and C_o . A lower k_2 value means a longer time needed to reach the equilibrium biosorption capacity under the given experimental condition, because k_2 plays the key role of a time-scaling factor in the corresponding kinetic equation [45]. The same trend was observed in other dye biosorption systems [16, 18, 41, 44].

DY27 biosorption isotherms and their mathematical modeling

Fig 7 illustrates the biosorption isotherms of DY27 by corncob at pH = 1.5 and different temperatures (18, 35, 50, and 60°C). The obtained isotherms are regular, smooth, and concave to

PLC	DS ONE			Biosorption	of Direct Yellow 27	dye from aqueous	solutions by cornco
-							
Table 2. Kinetic	model parameters for	DY27 biosorption or	nto corncob at differer	nt solution pH levels.			
Solution pH	1.5	2.0	2.5	3.0	4.0	5.0	6.0
$q_{e\ exp}\ [\mathrm{mg\ g}^{-1}]$	35.1551 ± 0.8815	28.4783 ± 0.4683	$\bf 20.5694 \pm 0.4657$	12.6606 ± 0.9735	6.900 ± 0.9896	4.2121 ± 0.79	1.5235 ± 0.08962
			Pseudo-fi	irst-order			
q _{e1}	32.52 ± 1.0	25.65 ± 0.92	18.58 ± 0.66	11.51 ± 0.58	8.052 ± 0.409	4.716 ± 0.215	1.437 ± 0.059
k_{I}	0.647 ± 0.0689	1.279 ± 0.204	1.222 ± 1.191	1.115 ± 0.8759	1.020 ± 0.2164	1.075 ± 0.2047	1.152 ± 0.1893
r^2	0.9662	0.9238	0.9269	0.8724	0.8726	0.8950	0.9486
ASS	197.5	254.7	129.6	95.13	45.04	12.73	0.5502
Sy.x	2.008	2.280	1.626	1.393	0.9587	0.5098	0.1203
AIC	75.57	88.52	54.08	38.31	0.1721	-64.26	-164.8
			Pseudo-sec	cond-order			
q _{e2}	35.62 ± 0.87	28.02 ± 0.62	20.35 ± 0.46	12.68 ± 0.61	8.883 ± 0.394	5.195 ± 0.203	1.579 ± 0.061
k_2	0.0259 ± 0.0031	0.06507 ± 0.0084	0.08452 ± 0.0109	0.1197 ± 0.03242	0.1577 ± 0.382	0.2840 ± 0.0616	0.9573 ± 0.1953
r^2	0.9831	0.9794	0.9795	0.9162	0.9301	0.9443	0.9664
ASS	98.66	68.92	36.40	62.51	24.70	6.759	0.3595
Sy.x	1.419	1.186	0.8619	1.129	0.7100	0.3714	0.09727
AIC	40.16	21.86	-10.69	16.89	-30.46	-96.56	-181.8
			Elor	vich			
B_E	0.1921 ± 0.0226	0.2794 ± 0.0298	0.3802 ± 0.0403	0.5944 ± 0.089	0.8387 ± 0.1064	1.443 ± 1.3508	4.581 ± 0.771
A_E	157.1 ± 74.14	448.1 ± 249.6	290 ± 157.1	140.9 ± 102.79	87.40 ± 52.73	54.80 ± 33.94	12.71 ± 9.506
r^2	0.9181	0.9313	0.9316	0.8710	0.9035	0.9016	0.8806
ASS	479.0	229.8	121.2	96.22	34.12	11.94	1.279
Sy.x	3.126	2.165	1.573	1.401	0.8344	0.4936	0.1835
AlC	120.7	83.27	50.67	38.88	-13.99	-67.54	-131.0
			Intrapartic	le diffusion			
k _{id}	3.545 ± 0.887	2.523 ± 0.719	1.852 ± 0.52	1.18 ± 0.343	0.8446 ± 0.2268	0.4862 ± 0.1354	0.1504 ± 0.505
с	13.76 ± 2.87	13.97 ± 2.32	9.961 ± 1.68	5.951 ± 1.109	4.053 ± 0.733	2.415 ± 0.438	0.6664 ± 0.1654
r ²	0.5688	0.5038	0.5117	0.4941	0.5340	0.5157	0.4887
ASS	2521	1659	865.7	377.3	164.7	58.76	5.478
Sy.x	7.173	5.819	4.203	2.775	1.833	1.095	0.3797
AIC	205.4	184.1	150.9	108.6	66.30	13.73	-72.86
			Fraction	al power			
k _{fp}	18.41 ± 1.53	17.56 ± 1.06	12.57 ± 0.77	7.588 ± 0.598	5.211 ± 0.372	3.086 ± 0.222	0.9003 ± 0.0932
v	0.1851 ± 0.0305	0.1465 ± 0.0234	0.1495 ± 0.0237	0.1563 ± 0.0301	0.1612 ± 0.0271	0.1580 ± 0.0276	0.1649 ± 0.0377
r^2	0.8392	0.8669	0.8949	0.8665	0.8438	0.8371	0.7984

Tab

https://doi.org/10.1371/journal.pone.0196428.t002

939.9

4.380

155.1

445.2

3.014

117.0

221.4

2.194

79.93

ASS

Sy.x

AIC

the x-axis (concentration axis). The concave shape and initial slope indicate that as more biosorption binding sites are occupied, it becomes more difficult for the remaining dye molecules to find available active sites on the corncob surface, which suggests a progressive saturation of corncob with DY27 [16, 46].

55.21

1.061

10.56

19.76

0.6350

-41.85

2.160

0.2384 -110.1

236.8

2.198

84.81

The DY27 biosorption performance decreased with increasing temperature from 18 to 60°C, revealing the exothermic nature of this process. This lower biosorption at higher temperature could be attributed to many possible factors: increased solubility of DY27 in water, stronger dye-solvent interaction than that between DY27 molecules and corncob, increased Brownian movement of DY27 molecules in solution, the breaking of existing intermolecular hydrogen bonding between DY27 and corncob, and/or damaged active biosorption sites that

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Particle size [mm]	0.297-0.500	0.500-0.800	0.800-1.000	1.000-1.180	1.18-1.410	1.410-1.700	1.700-2.000
$q_{e \ exp} \ [\mathrm{mg \ g}^{-1}]$	35.1551 ± 0.8815	32.9090 ± 1.5308	33.5694 ± 1.4657	33.2058 ± 0.7393	31.2677 ± 0.9359	29.5313 ± 0.9634	26.7755 ± 0.04481
			Pseudo-f	first-order			
<i>q</i> _{e1}	32.52 ± 1.0	31.27 ± 1.02	31.57 ± 0.95	30.58 ± 0.94	29.68 ± 0.95	28.05 ± 0.99	26.27 ± 0.89
<i>k</i> 1	0.647 ± 0.0689	0.4052 ± 0.0386	0.2301 ± 0.0172	0.22124 ± 0.0249	0.1965 ± 0.0153	0.1519 ± 0.0133	0.1144 ± 0.0096
r ²	0.9662	0.9724	0.9819	0.9810	0.9795	0.9765	0.9803
ASS	197.5	149.2	97.94	94.75	96.59	95.80	68.85
Sy.x	2.008	1.745	1.428	1.405	1.404	1.413	1.185
AlC	75.57	61.25	40.14	38.48	39.08	39.04	21.81
			Pseudo-se	cond-order			
<i>q</i> _{e2}	35.62 ± 0.87	34.12 ± 0.77	34.78 ± 0.7	33.78 ± 0.66	32.93 ± 0.84	31.67 ± 0.98	30.52 ± 1.12
<i>k</i> ₂	0.0259 ± 0.0031	0.01665 ± 0.00162	0.008266 ± 0.0006	0.007761 ± 0.0006	0.007286 ± 0.0007	0.005537 ± 0.0006	0.004020 ± 0.0005
r^2	0.9831	0.9891	0.9936	0.9941	0.9905	0.9879	0.9857
ASS	98.66	58.96	34.36	29.17	44.94	49.22	50.07
Sy.x	1.419	1.097	0.8460	0.7796	0.9577	1.013	1.011
AIC	40.16	13.91	-12.24	-20.42	0.06425	5.740	5.568
			Elo	vich			
B_E	0.1921 ± 0.0226	0.1806 ± 0.018	0.1508 ± 0.0105	0.1528 ± 0.0092	0.1541 ± 0.0103	0.1495 ± 0.0088	0.1425 ± 0.0098
A_E	157.1 ± 74.14	63.58 ± 20.91	20.82 ± 3.6	17.87 ± 2.6	15.54 ± 2.43	9.606 ± 1.189	5.678 ± 0.712
<i>r</i> ²	0.9181	0.9445	0.9774	0.9831	0.9794	0.9857	0.9828
ASS	479.0	300.5	122.1	84.09	96.86	58.22	60.20
Sy.x	3.126	2.476	1.595	1.324	1.406	1.101	1.108
AlC	120.7	96.97	51.17	32.52	39.22	14.13	14.97
			Intrapartie	cle diffusion			
k _{id}	3.545 ± 0.887	3.696 ± 0.749	4.104 ± 0.551	3.993 ± 0.493	3.894 ± 0.466	3.74 ± 0.348	3.519 ± 0.263
с	13.76 ± 2.87	10.04 ± 2.421	5.642 ± 1.787	5.045 ± 1.598	4.619 ± 1.508	3.004 ± 1.13	1.640 ± 0.8494
r ²	0.5688	0.6677	0.8240	0.8472	0.8521	0.9072	0.9367
ASS	2521	1799	951.7	760.5	696.8	378.6	221.2
Sy.x	7.173	6.059	4.453	3.980	3.771	2.808	2.125
AIC	205.4	188.2	153.8	142.6	139.9	107.7	81.34
			Fraction	al power			
k_{fp}	18.41 ± 1.53	14.66 ± 1.33	10.51 ± 1.007	9.733 ± 0.888	9.134 ± 0.851	7.239 ± 0.652	5.508 ± 0.532
ν	0.1851 ± 0.0305	0.2243 ± 0.03143	0.2980 ± 0.0305	0.3074 ± 0.0286	0.3146 ± 0.0292	0.3526 ± 0.0272	0.3968 ± 0.0283
r ²	0.8392	0.8678	0.9209	0.9330	0.9317	0.9535	0.9592
ASS	939.9	715.6	427.5	333.2	321.8	189.7	142.5
Sy.x	4.380	3.822	2.984	2.635	2.563	1.988	1.706
AIC	155.1	141.2	113.8	101.4	100.5	73.20	58.93

Table 3. Kinetic model parameters for DY27 biosorption onto corncob at different corncob particle sizes.

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decrease the surface activity [1, 47]. A decrease in the dye biosorption performance at higher temperatures has also been reported for the biosorption of FD&C red No. 40 and acid blue 9 by *Spirulina platensis* [48]; acid blue 9, food yellow 3, and FD&C yellow No. 5 by chitosan [49]; methylene blue, eriochrome black T, and alizarin S by prickly pear cactus (*Opuntia ficus indica*) cladodes [50]; and astrazon red by the leaves of *Posidonia oceanica* (Linnaeus) Delile [51], among others.

The DY27 biosorption isotherms were modeled with several two-parameter (Langmuir, Freundlich, Temkin, Halsey, and Dubinin-Radushkevich) and three-parameter (Sips, Radke-Prausnitz, Redlich-Peterson, and Toth) models. The parameters obtained from nonlinear regression analysis for different temperatures, along with their corresponding r^2 , *ASS*, *Sy.x*,

				Initia	l concentration [mg L ⁻¹]			
	7	13	28	50	74	85	100	145	293
$q_{e\ exp}\ [\mathrm{mg\ g}^{-1}]$	6.519 ± 0.20	11.762 ± 0.25	19.897 ± 0.07	$\textbf{34.918} \pm \textbf{0.88}$	$\textbf{48.425} \pm \textbf{0.65}$	57.56 ± 0.268	67.14 ± 1.273	72.996 ± 1.40	$\textbf{73.04} \pm \textbf{4.08}$
				Pseudo-j	first-order				
q _{e1}	6.085 ± 0.216	10.43 ± 0.28	17.74 ± 0.88	31.51 ± 1.62	44.70 ± 1.92	54.48 ± 2.66	63.13 ± 3.32	69.35 ± 4.36	71.51 ± 3.43
k_1	0.987 ± 0.142	0.825 ± 0.083	0.490 ± 0.076	0.510 ± 0.083	0.370 ± 0.045	0.365 ± 0.050	0.336 ± 0.052	0.351 ± 0.059	0.371 ± 0.049
r^2	0.9362	0.9714	0.9238	0.9195	0.9538	0.9372	0.9373	0.9355	0.9552
ASS	12.29	18.08	127.3	435.8	501.0	957.1	1245	1291	929.3
Sy.x	0.5008	0.6074	1.612	2.982	3.198	4.420	5.319	5.829	5.012
AIC	-66.08	-46.39	53.18	115.9	123.0	156.0	158.3	145.6	130.4
				Pseudo-se	cond-order				
<i>q</i> _{e2}	6.560 ± 0.191	11.46 ± 0.23	19.44 ± 0.6	34.58 ± 0.99	48.96 ± 1.05	58.90 ± 2.37	67.96 ± 2.58	74.87 ± 3.56	77.5 ± 3.33
<i>k</i> ₂	0.238 ± 0.039	0.101 ± 0.010	0.036 ± 0.005	0.021 ± 0.003	0.010 ± 0.001	0.009 ± 0.002	0.007 ± 0.001	0.007 ± 0.001	0.007 ± 0.001
r^2	0.9683	0.9879	0.9763	0.9795	0.9903	0.9644	0.9723	0.9685	0.9711
ASS	6.109	7.666	39.71	111.0	105.2	541.7	549.0	630.7	600.4
Sy.x	0.3531	0.3955	0.9002	1.505	1.465	3.325	3.532	4.074	4.028
AIC	-101.7	-90.14	-6.254	46.17	43.45	127.0	120.6	117.0	113.3
				Elo	wich				
B_E	1.175 ± 0.165	0.604 ± 0.076	0.332 ± 0.023	0.186 ± 0.012	0.122 ± 0.008	0.106 ± 0.009	0.091 ± 0.007	0.0809 ± 0.007	0.081 ± 0.011
A_E	82.35 ± 57.66	65.04 ± 34.79	53.84 ± 13.5	97.00 ± 22.43	75.57 ± 16.23	110.4 ± 33.78	116.4 ± 33.14	120.6 ± 34.1	148.5 ± 56.41
<i>r</i> ²	0.8862	0.9064	0.9707	0.9754	0.9737	0.9510	0.9699	0.9650	0.9487
ASS	21.91	59.11	48.94	133.4	285.0	746.9	597.1	701.0	1064
Sy.x	0.6686	1.098	0.9994	1.650	2.412	3.904	3.684	4.295	5.362
AlC	-36.59	14.04	4.404	55.54	94.26	143.4	124.5	121.2	135.6
				Intraparti	cle diffusion				
k _{id}	0.5787 ± 0.18	1.120 ± 0.305	2.100 ± 0.397	3.757 ± 0.723	5.500 ± 0.943	6.402 ± 1.17	7.574 ± 1.328	8.793 ± 1.722	8.683 ± 1.915
с	3.166 ± 0.582	4.774 ± 0.988	6.434 ± 1.283	11.52 ± 2.335	13.25 ± 3.04	17.04 ± 3.79	18.3 ± 4.47	18.61 ± 5.17	21.85 ± 6.04
<i>r</i> ²	0.4602	0.5258	0.6981	0.6902	0.7377	0.7120	0.7505	0.7379	0.6954
ASS	103.9	299.6	504.8	1677	2846	4387	4953	5250	6319
Sy.x	1.456	2.473	3.210	5.850	7.621	9.462	10.61	11.75	13.07
AIC	42.82	96.81	123.4	184.6	211.6	233.7	221.8	201.8	205.1
				Fraction	ial power				
k _{fp}	3.980 ± 0.302	6.276 ± 0.536	9.080 ± 0.604	16.27 ± 1.1	19.99 ± 1.53	24.93 ± 1.98	27.97 ± 2.25	30.15 ± 2.65	33.14 ± 3.14
v	0.149 ± 0.029	0.175 ± 0.032	0.210 ± 0.024	0.210 ± 0.024	0.237 ± 0.026	0.227 ± 0.028	0.234 ± 0.026	0.2422 ± 0.030	0.2248 ± 0.033
r^2	0.8111	0.8186	0.9116	0.9096	0.9104	0.8953	0.9188	0.9111	0.8935
ASS	36.37	114.6	147.7	489.0	972.0	1595	1612	1782	2209
Sy.x	0.8616	1.529	1.736	3.159	4.454	5.705	6.052	6.847	7.726
AIC	-10.73	47.80	60.75	121.8	156.8	182.1	170.2	158.5	164.1

Table 4. Kinetic model parameters for DY27 biosorption onto corncob at different initial DY27 concentrations.

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and *AIC* values are shown in Tables 5–8. The equilibrium biosorption of DY27 onto corncob is best described by the Redlich-Peterson model, as it yielded a higher r^2 value and lower *ASS*, *Sy.x* and *AIC* values at each assayed temperature compared to the other isotherm models. The good agreement between this model and experimental results are shown in Fig 7 (lines and points, respectively).

The Redlich-Peterson isotherm model is a hybrid model of Langmuir and Freundlich models, and it incorporates three parameters into a single empirical equation. This versatile model has been successfully used to represent biosorption equilibria over a wide concentration range, and can be applied to both homogeneous and heterogeneous systems [52]. Examples include

Two-parameter isotherms Base 100		Thre	Three-parameter isotherms		
	Langmuir		Sips	_	
$\overline{\mathrm{K}_{L} [\mathrm{L} \mathrm{mg}^{-1}]}$	0.05430 ± 0.01092	$K_{s}[(mg L^{-1})^{-1/ns}]$	0.05871 ± 0.01076	_	
$q_m [mg g^{-1}]$	81.75 ± 4.74	q _m [mg g ⁻¹]	79.61 ± 5.87	_	
r^2	0.9728	n _s	0.8127 ± 0.1872	_	
ASS	295.5	r ²	0.9748	_	
<u>Sy.x</u>	4.252	ASS	450.2	_	
AIC	85.60	Sy.x	4.331	_	
	Freundlich	AIC	85.79		
$\overline{K_F [(mg g^{-1}) (mg L^{-1})^{-1/nF}]}$	16.00 ± 4.21		Radke-Prausnitz	_	
n _F	3.267 ± 0.619	$A_R [L g^{-1}]$	5.029 ± 2.35	_	
r ²	0.8922	$R_R [L mg^{-1}]$	83.16 ± 5.32	_	
ASS	1913	B _R	0.0029 ± 0.1801		
<u>Sy.x</u>	8.748	r ²	0.9700		
AIC	122.1	ASS	7691		
Temkin		Sy.x	2.777		
$\overline{\mathbf{A}_T \left[\mathbf{L} \mathbf{mg}^{-1} \right]}$	1.071 ± 0.1061	AIC	2048	_	
$B_T [J \text{ mol}^{-1}]$	72.45 ± 1.56		Redlich-Peterson		
r ²	0.8919	$K_{RP} [L g^{-1}]$	3.245 ± 0.033	_	
ASS	26223	$A_{RP} \left[\left(L mg^{-1} \right)^{BRP} \right]$	0.01415 ± 0.0001		
<u>Sy.x</u>	5.126	B _{RP}	0.991 ± 0.009		
AIC	3273	r ²	0.9965	_	
	Halsey	ASS	282.7		
$\overline{K_H [(L g^{-1})^{-1/nH}]}$	1.218E-006 ± 0.888E-006	Sy.x	0.9672		
n _H	-0.2318 ± 0.0091	AIC	-61.61		
r ²	0.7992		Toth		
ASS	47971	$q_m [mg g^{-1}]$	75.03 ± 3.47		
Sy.x	6.933	$b_T [(L mg^{-1})^{-1/nT}]$	0.03345 ± 0.00559		
AIC	3877	n _T	0.4529 ± 0.1801		
Dul	binin-Radushkevich	r ²	0.9826		
$\overline{\mathbf{q}_m [\mathrm{mg}\mathrm{g}^{-1}]}$	73.73 ± 0.23	ASS	302.9		
$B_{DR} [mol^2 J^{-2}]$	0.0001679 ± 0.0015159	Sy.x	4.552		
r ²	0.9573	AIC	75.09		
ASS	9281				
Sy.x	4.350				
AIC	2234				

Table 5. Parameters of the isotherm models for the biosorption at 18°C [$q_{m exp}$: 73.71 ± 4.8 mg g⁻¹].

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the biosorption of FD&C red No. 40 onto chitosan [53], acid red 18 and FD&C blue No. 2 onto chitosan films [54], reactive blue 19 onto cross-linked chitosan/oil palm ash composite beads [55], Gemazol turquoise blue-G reactive dye onto dried sugar beet pulp [56], and methylene blue onto activated carbons [57].

The maximum experimental biosorption capacity $(q_{m exp})$ was found to be 73.7 mg g⁻¹ at 18°C (Table 5). As mentioned above, to the best of our knowledge, this is the first report of using a biosorbent to remove DY27 from aqueous solutions. The only previous work on DY27 removal used an inorganic adsorbent, namely poorly crystalline hydroxyapatite prepared via a precipitation method [26]. In that study, the maximum adsorption occurred at pH 3.5 to be 6.8 mg g⁻¹, while the corresponding value predicted by the Langmuir model was 89.26 mg g⁻¹ [26].



Two-parameter isotherms Base 100		Three-parameter isotherms			
Langn	nuir		Sips		
K _L [L mg ⁻¹]	0.04341 ± 0.0379	$K_{S}[(mg L^{-1})^{-1/ns}]$	0.04834 ± 0.01061		
$q_m [mg g^{-1}]$	65.03 ± 5.08	$q_m [mg g^{-1}]$	60.65 ± 5.67		
r ²	0.9425	n _s	0.7726 ± 0.2039		
ASS	419.6	r ²	0.9501		
Sy.x	4.025	ASS	424.7		
AIC	85.28	Sy.x	4.207		
Freund	llich	AIC	84.22		
$\frac{1}{K_F [(mg g^{-1}) (mg L^{-1})^{-1/nF}]}$	11.63 ± 3.822		Radke-Prausnitz		
n _F	3.210 ± 0.745	$A_R [L g^{-1}]$	3.400 ± 2.35		
r ²	0.822	$R_R [L mg^{-1}]$	15.38 ± 0.76		
ASS	1516	B _R	0.02599 ± 0.0103		
Sy.x	7.787	r ²	0.7939		
AIC	115.8	ASS	29491		
Teml	kin	Sy.x	5.439		
A _T [L mg ⁻¹]	0.5905 ± 0.0522	AIC	3392		
$B_T[J \text{ mol}^{-1}]$	84.60 ± 1.86		Redlich-Peterson		
r ²	0.8879	$K_{RP} [Lg^{-1}]$	1.816 ± 0.02		
ASS	16040	$A_{RP} \left[(L mg^{-1})^{BRP} \right]$	0.005964 ± 0.0001		
Sy.x	4.009	B _{RP}	0.999 ± 0.001		
AIC	2781	r ²	0.9947		
Hals	ey	ASS	358.8		
K _H [(L g ⁻¹) ^{-1/nH}]	2.710E-005 ± 1.637E-005	Sy.x	0.8724		
n _H	-0.2599 ± 0.0123	AIC	-268.0		
r ²	0.7939		Toth		
ASS	29491	$q_m [mg g^{-1}]$	56.6 ± 3.25		
Sy.x	5.436	$b_T [(L mg^{-1})^{-1/nT}]$	0.0252 ± 0.00393		
AIC	3390	n _T	0.3625 ± 0.1617		
Dubinin-Rad	lushkevich	r ²	0.9646		
q_m [mg g ⁻¹]	56.27 ± 0.23	ASS	431.0		
$B_{DR} [mol^2 J^{-2}]$	0.0002555 ± 0.0015159	Sy.x	4.541		
r ²	0.9624	AIC	74.92		
ASS	5382				
Sy.x	4.322				
AIC	1689				

Table 6. Parameters of the isotherm models for the biosorption at 35°C [$q_{m exp}$: 55.79 ± 4.29 mg g⁻¹].

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The above results indicate that corncob may be used as a low-cost, abundant, and green biosorbent with high capacity to remove DY27 from contaminated water and wastewater.

Thermodynamics of DY27 biosorption

The DY27 biosorption isotherm data at different temperatures were used to calculate important thermodynamic parameters, namely changes in Gibbs free energy (ΔG°), enthalpy (ΔH°), and entropy (ΔS°), in order to provide in-depth information about the inherent energetic changes during the biosorption process [37, 58]. According to Table 9, the ΔG° values were negative at all the temperatures (18–60°C), indicating that the biosorption of DY27 onto

Two-parameter isotherms Base 100		Three-parameter isotherms			
	Langmuir		Sips		
$\overline{\mathrm{K}_{L} [\mathrm{L} \mathrm{mg}^{-1}]}$	0.07311 ± 0.0379	$K_{S}[(mg L^{-1})^{-1/ns}]$	0.09408 ± 0.011		
$q_m [mg g^{-1}]$	43.19 ± 1.01808	$q_m [mg g^{-1}]$	38.73 ± 1.35		
\mathbf{r}^2	0.9524	n _s	0.5929 ± 0.0961		
ASS	216.4	r ²	0.9834		
<u>Sy.x</u>	2.042	ASS	275.44		
AIC	63.23	Sy.x	2.773		
	Freundlich	AIC	37.56		
$\overline{K_F [(mg g^{-1}) (mg L^{-1})^{-1/nF}]}$	10.09 ± 3.476		Radke-Prausnitz		
n _F	3.681 ± 1.022	$A_R [L g^{-1}]$	4.308 ± 0.115		
r ²	0.7713	$R_R [L mg^{-1}]$	41.12 ± 0.15		
ASS	1039	B _R	1.541E-016 ± 0.0103		
<u>Sy.x</u>	6.445	r ²	0.9422		
AIC	105.6	ASS	1886		
	Temkin	Sy.x	2.375		
$\overline{\mathbf{A}_T [\mathrm{L} \mathrm{mg}^{-1}]}$	6.114 ± 1.351	AIC	642.3		
$\overline{\mathbf{B}_T\left[\mathbf{J}\ \mathbf{mol}^{-1}\right]}$	188.6 ± 6.6		Redlich-Peterson		
r ²	0.76	$K_{RP} [L g^{-1}]$	2.685 ± 0.037		
ASS	7827	$A_{RP} [(L mg^{-1})^{BRP}]$	0.03161 ± 0.00114		
<u>Sy.x</u>	2.800	B _{RP}	0.995 ± 0.004		
AIC	2064	r ²	0.9901		
	Halsey	ASS	214.1		
$\overline{K_H [(L g^{-1})^{-1/nH}]}$	8.976E-009 ± 1.266E-009	Sy.x	0.5702		
n _H	-0.1553 ± 0.0123	AIC	-1119		
r ²	0.6827		Toth		
ASS	10347	$q_m [mg g^{-1}]$	38.32 ± 1.39		
Sy.x	3.220	$b_T [(L mg^{-1})^{-1/nT}]$	0.04885 ± 0.00709		
AIC	2343	n _T	0.4321 ± 0.1421		
Dut	binin-Radushkevich	r ²	0.9817		
$\overline{\mathbf{q}_m [\mathrm{mg}\mathrm{g}^{-1}]}$	38.04 ± 0.09	ASS	283.07		
$B_{DR} [mol^2 J^{-2}]$	8.049E005± 0.195E005	Sy.x	2.460		
r ²	0.9547	AIC	40.16		
ASS	1478				
Sy.x	2.217				
AIC	397.0				

Table 7. Parameters of the isotherm models for the biosorption at 50 °C [$q_{m exp}$: 38.99 ± 3.9 mg g⁻¹].

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corncob was spontaneous and thermodynamically favorable (i.e., there is an affinity of corncob towards the DY27 dye) [37, 44]. These ΔG° values are within the general range of -20 to 0 kJ mol⁻¹ for physisorption processes [37, 55].

The ΔG° value was more negative at 18°C, indicating that the biosorption was more energetically favorable at lower temperatures [1, 37]. This is due to the positive ΔS° value that indicates increased randomness at the solid-liquid interface after biosorption [37]. The positive ΔS° value also suggests the possibility of some structural changes or readjustments in the DY27-corncob complex [1].

The negative ΔH° values indicate that the biosorption is an exothermic process, in agreement with the observed decrease in biosorption as the temperature increases. The magnitude



Fig 7. Isotherms of DY27 biosorption by corncob at different temperatures.

of ΔH° value confirms that it is a physical sorption process (physisorption), for which the values of $|\Delta H^{\circ}|$ are usually lower than 84 kJ mol⁻¹ [37]. It is well known that the physisorption is exothermic and enhanced at lower temperature, because the interaction between adsorbates and active sites is weakened at higher temperatures [59].

FTIR analysis

FTIR spectra were collected for native and DY27-loaded corncob samples, in order to examine the interactions between dye molecules with the functional groups on the corncob. Fig 8 displays the spectra for the corncob samples and that of DY27 alone, and Table 10 lists the peak positions of the major absorption bands.

Cellulose, hemicellulose, and lignin are the main constituents of corncob. They contain functional groups that exhibit common absorption bands, including the OH stretching at 3428 cm⁻¹ [60] and the symmetric stretching of aliphatic chains (-CH) at 2916 cm⁻¹ [61]. Absorption bands related mainly to cellulose were observed at 1374, 1160, and 900 cm⁻¹, which are associated with the symmetric C-H deformations, C-O-C stretching of glycosidic bonds, and antisymmetric out-of-plane ring stretching, respectively [66, 67, 70]. Two characteristic absorption bands of hemicellulose were identified, the first one at 1732 cm⁻¹ is associated with the stretching vibration of C = O group [62], and the second at 1250 cm⁻¹ is attributed to the stretching of C-O bond [68, 71]. Representative absorption bands of lignin were found at

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Two-parameter isotherms Base 100		Th	Three-parameter isotherms			
	Langmuir		Sips			
$\overline{\mathbf{K}_{L}\left[\mathbf{L}\mathbf{mg}^{-1}\right]}$	0.1615 ± 0.0409	$K_{s}[(mg L^{-1})^{-1/ns}]$	0.1711 ± 0.0304			
$q_m [mg g^{-1}]$	29.30 ± 1.28	$q_m [mg g^{-1}]$	27.80 ± 1.15			
r ²	0.9490	n _s	0.7354 ± 0.1472			
ASS	42.63	r ²	0.9657			
Sy.x	1.425	ASS	63.33			
AIC	30.49	Sy.x	1.697			
	Freundlich	AIC	23.89			
$\overline{K_F [(mg g^{-1}) (mg L^{-1})^{-1/nF}]}$	11.70 ± 3.213		Radke-Prausnitz			
n _F	5.543 ± 1.956	$A_R [L g^{-1}]$	5.554 ± 0.158			
r ²	0.6886	$R_R [L mg^{-1}]$	28.6 ± 0.49			
ASS	386.8	B _R	1.717E-016 ±000.0103			
Sy.x	4.193	r ²	0.9757			
AIC	73.92	ASS	212.6			
	Temkin	Sy.x	0.4617			
$\overline{\mathbf{A}_T \left[\mathbf{L} \mathbf{mg}^{-1} \right]}$	116.6 ± 39.67	AIC	-1540			
$\overline{\mathbf{B}_T\left[\mathbf{J}\ \mathbf{mol}^{-1}\right]}$	377.3 ± 13.8		Redlich-Peterson			
r ²	0.7404	$\mathrm{K}_{RP}[\mathrm{L}\mathrm{g}^{-1}]$	4.158 ± 0.037			
ASS	2266	$A_{RP} [(L mg^{-1})^{BRP}]$	0.113 ± 0.0017			
Sy.x	1.507	B _{RP}	0.989 ± 0.001			
AIC	824.0	r ²	0.9961			
	Halsey	ASS	34.04			
$K_H [(L g^{-1})^{-1/nH}]$	4.236E-012 ± 2.32E-012	Sy.x	0.1848			
n _H	-0.1066 ± 0.0051	AIC	-3372			
r ²	0.6833		Toth			
ASS	2765	$q_m [mg g^{-1}]$	27.71 ± 1.29			
Sy.x	1.664	$b_T [(L mg^{-1})^{-1/nT}]$	0.11 ± 0.03423			
AIC	1023	n _T	0.6536 ± 0.2326			
Dubir	nin-Radushkevich	r ²	0.9636			
$\mathbf{q}_m [\mathbf{mg}\mathbf{g}^{-1}]$	27.15 ± 0.06	ASS	45.19			
$\mathbf{B}_{DR} \left[\mathbf{mol}^2 \mathbf{J}^{-2} \right]$	2.789E005± 0.095E005	Sy.x	1.467			
r ²	0.9052	AIC	25.29			
ASS	827.4					
Sy.x	2.9105					
AIC	-183.5					

Table 8. Parameters of the isotherm models for the biosorption at 60 °C [$q_{m exp}$: 28.23 ± 1.0 mg g⁻¹].

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1515, 1603, 1426, and 1070 cm⁻¹, which correspond to the aromatic skeletal vibration, the aromatic skeletal vibration breathing with C = O stretching [64, 66, 72], the C-H in-plane deformation with aromatic ring stretching, and the C-O deformations of secondary alcohols and aliphatic ethers [64], respectively. The characteristic absorption band of amide I C = O stretching at 1653 cm⁻¹ [63] was also detected in the spectrum of native biosorbent, and this band most probably corresponds to the amide groups in the proteins.

After loading DY27 onto the native biosorbent, the FTIR spectrum showed a few changes. The relative intensity decreased for the bands at 1653, 1603, 1374, and 1070 cm⁻¹. Based on the assignments described above, these changes indicate the involvement of the proteins, cellulose, and lignin of corncob in the biosorption of DY27 dye molecules. Additionally, the absorption



T [°C]	ΔG° [J mol ⁻¹]	ΔS° [J mol ⁻¹ K ⁻¹]	ΔH° [J mol ⁻¹]
18	-2135.00 ± 39.03	25.956 ± 0.6311	-9692.09 ± 39.03
35	-1522.24 ± 34.97		-9520.58 ± 34.97
50	-1136.06 ± 29.24		-9523.74 ± 29.24
60	-1074.53 ± 26.87		-9721.77 ± 26.95

Table 9. Thermodynamic parameters for the biosorption.

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band at 1514 cm⁻¹ became more intense, which can be explained by the combination between the band corresponding to the aromatic skeletal vibration of lignin at 1514 cm⁻¹ and a characteristic absorption band of DY27 (the stretching of the azo functional group) at 1507 cm⁻¹ [65]. A new absorption band appeared at 1458 cm⁻¹, which could be related again to the presence of the DY27 dye, because this wavenumber is close to the C = C stretching frequency of its aromatic ring at 1464 cm⁻¹ [65]. The presence of these functional groups belonging to DY27 in the dye-loaded corncob may come from the dye molecules that either do not participate directly in the biosorption process or were biosorbed through weak forces.

Dyes with linear molecular structures (including DY27) are used mainly for dyeing paper. The paper dyeing process basically involves the penetration of dye molecules into the capillary spaces of cellulose, followed by dye adsorption on the fiber surface. The adsorption is driven



Fig 8. FTIR spectra of (A) native corncob, (B) DY27-loaded corncob, and (C) DY27 dye.

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IR band frequency [cm ⁻¹]		Functional group assignment	Reference	
Native corncob	DY27-loaded corncob	DY27 dye		
3428	3421	3498	O-H stretching	[60]
2916	2910	-	Symmetric stretching of C-H	[61]
1732	1734	-	C = O stretching vibration of hemicellulose	[62]
1653	1654	1664	C = O stretching of amide I of proteins	[63]
1603	1603	1598	Aromatic skeletal vibration + C = O stretching	[64]
1515	1514	-	Aromatic skeletal vibration of lignin	[64]
-	-	1507	N = N stretching	[65]
-	1458	1464	C = C stretching of aromatic rings of dye	[65]
1426	1430	1438	C-H in plane deformation with aromatic ring stretching	[66]
1374	1375	-	Symmetric C-H deformation of cellulose	[67]
1250	1248	-	C-O stretching of hemicellulose	[68]
-	-	1211, 1188	Amine C-N stretching	[69]
1160	1163	-	C-O-C stretching of glycosidic linkages	[70]
1070	1085	-	C-O deformation of secondary alcohols and aliphatic ethers	[64]
1049 shoulder	1051 shoulder	1049	C-O stretching vibrations of cellulose S = O stretching of sulfonate groups	[70] [63]
900	898	-	Antisymmetric out-of-plane ring stretching of cellulose	[66]

Table 10. Summary of infrared spectral bands observed in the native corncob, DY27-loaded corncob, and DY27 dye.

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by the effects of charge, precipitation, and intermolecular forces. It is known that lignin has a considerable influence on the dyeability, but not the hemicellulose [73]. This finding in paper dyeing is consistent with the analysis of the FTIR spectra of DY27 biosorbed on corncob.

CSLM analysis

CSLM is a powerful optical technique for visualizing the structure of biopolymer mixtures. In CSLM, optical contrast is obtained by differences in fluorescence, either by auto-fluorescence of the material or by the addition of specific fluorescent dyes (fluorochromes) [74]. Each fluorochrome has its own particular emission wavelength [75]. In the present work, two fluorochromes were used: fluorescein and calcofluor white M2R.

The auto-fluorescence of plant cells is mainly due to the presence of lignin and chlorophylls [76]. In the present work, the maximum emission wavelength ($\lambda_{em max}$) for lignin auto-fluorescence was observed at 501 nm (data not shown), which agrees with that reported by Rosas-Hernández (500 nm) [77]. Likewise, the peak of maximum emission for DY27 was found at 598 nm (data not shown).

The proteins were examined by staining DY27-loaded corncob samples with fluorescein, which binds covalently to the primary amine groups on the proteins. The $\lambda_{em max}$ for fluorescein was found at 519 nm, which is close to those reported by van de Velde et al. (518 nm) [74], Becker et al. (530 nm) [78], and the datasheet from the manufacturer (514 nm) [79].

Fig 9 shows CSLM images of DY27-loaded corncob stained with fluorescein. The auto-fluorescence of lignin (green color, Fig 9A), the fluorescence of DY27 dye on the corncob surface (red color, Fig 9B), and the fluorescein-induced fluorescence of corncob proteins (blue color, Fig 9C) can be observed. Furthermore, the interaction between DY27 with the corncob proteins was evidenced by a violet color in the overlay image (Fig 9D) produced by the combination of the red and blue colors from the DY27 dye and the fluorescein-induced fluorescence of proteins, respectively. These results clearly demonstrate that the DY27 dye binds to the proteins present in corncob.



Fig 9. CSLM images of a) lignin (green); b) lignin and DY27 dye in green and red colors, respectively; and c) fluorescein-induced fluorescence of corncob proteins in blue color. d) An overlay image showing the interaction between the DY27 dye and the corncob proteins in violet color.

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Fig 10. CSLM images of a) lignin (green); b) DY27 dye on the corncob surface in red color; and c) calcofluor white-induced fluorescence of corncob cellulose in blue color. d) An overlay image showing the interaction between the DY27 dye and the corncob cellulose in violet color.

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The cellulose was examined by staining DY27-loaded corncob samples with calcofluor white M2R solution. This fluorochrome emits fluorescence when activated by ultraviolet radiation and has an affinity for the cellulose and chitin in cell walls [80, 81]. The $\lambda_{em max}$ for calcofluor white was observed at 463 nm (data not shown), which is close to that reported in the datasheet from the manufacturer (433 nm) [82].

Fig 10 shows CSLM images of DY27-loaded corncob stained with calcofluor white. The auto-fluorescence of lignin is shown in green color (Fig 10A), the fluorescence of DY27 on the corncob surface in red (Fig 10B), and the calcofluor-induced fluorescence of corncob cellulose in blue (Fig 10C). The binding of DY27 onto the corncob cellulose was revealed by a violet color in the overlay image (Fig 10D), which is also due to the combination of the red and blue colors from DY27 and the calcofluor-induced fluorescence of cellulose, respectively. These results indicate that the DY27 dye also binds to the corncob cellulose.

The above findings clearly show that proteins and lignocellulose in the corncob play a crucial role in DY27 biosorption from aqueous solutions, which is consistent with the FTIR findings in this study.

DY27 desorption studies

Desorption of the dye from the DY27-loaded corncob was tested using several chemical agents, including acid, alkali, neutral salt, and organic solvent solutions. The performances of these eluent solutions are shown in Fig 11A. The highest DY27 desorption percentage was observed for strong alkali solutions, followed by weak alkaline, neutral salt, weak acid, organic solvent, and lastly strong acid solutions. Therefore, the DY27 desorption percentage depends strongly on the eluent pH: the higher the pH, the higher the desorption percentage. This trend may be explained on the basis of electrostatic repulsion between the negatively charged sites on the corncob biomass and the anionic DY27 molecules [16, 83]. The strong acid solutions performed poorly in the desorption, eluting only negligible amounts (< 4%) of biosorbed DY27. Therefore, we selected the alkaline solutions (NaOH, KOH, and Na₂CO₃) for further kinetic studies of the desorption process.

Fig 11B displays the DY27 desorption percentage as a function of desorption time. It is evident that for all three alkaline solutions, the desorption was initially very fast and became slower towards equilibrium. The highest rate and extent of DY27 desorption were achieved with 0.1 M NaOH solution, with a desorption efficiency of approximately 98%. These results clearly indicate that 0.1 M NaOH is the best eluent for DY27 desorption from the corncob-based biosorbent.

Conclusions

The present work clearly demonstrates the feasibility of utilizing corncob as a novel, effective, efficient, eco-friendly, and inexpensive biosorbent for the removal of DY27 dye from aqueous solutions. The DY27 biosorption performance of corncob was affected significantly by solution pH, corncob particle size, shaking contact time, and initial dye concentration. The kinetic data of biosorption were accurately described by the pseudo-second-order model, and the biosorption isotherms were well predicted by the Redlich-Peterson model. Thermodynamic study indicated that the DY27 biosorption process was spontaneous and exothermic. FTIR and CSLM results confirmed that lignocellulose and proteins are involved in the biosorption process. Finally, a 0.1 M NaOH solution successfully eluted most of the DY27 dye from the DY27-loaded corncob biomass.

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