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# Bacterial conversion of depolymerized Kraft lignin



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## **Abstract**

**Background:** Lignin is a potential feedstock for microbial conversion into various comicals. However, the degradation rate of native or technical lignin is low, and depolymerization is needed to obtain it asonable conversion rates. In the current study, base-catalyzed depolymerization—using NaOH (5 wto —o coftwood Kraft lignin was conducted in a continuous-flow reactor system at temperatures in the range 190–240 and residence times of 1 or 2 min. The ability of growth of nine bacterial strains belonging to the genera *P. sudomona*, and *Rhodococcus* was tested using the alkaline-treated lignin as a sole carbon source.

**Results:** *Pseudomonas fluorescens* and *Rhodococcus opacus* showed the best growth of the tested species on plates with lignin. Further evaluation of *P. fluorescens* and *R. opacus* was made in liquid cultivations with depolymerized lignin (DL) at a concentration of 1 g/L. Size exclusion chromat ograp. (SEC) showed that *R. opacus* consumed most of the available lower molecular weight compounds (approximately 0.1–0.4 kDa) in the DL, but the weight distribution of larger fractions was almost unaffected. Importantly, the continued compounds included guaiacol—one of the main monomers in the DL. SEC analysis of *P. fluorescena* by the proth, in contrast, did not show a large conversion of low molecular weight compounds, and guaiacol remain all unconsumed. However, a significant shift in molecular weight distribution towards lower average weights. Tas seen

**Conclusions:** Rhodococcus opacus ar \*1.P. fluore arms were identified as two potential microbial candidates for the conversion/consumption of base-catalyzed depolymerized lignin, acting on low and high molecular weight lignin fragments, respectively. These find ags will be of relevance for designing bioconversion of softwood Kraft lignin.

**Keywords:** Base-catalyzed depolymentation, Indulin AT, Microbial conversion, *Pseudomonas, Rhodococcus*, Guaiacol, Biorefineries

#### **Background**

Lignin is one of the mo. bundant biopolymers on Earth and constitutes [2-32% [x]] of woody biomass by weight and 40% by energy [1]. Its natural abundance, high calorific value and the fact that it is one of the few available renewal. Source's of aromatic chemicals in nature make it a prime eastock for targeted valorization towards lifety in the case of the few available can be called the polymer composites, and valuable can be called a precursors [3, 4]. Despite this, most large-scale

industrial plants that exploit plant polysaccharides have almost exclusively burned the co-produced lignin to generate heat and power desired for biomass conversion and/or product drying, and have not aimed for higher value products [5]. The emergence of biorefineries, which predominately convert the carbohydrate part of cellulosic biomass into liquid fuels, will generate substantially more lignin than needed for process heat or for power generation to the operation, which will add to lignin already generated in the pulp and paper industry. Hence, efforts are underway for lignin conversion into higher value products [6, 7].

Among the three principal polymers in plants, lignin has the most heterogeneous and complex composition

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and structure, turning its efficient utilization into a major technological challenge. The lignin biopolymer is built from phenyl propanoid units which are substituted at different positions and connected together by ether and C–C cross-links/bonds [8, 9]. It should be noted in this context that the relative contribution of these building blocks is very different between different types of biomass [10]. In addition, there are different techniques for the separation of lignin from lignocellulosic biomass and the resultant lignin vary in terms of bond structures, building block composition, added functional groups, and molecular weight distribution depending on the separation process. Hence, the term "lignin" is per se not very descriptive.

A central process step needed for biological lignin valorization is the breakdown of the large lignin macromolecule into smaller compounds that can be converted by microorganisms [1]. This depolymerization process is essential, as only low molecular weight compounds (mono- or possibly oligomers) can pass the cellular membranes of microbes and enter the cellular metabolism. The depolymerization is quite challenging, owing to the distribution of bond strengths in the different C-O and C-C bonds. Ideally, pretreated lignin streams for back rial conversion should consist of water-soluble monomers. For chemical depolymerization, in partic 'ar, char formation and repolymerization of produced phaselicradicals to large water-insoluble comple s thus h ed to be avoided. High concentrations are an normally desirable—a value of 100 g/L [5] has been men .oned although tolerance of the microbial catalyst to the monomers will likely limit the allowable oncentration in the process. Until now, obtained monome... species in lignin streams are still in the range few g/L [11]. The monomers resulting from chemical depolymerization are quite diverse. In a bic ogical conversion process, it is thus important to companisms that have the ability to funnel a large range of the lignin-derived aromatic compounds to tral intermediates (mainly protocatechuate and cateca. Via the upper-funneling catabolic pathways [1]. The principal intermediates are then further charged to valuable products using natural or ger cally sineered microorganisms [12-14]. In preous studies, microorganisms which are capable of utiliz r lignin/model compounds a sole source of carbon have een identified, including Pseudomonas putida, Rhodococcus opacus, Rhodococcus jostii, Amycolatopsis sp., Sphingobium sp., and Cupriavidus necator [15–18]. Successful proof-of-principle studies have been made for the bioconversion of lignin from alkaline pretreated corn stover and organosolv lignin into various products such as PHA, muconate and lipids [11, 19-25]. Apart from monomers, the depolymerization of lignin also results

in a heterogeneous mixture of oligomers and higher molecular weight compounds and ability of the selected host organisms to break down these larger fragments represents one process option in which a complete depolymerization is not necessary. This, however, requires the use of organisms which secrete extracer or lightlytic enzymes (e.g. laccases and DyP-type peroxida.

In the present study, we explore the possibilities of utilizing depolymerized softwood Fraft nin (indulin AT) as a substrate for bacterial conversion. oftwood Kraft lignin is potentially available large quantities from pulp mills in Scandinavia [26] It is a mylex substrate with a broad molecular weight a sibution and a high average molecular weight, Illing for depolymerization in bioprocessing. The lka. depolymerization of the lignin was performe a cont uous-mode operation without incorporation are capping agents or hydrogen donors, supporting it urtner scalability and commercial applicability Nine becerial species with known aromaticmetabour. expacity were evaluated for their ability to grow on plates using the depolymerized product, with the objective of finding organisms capable of both extrallular lignin breakdown along with the intracellular at matic metabolism. Further evaluation of selected strains was carried out in liquid shake flask cultivations using alkaline treated partly depolymerized lignin as a substrate. Additional cultivations were also made with a mixture of selected model compounds representing monomers found in the depolymerized lignin.

#### **Methods**

## Lignin substrate preparation

A commercial Kraft lignin, Indulin AT, which is a pine softwood lignin precipitated from the black liquor of linerboard-grade pulp [27], was obtained as a dry brown powder from MeadWestvaco Corporation (Charleston Heights, SC, USA). This lignin fraction has a typical moisture content of 4.5 wt% and ash content of 3.5 wt%. The feed of the chemical depolymerization setup was prepared before each experimental run and was characteristically comprised 5 wt% lignin substrate, 5 wt% NaOH, and 90 wt% deionized water. Vacuum filtration for the reactor feed was performed to ensure that no precipitation could take place in the pump and to counteract clogging problems within the pressure valve post reaction.

#### Depolymerization apparatus and operation

The base-catalyzed (chemical) lignin depolymerization experimental runs were conducted using a tailored bench-scale continuous flow reactor (CFR) as previously described in Abdelaziz et al. [28]. In short, the continuous plug flow reactor system consisted of

a Gilson 307 HPLC pump (Middleton, WI, USA) and Salamander tubular CFR reactor (Cambridge Reactor Design Ltd., Cottenham, UK) with a typical volume of 10 mL. The reactor was equipped with a preheater (8 mL) that ensures bringing the reactant mixture to the desired reaction temperatures. The product stream exiting the reactor is further condensed with the aid of a Julabo circulating water bath, and a pressure control valve made from nickel-molybdenum-chromium superalloy (Hastelloy C276; UNS N10276) was installed to maintain the pressure throughout the reaction. A filter was connected directly after the condensation step and attached to the inlet of the pressure control valve to protect it from any char depositions if occurred. The total operating system volume was about 50 mL. The setup was heated up and pressurized to the targeted working conditions, before starting each experiment and then the feed was pumped continuously to the plug flow reactor at volumetric flowrates of 10 and 5 mL/ min, respectively. These mixtures flowing through the continuous reactor system correspond to residence times in the heating zone of about 1 and 2 min, respectively. The operating conditions were chosen according to previous work by Abdelaziz et al. [28] i.e. the tel perature was below the coking point but high enough to give as high fraction of low molecular weight aromatic compounds as possible in the continuous lymerization process. The experimental conditions and parameters are summarized in Table 1. For operating parameters, the temperature was djusted using a Eurotherm temperature controller (Eurotherm, Ashburn, Virginia, USA) and the ressure was tuned through a backpressure regulator , ... nkhorst High-Tech B.V., Ruurlo, Netherlands, Ger exiting the reaction zone and throttling the product mixture/effluent to atmospheric pressure, to liquid depolymerized products (bio-oils) were style sly collected and samples were taken for analysi. The collected depolymerized product sample at three different temperatures (190, 220, 240 °C) were used as carbon sources in plate

ble Rase-catalyzed	lignin	depolymerization
eλ, rimental conditions		

Paramy cers	Range tested			
NaOH (wt%)	5			
Lignin loading (wt%)	5			
pH measure	12–14			
Temperature (°C)	190, 220, and 240			
Pressure (bar)	120-130			
Flowrate (mL/min)	5 and 10			

screening experiments. For liquid cultures, the collected lignin products treated at 220 °C and a flow rate of 5 mL/min were used.

#### **Bacterial strains**

Nine bacterial strains were used in this domonas putida KT2440 (DSM 6125), Pse. omonas fluorescens (DSM 50090), Rhoa ccus rythropolis (DSM 43066) and Rhodococcus opacus SM 1069) were purchased from the Deutsch Sammlung von Mikroorganismen und Zellkulturen Braunschweig, Germany. Pseudomonas putida L. 12 w. generously provided by Esteban Martínez-Garca t the Centro Nacional de Biotecnología, Maa Spain Four previously reported isolates (B, C, 9.1 and with tentative identifications, were used. Ison e B (Pseudomonas sp. strain B; DSM 104484) an is C (Pseudomonas plecoglossicida strain C; DSM 94486) were isolated from mature compost s les [29] and isolate 9.1 (Pseudomonas deceptionensis) and isolate 19 (Rhodococcus erythropolis) were isolated from sediments of the Baltic Sea [30].

#### Cure media and carbon source

M) medium was used for all the solid and liquid culture experiments. The media consists of M9 salts (containing per L: 6 g Na<sub>2</sub>HPO<sub>4</sub>, 3 g KH<sub>2</sub>PO<sub>4</sub>, 1 g NH<sub>4</sub>Cl, 0.5 g NaCl), 2 mM MgSO<sub>4</sub>, 100  $\mu$ M CaCl<sub>2</sub> and 10 mL/L trace element mixture [31, 32]. All the media components were either autoclaved or sterile filtered. The final pH was tuned to 7.

If lignin or depolymerized lignin ( $\overline{D}L$ ) was used as a carbon source, the pH was adjusted approximately to 7 using 50%  $H_2SO_4$ , before adding it to the M9 medium. At pH 7, the solubility of lignin was poor and hence precipitates were formed; although at the final concentration of 1 g/L, a complete solubility was observed. Henceforth, for liquid culture experiments, a concentration of not more than 1 g/L was used to enable biomass measurements.

If lignin model compounds were used as a carbon source, 50 mM stock solutions were prepared and stored (4 °C) for use within 2 weeks. Model compounds, if not soluble in water, were dissolved using a few drops of 5 N NaOH. All the chemicals, reagents and materials used were obtained from either VWR (West Chester, PA, USA) or Sigma-Aldrich (St. Louis, USA), unless specified.

#### **Growth on plates**

Plates were prepared using a final concentration of 1.5% agar in M9 medium. The carbon source utilized was either 4 g/L glucose, 5 mM vanillin/guaiacol or 1–5 g/L DL. The prepared plates were stored (4 °C) until further use. For screening of several organisms in a single plate, the plates were equally gridded and each grid was

inoculated with a single strain. The inoculated plates were sealed (parafilm) and incubated (30 °C).

The lignin (or lignin model compounds) plates were initially inoculated from a freshly grown glucose plate (in M9 medium). Later, when a few colonies were visible, it was re-streaked on to the fresh plates containing the same carbon source. This was performed to maintain the microbial adaptation to a particular carbon source.

#### Liquid culture experiments

Liquid culture experiments were conducted in 250 mL shake flasks containing 50 mL culture media. The carbon source added was either 10 g/L glucose (pre-culture), 5 mM guaiacol, 3 mM each of guaiacol, vanillin and 4-HBA (4-hydroxybenzoic acid), 1 g/L DL supplemented with 5 g/L glucose (high cell density experiments), or 1 g/L lignin/DL (low cell density experiments).

For the experiments with DL (or un-processed lignin), the flasks were inoculated using a single colony of the corresponding microorganism from 1 g/L DL plates. Shake flasks with model compounds as a carbon source were inoculated with a fixed amount of biomass, using 10 g/L glucose as pre-culture, to achieve an initial OD of around 0.2 or 0.5. All experiments were carried in duplicates. The flasks were incubated at 27 °C with agitation (180 rpm). Samples were withdrawn to sheady intervals to monitor the biomass density OD stical density), change in molecular weight of lignin (a construction of monomers (UHPLC-ultra-high performance liquid chromatography).

#### **Biomass measurements**

Biomass growth was measure actrophotometrically by optical density at 620 nm ( $\mathrm{OD}_{620}$ ). The color of lignin is dark and it significant, affect the absorbance measurements; hence, the rule as centrifuged to remove cells and the obtained apernatant was used as a blank prior biomass consurements. The samples, whenever required, were diluct with water/saline to stay in the linear range of optical density (0.03–0.3), in which case, the supernative was also diluted with the same factor. Subsequently, the rules were removed by centrifugation (3 min 12,  $0.0 \times g$ ) and the supernatants were kept at  $-20~\mathrm{C}$  for THPLC and SEC analysis.

## Calculation of yield and rates

Growth rate, uptake rate and yield were calculated for R. opacus on 5 mM guaiacol. Biomass dry weight was measured at the end of cultivation. This was used to convert the optical density to biomass concentration with a response factor of 0.4. Biomass yield ( $Y_{SX}$ ), expressed in both g/g and g/mmol, was calculated using the phase

plane plot of biomass produced and substrate utilized. The maximum specific growth rate ( $\mu$ ) (1/h) was determined from the plot of natural logarithm of biomass in the broth over time. The specific substrate uptake rate ( $q_{Substrate}$ ) was calculated by dividing  $\mu$  with  $T_{SX}$ .

#### **UHPLC** analysis

The frozen samples were thawed mixed and filtered (0.2  $\mu$ m pore size) before analysis. Waters Acquity UPLC system connected with a photod, as array detector (Waters, Milford, MA, SA) was operated for the analysis of phenolic cor pour. The column used for separation was Ethylene widged Hybrid C18 with a length of 100 mm, 1 mm in anal diameter and 1.7  $\mu$ m particle size. San ples were injected at a volume of 2.5  $\mu$ L and the temperature of the column was kept at 47 °C. The mobile phas was composed of 3% acetonitrile, 95% water, 2% acceptance (fraction A) and 85% acetonitrile, 13% water, 2% acceptance in a calcumate of the column was set of the column w

The analysmethod was obtained from Schwarz et al. [33] with minor modifications. A flow rate of 0.6 mL/min was used. The LC gradient elution method is as follows: '0% A at time 0, decreased to 90% A in 5 min, held at 9th A for 2 min and decreased to 25% A in 4.5 min. Following the gradient, the column was washed for 5 min with 100% B and was equilibrated for 5 min with 100% A. The chromatographic system was controlled by Acquity UPLC Console and the data were processed using Empower 3 software.

If unknown peaks appeared as a result of metabolic intermediates excretion, MS–MS analysis [30] was performed to identify and confirm the respective compounds. Later, the identified peaks were quantified using standards in UHPLC.

## Size exclusion chromatography

The molecular weight distributions (MWDs) of different lignin samples were determined employing a size exclusion chromatography (SEC) system, following an established method [34]. The setup adopted for the size measurements was a Waters 600E high-performance liquid chromatography (HPLC) system (Waters, Milford, MA, USA) equipped with a Waters 2414 refractive index detector, a Waters 486 ultraviolet (UV) tunable absorbance detector, and an analytical column packed with 30 cm of Superdex 30 and 30 cm of Superdex 200 (GE Healthcare, Uppsala, Sweden). The column was operating at ambient temperature and eluted with 125 mM NaOH solution (analytical grade) as the mobile phase at a flowrate of 1.0 mL/min. Calibration was carried out using polyethylene glycol (PEG) standards ranging from 400 to 35,000 g/mol in the eluent (Merck Schuchardt OHG, Hohenbrunn, Germany). The samples were diluted at concentrations of 1 mg/mL in the eluent and the solutions were filtered using a 0.2  $\mu m$  filter (Schleicher and Schuell, Dassel, Germany) to get rid of any suspended matter. Finally, about 500  $\mu L$  from the filtered solution was injected into the SEC system for data acquisition. Due to comparison with incorporated PEG standards, both the molecular weight and the molecular number should be interpreted relatively.

#### **Results**

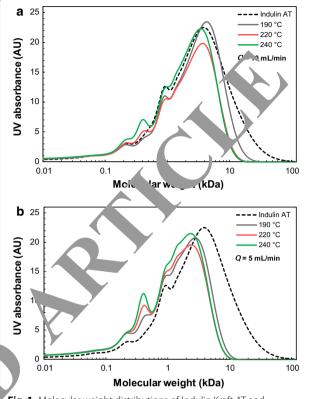
## Characterization of depolymerized lignin streams produced at different temperatures and residence times

The molecular weight distribution (MWD) of the depolymerized Kraft lignin as assessed by SEC was in good agreement with previously reported measurements by Abdelaziz et al. [28]. The analyses of samples obtained at three different temperatures, as well as the original (untreated) lignin are shown in Fig. 1.

The untreated Indulin Kraft lignin exhibited a rather wide-range distribution from 0.1 to 100 kDa with a peak molecular weight average (Mp) around 4 kDa—estimated based on the used PEG standards—and most of the starting lignin material had a molecular mass larger than 1 kDa (Fig. 1a). The MWDs of the DL products, on t. other hand, gradually shifted towards lower melecular mass components with increasing reaction terrora ure, and the fraction with a molecular weight higher than 10 kDa was very small at a depolymerization temperature of 240 °C. A bimodal curve distribution, when a second peak at 0.2-0.4 kDa—representing mainly me lomeric and dimeric fractions was shown at 220 °C. Increased lignin solubilization was obtained a be higher depolymerization temperatures, and higher reaction temperatures gradually yielded more monon, and dimeric compounds, in agreement with our previous work [28]. The shift of the MWD y as n are pronounced at the higher residence time (Fig. 1 and the fraction of components larger than 10 l Da was a set depleted already at a temperature of '90 The peak around 0.4 kDa was more distinct ard higher th a longer residence time (2 min), and inc eased more distinctly with reaction temperature. UHPLC Tysis of fractions collected after SEC revealed the esent of aromatic monomers in the 0.4 kDa n (Additional file 1: Figure S1). The monome were subsequently identified and quantified using UHP<sub>1</sub> C (see: Shake flask experiments with depolymerized lignin-Monomer analysis).

## Bacterial screening on agar plates with depolymerized lignin

To identify bacteria able to grow on DL, growth on plates was tested for nine bacteria/isolates previously known to metabolize aromatic compounds or isolated on



**Fig. 1** Molecular weight distributions of Indulin Kraft AT and depolymerized lignin products, measured as UV absorbance at 280 nm at two different residence times. The depolymerization reaction sets were carried out at temperatures 190, 220 and 240 °C with the flowrates of **a** 10 mL/min (residence time of 1 min) and **b** 5 mL/min (residence time of 2 min)

lignin-media [17, 29, 30, 35, 36]. The lignin used in this screening had been depolymerized at two different flow rates (5 and 10 mL/min) and three different temperatures (190, 220 and 240 °C) for each flow rate, i.e. at six different conditions. The concentration of DL used was 5 g/L, but there was no visible growth on plates for the first 3 weeks. Hence, another set of plate screening experiments were made with a reduced DL concentration (2 g/L). Within the first 2 weeks, colonies were formed by *R. opacus* on almost all plates—although only a few at each plate. After one additional week, some colonies were also seen for a few other organisms (Table 2). The growth on plates was somewhat ambiguous for some of the organisms, but was clearly seen when those organisms were re-streaked onto fresh plates with 1 g/L DL (not shown).

#### Shake flask experiments with depolymerized lignin

Rhodococcus opacus showed growth on the largest number of plates (12), followed by *P. fluorescens* (8). These two organisms were chosen for experiments with liquid

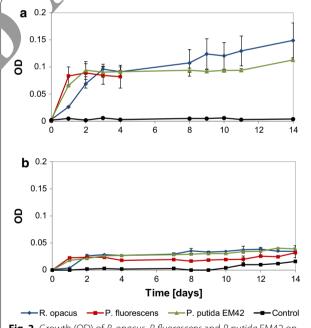
Organisms		5 mL/min				10 mL/min					No. of '+'		
	190 °C		220 °C		240 °C		190 °C		220 °C		240 °C		for each organism
	1	2	1	2	1	2	1	2	1	2	1	2	7
R. opacus	+	+	+	+	+	+	+	+	+	+	+	+	
R. erythropolis	+	_	-	-	+	+	_	+	+	_	-	( -	6
P. fluorescens	_	+	-	+	+	_	+	-	+	+	+	+	3
P. putida KT2440	-	-	-	+	-	-	-	-	-	+		7	3
P. putida EM42	_	_	+	+	-	+	+	-	+	- (	_	- 7	5
solate B	_	_	+	+	-	+	+	_	-	(+	-	<b>/</b> -	5
solate C	_	_	-	+	+	+	+	+	- ^			-	6
Isolate 9.1	+	-	-	-	-	+	_	+	-	-	<b>_</b>	+	4
Isolate 19	-	_	_	+	_	+	_	+		- )	_	_	4
No. of species showing growth on depo- lymerized lignin with different severities	4		7		8		8		8		4		

Table 2 Growth of bacterial strains on agar plates with 2 g/L depolymerized lignin in M9 medium

The lignin used was depolymerized with two different flow rates (5 and 10 mL/min) and three different flow rate. The plates were incubated for 3 weeks at 30 °C. The isolates were previously tentatively reported as *Pseudomonas* sp. (B); F. orglossicida (C); P. deceptionensis (9.1); and Rhodococcus erythropolis (19). Biological duplicates are indicated as 1 and 2. (+ growth; – no growth)

cultures on DL. In addition, P. putida EM42 was included due to the well-known robustness of its parental strair KT2440 [29]. Furthermore, the strain EM42 is stream lined for industrial applications by heavy genome editing and hence of interest as a host organism [37], M9 medium with DL 1 g/L, produced at a flow rate min and a temperature of 220 °C was used in these eximents. The yield of lower molecular weight anin compounds appear to be somewhat higher at 240 °C sed on the SEC analyses. However, this to perature is close to the coking point (250 °C) [28], and for reasons of operational stability temperature of 220 schosen. Low inoculum flask experiments (  $^{1}$  OD < 0.01) were conducted with both depolymerized and non-depolymerized lignin. A non-inoculated ontro lask was also included c in OD due to lignin to monitor potent. precipitation.

res of *I. fluorescens* reached a plateau The OD in a after 1 day at value of about 0.1, and similar trend was seen for P. putida EM42 and R. opacus, but on day 2 and 3, respectively (Fig. 2a). P. fluorescens formed aggregates after 4 day, and thereafter it was not possible to measure he call density by OD. The aggregates were recognized cuernal clusters from microscopic inspection. For all orga. Ins, formation of biofilms was seen at the air-liquid interface of the flask after 4 days. The biofilms were re-suspended before sampling. In case of non-depolymerized lignin, the increase in OD was lower, but a small growth was detected for all three organisms in the first 2 days, after which the OD stayed constant until day 14 (Fig. 2b). The maximum OD reached for non-depolymerized was approximately only one-third of that obtained



**Fig. 2** Growth (OD) of *R. opacus*, *P. fluorescens* and *P. putida* EM42 on 1 g/L **a** depolymerized (at 220 °C with 2 min residence time) lignin **b** non-depolymerized lignin as the only carbon source in M9 medium. A non-inoculated control is shown with a black line. Same scale on the *Y*-axis is maintained for easy comparison between the graphs. All experiments were performed in duplicates

with depolymerized lignin, which is in line with the lower availability of monomers/oligomers for direct consumption in the unprocessed lignin.

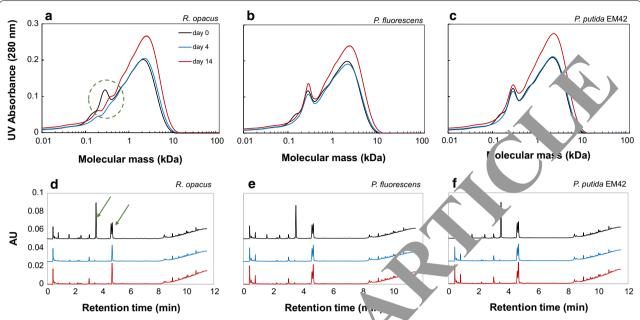


Fig. 3 SEC (a-c) and UHPLC (d-f) chromatograms of 1 g/L depolymerization lignin on day 0,4 and 14 after bacterial conversion by *R. opacus*, *P. fluorescens* and *P. putida* EM42. The bacterial cultures, which were inoculated with a single colony, were grown using depolymerized lignin as the only carbon source in M9 medium. In the UHPLC chromatograms (d-f) consults at retention time 1.6, 2.4, 3.5, 4.6 and 4.7 min were identified as 4-HBA, vanillate, vanillin, guaiacol and acetovanillone, respectively. The consultation of low molecular weight fraction (0.1–0.4 kDa) by *R. opacus* within 4 days is highlighted with a green dashed circle. The consumption of molecular weight fraction (0.1–0.4 kDa) by *R. opacus* is highlighted with green arrows

From SEC analyses, it was found that the low more ular weight fraction (0.1–0.4 kDa) of the DL cappeared within 4 days in the cultures with *P. opacus*, it dicating a possible consumption of these compounds (Fig. 3a). In the cultures of *P. fluorescens*, the cak for compounds between 0.1 and 0.2 kDa disappeared, and minor effects were possibly seen also for care fractions in the first 4 days (Fig. 3b). In the case of *P. putida* EM42, only compounds between 0.1 and 0.2 kDa disappeared, with no other change in the Case of *P. putida* EM42, only compounds between 0.1 and 0.2 kDa disappeared, with no other change in the Case of *P. putida* EM42, only compounds between 0.1 and 0.2 kDa disappeared, with no other change in the Case of *P. putida* EM42, only compounds between 0.1 and 0.2 kDa disappeared, with no other change in the Case of *P. putida* EM42, only compounds between 0.1 and 0.2 kDa disappeared, with no other change in the Case of *P. putida* EM42, only compounds between 0.1 and 0.2 kDa disappeared, with no other change in the Case of *P. putida* EM42, only compounds between 0.1 and 0.2 kDa disappeared, with no other change in the Case of *P. putida* EM42, only compounds between 0.1 and 0.2 kDa disappeared, with no other change in the Case of *P. putida* EM42, only compounds between 0.1 and 0.2 kDa disappeared, with no other change in the Case of *P. putida* EM42, only compounds between 0.1 and 0.2 kDa disappeared, with no other change in the Case of *P. putida* EM42, only compounds between 0.1 and 0.2 kDa disappeared, with no other change in the Case of *P. putida* EM42, only compounds between 0.1 and 0.2 kDa disappeared, with no other change in the Case of *P. putida* EM42, only compounds between 0.1 and 0.2 kDa disappeared, with no other change in the Case of *P. putida* EM42, only compounds between 0.1 and 0.2 kDa disappeared, with no other change in the Case of *P. putida* EM42, only compounds between 0.1 and 0.2 kDa disappeared, with no other change in the Case of *P. putida* EM42, only compounds be

#### Monome. aysi in depolymerized lignin

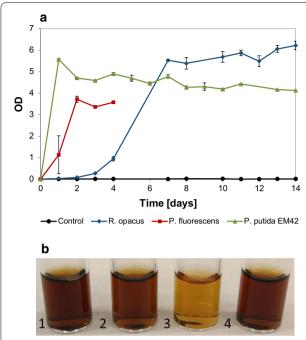
UPC and is showed the presence of several monoers some of which have previously been identified by su, critical fluid chromatography—mass spectrometry [28, 5]. The previously identified compounds found in the DL were 4-HBA (retention time 1.6 min), vanillate (2.4 min), vanillin (3.5 min), guaiacol (4.6 min) and acetovanillone (4.7 min). These monomer peaks were quantified by UHPLC and the yields based on initial lignin were found to be 3.5, 1.3, 0.95, 0.31, and 0.27 wt% respectively for guaiacol, vanillin, acetovanillone, vanillate and 4-HBA, which adds up to a total monomer yield

of 6.3 wt%. From the UHPLC chromatograms, it appears that all three organisms consumed 4-HBA, vanillate and vanillin within 4 days (Fig. 3d–f). In addition, *R. opacus* consumed guaiacol and an unknown compound giving a peak eluting at 0.8 min. None of the organisms consumed acetovanillone.

#### Flask experiments with depolymerized lignin and glucose

Experiments were also made in which glucose (5 g/L) was added as a second carbon source, allowing an initial growth phase and a higher biomass concentration for conversion of the DL (1 g/L). All organisms showed growth due to glucose (Fig. 4a), and the fastest growth was found for P. putida EM42 which obtained its maximum OD within 24 h. P. fluorescens achieved its maximum OD within 2 days. For *R. opacus* the initial growth phase lasted 5-7 days, but the OD kept increasing slowly throughout the entire experiment (Fig. 4a). Similar to the experiments with low cell density, P. fluorescens formed aggregates after day 4, and hence it was not possible to monitor the cell density beyond that time. Interestingly, there was a noticeable change in media color and a significant reduction in color was visible for P. fluorescens (Fig. 4b).

SEC analyses showed that the lower molecular weight fraction (between 0.1 and 0.4 kDa) was consumed by



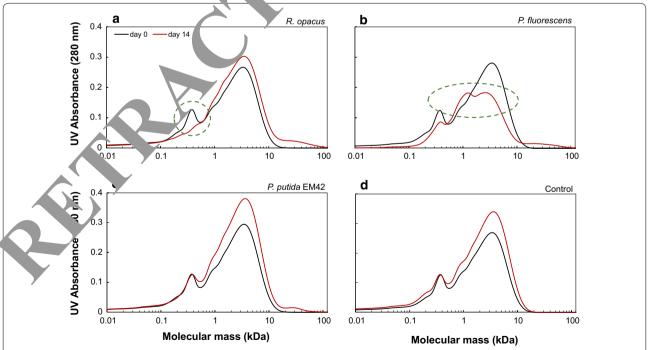
**Fig. 4** a Growth (OD) of *R. opacus*, *P. fluorescens* and *P. putida* EM42 on 1 g/L depolymerized (at 220 °C with 2 min residence time) lignin supplemented with 5 g/L glucose in M9 medium. A non-inoculated control is shown with a black line. **b** Change in media color on day 14 for 1-non-inoculated control; 2-*R. opacus*; 3-*P. fluorescens*; 4-*P. rutida* EM42

R. opacus, similar to the experiments without glucose (Fig. 5a). There was, in contrast, a large change in the SEC chromatograms in the cultivations of P. fluorescens. The peak height of the largest fraction (2–7 kDa) was markedly reduced. Part of the lower molecular weight fraction also remained unconsumed. The FC chromatogram of P. putida EM42 was almost the some as the control, i.e. no major conversion of any fraction was found. An overall increase in peak he but between 1 and 10 kDa was observed also for the control, which shows that this change is not due to microbial activity. A slight increase in molecular weight two n 15 and 100 kDa was observed for all the incomplete samples (Fig. 5).

The UHPLC chromatogram, show similar patterns for the conversion of the conomers by all the three organisms as in the periment without added glucose (Fig. 6). However, one significant difference was the occurrence of new large permanents at the property of the property of the experiments of the property of the proper

## Shake flack experiments with aromatic compounds

The UHILC analyses of shake flask experiments using L, indicate consumption of guaiacol by *R. opacus* (c. Figs. 3d, 6a), a not very common feature among microbes. Consumption of guaiacol was, therefore, tested in shake flask experiments of *R. opacus* using guaiacol (5 mM) as the only source of carbon and energy, and



**Fig. 5** SEC chromatograms of 1 g/L depolymerized lignin on day 0 and 14 after high cell density bacterial conversion by **a** *R. opacus* **b** *P. fluorescens* **c** *P. putida* EM42 and **d** non-inoculated control in M9 medium. The consumption of low molecular weight fraction (0.1–0.4 kDa) by *R. opacus* and breakdown of high molecular weight fraction (1–10 kDa) by *P. fluorescens* are highlighted with green dashed circles

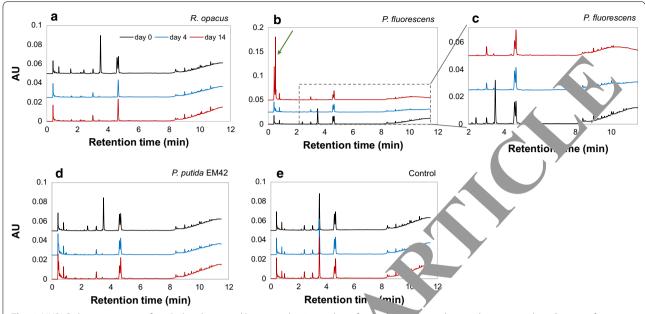
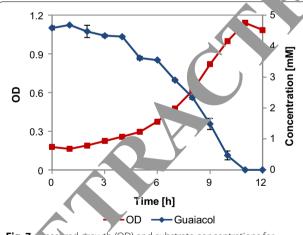


Fig. 6 UHPLC chromatograms of 1 g/L depolymerized lignin on day 0, 4 and 14 after high cere density bacterial conversion by **a** *R. opacus* **b**, **c** *P. fluorescens* **d** *P. putida* EM42 **e** non-inoculated control in M9 medium. The peaks at retention time 1.6, 2.4, 3.5, 4.6 and 4.7 min were identified as 4-HBA, vanillate, vanillin, guaiacol and acetovanillone, respectively. The occurrence of 1 ew large peaks at retention times 0.4–0.5 min for *P. fluorescens* on day 14 are highlighted with a green arrow. For a better value. If the newly formed peaks on day 14, the chromatograms are arranged in reverse order (bottom to top) for **b**, **c** 

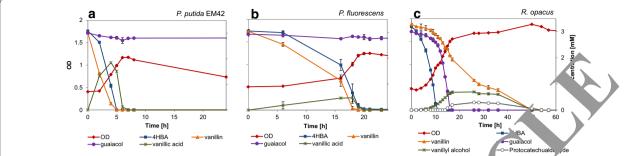


**Fig. 7** past red growth (OD) and substrate concentrations for cultures a pace grown on guaiacol as the only source of carbon mediu. The standard deviation of duplicate experiments is indicated with error bars

growth on guaiacol was indeed confirmed (Fig. 7). Inoculum was taken from agar plates with guaiacol and no lag phase was observed in the shake flask cultures. A specific growth rate of 0.2  $(\pm\,0.001)~h^{-1}$  and a biomass yield of 0.7  $(\pm\,0.02)~g_{cdw}/g$  was obtained and a complete conversion of guaiacol was achieved with a specific conversion rate of 2.4 (mmol/g<sub>cdw</sub>/h).

The main identified low molecular weight aromatic compounds in the DL apart from guaiacol were vanillin, vanillate and 4-HBA (4-hydroxybenzoate). Additional shake flask experiments were made, in which a mixture of 3 mM each of guaiacol, vanillin and 4-HBA served as carbon and energy source for growth of *R. opacus*, *P. fluorescens* and *P. putida* EM42. These three compounds also represent different branches of the upper funneling/β-ketoadipate pathways [1].

The fastest growth was obtained for P. putida EM42, which reached its maximum OD in 6 h (Fig. 8a). Vanillin and 4-HBA were converted/consumed almost simultaneously. Vanillin was converted to vanillic acid and the excreted vanillic acid was subsequently taken up (Fig. 8a). Guaiacol, however, remained unconsumed even after 200 h. P. fluorescens showed a somewhat longer lag phase, but after this, vanillin and 4-HBA were simultaneously converted/taken up (Fig. 8b). The conversion of vanillin to vanillic acid was similar to that of P. putida EM42 and guaiacol remained unconsumed for 200 h. Multiauxic growth was observed in case of R. opacus, which reached its maximum OD in around 50 h (Fig. 8c). The substrate consumption was sequential starting with 4-HBA, which was consumed in 11 h. After the depletion of 4-HBA, guaiacol was consumed within 17 h. Interestingly, R. opacus converted vanillin to vanillyl alcohol—partly concomitant with the conversion of 4-HBA and guaiacol. A



**Fig. 8** Growth (OD) and consumption of model compounds by **a** *P. putida* EM42 **b** *P. fluorescens* **c** *R. opacus* in M9 edium. 3 mM each of guaiacol, 4-HBA and vanillin were provided as the only source of carbon. The standard deviation of duplicate experiments is a sated with error bars

slight excretion of protocatechualdehyde was observed in the medium. The remaining vanillin and the excreted protocatechualdehyde and vanillyl alcohol were all consumed around 50 h (Fig. 8c).

#### Discussion

The present study aimed to identify bacterial strains capable of bioconversion of Indulin AT-a technical softwood Kraft lignin. Kraft lignin is a complex substrate with high molecular weight making it a difficult si strate for direct microbial uptake, and therefore, a reviously developed alkaline depolymerization method was employed to break down a part of the higher n. weight lignin fragments into compounds of lower m ular weight [28]. The nine bacterial strain. rolates (cf. Table 2) examined here have been priviously ported to metabolize lignin-related compounds [17, 21, 29, 35, 36, 39], and fall into two phyla (Pre-eobacteria and Actinobacteria), with a significant evaluation distance between them. During the p screening with DL, an initial concentration of 5 g/L was an eady inhibitory and hence no growth was or rved within the first 3 weeks. It is well-known the sever compounds, e.g. vanillin, are inhibnory microbial growth and metabolism of many begin and yeasts [40]. However, the total concentration of q artified aromatic monomers was only in the range of few nullimoles (2.3 mM) in 5 g/L DL, well concentrations of the defined medium tested at which I. vti aa EM42, P. fluorescens, and R. opacus all rew Fig. 8). This suggests that the inhibition is due to presence of other unknown toxic compounds present the depolymerisate. When the concentration was decreased to 2 g/L DL, a few colonies appeared in 2 weeks. An adaption of at least 2 weeks was essential for the organisms to initiate growth on DL plates (Table 2). Upon re-streaking to new lignin plates, the growth was rather quick (3 days).

In liquid cultures of *R. opacus, P. fluorescens* and *P. putida* EM42, the initial increase in OD was due to the

uptake of the r ao. available low molecular weight lignin compounds in the L. Later, there was a stationary phase for 8 10 c.ys, after which a slight further growth could be see (1.5. ...). Possibly, enzymes for the breakdown of HMV 'higher molecular weight) lignin were uring this time and LMW (lower molecular expres weight) fragnients were obtained. It was not possible to measure be growth of *P. fluorescens* after the first 3 days he to the aggregation of cells in the culture medium. Sh behavior of cell aggregation and increased hydroph bicity has previously been observed in *P. putida* CP1 [41, 42] and P. fluorescens [21] when grown under stressful conditions. Due to the reduction in size of cells under these conditions, the increase in cell counts were in disagreement with the cell dry weight and optical density

The SEC analyses at day 14 indicated an increase in the average molecular weight (Fig. 3). The main change—at 1-10 kDa-was observed not only for the inoculated cultures, but also for the non-inoculated lignin (cf. Figure 5d). This phenomenon is, therefore, probably not a result of microbial action, but may be due to instability of lignin fragments, which undergoe reconfiguration and results in higher UV absorbance. The slight increase in molecular weight (in the range of 15–100 kDa) for all the inoculated flasks in high cell density cultures (Fig. 5a-c) is probably due to the secretion of extracellular laccases. Even though the secreted extracellular enzymes are expected to break down the higher molecular weight lignin, they are also known for polymerizing phenolic compounds [43]. The balance between the depolymerizing and polymerizing abilities of these enzymes depends on several factors (reaction temperature, origin of enzyme, feedstock structure, etc.) and have been previously reported [44].

The use of cultures with glucose as a co-substrate together with DL revealed the ability of *P. fluorescens* to break down the HMW fraction (Fig. 5b). Both the reduction in color of the DL media (Fig. 4b) and the

reduction of peak size in the UHPLC chromatograms after 10 min (Fig. 6c) clearly showed that this organism degrades HMW lignin fragments. It was not determined to what extent the degradation products were consumed. Additional peaks appearing in the UHPLC chromatograms indicated that at least not all the generated compounds were consumed (Fig. 6b). This is in agreement with reports by Salvachúa et al. on *P. fluorescens* Pf-5, which was found able to depolymerize HMW lignin, but was not able to consume the produced monomers [21]. In contrast to a few previous reports on the ability of P. putida KT2440 to breakdown HMW lignin [20, 21], no such behavior was observed for EM42 strain, despite its genetic similarities with KT2440 [37]. This could be due to the different lignin source used in this study, since the lignin obtained from corn stover differs much from the technical lignin (Indulin AT) in terms of the structure, primary building blocks and successive breakdown products [45, 46]. R. opacus has previously been reported to break down HMW corn stover lignin to some extent, but a larger conversion was obtained with *R. jostii* in the same study [24]. Co-fermentation using both these strains furthermore proved to increase the conversion significantly [24].

Laccases and peroxidases are known to be responsible for higher molecular weight lignin breakd wn [47] 48]. Hence, BLAST (Basic Local Alignment Searce Cool) searches [49] were performed using well haracter, ed DyP-type peroxidases [50, 51] and lace ses [2] against the genome of P. fluorescens, P. putiau KT244 and R. opacus to discover if there are any putative enzymes responsible for lignin breakdow. (Additional file 1: Table S1 and S2). Two enzymes from the subfamily of DyPA and three from DyPB \ lected from R. jostii RHA1 and *P. fluorescens* Pf-5 [16, 53, 54]. As the enzymes from subfamily DyPC an DyPD were not found/annotated either in Pseu w or Rhodococcus group of organisms, those were lected from the well-studied Amycolatopsis s. 75iv2 and Bjerkandera adusta, respectively [50, 55]. Additionally, two laccases were selected, one from P. fluorescens and another from R. opacus PD630. seem that laccases, DyPA and DyPB are with pres + (Note the absence of DyPA in KT2440) all he three organisms used in this study (Additional Table S1 and S2). However, the inability of R. opacus a. P. putida to depolymerize the HMW lignin to the expected level might be due to the inefficiency of enzyme secretion or due to the lack of gene expression in the given environment. P. fluorescens was the only organism in which proteins similar to DyPC and DyPD were found (Additional file 1: Table S1). Steady-state enzyme kinetics of the DyP subclasses has revealed that C and D type DyPs have higher peroxidase activities than A and B type

DyPs [54], possibly explaining the much more efficient degradation of high molecular weight lignin fractions by *P. fluorescens* than by the other organisms studied.

The growth of organisms on the mixture of model compounds was in good agreement with their growth on DL—regarding the consumption r.ono ners. Depending on both the origin of lignin, its means of separation and the method of depolyr rization a very different distribution between the formal low molecular weight compounds can be obtained. Prevously identified compounds in the depolyme ized softwood lignin samples using the depolyme zati method employed here, have been 4-HBA, venilla vanillin and guaiacol [38]. Similar compounds were also eported in other related substrates (Table 3). •se compounds belong to different branches the up er funneling pathways, which mostly con orge of either protocatechuic acid or catechol that eventua. enter the beta-ketoadipate pathway and promote cell greath [56]. In this study, 4-HBA, vanillin and value were consumed by all the three organisms, shown 15th from liquid cultures on DL and in model media. 4-VBA, which belongs to the p-coumaryl branch the upper funneling pathway, has been reported to be co verted by a number of microbes, including Acinetobocter baylyi, Cupriavidus necator, P. aeruginosa, and is apparently not very toxic [29, 57–59]. In contrast, vanillin and guaiacol (belonging to the coniferyl branch of the funneling pathway) have been reported to be quite toxic to many organisms [60-65]. Microbial detoxification mechanisms of vanillin are related to conversion of its aldehyde group either by oxidation to vanillic acid or by reduction to vanillyl alcohol [66]. The conversion of vanillin towards the less toxic intermediate vanillic acid is common in *Pseudomonas* species [29, 67]. The vanillic acid is—possibly following a period of excretion and accumulation in the medium-further converted to protocatechuic acid entering the beta-ketoadipate pathway. This was observed in our liquid cultivations for P. fluorescens and P. putida EM42 (Fig. 8a, b). Strong effects on gene expression have been reported when vanillin was provided as a sole carbon source to P. putida KT2440. The expression of more than 600 genes was changed, including the beta-ketoadipate pathway but also the central energy metabolism and genes associated to solvent tolerance [68].

Interestingly, in our study *R. opacus* converted vanillin to vanillyl alcohol in the presence of other carbon sources (guaiacol and 4-HBA) (Fig. 8c). The detoxification of vanillin to vanillyl alcohol is mainly reported in yeasts [64], but also in some bacterial species such as *Gluconacetobacter xylinus*, which converted vanillin (0.5 mM) to vanillyl alcohol at 80% yield, in the presence of 25 g/L glucose [63]. In the current study, the excretion of vanillyl

Table 3	Some studies reporting	g guaiacol, vanillii	n and related compou	unds in depolymerized lignin
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Substrate	Depolymerization method	Major monomers	References
Steam treated softwood and hemp lignin	Base-catalyzed depolymerization	Guaiacol, vanillin, catechol	[69]
Organosolv lignin	Base-catalyzed depolymerization	Guaiacol, syringol, syringaldehyde,	[70]
Spruce lignin	Hydrogenolysis/extraction	Ethylguaiacol, propylguaiacol,	[71]
Dealkaline lignin	Solid acid catalysts	Vanillin, guaiacyl acetone, homovanillic acid	[72]
Birch wood lignin	Hydrogenolysis with Ni-based catalysts	Propyl guaiacol and propyl syrings	[/3]
Organosolv switchgrass lignin	Depolymerization and hydrodeoxygenation with formic acid and Pt/C	Guaiacol, 4-methylguaiacol, homo illyl alcohol	[74]
Organosolv hardwood lignin	Depolymerization in supercritical carbon dioxide/acetone/water fluid	Guaiacol, 2-methoxy-4- ethyl-phenol, syringol	[75]
Kraft lignin	Oxidation in aqueous methanol at acidic pH	Vanillin, methyl va. te	[76]
Kraft lignin (Indulin AT)	Base (NaOH) catalyzed depolymerization at 220 $^{\circ}\mathrm{C}$	Guaiacol, vanillin, aceta nillone, vanillate, 4-hydroy, nzoate	This study

alcohol stopped after exhaustion of 4-HBA and guaiacol and the remaining vanillin was metabolized within 50 h. Enzymes responsible for vanillin degradation (vanillin dehydrogenase, vanillate O-demethylase and vanillylalcohol oxidase) have been identified in *Rhodococcus*, but during growth on vanillin, vanillate O-demethylase was found to be more upregulated than vanillin dehydro nase [77]. The inefficient expression of *vdh* (vanillin dehy drogenase) is a possible reason for the detoxification of vanillin to vanilly alcohol instead of the more vanillic acid. The slight generation of protocated dehyde could be due to the action of the basical demethylase system, which must be highly express in this mixture. This system could recognize vanillin with lower affinity removing its methoxy moley and giving rise to small amounts of protocatechualdeh

Guaiacol is one of the ma depolymerization products in alkaline treatment of softy of a lignin [38]. R. opacus was the only organitested which showed growth on guaiacol plates ddi file 1: Table S3). When grown in liquid cultu. the growth rate on guaiacol was  $0.2 \text{ h}^{-1}$ , ich is sightly higher than the growth of the same strangen vanillin (0.127  $h^{-1}$ ) and 4-HBA (0.126 h<sup>-/</sup>) [17]. Some species of *Rhodococcus* have previously 2 n poi ted to be able to grow on guaiacol [78– 80] or poses seenes related to guaiacol breakdown [81]. The quence of the enzyme (Cytochrome P450, Uni-4A0P3) responsible for the conversion of guaiacol catechol from R. rhodochrous (ATCC 21198) [81] was blasted against the genome of the R. opacus (taxid: 37919) and an identity of 81% was found, which strongly indicates the presence of a similar protein. When the same enzyme was blasted against the genome of KT2440, no similar protein was found and with P. fluorescens (taxid: 294) only a protein with low identity (28%) was found. Pseudomonas sp. strain PN-1 has been reported to convert a piace of catechol under *anaerobic* conditions [82]. It was suggested that the anaerobic demethylation sy of this species has a broader specificity in the degradation of lignin molecules than the aerobic enzyme [82]. *P. putida* KT2440 possesses genes encoding proteins necessary to metabolize catechol [18], and the addituor of genes responsible for the guaiacol-demethylating cy ochrome P450 might be very interesting to expand the ange of substrates which the organism can utilize. As the softwood lignin contains a significant amount of G-type lignin, the utilization of guaiacol remains essential.

## **Conclusions**

Lignins of different origin vary significantly in their composition, and the fragments generated after depolymerization are strongly affected by the type and severity of the treatment. Softwood Kraft lignin is a rather complex substrate, which has been less studied than corn stover lignin for biological valorization. Here, microbial conversion of alkalinetreated Indulin AT—a softwood-based Kraft lignin—was demonstrated. Bacterial growth on the pretreated (partly depolymerized) lignin was clearly better than the growth on unprocessed lignin. Of the bacterial strains tested, R. *opacus* was able to consume most of the LMW compounds and HMW lignin was also converted to some extent. P. fluorescens showed a considerable ability for breakdown of HMW lignin. Even more interesting was the ability of R. opacus to consume guaiacol, which is monomer present in appreciable amounts after the depolymerization of softwood-based lignin. The yields of monomeric compounds obtained after depolymerization of softwood Kraft lignin typically amount to only a few percent, and this would need to be substantially increased for any commercial process. However, the organisms of this study are certainly of interest—either as potential host organisms, or sources of genetic material, in development of microbes for the production of fine/bulk chemicals from softwood lignin.

#### **Additional file**

Additional file 1: Figure S1. (a) The SEC chromatogram of 1 g/L depolymerized (at 220 °C, 5 mL/min) lignin. Red dotted lines represent the fractions collected. The collected fraction at 95-108 min corresponds to the 0.2-0.4 kDa peak in the SEC chromatograms calibrated with PEG standards. (b) UHPLC chromatograms of the fractions obtained from SEC. The peaks in the fraction 95-108 min correspond to aromatic monomers (Vanillin-3.5 min; guaiacol-4.6 min; acetovanillone-4.7 min). Table S1. Results of homology BLAST with the previously well-characterized DyP proteins against the genome of the organisms used in this study. P. putida EM42 strain used in this study is the modified version of KT2440 and hence the genome of KT2440 (parental strain) was used for BLAST searches. Proteins with identity more than 75% are emphasized in green. Proteins that were found absent and the ones with less than 30% query are highlighted in red. P. fluorescens highlighted in blue is the only organism with proteins similar to DyPC and DyPD. Table S2. Results of homology BLAST with laccases against the genome of the organisms used in this study. P. putida EM42 strain used in this study is the modified version of KT2440 (parental strain) and hence the genome of KT2440 was used for BLAST searches. Proteins with identity more than 75% are emphasized in green. Table S3. Growth of bacterial strains on agar plates with 5 mM vanillin/guaiacol as the only source of carbon in M9 medium, incubated for 2 weeks at 30 °C. The isolates were previously tentatively reported as *Pseudomonas* sp. (B): P. plecoglossicida (C); P. deceptionensis (9.1); and Rhodococcus erythropolis (19). Duplicate experiments were performed and the growth results were similar (++ abundant growth; + growth; - no growth). R. opacus x only organism to show growth on guaiacol (highlighted).

#### **Abbreviations**

DL: depolymerized lignin; SEC: size exclusion chromator aphy. O: optical density; MWD: molecular weight distribution; UHPLC: litra-high; commance liquid chromatography; UV: ultraviolet; LMW: low nolecular weight, HMW: high molecular weight; BLAST: Basic Local Alignic ent Search Tool; CFR: continuous flow reactor; 4-HBA: 4-hydroxybenzo o: PEG: polyethylene glycol; HPLC: high performance liquid chromatography.

## Authors' contributions

GL, CH and MGG conceived the project. Kk, CA, JG, H and GL designed the study. KR, OA and MN performed experiments. All authors discussed and interpreted the results. KR and OA confed the manuscript. GL, CH and JGH revised the manuscript. All authors disproved the final manuscript.

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### Competing interests

The authors declare that they have no competing interests.

#### Availability of supporting data

The datasets used and/or analyzed during the current study are available from the corresponding author on reasonable request.

#### Consent for publication

Not applicable

#### Ethics approval and consent to participate

This article does not contain any studies with human participal performed by any of the authors.

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#### Referen

- Abde azızınınk DP, Prothmann J, Ravi K, Sun M, García-Hidalgo J, Sancanl M, Hulteberg CP, Turner C, Lidén G, Gorwa-Grauslund MF. Biological valorization of low molecular weight lignin. Biotechnol Adv. 2016;34:1318–46.
- Zakzeski J, Bruijnincx PC, Jongerius AL, Weckhuysen BM. The catalytic valorization of lignin for the production of renewable chemicals. Chem Rev. 2010;110:3552–99.
- Rinaldi R, Jastrzebski R, Clough MT, Ralph J, Kennema M, Bruijnincx PCA, Weckhuysen BM. Paving the way for lignin valorisation: recent advances in bioengineering, biorefining and catalysis. Angew Chem Int Ed. 2016;55:8164–215.
- Upton BM, Kasko AM. Strategies for the conversion of lignin to high-value polymeric materials: review and perspective. Chem Rev. 2016:116:2275–306.
- Beckham GT, Johnson CW, Karp EM, Salvachúa D, Vardon DR. Opportunities and challenges in biological lignin valorization. Curr Opin Biotechnol. 2016;42:40–53.
- Ragauskas AJ, Beckham GT, Biddy MJ, Chandra R, Chen F, Davis MF, Davison BH, Dixon RA, Gilna P, Keller M, et al. Lignin valorization: improving lignin processing in the biorefinery. Science. 2014;344:1246843.
- Schutyser W, Renders T, Van den Bosch S, Koelewijn SF, Beckham GT, Sels BF. Chemicals from lignin: an interplay of lignocellulose fractionation, depolymerisation, and upgrading. Chem Soc Rev. 2018;47:852–908.
- Sjostrom E. Wood chemistry. 2nd ed. San Diego: Academic Press; 1993. p. 71–89.
- Chakar FS, Ragauskas AJ. Review of current and future softwood kraft lignin process chemistry. Ind Crops Prod. 2004;20:131–41.
- Bugg TD, Rahmanpour R. Enzymatic conversion of lignin into renewable chemicals. Curr Opin Chem Biol. 2015;29:10–7.
- Linger JG, Vardon DR, Guarnieri MT, Karp EM, Hunsinger GB, Franden MA, Johnson CW, Chupka G, Strathmann TJ, Pienkos PT, Beckham GT. Lignin valorization through integrated biological funneling and chemical catalysis. Proc Natl Acad Sci USA. 2014;111:12013–8.
- 12. Shi Y, Yan X, Li Q, Wang X, Liu M, Xie S, Chai L, Yuan J. Directed bioconversion of Kraft lignin to polyhydroxyalkanoate by Cupriavidus basilensis B-8 without any pretreatment. Process Biochem. 2017;52:238–42.
- Zhao C, Xie S, Pu Y, Zhang R, Huang F, Ragauskas AJ, Yuan JS. Synergistic enzymatic and microbial lignin conversion. Green Chem. 2016;18:1306–12.
- Barton N, Horbal L, Starck S, Kohlstedt M, Luzhetskyy A, Wittmann C. Enabling the valorization of guaiacol-based lignin: integrated chemical and biochemical production of cis, cis-muconic acid using metabolically engineered Amycolatopsis sp ATCC 39116. Metab Eng. 2018;45:200–10.

- Shi Y, Chai L, Tang C, Yang Z, Zhang H, Chen R, Chen Y, Zheng Y. Characterization and genomic analysis of kraft lignin biodegradation by the beta-proteobacterium *Cupriavidus basilensis* B-8. Biotechnol Biofuels. 2013;6:1.
- Ahmad M, Roberts JN, Hardiman EM, Singh R, Eltis LD, Bugg TD. Identification of DypB from *Rhodococcus jostii* RHA1 as a lignin peroxidase. Biochemistry. 2011;50:5096–107.
- Kosa M, Ragauskas AJ. Bioconversion of lignin model compounds with oleaginous *Rhodococci*. Appl Microbiol Biotechnol. 2012;93:891–900.
- Jiménez JI, Miñambres B, García JL, Díaz E. Genomic analysis of the aromatic catabolic pathways from *Pseudomonas putida* KT2440. Environ Microbiol. 2002;4:824–41.
- Vardon DR, Franden MA, Johnson CW, Karp EM, Guarnieri MT, Linger JG, Salm MJ, Strathmann TJ, Beckham GT. Adipic acid production from lignin. Energy Environ Sci. 2015;8:617–28.
- Rodriguez A, Salvachua D, Katahira R, Black BA, Cleveland NS, Reed M, Smith H, Baidoo EEK, Keasling JD, Simmons BA, et al. Base-catalyzed depolymerization of solid lignin-rich streams enables microbial conversion. Acs Sustain Chem Eng. 2017;5:8171–80.
- Salvachúa D, Karp EM, Nimlos CT, Vardon DR, Beckham GT. Towards lignin consolidated bioprocessing: simultaneous lignin depolymerization and product generation by bacteria. Green Chem. 2015;17:4951–67.
- Le RK, Wells T Jr, Das P, Meng X, Stoklosa RJ, Bhalla A, Hodge DB, Yuan JS, Ragauskas AJ. Conversion of corn stover alkaline pre-treatment waste streams into biodiesel via Rhodococci. RSC Adv. 2017;7:4108–15.
- Kosa M, Ragauskas AJ. Lignin to lipid bioconversion by oleaginous Rhodococci. Green Chem. 2013;15:2070.
- He Y, Li X, Ben H, Xue X, Yang B. Lipid production from dilute alkali corn stover lignin by rhodococcus strains. ACS Sustain Chem Eng. 2017;5:2302–11.
- Numata K, Morisaki K. Screening of marine bacteria to synthesize polyhydroxyalkanoate from lignin: contribution of lignin derivatives to biosynthesis by *Oceanimonas doudoroffii*. ACS Sustain Chem F 2015:3:569–73.
- Gellerstedt G. Softwood kraft lignin: raw material for the future of ps Prod. 2015;77:845–54.
- Hu Z, Du X, Liu J, Chang H-M, Jameel H. Structural characterization opine kraft lignin: bioChoice lignin vs indulin AT. J W on m Technøl. 2016;36:432–46.
- 28. Abdelaziz OY, Li K, Tunå P, Hulteberg CP. Continuous catalytic expolymerisation and conversion of industrial kraft ligran into low-molecular-weight aromatics. Biomass Convers Biorefinery. 20 3:455–70.
- Ravi K, Garcia-Hidalgo J, Gorwa-Grauslund M, Jen G Conversion of lignin model compounds by *Pseudomonas putiua* x12440 and isolates from compost. Appl Microbiol Biote. 2017;101:5059–70.
- Ravi K, Garcia-Hidalgo J, Nobel M, Gonva Gra, slund MF, Liden G. Biological conversion of aromatic collignol compounds by a Pseudomonas isolate from sediments of the litic Sea. AMB Express. 2018;8:32.
- Sambrook J, Russell D. Porting a laboratory manual. Cold Spring Harbor: Cold Spring aboratory Press; 2001.
- 32. Pfennig N, Lipp KD. Über a itamin B12-Bedürfnis phototropher Schwefelbal erie. Tchiv für Mikrobiologie. 1966;55:245–56.
- 33. Schwarz M, Rodrígue C Guillén DA, Barroso CG. Development and validation of UPLC for tye determination of phenolic compounds and furar oder vatives in *Brandy de Jerez*. J Sep Sci. 2009;32:1782–90.
- 34. Abdel. 17, Flul Loerg CP. Physicochemical characterisation of technilignins. Their potential valorisation. Waste and Biomass Valoriz. 20, 78:859–03.
- W g G, Huang F, Kosa M, Huang D, Ragauskas AJ. Bioconversion of vgen-pretreated Kraft lignin to microbial lipid with oleaginous *Rhodocoe opacus* DSM 1069. Green Chem. 2015;17:2784–9.
- Taylor CR, Hardiman EM, Ahmad M, Sainsbury PD, Norris PR, Bugg TD. Isolation of bacterial strains able to metabolize lignin from screening of environmental samples. J Appl Microbiol. 2012;113:521–30.
- Martinez-Garcia E, Nikel PI, Aparicio T, de Lorenzo V. Pseudomonas 2.0: genetic upgrading of P. putida KT2440 as an enhanced host for heterologous gene expression. Microb Cell Fact. 2014;13:159.
- Prothmann J, Sun M, Spegel P, Sandahl M, Turner C. Ultra-highperformance supercritical fluid chromatography with quadrupoletime-of-flight mass spectrometry (UHPSFC/QTOF-MS) for analysis of

- lignin-derived monomeric compounds in processed lignin samples. Anal Bioanal Chem. 2017;409:7049–61.
- 39. Nelson KE, Weinel C, Paulsen IT, Dodson RJ, Hilbert H, Martins dos Santos VA, Fouts DE, Gill SR, Pop M, Holmes M, et al. Complete genome sequence and comparative analysis of the metabolically versatile *Pseudomonas putida* KT2440. Environ Microbiol. 2002;4:799–508.
- Franden MA, Pilath HM, Mohagheghi A, Pienkos PT, Z'ang M. Inhibition of growth of Zymomonas mobilis by model compound of and in gnocellulosic hydrolysates. Biotechnol Biofuels. 2013;6:99.
- 41. Fakhruddin ANM, Quilty B. Measurement of the growth of an ac forming bacterium *Pseudomonas putida* CP1. Biodeg. https://doi.org/18.189-97.
- Farrell A, Quilty B. Substrate-dependent toag nation of Pseudomonas putida CP1 during the degradation of many-chlorophenols and phenol. J Ind Microbiol Biotecthol. 2002;28;316–24.
- 43. Ghoul M, Chebil L. Enzymatic polyresization of phenolic compounds by oxidoreductases. Dordrecht singe. 1–46.
- 44. Brzonova I, Kozliak EI, Andriano A, LaVallie A, Kubátová A, Ji Y. Production of lignin based in stuble poly. (anionic hydrogels) by *C versicolor*. Sci Rep. 2017;7:1750/.
- 45. Constant S, Wienk, LJ, Fn. AE, de Peinder P, Boelens R, van Es DS, Grisel RJH, Weckhovsen BM, J, gen WJJ, Gosselink RJA, Bruijnincx PCA. New insights into the structure and composition of technical lignins: a comparation of harmonic on study. Green Chem. 2016;18:2651–65.
- Katahira R, Mr. A, McKinney K, Chen XW, Tucker MP, Johnson DK, Beckham GT. Basa dalyzed depolymerization of biorefinery lignins. ACS Sust page 2016;4:1474–86.
- 47. de Gonza Colpa DI, Habib MH, Fraaije MW. Bacterial enzymes involve i in lignin degradation. J Biotechnol. 2016;236:110–9.
- 48. Brown N.C, Chang MC. Exploring bacterial lignin degradation. Curr Opin Chem Biol. 2014;19:1–7.
- Altschul SF, Gish W, Miller W, Myers EW, Lipman DJ. Basic local alignment search tool. J Mol Biol. 1990;215:403–10.
- 50 Yoshida T, Sugano Y. A structural and functional perspective of DyP-type peroxidase family. Arch Biochem Biophys. 2015;574:49–55.
- 51. Colpa DI, Fraaije MW, van Bloois E. DyP-type peroxidases: a promising and versatile class of enzymes. J Ind Microbiol Biotechnol. 2014;41:1–7.
- 52. Chauhan PS, Goradia B, Saxena A. Bacterial laccase: recent update on
- production, properties and industrial applications. 3 Biotech. 2017;7:323.

  53. Rahmanpour R, Bugg TD. Characterisation of Dyp-type peroxidases from
- Pseudomonas fluorescens Pf-5: oxidation of Mn(II) and polymeric lignin by Dyp1B. Arch Biochem Biophys. 2015;574:93–8. 54. Roberts JN, Singh R, Grigg JC, Murphy MEP, Bugg TDH, Eltis LD. Charac-
- terization of dye-decolorizing peroxidases from *Rhodococcus jostii* RHA1. Biochemistry. 2011;50:5108–19.
- Brown ME, Barros T, Chang MC. Identification and characterization of a multifunctional dye peroxidase from a lignin-reactive bacterium. ACS Chem Biol. 2012;7:2074–81.
- The eLignin Microbial Database. http://www.elignindatabase.com/. Accessed 12 Apr 2018.
- Fischer R, Bleichrodt FS, Gerischer UC. Aromatic degradative pathways in Acinetobacter baylyi underlie carbon catabolite repression. Microbiology. 2008;154:3095–103.
- Pérez-Pantoja D, De la Iglesia R, Pieper DH, González B. Metabolic reconstruction of aromatic compounds degradation from the genome of the amazing pollutant-degrading bacterium *Cupriavidus necator JMP134*. FEMS Microbiol Rev. 2008;32:736–94.
- Kuhnigk T, Konig H. Degradation of dimeric lignin model compounds by aerobic bacteria isolated from the hindgut of xylophagous termites. J Basic Microbiol. 1997;37:205–11.
- Zaldivar J, Martinez A, Ingram LO. Effect of alcohol compounds found in hemicellulose hydrolysate on the growth and fermentation of ethanologenic *Escherichia coli*. Biotechnol Bioeng. 2000;68:524–30.
- Zaldivar J, Martinez A, Ingram LO. Effect of selected aldehydes on the growth and fermentation of ethanologenic *Escherichia coli*. Biotechnol Bioeng. 1999:65:24–33.
- Kitahara Y, Yin T, Zhao X, Wachi M, Du W, Liu D. Isolation of oleaginous yeast (*Rhodosporidium toruloides*) mutants tolerant of sugarcane bagasse hydrolysate. Biosci Biotechnol Biochem. 2014;78:336–42.
- Zhang S, Winestrand S, Guo X, Chen L, Hong F, Jönsson LJ. Effects of aromatic compounds on the production of bacterial nanocellulose by Gluconacetobacter xylinus. Microb Cell Fact. 2014;13:62.

- Wang X, Liang Z, Hou J, Bao X, Shen Y. Identification and functional evaluation of the reductases and dehydrogenases from Saccharomyces cerevisiae involved in vanillin resistance. BMC Biotechnol. 2016;16:31.
- Capasso R, Evidente A, Schivo L, Orru G, Marcialis MA, Cristinzio G. Antibacterial polyphenols from olive oil mill waste waters. J Appl Bacteriol. 1995;79:393–8.
- Shen Y, Li H, Wang X, Zhang X, Hou J, Wang L, Gao N, Bao X. High vanillin tolerance of an evolved *Saccharomyces cerevisiae* strain owing to its enhanced vanillin reduction and antioxidative capacity. J Ind Microbiol Biotechnol. 2014;41:1637–45.
- Overhage J, Priefert H, Steinbüchel A. Biochemical and genetic analyses of ferulic acid catabolism in *Pseudomonas* sp. strain HR199. Appl Environ Microbiol. 1999;65:4837–47.
- Simon O, Klaiber I, Huber A, Pfannstiel J. Comprehensive proteome analysis of the response of *Pseudomonas putida* KT2440 to the flavor compound vanillin. J Proteomics. 2014;109:212–27.
- 69. Lavoie JM, Bare W, Bilodeau M. Depolymerization of steam-treated lignin for the production of green chemicals. Bioresour Technol. 2011;102:4917–20.
- Roberts VM, Stein V, Reiner T, Lemonidou A, Li X, Lercher JA. Towards quantitative catalytic lignin depolymerization. Chemistry. 2011:17:5939

  –48.
- Shuai L, Amiri MT, Questell-Santiago YM, Héroguel F, Li Y, Kim H, Meilan R, Chapple C, Ralph J, Luterbacher JS. Formaldehyde stabilization facilitates lignin monomer production during biomass depolymerization. Science. 2016;354:329–33.
- Deepa AK, Dhepe PL. Lignin depolymerization into aromatic monomers over solid acid catalysts. ACS Catalysis. 2015;5:365–79.
- Song Q, Wang F, Cai JY, Wang YH, Zhang JJ, Yu WQ, Xu J. Lignin depolymerization (LDP) in alcohol over nickel-based catalysts via a fragmentation hydrogenolysis process. Energy Environ Sci. 2013;6:994–1007.

- Xu W, Miller SJ, Agrawal PK, Jones CW. Depolymerization and hydrodeoxygenation of switchgrass lignin with formic acid. ChemSusChem. 2012;5:667–75.
- 75. Gosselink RJ, Teunissen W, van Dam JE, de Jong E, Gellerstedt G, Scott EL, Sanders JP. Lignin depolymerisation in supercritical carbon dioxide/acetone/water fluid for the production of aromatic chemicals. Bioresour Technol. 2012;106:173–7.
- Voitl T, von Rohr PR. Reply to Comments on Demonstruction a process for the conversion of Kraft lignin into vanillin and thy vanillate by acidic oxidation in aqueous methanol. Ind Englemen Res. 2010;49:3501–3.
- 77. Chen HP, Chow M, Liu CC, Lau A, Liu J, Fl LD. Uin co abolism in Rhodococcus jostii RHA1. Appl Enviro 1 Microbiol. 22. //8:586–8.
- Haggblom M, Apajalahti J, Salkinoj alonen M. Metabolism of chloroguaiacols by *Rhodococcus Chloro*, nolicus. A ppl Microbiol Biotechnol. 1986;24:397–404.
- Acevedo C, Brezny R, Joyce TVv. pzález B. Metabolism of mono-and dichlorinated guaiacol by Rhodos pur ruber CA16. Curr Microbiol. 1995;30:63–7.
- Andreoni V, Bernas oni S, tetti P, Villa M. Metabolism of lignin-related compounds by Stadococcus Adochrous: bioconversion of anisoin. Appl Microbiol Biotech. Jl. 1991;36:410–5.
- 81. Shields-Mc of S SD, Klingeman DM, Indest K, Hancock D, Wewalwela D, nich WT, Donaldson JR. Draft genome sequence of *Rhodococcus n. ochrous* strain ATCC 21198. Genome Announc. 201. 2054–114
- 82. Taylo BF As and anaerobic catabolism of vanillic acid and some other nethoxy-aromatic compounds by Pseudomonas sp. strain Pn-1. Appl En iron Microbiol. 1983;46:1286–92.

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