1 Challenges and progress towards industrial recombinant protein production in

- 2 yeasts: A review
- 3 Pieter De Brabander^{a,b}, Evelien Uitterhaegen^b, Tom Delmulle^a, Karel De Winter^b, and Wim Soetaert^{a,b}
- ^aCentre for Industrial Biotechnology and Biocatalysis (InBio.be), Department of Biotechnology, Faculty
 of Bioscience Engineering, Ghent University, Coupure Links 653, 9000 Ghent, Belgium
- 6 ^bBio Base Europe Pilot Plant (BBEPP), Rodenhuizekaai 1, 9042 Ghent (Desteldonk), Belgium
- 7
- 8 *corresponding author
- 9 karel.de.winter@bbeu.org
- 10 Bio Base Europe Pilot Plant (BBEPP)
- 11 Rodenhuizekaai 1,
- 12 9042 Ghent (Desteldonk), Belgium.
- 13 Phone: +32 9 335 70 01

14 Abstract

15 Recombinant proteins (RP) are widely used as biopharmaceuticals, industrial enzymes, or sustainable 16 food source. Yeasts, with their ability to produce complex proteins through a broad variety of cheap 17 carbon sources, have emerged as promising eukaryotic production hosts. As such, the prevalence of 18 yeasts as favourable production organisms in commercial RP production is expected to increase. Yet, 19 with the selection of a robust production host on the one hand, successful scale-up is dependent on a 20 thorough understanding of the challenging environment and limitations of large-scale bioreactors on 21 the other hand. In the present work, several prominent yeast species, including Saccharomyces 22 cerevisiae, Pichia pastoris, Yarrowia lipolytica, Kluyveromyces lactis and Kluyveromyces marxianus are 23 reviewed for their current state and performance in commercial RP production. Thereafter, the impact 24 of principal process control parameters, including dissolved oxygen, pH, substrate concentration, and 25 temperature, on large-scale RP production are discussed. Finally, technical challenges of process scale-26 up are identified. To that end, process intensification strategies to enhance industrial feasibility are 27 summarized, specifically highlighting fermentation strategies to ensure sufficient cooling capacity, 28 overcome oxygen limitation, and increase protein quality and productivity. As such, this review aims 29 to contribute to the pursuit of sustainable yeast-based RP production.

30

31 Keywords:

32 recombinant protein, industrial enzyme, biopharmaceutical, protein transition, yeast, scale-up,

33 process intensification, Saccharomyces cerevisiae, Pichia pastoris, Yarrowia lipolytica

34 **1. Introduction**

35 Recombinant protein (RP) production is a fast-growing field. In 2020, the industrial enzyme market was 36 valued at 5.96 billion USD and is expected to grow at a CAGR of 7.66% (Verfied Market Research, 2021) 37 while the entire RP market was valued at over 125 billion USD with an expected CAGR of 11.2% 38 (Mordor Intelligence, 2021). This can be attributed to the maturation of several factors within the field 39 of RP production such as the increased number of available production hosts, the availability of 40 advanced genetic manipulation tools, the possibility to perform high-throughput screening of newly 41 developed strains and the growing amount of process optimization studies which have led to an overall 42 increase in process efficiency and industrial feasibility (Madhavan et al., 2021). Furthermore, the so-43 called protein transition has incentivized the search for alternative, more sustainable ways of protein 44 production and could result in up to 20% of the emission mitigations required to stay below a 45 temperature increase of 1.5 °C by 2050 (Roe et al., 2019; Stehfest et al., 2019). Besides the well-known 46 plant-based protein alternatives, the importance of microbial protein has been steadily growing as 47 well. Indeed, the alternative protein market was valued at 49.7 million USD and is expected to grow at 48 an impressive CAGR of 16.9% to reach more than 126 million USD by 2028 (Vantage Market Research, 49 2022). This is further demonstrated by the multitude of recent microbial-based protein start-ups such 50 as Those Vegan Cowboys, Nature's Fynd and Perfect Day.

51 Today, there is a microbial pathway available for the conversion of almost any carbon source into 52 almost any desired target product, and significant progress has been made regarding the expression 53 of a large variety of RP. Yet, these proofs-of-concept often remain within the laboratory environment. 54 To successfully translate these findings to an industrial level, research toward scale-up is crucial. 55 Indeed, scaling a process involves multiple technical challenges such as achieving a sufficient oxygen 56 supply and cooling capacity and dealing with heterogeneous environments, while important capital 57 investments with a high risk level are required (de Lorenzo and Couto, 2019). An increased 58 understanding of these challenges that are inherent to large-scale biotechnological processes is a key 59 aspect for efficient scale-up and to lower production prices to obtain market competitive products. 60 There has been increasing interest in yeasts as versatile production hosts for a wide range of RP such 61 as industrial enzymes, biopharmaceuticals and microbial proteins for food and feed industries (Anchel, 62 2016; Mattanovich et al., 2012). Therefore, this review aims to provide a comprehensive overview of 63 the state of the art on yeast-based RP production, the remaining challenges and hurdles that limit 64 widespread industrialization, and innovative process strategies to meet these challenges.

65 2. Yeast production hosts

66 Yeasts are an interesting and versatile class of RP production hosts. Their eukaryotic nature, which 67 allows these organisms to perform post-translational modifications, makes them applicable for the 68 production of a wide range of RP, a significant advantage over bacteria such as Escherichia coli 69 (Mattanovich et al., 2012; Patra et al., 2021; Waegeman and Soetaert, 2011). In addition, as compared 70 to higher eukaryotes such as mammalian or insect cell lines, they are significantly easier to genetically 71 modify, show higher growth rates, obtain higher cell densities, have a lower substrate cost, and are 72 more robust to large-scale fluctuations (Gomes et al., 2018; Potvin et al., 2012). Besides, yeast species 73 are generally known for their higher RP secretion efficiency as compared to bacteria. Extracellular 74 production can be more advantageous as compared to intracellular production hosts in terms of the 75 downstream purification process, an important cost driver that can amount to up to 80 % of the overall 76 RP production process (Labrou, 2014). Indeed, eliminating a cell disruption step avoids the release of 77 intracellular components in the medium facilitating the downstream processing. Additionally, yeasts 78 are a dominating group regarding the production of microbial proteins, such as single cell protein, and 79 have seen wide acceptance on the market, as recently reviewed by Ritala et al. (2017). Within this 80 section, the most prominent yeast species and their relevant characteristics regarding industrial RP 81 production are discussed. This will be confined to Saccharomyces cerevisiae, Pichia pastoris (or 82 Komagataella phaffii), Yarrowia lipolytica, and two Kluyveromyces species, i.e., K. lactis and K. 83 marxianus (Table 1). All have been given the "generally recognized as safe" (GRAS) status by the US 84 Food and Drug Agency (FDA).

85 2.1. Saccharomyces cerevisiae

86 Owing to its robustness for large-scale production, S. cerevisiae has been widely used for the 87 production of bioethanol, platform chemicals and single-cell protein as well as in bakery, brewing and 88 wine applications and to a lesser extent in RP production (Deparis et al., 2017). Because of its 89 robustness and ease of genetic manipulation, this well-researched model yeast was originally 90 developed as an alternative host for the production of RPs which could not be produced by bacterial 91 hosts (Baghban et al., 2019; Huang et al., 2010). Expression of these RP is typically accomplished using 92 either strong constitutive promoters such as PTEF1 and PGADPH or inducible promoters like the galactose-93 inducible P_{GAL1} (Partow et al., 2010). For a more comprehensive review on the available genetic tools 94 and secretion factors, the reader is referred to Gomes et al. (2018). Nonetheless, S. cerevisiae is less 95 attractive for the production of complex proteins due to its characteristic hyper-glycosylation patterns. 96 This can lead to protein heterogeneity, reducing the therapeutic applicability of these proteins and 97 complicating their downstream purification (Jacobs et al., 2009). In addition, challenging protein 98 folding of complex proteins can lead to insufficient protein secretion causing partially degraded 99 products, hereby lowering product yield and further complicating the purification process. To that end, 100 a large amount of research efforts has been dedicated to the improvement of protein secretion of 101 S. cerevisiae, and to extension yeasts in general, and has recently been reviewed by Madhavan et al. 102 (2021). Besides, fermentations using this Crabtree positive yeast can lead to undesired carbon losses 103 due to the production of ethanol, thereby lowering the overall process efficiency. These limitations 104 have motivated the investigation of alternative production hosts. As a result, since 2000, a trend 105 reversal in the number of published research papers has been observed with respect to the use of the 106 methylotrophic yeast P. pastoris as the expression host for RP production as compared to S. cerevisiae 107 (Figure 1). Furthermore, specifically for newly approved pharmaceuticals and from 2018 to 2022, 108 P. pastoris was used more frequently as the expression host as compared to S. cerevisiae. (Walsh and 109 Walsh, 2022). Yet, owing to its well-known genome and beneficial characteristics listed above, several 110 classes of RP are currently still produced on a commercial scale using S. cerevisiae. Examples include insulin, one of the first and most well-known RP, a wide range of hormones, vaccines and fusionproteins as well as industrial enzymes (G. Wang et al., 2017).

113 2.2. Pichia pastoris

114 In the past two decades, P. pastoris, also known as Komagataella phaffii, has seen a surge in strain 115 development (Pan et al., 2022) and process intensification efforts to boost RP expression (Looser et 116 al., 2014; Yang and Zhang, 2018). Moreover, recent advances in glycoengineering, a tool to improve 117 protein properties through the addition or removal of glycans, have enabled the yeast-based 118 production of complex humanized RP, as extensively reviewed by Ma et al. (2020). The latter include 119 monoclonal antibodies, that are currently mostly produced using mammalian cell lines (Jacobs et al., 120 2009; Walsh and Walsh, 2022). For P. pastoris, many successful glycosylation efforts have been 121 performed in the past couple of decades, turning this methylotrophic yeast into one of the most 122 commonly used yeast hosts for protein expression (Shenoy et al., 2021). Yet, P. pastoris' most 123 commonly used promoter system enabling the expression of foreign proteins, i.e., the highly efficient alcohol oxidase 1 promoter (PAOX1), requires methanol as an inducer. The use of methanol has 124 125 considerable drawbacks, of which the risk of storing and handling large amounts of methanol, as well 126 as the increased heat production and oxygen demands are the most prominent. To address this issue, 127 novel *P. pastoris* strains have been developed. For instance, Mut^s strains, where the AOX1 gene was 128 knocked out, exhibit a lower methanol consumption rate. As a result, these strains are more desired 129 as compared to Mut⁺ wild-type strains. A study comparing both phenotypes for the production of two 130 recombinant enzymes found that the Mut^s strain showed a three-fold higher volumetric productivity 131 and was seven times more efficient in converting methanol into the product (Krainer et al., 2012). 132 Alternatively, methanol-free systems have been developed of which the constitutive glyceraldehyde-133 3-phosphate dehydrogenase (P_{GAP}) system has been gaining the most attention (Cankorur-Cetinkaya 134 et al., 2018; Vogl and Glieder, 2013). Moreover, novel promoters, such as the innovative methanol-135 free P_{PDF} and P_{UFF}, are continuously being developed and investigated (Garrigós-Martínez et al., 2021). 136 Several studies have already improved RP production using these alternative promoters as compared 137 to the methanol-inducible PAOXI (Wang et al., 2012a; Zhu et al., 2014). Nevertheless, while constitutive 138 expression can omit the handling of potentially hazardous inducers, toxic proteins are still best expressed by a strong inducible promoter because of the decoupled growth and production phase 139 140 hence avoiding limitations in end titer and productivity due to product inhibition. For a more 141 comprehensive understanding of promoter engineering for P. pastoris, the reader is referred to 142 reviews by Fisher and Glieder (2019) and Ahmad et al. (2014).

143

144 The development of humanized RP combined with the availability of strong promoters and efficient 145 cultivation on simple and inexpensive media, has positioned P. pastoris as a powerful industrial 146 workhorse. Indeed, *P. pastoris* has a lower maintenance coefficient (m_s) (0.013 g g⁻¹ h⁻¹) as compared 147 to *E. coli* (0.04 g g⁻¹ h⁻¹), a conventional bacterial RP production host, or *S. cerevisiae* (0.16 g g⁻¹ h⁻¹) (Jahic et al., 2002; Rebnegger et al., 2016; Uribelarrea et al., 1990). While, as compared to S. cerevisiae, 148 149 P. pastoris may be a more optimal choice for RP production, this yeast has not been studied as 150 extensively and may therefore result in a more extensive process development. For example, 151 additional research focusing on enhancing protein folding and secretion efficiency is required to 152 further increase RP productivity (Yu et al., 2017). In 2009, the first biopharmaceutical RP produced by 153 P. pastoris was approved (Çelik and Çalik, 2012; Corchero et al., 2013). Since then, regulatory agencies 154 such as the FDA have approved the production by companies including Sanofi and Merck of over 70 industrial and biopharmaceutical RPs by P. pastoris (Lee et al., 2021). A recent overview of industrial 155 156 enzymes produced by *P. pastoris* in bioreactors up to 50 L scale and RP products on the market is 157 provided by Duman-Özdamar and Binay (2021) and Baghban et al. (2019), respectively.

158

159 2.3. Yarrowia lipolytica

160 Y. lipolytica is another emerging yeast RP expression host for which research interest has been steadily 161 increasing over the past decade (Figure 1). Much like P. pastoris, this dimorphic yeast has been gaining 162 popularity as a RP production host owing to its capacity to grow to high cell densities on low-cost 163 substrates and efficient excretion of RP. Its ability to use a wide range of carbon sources including plant 164 oils, alkanes and starch hydrolysates has further sparked industrial interest (Coelho et al., 2010; Park 165 and Ledesma-Amaro, 2022). However, its uncontrolled dimorphism, which is not fully understood, and 166 high oxygen requirements are limiting its industrial implementation (Madzak and Beckerich, 2013; 167 Vandermies and Fickers, 2019). RP expression in Y. lipolytica is frequently done with the strong 168 constitutive P_{TEF}. In addition, several lipid-regulated promoters have also been researched and a 169 comprehensive review of Y. lipolytica promoters was recently performed by Sun et al. (2022). Major 170 RP groups which have been produced by Y. lipolytica include enzymes such as lipases, proteases, 171 phosphatases, RNases and esterases as reviewed by Darvishi Harzevili (2014). For example, First Wave 172 BioPharma has recently issued a press release on the clinical phase trials of Adrulipase, a recombinant 173 lipase for the treatment of exocrine pancreatic insufficiency (First Wave BioPharma, 2022; Leblond et 174 al., 2012). An important step towards further exploitation of Y. lipolytica for biopharmaceutical RP 175 production was the technology development to produce humanized glycoproteins (de Pourcq et al., 176 2012), patented by Oxyrane (US8026083B2 and EP2508612B1), a UK-based company focussing on 177 enzyme therapy. Nevertheless, as compared to P. pastoris or S. cerevisiae, industrial-scale RP processes 178 with Y. lipolytica are still scarce, though the steady rise in published research papers and filed patent 179 applications in the last decade on Y. lipolytica indicates an upcoming increase in this area.

180 2.4. Kluyveromyces spp.

181 K. lactis is an emerging lactose-consuming yeast with high levels of protein secretion. The principal 182 value of the Kluyveromyces genus lays in the fact that it is one of the few yeast genera that can utilize 183 lactose as a sole carbon and energy source which has been particularly interesting for dairy waste 184 stream valorization. Further owing to its high growth rate and RP yield, a wide range of enzymes, 185 antibodies and virus particles have been produced in large-scale cultivations up to 140 m³. Examples 186 are the large-scale production of chymosin and β -galactosidase by DSM in the Netherlands (Guimarães 187 et al., 2010; Spohner et al., 2016). With respect to promoter utilization for RP expression in K. lactis, 188 the commonly employed S. cerevisiae promoters P_{GAL1} or P_{PGK} have been frequently used (Gomes et al., 189 2018). The lesser studied K. marxianus has similar beneficial properties while also being able to grow 190 at temperatures of 45 °C and above and utilize a broad amount of substrates such as lactose, xylose, 191 cellobiose, arabinose and polysaccharides such as pectin and inulin (Karim et al., 2020). The most 192 important industrial applications of K. marxianus include the production of endogenous enzymes such 193 as β -galactosidase, β -xylosidase, β -glucosidase, and inulase, often through the use of the *INU1* 194 promoter (Lane and Morrissey, 2010; Zhou et al., 2018). In conclusion, this non-conventional yeast 195 holds great potential for industrial applications and recent advancements regarding the available 196 biotechnological tools have been reviewed by Nurcholis et al. (2020). Furthermore, the ability of 197 K. marxianus to valorise waste streams is promising from an economic point of view and can aid in the 198 transition towards a more sustainable bioeconomy.

199

Placeholder Figure 1.

Table 1. Yeast expression systems for recombinant protein production.

	Saccharomyces cerevisiae	Pichia pastoris	Yarrowia lipolytica	Kluyveromyces lactis	Kluyveromyces marxianus
GRAS status	Approved	Approved	Approved	Approved	Approved
Crabtree effect	Positive	Negative	Negative	Negative	Negative
Oxygen metabolism	Facultative anaerobic	Facultative anaerobic	Obligate aerobic	Obligate aerobic	Obligate aerobic
Optimal growth temperature	30-32 °C	28-30 °C	30-34 °C	25-28 °C	40 °Cª
Native carbohydrate substrates	Hexoses Disaccharides (maltose, sucrose) Alcohols (glycerol, ethanol)	Hexoses Alcohols (glycerol, ethanol, methanol)	Hexoses Pentoses Disaccharides Alcohols (glycerol, ethanol) Lipids Alkanes	Hexoses Pentoses Disaccharides Polysaccharides Alcohols (glycerol, ethanol)	Hexoses Pentoses Disaccharides (sucrose, lactose, cellobiose) Polysaccharides Alcohols (glycerol, ethanol)
Humanized strains available	Yes	Yes	Yes	No	Yes

^a high growth rates up to 47 °C have been reported (Lehnen et al., 2019)

202 3. Principal process parameters and scale-up considerations

203 The recent surge of interest in yeasts as attractive RP production hosts has led to the development of 204 many yeast-based RP production processes for a variety of proteins. Yet, to translate these findings to 205 a commercial scale, it is crucial to understand important scale-up parameters and the effects of the 206 inherent fluctuating environmental conditions in large-scale fermentations and to anticipate the 207 consequences during an early stage. Indeed, while the occurrence and impact of heterogenic zones in 208 terms of nutrient concentrations, oxygen availability, pH, pressure, shear stress etc. are limited on a 209 laboratory scale, on an industrial scale they often negatively impact biomass accumulation, substrate 210 yield, productivity, and in case of more complex products such as RP, product quality. Such zones are 211 often induced by the increased mixing time in large-scale fermentation, which can vary from less than 212 a second at lab scale to over 100 seconds for 30 m³ stirred tank reactors (STR) for microbial production 213 hosts and over 1000 seconds for mammalian cells (Lara et al., 2006). While ideally processes are 214 directly developed on a relevant scale, this is generally not possible due to restraints in the availability 215 of equipment and the labour- and cost-intensiveness of these trials. Furthermore, when processes are 216 scaled, this is often done by companies who are reluctant to publish their results. As a consequence, 217 most available knowledge on scale-up is obtained through scale-down reactors (SDR), an alternative 218 and more practical methodology to replicate the effects of large-scale fermentations in a laboratory 219 environment (Nadal-Rey et al., 2021; Olughu et al., 2019). Here, the bioreactor setup is commonly split 220 up into two separate parts, usually one STR and a plug flow reactor (PFR), allowing the replication of 221 fluctuations in substrate, oxygen, or pH similar to those experienced at large-scale fermentations. 222 Within this section, the most important process parameters and their impact on process performance 223 are discussed in the light of scale-up considerations for yeast-based RP production based on SDR 224 studies and available large-scale fermentation data. Alternatively, mathematical models and so-called 225 digital twins have been used to increase scientific understanding of industrial fermentations as 226 extensively discussed in Haringa et al. (2018) but are beyond the scope of this review.

227 3.1. Dissolved oxygen

228 The dissolved oxygen (DO) concentration has been described as the most influential parameter on 229 overall process performance for RP production (Li et al., 2007). Indeed, for aerobic production hosts, 230 anaerobic conditions can reduce biomass or product yield and trigger by-product formation as a 231 consequence of metabolic shifts. In this respect, obligate aerobic yeasts such as Y. lipolytica, K. lactis 232 and K. marxianus are more sensitive to anaerobic conditions as compared to S. cerevisiae and P. 233 pastoris. Particularly for the production of cost-sensitive industrial enzymes or microbial protein, the 234 substrate is one of the largest cost factors and optimal process efficiency is highly desirable (Puetz and 235 Wurm, 2019). In contrast, speciality biopharmaceutical production processes have higher profit 236 margins and are more driven by patent lifetime restraints and will thus be less inclined to intensively 237 optimize process efficiency. Yet in this case, unforeseen by-products can negatively impact product 238 quality and complicate downstream purification. In general and despite its increase in production cost, 239 high oxygen availability is desired as it increases RP production, substrate consumption and biomass 240 accumulation (Gorczyca et al., 2020; Signori et al., 2014). Furthermore, the degree of oxygen 241 availability also impacts the morphology of Y. lipolytica, where high oxygen availability promotes the 242 prevalence of an ovoid morphology over filaments (Gorczyca et al., 2020). Due to the increased shear 243 stress and negative impact on rheology, the latter is less desired on an industrial scale. However, the 244 low solubility of oxygen in the liquid phase, the high oxygen demand of high-cell density fermentations, 245 and increased difficulty in supplying oxygen in large bioreactors makes it challenging to attain high 246 oxygen availability at an industrial scale. Applicable process intensification strategies to tackle this are 247 discussed Section 4.

248 In addition to the challenges of maintaining high oxygen availability, oxygen fluctuations, with zones 249 ranging from high oxygen concentrations to microaerobic or anaerobic, are commonly encountered 250 and may reduce the process performance of large-scale fermentations. For instance, oxygen 251 fluctuations have been shown to reduce lipase gene expression rates for Y. lipolytica (Kar et al., 2010) 252 and decrease biomass production of a S. cerevisiae fermentation while showing a variable effect on 253 ethanol, acetate and glycerol accumulation (Lara et al., 2006). Lorantfy et al. (2013) observed that 254 biomass production was reduced and by-product formation was increased as a consequence of larger 255 oxygen fluctuations, prolonging the exposure of the baker's yeast to low DO concentrations. In 256 essence, DO fluctuations are more difficult to control at an industrial scale as compared to DO 257 limitations, emphasizing the importance of working with a robust production host. For instance, for K. 258 marxianus, DO fluctuations between 0-20% did not notably influence the volumetric β -galactosidase 259 production rate, biomass production or biomass yield as compared to constant DO concentrations in 260 the same range, showing the robustness of the evaluated yeast to DO variations (Cortés et al., 2005).

261 *3.2. Pressure*

262 Because of the increased hydrostatic pressure in large-scale reactors (approximately 1 bar for every 10 263 meters of height), elevated reactor pressures are inextricably linked to large-scale fermentations. 264 Additionally, elevated pressures of up to 1.5 barg are commonly intentionally applied to increase 265 oxygen supply and ensure sterility. This means that microbial organisms will move through different 266 pressure zones and potential consequences must be considered. While on the one hand, these 267 moderately increased pressures do not lead to a notable reduction in cell viability of yeasts (Fernandes, 268 2005), on the other hand, an increased reactor pressure will increase the solubility of gaseous 269 compounds in the media as a consequence of Henry's law. For instance, CO₂, the main by-product of 270 aerobic fermentation, has been shown to affect microbial physiology as it can easily enter the cell, 271 even more so at an elevated pressure, and lower intracellular pH. As a result of elevated CO₂ 272 concentrations by increasing the CO₂ concentration in the ingoing gas from 0.05 to 50%, a 60% increase 273 in maintenance energy requirements was registered for S. cerevisiae which would lead to a reduced 274 biomass and product yield (Eigenstetter and Takors, 2017). Yet, another study on similarly elevated 275 CO₂ concentrations, observed no effects on the biomass yield or viability of *S. cerevisiae* (Hakkaart et 276 al., 2020). Here, the authors hypothesized that these contradicting results were caused by the lower 277 growth rate applied in the latter study. Nevertheless, as many metabolic pathways are related to the 278 CO₂ concentration, metabolic changes cannot be excluded and should be the subject of further 279 research in a case-by-case evaluation. Nonetheless, increasing total air pressure as a strategy for 280 process intensification has been frequently applied to meet high oxygen demands and will be more 281 elaborately discussed in Section 4.2.2.

282 *3.3. Substrate gradients*

283 As a consequence of the usual single substrate addition point at the top of a reactor, substrate gradients are present during large-scale fermentations. Indeed, during a 30 m³ fed-batch fermentation 284 285 with S. cerevisiae, a 2.75-fold difference in substrate concentration was estimated between the 286 addition point and the bulk medium (Larsson et al., 1996). These varying substrate concentration zones could trigger unwanted metabolic changes such as overflow metabolisms or the Crabtree effect 287 288 resulting in the production of by-products. Studies evaluating the impact of increased mixing times and 289 substrate gradients during S. cerevisiae cultivations have observed up to 30% lower biomass yields due 290 to increased ethanol production (George et al., 1993; Lejeune et al., 2010; Risager Wright et al., 2016). 291 This highlights the impact of substrate gradients on process performance and the need for further 292 research towards large-scale production environments, where substrate costs often dictate the final 293 product price.

294 3.4. pH

295 pH is a critical process parameter because of its impact on yeast growth, product formation, RP 296 stability, and protease activity. Additionally, the optimal pH varies among production strains and target 297 RPs. While in a laboratory environment buffer solutions are typically used, on an industrial scale highly 298 concentrated acid or base solutions must be added to maintain continuous control over the pH 299 setpoint resulting in important pH fluctuations at the point of addition. Indeed, for a pilot-scale 300 fermentation of 100 L, a pH range of the setpoint of 4 in the bulk medium to 9 at the point of addition 301 has been predicted (Reuss et al., 1994). Nonetheless, for Y. lipolytica, extended exposure to pH 302 perturbations in a 1.6 L bioreactor affected the cell morphology but process yields and cell viability 303 remained largely unchanged (Timoumi et al., 2017). While yeasts can grow at a wide pH range and are 304 generally robust to fluctuations, protein quality and productivity can be negatively affected. For 305 instance, for the production of recombinant erythropoietin by P. pastoris, the highest product 306 concentration was obtained at pH 4.5 while this was reduced to less than 50% at pH 5.0 (Soyaslan and 307 Çalik, 2011). Furthermore, excessive pH fluctuations could result in protein aggregation or degradation 308 (Zheng and Janis, 2006). Investigating the effects of, and sensitivity to, different pH values on process 309 performance is therefore paramount to gain early insight into the potential impact of pH fluctuations 310 on an industrial scale.

311 *3.5. Temperature*

312 Process temperature plays an important role in RP productivity and product stability and quality. 313 However, as opposed to the previously discussed parameters, no temperature gradients have been 314 observed in an industrial-scale fermenter. This is because of the larger cooling surface, usually in the 315 form of a cooling jacket or cooling spirals, as opposed to the single addition points of acid or base for 316 pH control, of substrate for feeding, or bottom sparging for oxygen addition. Yet, sufficient heat 317 dissipation can become a limiting factor during scale-up, especially for high-cell density fermentations. 318 The heat transfer rate (HTR) can be calculated and is a measure of the cooling capability of a fermenter 319 (Yang, 2010):

$$HTR = hA(T - T_i)$$

Where *h* is the heat transfer coefficient, related to the specific heat transfer efficiency of the cooling jacket material and configuration, *A* is the vessel jacket area, *T* is the culture temperature and T_j is the temperature of coolant inside the jacket. During scale-up, the ratio of jacket area A to the volume of the reactor decreases, consequently lowering cooling capacity. To overcome inadequate cooling capacity during large-scale high-cell density fermentation, process intensification strategies can be applied and are discussed throughout Section 4.

327

4. Process intensification strategies

328 Process intensification (PI) aims to improve process performance and reduce the resources required 329 through intensified equipment or operating strategies and has been described as one of the most 330 promising development paths to tackle techno-economic challenges and enhance industrial feasibility 331 (Górak and Stankiewicz, 2018). As an example, the intensification of fermentation processes through 332 in situ product removal can increase product titer and overall productivity for yeast-based production 333 of industrial platform chemicals (De Brabander et al., 2021). Likewise, yeast-based RP processes often 334 require a further increase in efficiency or productivity to compete with well-established RP production 335 systems using E. coli or mammalian cells. Additionally, PI strategies can be crucial to overcome 336 commonly observed large-scale limitations in cooling capacity or oxygen supply and improve process 337 scalability. In recent years, multiple innovative PI strategies have been designed and evaluated on lab 338 scale to boost process efficiency and increase industrial feasibility. Yet, due to the often risk-averse 339 approach of industrial processing and limited fundamental understanding of PI strategies, PI 340 innovation is rather slow (Górak and Stankiewicz, 2018). To enhance industrial implementation of PI 341 strategies, this review aims to increase understanding and provide a critical overview of potential PI 342 strategies for yeast-based RP production (Table 2). Because of the well-established nature of 343 S. cerevisiae and the surge of interest in P. pastoris for RP production, most RP-focussed PI strategies 344 have been applied to either or both of these yeasts. However, the strategies discussed within this 345 section could likely also be applied to other RP production hosts, but this would require further 346 validation.

347 4.1. Feed-related PI strategies

The applied feed strategy has a major impact on the process performance. For the production of RP with *P. pastoris*, well-known protocols such as the ones from Pichia Protocols (Stratton et al., 1998), Invitrogen (2002) or Nature Protocols (Tolner et al., 2006) form a solid starting point for process development. Yet, the implementation of PI strategies can significantly increase process performance which is often required to attain industrially feasible production. Feed-related PI strategies that have increased process performance while also tackling scale-up challenges and limitations are discussed below.

355 4.1.1. <u>Growth rate-controlled feeding</u>

356 With respect to RP production, studies have shown a strong correlation between growth rate (μ) and 357 productivity. For *P. pastoris*, this correlation was shown to be strain-dependent and the role of μ and 358 its impact on RP production has recently been investigated (Garrigós-Martínez et al., 2019). In general, 359 it has been shown that a reduced growth rate, as compared to μ_{max} , can significantly enhance RP 360 productivity when working with methanol-inducible *P. pastoris* strains. Conversely, when utilizing 361 P. pastoris strains that do not require methanol for RP production (e.g. P_{GAP}), a growth-coupled RP 362 production is typically observed (Looser et al., 2014). In S. cerevisiae, protein-dependent correlations 363 have been observed (Liu et al., 2013). Furthermore, growth rate control is an important factor to 364 maintain process stability, increase reproducibility and control oxygen demands (Egli, 2015). The latter will be discussed more elaborately in the Section 4.2. Constant growth rates can be achieved by 365 applying an exponential feed rate. To accurately and continuously control this process parameter at a 366 367 fixed level and avoid deviations and time-consuming offline measurements, advanced feed rate 368 control strategies based on online process monitoring, e.g., through detailed off-gas analysis, inline 369 biomass probes, or mathematical models are becoming more prevalent (Lisa Mears et al., 2017; Ponte 370 et al., 2018). For instance, for a *P. pastoris* process, controlling μ at an optimal level of 0.03 h⁻¹ based on the online measurement of the ammonia consumption rate, could increase the end titer of alpha 371 372 1-antitrypsin up to 2 times as compared to conventional control strategies such as DO-stat (Tavasoli et 373 al., 2019). For S. cerevisiae and K. marxianus, μ was accurately controlled by regulating a glucose feed 374 based on online measurements of a biomass probe (Dabros et al., 2010; Xiong et al., 2015). In 375 conclusion, a growth rate-controlled feed regime at a predetermined and optimized μ has been a 376 proven strategy to enhance process performance. Hence, accurate growth rate control through process automation based on online measurement of key parameters will play a key role in the 377 378 development of intensified industrial processes with improved performance and stability.

379 4.1.2. Mixed substrate feeding

380 Whereas conventional fed-batch systems use a single carbon feed for either growth or induction, a 381 mixed feed strategy combines two or more. The initial goal of this strategy was to enhance the 382 volumetric productivity by increasing biomass accumulation throughout the induction phase of 383 methanol-induced *P. pastoris* fermentations. This PI strategy has been successfully applied for both 384 Mut⁺ (Jungo et al., 2007a) and Mut^s phenotypes (Zalai et al., 2012). Doing this, a mixed feed with 385 methanol and glycerol led to a 4.5-fold improvement in volumetric productivity of bovine lysozyme by 386 P. pastoris (Brierley et al., 1990). In addition, follow-up research observed that oxygen consumption 387 and heat production could thereby also be reduced as a consequence of the partial substitution of 388 methanol by an alternative carbon source (Niu et al., 2013). This results from the higher oxygen 389 requirement and heat production during methanol metabolization as compared to commonly used co-390 substrates such as glycerol, mannitol or sorbitol (Jungo et al., 2007a; Zalai et al., 2012). The ratio of 391 methanol over the co-substrate in the mixed feed plays an important role in RP productivity. While at 392 suboptimal ratios RP productivity dropped to zero, for an optimized ratio RP productivity could be 393 increased by more than 40% compared to a methanol-only benchmark (Niu et al., 2013). However, 394 because of the additional feeds, process complexity will increase, undesired protein variation could be 395 introduced (Ye et al., 2011), and repression of the induction system has been described, thereby 396 lowering process efficiency (J. Wang et al., 2017). Yet, the latter has been circumvented by using non-397 repressive carbon sources such as sorbitol, mannitol or ascorbic acid (Vuree, 2020; Zavec et al., 2020). 398 As an example, a methanol/sorbitol mixed feed strategy with P. pastoris reduced heat production and 399 oxygen consumption by almost 40% while increasing RP productivity by 30% as compared to the 400 conventional process with a sole methanol feed (Jungo et al., 2007b). In conclusion, for the widely used 401 methanol-inducible *P. pastoris* processes, a mixed feed strategy could be an attractive PI strategy to 402 lower oxygen and cooling demands but requires careful optimization of all carbon fractions in the feed 403 to avoid a reduction in RP productivity. Additionally, the associated increased substrate cost should be 404 justified by, on the one hand, a proportional increase in RP formation or, on the other hand, a reduction 405 in costs related to cooling or oxygen supply.

406 4.1.3. Intermittent feeding

407 An intermittent feeding strategy applies successive substrate feeding breaks during which the yeast is 408 deprived of any carbon source. As a response to these applied stress conditions, higher RP synthesis 409 rates have been observed for P_{GAP} -controlled *P. pastoris* processes. It has been hypothesized that this 410 increase could be caused by the upregulation of glycolytic pathways (Garcia-Ortega et al., 2016). By 411 optimizing the duration of the carbon starvation periods, this rather simple to implement fed-batch 412 strategy has led to an increased biomass-specific yield and could increase the substrate yield by 50% 413 while overall process productivity could be increased by 30% (Garcia-Ortega et al., 2016). Because of 414 the higher biomass-specific yield, lower biomass concentrations could be targeted while still 415 maintaining similar end product concentrations which is beneficial for lowering oxygen demand and 416 cooling requirements. However, repeated carbon starvation cycles are known to induce stress on the 417 production organism and consequentially lower RP yield or quality due to proteolytic degradation (Heo 418 et al., 2008). Nevertheless, this readily scalable strategy appears promising for large-scale 419 fermentation processes owing to the increased cost-effectiveness following the higher substrate yield. 420 To date, industrial feasibility of the strategy has not been investigated and further research is necessary 421 to increase insight into the consequences of repeated carbon starvation cycles on protein breakdown 422 and evaluate the effectiveness of this strategy for other types of RP as well as production hosts.

423 4.2. Oxygen-related PI strategies

One of the most commonly observed challenges during scale-up involves insufficient oxygen availability for the cells. Indeed, as discussed in Section 3.1, the oxygen concentration and its fluctuations in a bioreactor can significantly influence process performance. To overcome these limitations, PI strategies aiming to increase the oxygen transfer or lower oxygen requirements can be applied, hereby increasing technical feasibility as well as process performance. Alternatively, process temperature, choice of the production host, promoter, and strain phenotype can also considerablyinfluence oxygen requirements, which will be more elaborately discussed in the next sections.

431 4.2.1. Lowering growth rate

432 A drawback of high-cell density fermentations, as commonly observed with yeast, is the high OTR 433 requirement. As discussed above, the applied growth rate is known to control oxygen demands. While 434 generally regarded as disadvantageous due to the increased processing time and lower RP 435 productivity, operating at a lower cell growth rate by limiting the substrate feed rate is an interesting 436 strategy for RP production (Looser et al., 2014). The reduced cell growth results in lower oxygen and 437 maintenance energy requirements, thereby increasing cell viability and minimizing cell lysis and the 438 consequential release of intracellular proteases into the medium causing RP product breakdown (Zhao 439 et al., 2008). Therefore, low growth rates could allow extended accumulation of the desired proteins 440 and has successfully been applied on pilot-scale. As an example, limiting the growth rate of a PGAP-441 controlled P. pastoris strain resulted in a 13% increase in specific lipase activity and over 50% reduction 442 of the cell maintenance energy consumption upon scale-up to 800 L. However, as a consequence of 443 lower biomass accumulation, lipase productivity was decreased by about 10% (Zhao et al., 2008). 444 Nevertheless, the accumulation of 20% less biomass is expected to reduce downstream purification 445 costs.

446 4.2.2. Increasing total air pressure

447 Oxygen limitation can also be avoided by increasing the oxygen transfer rate (OTR). The OTR is a 448 measure of the rate at which oxygen is transferred from a gaseous phase (e.g. air) to a liquid phase 449 (e.g. cultivation medium) and can be optimized by reactor type and design, which has been extensively 450 reviewed elsewhere (Garcia-Ochoa and Gomez, 2009). Yet, most advanced bioreactor types able to 451 achieve high OTR values are still within a research stage. Alternatively, the OTR can be improved by 452 increasing the partial pressure of oxygen through the supply of oxygen-enriched air, or by increasing 453 the total reactor pressure. While in a laboratory environment, further increase of the oxygen supply 454 after maximizing both stirrer speed and aeration rate is often achieved through pure oxygen 455 supplementation, the operational risks and high cost of pure oxygen limit this strategy for large-scale 456 fermentations. Contrastingly, increasing the total air pressure is a viable and well-known strategy for 457 industrial fermentations which are commonly performed in pressure-resistant stainless-steel reactors 458 (Knoll et al., 2007). Because of the higher solubility of oxygen in the medium phase at an increased 459 pressure, pure oxygen supplementation can hereby be avoided (Liu et al., 2016). As an example, an 460 enhanced oxygen uptake rate by increasing the reactor pressure from 0.2 to 0.9 barg resulted in a 50% 461 increased β-glucosidase yield for a *P. pastoris* fermentation process (Charoenrat et al., 2006). Also, by 462 increasing the operating pressure up to 7 barg for a Y. lipolytica process, lipase production was increased by over 500% (Lopes et al., 2008). Yet, a pressure above approx. 1.7 barg requires specialized 463 464 equipment, further increasing capital investments.

465 4.2.3. <u>Hypoxic conditions</u>

466 The driving force of oxygen transfer to the medium phase can be increased by lowering the DO setpoint (e.g., < 5%) (Charoenrat et al., 2005). As a result, a reduced amount of energy in the form of agitation 467 468 or aeration is required for similar oxygen demands which is beneficial from both a technical and 469 economic point of view. However, under these low oxygen or hypoxic conditions, varying results have 470 been obtained depending on the specific yeast species. By reducing DO of a *P. pastoris* fermentation 471 process, a 2.3-fold higher volumetric RP productivity could be attained and product purity could be 472 increased because of a reduced amount of homogeneous proteins secreted in the medium (Baumann 473 et al., 2008; Garcia-Ortega et al., 2017; Gasset et al., 2022). Even more, lower biomass accumulation 474 was observed, which facilitates cell removal during downstream processing. Contrastingly, for 475 S. cerevisiae RP productivity was lowered and cell growth was negatively affected (Aon et al., 2018; 476 Baumann et al., 2011). Besides, monitoring and controlling the desired hypoxic conditions is 477 challenging. A commonly applied strategy for hypoxic process control involves monitoring the specific 478 ethanol production rate. Yet, this requires intensive sampling and off-line quantification of the ethanol 479 concentration in the culture medium. Alternatively, respiratory quotient (RQ) monitoring, based on O₂ 480 and CO₂ data from off-gas analysis which indicates the physiological state of the yeast, has been used 481 as a continuous monitoring method for the degree of oxygen limitation (Gasset et al., 2022).

482 *4.3. Continuous cultivation*

483 Progressing towards continuous production has led to an increased process efficiency and subsequent 484 improved cost-competitiveness within several industries. For industrial yeast-based RP production, 485 continuous cultivation presents a highly promising PI strategy to improve its industrial feasibility as it 486 allows extension of the production phase, thereby reducing process downtime and increasing space-487 time yield. Increased product consistency and manufacturing flexibility as well as equipment size 488 reduction are some of the major cost reduction factors that can be obtained through continuous 489 processing (Peebo and Neubauer, 2018; Walther et al., 2015). Furthermore, higher RP yields have been 490 obtained through continuous cultivation, which could be attributed to the higher fraction of younger, 491 more efficient cells as compared to traditional batch or fed batch processes (Nieto-Taype et al., 2020) 492 or the lower stress levels observed in these cells (Zahrl et al., 2017). For example, a recent P. pastoris 493 study demonstrated that within a timeframe of six weeks, almost six times more recombinant lipase B 494 could be produced in a continuous mode as compared to a traditional fed-batch mode, thereby 495 enabling lower production costs (de Macedo Robert et al., 2019). Another P. pastoris study was able 496 to increase the volumetric productivity for a recombinant Hepatitis B antigen by roughly 50% by 497 employing a continuous production mode. The authors reported high genetic stability and the absence 498 of contaminants after two weeks of cultivation (Rahimi et al., 2019). Through the continuous 499 cultivation of a β -galactosidase enzyme using a lactose-consuming *S. cerevisiae* strain, productivity was 500 increased up to 11-fold as compared to the batch benchmark process (Domingues et al., 2005).

501 Yet, continuous cultivation often involves increased complexity for process control, requires high 502 genetic stability of the production strain, is more prone to contamination, and faces regulatory 503 challenges with regard to the production of biopharmaceutical RP (Fisher et al., 2019). For the latter, 504 additional challenges are observed as compared to industrial enzymes due to their typically lower 505 production quantities, the increased importance of time to market and the required time for 506 regulatory approval. Yet, with regard to regulatory approval, the rise of continuous (perfusion) 507 cultivations for biopharmaceutical RP production with mammalian cell cultures could also accelerate 508 the transition towards yeast-based continuous cultivation (L. Mears et al., 2017). Currently, continuous 509 industrial production of insulin, one of the first produced RP in yeast, is being performed by Novo 510 Nordisk using S. cerevisiae (Peebo and Neubauer, 2018). To the best of our knowledge, no reports on 511 commercial continuous RP production have been described for the other yeasts discussed within this 512 review, rendering fed-batch processing still the most prevalent mode of operation to date. The lack of 513 such processes could be caused by the low risk tolerance, short process development times, the 514 relatively young industry of biotechnology and the initial focus on the development of innovative RP 515 rather than PI and cost reduction strategies (Cankorur-Cetinkaya et al., 2018; Walther et al., 2015). Yet, 516 the pressure to continuously lower production costs of RP will push researchers to tackle current 517 challenges of continuous yeast-based cultivation and it is expected that with the maturation of the 518 biotechnology industry, the implementation of continuous processes on an industrial scale will 519 increase.

520 *4.4. Lowering process temperature*

521 Lowering the process temperature setpoint is a convenient PI strategy that has led to 1) reduced 522 proteolytic degradation, 2) increased product quality, and 3) increased protein productivity. The 523 former two can be explained by reduced thermodynamic protease activity at lower temperatures and 524 reduced release of intracellular proteases into the medium as a consequence of decreased metabolic 525 stress or cell death (Jahic et al., 2003b). Lower temperatures can furthermore increase protein stability 526 thereby reducing protein refolding requirements and consequently increasing RP productivity 527 (Dragosits et al., 2009). For example, Jahic et al. (2003b) could double the final RP concentration, yet 528 this required the temperature to be lowered to 12 °C as compared to 30 °C. Other studies have 529 reported specific productivity improvements of up to 100-fold by lowering the temperature to 20 °C 530 (Dragosits et al., 2009; Jin et al., 2011; Li et al., 2013; Zhong et al., 2014). Additionally, at lower 531 temperatures oxygen demand is lowered because of a reduced metabolic activity, while oxygen 532 solubility is increased (Trentmann et al., 2004). However, a temperature close to optimal is desired to 533 maximize the growth rate of the yeast and, as a consequence, the abovementioned benefits are usually 534 at a cost of a lower growth rate. Furthermore, care must be taken when scaling these processes since 535 sufficient heat dissipation will become more challenging and cooling costs will rise with an increasing 536 bioreactor size. This especially holds for high-cell density fermentations and those with high heat 537 generation such as methanol-induced P. pastoris fermentations. Yet, other PI strategies can be applied 538 in parallel to lower heat production and decrease oxygen requirements (Table 2). Hence, during 539 process development an important trade-off must be made in which increased protein production 540 and/or quality is balanced with higher cooling capacity requirements and their related processing costs 541 while also considering the technical limitations of a large-scale bioreactor.

542 4.5. Lowering process pH

543 Similar to temperature, the pH can be optimized to reduce proteolytic degradation and increase overall 544 process efficiency (Kong et al., 2009). In general, conditions of low pH reduce proteolytic activity 545 because of the associated partial or complete inactivation of proteases (Jahic et al., 2003a). While in 546 contrast to lowering the temperature, adjusting pH is not limited by technical limitations. Yet, strain-547 and protein-specific limitations will apply and must be accounted for. This strategy was found effective 548 for the production several types of RP including a cellulose-binding module fused to a Candida 549 antarctica lipase (CBM-CALB) where the fraction of full-length protein was increased from 40 to 90% 550 by decreasing the pH from 5.0 to 4.0 (Jahic et al., 2003a), and a proteolysis reduction of a growth 551 hormone (IGF-1) by lowering the pH to 3.0 (Baghban et al., 2019). Additionally, processes at a lower 552 pH setpoint are less prone to contamination.

553 4.6. Strain selection and development

554 Besides process-related intensification strategies, strain selection and engineering can significantly 555 improve process performance and technical feasibility. While an in-depth review of strain 556 development is beyond the scope of this review, some prominent examples are provided. For instance, 557 the use of thermotolerant organisms, such as K. marxianus, can improve cooling efficiency because of 558 the increased driving force between the coolant and fermentation medium, thereby lowering process 559 costs related to temperature control (Gombert et al., 2016). A higher temperature furthermore 560 reduces the risk of contamination. Yet, the optimal temperature setpoint will also be determined by 561 the stability of the target protein at these elevated temperatures. Alternatively, strain phenotypes 562 have been developed with reduced heat generation. For instance, for P. pastoris, this has led to the 563 development of strains where both AOX1 and AOX2 genes are knocked out, resulting in a Mut⁻ strain. 564 Recent progress has shown that important scale-up parameters including the oxygen demand, heat 565 output and specific methanol uptake rate were reduced by more than 80% as compared to the more commonly used Mut^s or Mut⁺ alternatives, while high viability could be maintained throughout the 566 567 fermentation (Zavec et al., 2020). Additionally, because of the lower methanol uptake rate, which was 93% lower as compared to a Mut^s strain, lower volumes of methanol are required and hence the 568 569 inherent drawbacks of having to store large amounts of methanol can be minimized, while also 570 lowering feedstock costs. An important drawback is the currently lower (42%) product titer as 571 compared to a Mut^s phenotype. Yet, while still in a research stage, new methanol-free promoter 572 systems are continuously being developed aiming to lower heat production and oxygen demand 573 (Garrigós-Martínez et al., 2021). As an alternative to lowering oxygen demand, the oxygen uptake rate 574 (OUR) in yeasts can be enhanced through modification of the cell physiology. This has been 575 demonstrated in Y. lipolytica and P. pastoris through co-expression of Vitreoscilla hemoglobin, 576 resulting in nearly a two-fold increase in the activity of a recombinant lipase produced using P. pastoris 577 (Bhave and Chattoo, 2003; Wang et al., 2012b). Finally, recent advancements in synthetic biology 578 techniques, including the latest CRISPR/Cas9 technology, are expected to play a major role in 579 improving homologous recombination efficiency, protein folding, and optimizing promoters, 580 terminators, and transcription factors. For a more comprehensive view on the subject, the reader is 581 referred to recent reviews (Pan et al., 2022; Patra et al., 2021; Raschmanová et al., 2021, 2018).

Table 2. Process intensification strategies to overcome principal scale-up challenges.

Scale-up challenge	Process strategies
Oxygen limitation	Increase oxygen supply by
	Reactor design
	Increasing total operating air pressure
	Lowering the DO setpoint to increase the driving force
	Lowering the temperature to increase the oxygen solubility
	Enhancing OUR through cell physiology modification
	Reduce oxygen demand by
	Lowering feed rate to maintain lower growth rate
	Mixed feed strategies
	Intermittent feeding strategy
	Lower temperature setpoint to reduce metabolic activity
	Development of alternative promoter systems
Insufficient cooling capacity	Increase cooling efficiency by
	Increasing temperature setpoint
	Reactor design
	Reduce heat production by
	Reducing working biomass concentration
	Applying a mixed feed strategy
	Applying an intermittent feeding
	Development of alternative promoter systems (e.g., Mut ⁻)
	Using alternative production hosts
Gradients (e.g., substrate, pH)	Optimize reactor design to reduce gradients
	Select robust strains to reduce impact of gradients
Low or inconsistent protein	Reduce proteolytic degradation by
quality	Lowering the temperature to avoid cell lysis and release of
	protease to the medium
	Lowering the temperature to reduce protease activity
	Lowering pH setpoint to reduce protease activity
	Developing protease deficient strains
	Reduce batch-to-batch variability by
	Implementing a continuous cultivation strategy
	Developing protease deficient strains
	Advanced process automatization with online monitoring and
	process control

584 5. Conclusions and prospects

585 Yeasts have demonstrated high potential as industrial hosts for RP production. While S. cerevisiae and 586 P. pastoris have been used to produce a variety of RP products at an industrial level, including microbial 587 protein, enzymes and biopharmaceutical proteins, research efforts towards the use of alternative or 588 non-conventional yeasts such as Y. lipolytica, K. lactis and K. marxianus have been increasing because of the ever-rising importance to use sustainable and low-cost feedstocks. The higher protein secretion 589 590 efficiency of yeasts as compared to bacterial production hosts allows to significantly reduce the 591 downstream processing costs of price-sensitive products, hereby attaining economically feasible 592 production. In addition, particularly for the production of biopharmaceutical RP, yeasts provide a fast 593 development timeline and can use cheap cultivation media while being able to produce complex 594 humanized proteins.

595 Yet, the road from laboratory-scale development to an industrial process is challenging and starts with 596 the selection of a suitable production host. In addition to the widely used selection criterium of 597 maximal productivity, the strain robustness towards the conditions of large-scale fermentations must 598 be considered and investigated during early process development. Indeed, in large-scale bioreactors, 599 fluctuations in terms of dissolved gaseous compounds, substrate concentrations, and pH, are not easily 600 controlled or overcome and hence their impact on the process performance should be taken into 601 account when scaling a fermentative production process.

- 602 Besides gradients, technical limitations of large-scale bioreactors are often the cause of commonly 603 encountered scale-up challenges such as insufficient oxygen supply or cooling capacity and must be 604 considered during the early stages of process development to establish crucial process boundaries. To 605 overcome these challenges and expand the feasibility window for process operation, multiple effective 606 process intensification strategies focussing on lowering oxygen demands and cooling requirements 607 while increasing product quality and reducing process costs have been investigated to enhance 608 industrial feasibility and subsequent commercialization. In addition, strain engineering efforts and the 609 development of novel promoter systems for P. pastoris could bring further process improvements, the 610 latter through the use of non-hazardous inducers with concurrent reduced oxygen demand or heat 611 production.
- Finally, up until now, most research has focused on the use of *S. cerevisiae* and *P. pastoris* for RP production, both from a strain engineering and fermentation process development point-of-view and has already allowed efficient industrial yeast-based RP production. Nonetheless, future process development and intensification efforts should also be directed towards other emerging yeast species, opening doors to a wide range of potential novel RP products with a variety of applications and to the
- 617 use of alternative abundant, economical and sustainable feedstocks.

618 6. Research funding

This work did not receive any specific grant from funding agencies in the public, commercial, or notfor-profit sectors.

621 **7. Declaration of competing interest**

622 The authors declare that there is no conflict of interest.

623 8. Acknowledgements

- 624 No acknowledgements are necessary as the work provided to create this manuscript was fully
- 625 performed by the authors themselves.

626 9. Refences

- Ahmad, M., Hirz, M., Pichler, H., Schwab, H., 2014. Protein expression in *Pichia pastoris*: Recent
 achievements and perspectives for heterologous protein production. Appl Microbiol Biotechnol.
 https://doi.org/10.1007/s00253-014-5732-5
- Anchel, D., 2016. Methods and compositions for egg white protein production. WO2016077457A1.
- Aon, J.C., Tecson, R.C., Loladze, V., 2018. Saccharomyces cerevisiae morphological changes and
 cytokinesis arrest elicited by hypoxia during scale-up for production of therapeutic recombinant
 proteins. Microb Cell Fact 17, 1–15. https://doi.org/10.1186/s12934-018-1044-2
- Baghban, R., Farajnia, S., Rajabibazl, M., Ghasemi, Y., Mafi, A.A., Hoseinpoor, R., Rahbarnia, L., Aria,
 M., 2019. Yeast Expression Systems: Overview and Recent Advances. Mol Biotechnol 61, 365–
 384. https://doi.org/10.1007/s12033-019-00164-8
- Baumann, K., Dato, L., Graf, A.B., Frascotti, G., Dragosits, M., Porro, D., Mattanovich, D., Ferrer, P.,
 Branduardi, P., 2011. The impact of oxygen on the transcriptome of recombinant *S. cerevisiae*and *P. pastoris* a comparative analysis. BMC Genomics 12, 1–16. https://doi.org/10.1186/14712164-12-218
- Baumann, K., Maurer, M., Dragosits, M., Cos, O., Ferrer, P., Mattanovich, D., 2008. Hypoxic fed-batch
 cultivation of *Pichia pastoris* increases specific and volumetric productivity of recombinant
 proteins. Biotechnol Bioeng 100, 177–183. https://doi.org/10.1002/bit.21763
- Bhave, S.L., Chattoo, B.B., 2003. Expression of *Vitreoscilla* Hemoglobin Improves Growth and Levels of
 Extracellular Enzyme in *Yarrowia lipolytica*. Biotechnol Bioeng 84, 658–666.
 https://doi.org/10.1002/bit.10817
- Brierley, R.A., Bussineau, C., Kosson, R., Melton, A., Siegel, R.S., 1990. Fermentation Development of
 Recombinant *Pichia pastoris* Expressing the Heterologous Gene: Bovine Lysozyme. Ann N Y Acad
 Sci 589, 350–362. https://doi.org/10.1111/j.1749-6632.1990.tb24257.x
- Cankorur-Cetinkaya, A., Narraidoo, N., Kasavi, C., Slater, N.K.H., Archer, D.B., Oliver, S.G., 2018. Process
 development for the continuous production of heterologous proteins by the industrial yeast,
 Komagataella phaffii. Biotechnol Bioeng 115, 2962–2973. https://doi.org/10.1002/bit.26846
- 653 Çelik, E., Çalik, P., 2012. Production of recombinant proteins by yeast cells. Biotechnol Adv 30, 1108–
 654 1118. https://doi.org/10.1016/j.biotechadv.2011.09.011
- Charoenrat, T., Ketudat-Cairns, M., Jahic, M., Veide, A., Enfors, S.O., 2006. Increased total air pressure
 versus oxygen limitation for enhanced oxygen transfer and product formation in a *Pichia pastoris*recombinant protein process. Biochem Eng J 30, 205–211.
 https://doi.org/10.1016/j.bej.2006.04.004
- Charoenrat, T., Ketudat-Cairns, M., Stendahl-Andersen, H., Jahic, M., Enfors, S.O., 2005. Oxygenlimited fed-batch process: An alternative control for *Pichia pastoris* recombinant protein
 processes. Bioprocess Biosyst Eng 27, 399–406. https://doi.org/10.1007/s00449-005-0005-4
- Coelho, M.A.Z., Amaral, P.F.F., Belo, I., 2010. *Yarrowia lipolytica* : an industrial workhorse. Applied
 microbiology and microbial biotechnology 2, 930–944.
- Corchero, J.L., Gasser, B., Resina, D., Smith, W., Parrilli, E., Vázquez, F., Abasolo, I., Giuliani, M., Jäntti,
 J., Ferrer, P., Saloheimo, M., Mattanovich, D., Schwartz, S., Tutino, M.L., Villaverde, A., 2013.

- 666 Unconventional microbial systems for the cost-efficient production of high-quality protein 667 therapeutics. Biotechnol Adv 31, 140–153. https://doi.org/10.1016/j.biotechadv.2012.09.001
- Cortés, G., Trujillo-Roldán, M.A., Ramírez, O.T., Galindo, E., 2005. Production of β-galactosidase by
 Kluyveromyces marxianus under oscillating dissolved oxygen tension. Process Biochemistry 40,
 773–778. https://doi.org/10.1016/j.procbio.2004.02.001
- Dabros, M., Schuler, M.M., Marison, I.W., 2010. Simple control of specific growth rate in
 biotechnological fed-batch processes based on enhanced online measurements of biomass.
 Bioprocess Biosyst Eng 33, 1109–1118. https://doi.org/10.1007/s00449-010-0438-2
- Darvishi Harzevili, F., 2014. Yarrowia lipolytica in Biotechnological Applications. Springer, Cham, pp.
 17–74. https://doi.org/10.1007/978-3-319-06437-6_2
- De Brabander, P., Uitterhaegen, E., Verhoeven, E., Cruyssen, C. Vander, De Winter, K., Soetaert, W.,
 2021. In situ product recovery of bio-based industrial platform chemicals: A guideline to solvent
 selection. Fermentation 7, 26. https://doi.org/10.3390/fermentation7010026
- de Lorenzo, V., Couto, J., 2019. The important versus the exciting: reining contradictions in
 contemporary biotechnology. Microb Biotechnol 12, 32–34. https://doi.org/10.1111/17517915.13348
- de Macedo Robert, J., Garcia-Ortega, X., Montesinos-Seguí, J.L., Guimaraes Freire, D.M., Valero, F.,
 2019. Continuous operation, a realistic alternative to fed-batch fermentation for the production
 of recombinant lipase B from Candida antarctica under the constitutive promoter PGK in *Pichia pastoris*. Biochem Eng J 147, 39–47. https://doi.org/10.1016/j.bej.2019.03.027
- de Pourcq, K., Tiels, P., van Hecke, A., Geysens, S., Vervecken, W., Callewaert, N., 2012. Engineering
 Yarrowia lipolytica to Produce Glycoproteins Homogeneously Modified with the Universal
 Man3GlcNAc2 N-Glycan Core. PLoS One 7, 39976.
 https://doi.org/10.1371/JOURNAL.PONE.0039976
- Deparis, Q., Claes, A., Foulquié-Moreno, M.R., Thevelein, J.M., 2017. Engineering tolerance to
 industrially relevant stress factors in yeast cell factories. FEMS Yeast Res 17, 36.
 https://doi.org/10.1093/femsyr/fox036
- Domingues, L., Lima, N., Teixeira, J.A., 2005. Aspergillus niger β-galactosidase production by yeast in a
 continuous high cell density reactor. Process Biochemistry 40, 1151–1154.
 https://doi.org/10.1016/j.procbio.2004.04.016
- Dragosits, M., Stadlmann, J., Albiol, J., Baumann, K., Maurer, M., Gasser, B., Sauer, M., Altmann, F.,
 Ferrer, P., Mattanovich, D., 2009. The effect of temperature on the proteome of recombinant *Pichia pastoris*. J Proteome Res 8, 1380–1392. https://doi.org/10.1021/pr8007623
- Duman-Özdamar, Z.E., Binay, B., 2021. Production of Industrial Enzymes via *Pichia pastoris* as a Cell
 Factory in Bioreactor: Current Status and Future Aspects. Protein Journal 40, 367–376.
 https://doi.org/10.1007/s10930-021-09968-7
- Figli, T., 2015. Microbial growth and physiology: A call for better craftsmanship. Front Microbiol 6, 287.
 https://doi.org/10.3389/fmicb.2015.00287
- Eigenstetter, G., Takors, R., 2017. Dynamic modeling reveals a three-step response of *Saccharomyces cerevisiae* to high CO2 levels accompanied by increasing ATP demands. FEMS Yeast Res 17, 8.
 https://doi.org/10.1093/femsyr/fox008

- Fernandes, P.M.B., 2005. How does yeast respond to pressure? Brazilian Journal of Medical and
 Biological Research 38, 1239–1245. https://doi.org/10.1590/S0100-879X2005000800012
- First Wave BioPharma, 2022. First Wave BioPharma provides update on adriulipase (FW-EPI) clinical
 program. https://www.firstwavebio.com/firstwavebio-news/58-2022-news/352 irstaveioharmarovidespdateondrulipase20220113120502.
- Fischer, J.E., Glieder, A., 2019. Current advances in engineering tools for *Pichia pastoris*. Curr Opin
 Biotechnol 59, 175–181. https://doi.org/10.1016/J.COPBIO.2019.06.002
- Fisher, A.C., Kamga, M.H., Agarabi, C., Brorson, K., Lee, S.L., Yoon, S., 2019. The Current Scientific and
 Regulatory Landscape in Advancing Integrated Continuous Biopharmaceutical Manufacturing.
 Trends Biotechnol. https://doi.org/10.1016/j.tibtech.2018.08.008
- Garcia-Ochoa, F., Gomez, E., 2009. Bioreactor scale-up and oxygen transfer rate in microbial processes:
 An overview. Biotechnol Adv 27, 153–176. https://doi.org/10.1016/j.biotechadv.2008.10.006
- Garcia-Ortega, X., Adelantado, N., Ferrer, P., Montesinos, J.L., Valero, F., 2016. A step forward to
 improve recombinant protein production in *Pichia pastoris*: From specific growth rate effect on
 protein secretion to carbon-starving conditions as advanced strategy. Process Biochemistry 51,
 681–691. https://doi.org/10.1016/j.procbio.2016.02.018
- Garcia-Ortega, X., Valero, F., Montesinos-Seguí, J.L., 2017. Physiological state as transferable operating
 criterion to improve recombinant protein production in *Pichia pastoris* through oxygen limitation.
 Journal of Chemical Technology and Biotechnology 92, 2573–2582.
 https://doi.org/10.1002/jctb.5272
- Garrigós-Martínez, J., Nieto-Taype, M.A., Gasset-Franch, A., Montesinos-Seguí, J.L., Garcia-Ortega, X.,
 Valero, F., 2019. Specific growth rate governs AOX1 gene expression, affecting the production
 kinetics of *Pichia pastoris* (*Komagataella phaffii*) P AOX1 -driven recombinant producer strains
 with different target gene dosage. Microb Cell Fact 18, 1–15. https://doi.org/10.1186/s12934 019-1240-8
- Garrigós-Martínez, J., Vuoristo, K., Nieto-Taype, M.A., Tähtiharju, J., Uusitalo, J., Tukiainen, P., Schmid,
 C., Tolstorukov, I., Madden, K., Penttilä, M., Montesinos-Seguí, J.L., Valero, F., Glieder, A., GarciaOrtega, X., 2021. Bioprocess performance analysis of novel methanol-independent promoters for
 recombinant protein production with *Pichia pastoris*. Microb Cell Fact 20, 1–12.
 https://doi.org/10.1186/s12934-021-01564-9
- Gasset, A., Garcia-Ortega, X., Garrigós-Martínez, J., Valero, F., Montesinos-Seguí, J.L., 2022. Innovative
 Bioprocess Strategies Combining Physiological Control and Strain Engineering of *Pichia pastoris* to Improve Recombinant Protein Production. Front Bioeng Biotechnol 10, 1–12.
 https://doi.org/10.3389/FBIOE.2022.818434
- George, S., Larsson, G., Enfors, S.O., 1993. A scale-down two-compartment reactor with controlled
 substrate oscillations: Metabolic response of *Saccharomyces cerevisiae*. Bioprocess Engineering
 9, 249–257. https://doi.org/10.1007/BF01061530
- Gombert, A.K., Madeira, J.V., Cerdán, M.E., González-Siso, M.I., 2016. *Kluyveromyces marxianus* as a
 host for heterologous protein synthesis. Appl Microbiol Biotechnol 100, 6193–6208.
 https://doi.org/10.1007/s00253-016-7645-y

- Gomes, A.M.V., Carmo, T.S., Carvalho, L.S., Bahia, F.M., Parachin, N.S., 2018. Comparison of yeasts as
 hosts for recombinant protein production. Microorganisms.
 https://doi.org/10.3390/microorganisms6020038
- 750 Górak, A., Stankiewicz, A., 2018. Intensification of Biobased Processes. The Royal Society of Chemistry.
- Gorczyca, M., Kaźmierczak, J., Steels, S., Fickers, P., Celińska, E., 2020. Impact of oxygen availability on
 heterologous geneexpression and polypeptide secretion dynamics in *Yarrowia lipolytica*-based
 protein production platforms. Yeast 37, 559–568. https://doi.org/10.1002/yea.3499
- Guimarães, P.M.R., Teixeira, J.A., Domingues, L., 2010. Fermentation of lactose to bio-ethanol by
 yeasts as part of integrated solutions for the valorisation of cheese whey. Biotechnol Adv 28,
 375–384. https://doi.org/10.1016/j.biotechadv.2010.02.002
- Hakkaart, X., Liu, Y., Hulst, M., el Masoudi, A., Peuscher, E., Pronk, J., van Gulik, W., Daran-Lapujade,
 P., 2020. Physiological responses of *Saccharomyces cerevisiae* to industrially relevant conditions:
 Slow growth, low pH, and high CO2 levels. Biotechnol Bioeng 117, 721–735.
 https://doi.org/10.1002/bit.27210
- Haringa, C., Mudde, R.F., Noorman, H.J., 2018. From industrial fermentor to CFD-guided downscaling:
 what have we learned? Biochem Eng J 140, 57–71. https://doi.org/10.1016/j.bej.2018.09.001
- Heo, J.H., Ananin, V., Kang, H.A., Rhee, S.K., Kim, C.H., 2008. Feeding strategies for the enhanced
 production of recombinant human serum albumin in the fed-batch cultivation of *Hansenula polymorpha*. Process Biochemistry 43, 918–924. https://doi.org/10.1016/j.procbio.2008.04.017
- Huang, C., Lowe, A.J., Batt, C.A., 2010. Recombinant immunotherapeutics: Current state and
 perspectives regarding the feasibility and market. Appl Microbiol Biotechnol 87, 401–410.
 https://doi.org/10.1007/s00253-010-2590-7
- 769 Invitrogen, 2002. *Pichia* fermentation process guidelines 1–11.
- Jacobs, P.P., Geysens, S., Vervecken, W., Contreras, R., Callewaert, N., 2009. Engineering complex-type
 N-glycosylation in *Pichia pastoris* using GlycoSwitch technology. Nat Protoc 4, 58–70.
 https://doi.org/10.1038/nprot.2008.213
- Jahic, M., Gustavsson, M., Jansen, A.K., Martinelle, M., Enfors, S.O., 2003a. Analysis and control of
 proteolysis of a fusion protein in *Pichia pastoris* fed-batch processes. J Biotechnol 102, 45–53.
 https://doi.org/10.1016/S0168-1656(03)00003-8
- Jahic, M., Rotticci-Mulder, J., Martinelle, M., Hult, K., Enfors, S.O., 2002. Modeling of growth and
 energy metabolism of *Pichia pastoris* producing a fusion protein. Bioprocess Biosyst Eng 24, 385–
 393. https://doi.org/10.1007/s00449-001-0274-5
- Jahic, M., Wallberg, F., Bollok, M., Garcia, P., Enfors, S.O., 2003b. Temperature limited fed-batch
 technique for control of proteolysis in *Pichia pastoris* bioreactor cultures. Microb Cell Fact 2, 6.
 https://doi.org/10.1186/1475-2859-2-6
- Jin, H., Liu, G., Dai, K., Wang, H., Li, Z., Shi, Z., 2011. Improvement of porcine interferon-α production
 by recombinant *pichia pastoris* via induction at low methanol concentration and low
 temperature. Appl Biochem Biotechnol 165, 559–571. https://doi.org/10.1007/s12010-011 9275-2

- Jungo, C., Marison, I., von Stockar, U., 2007a. Mixed feeds of glycerol and methanol can improve the
 performance of *Pichia pastoris* cultures: A quantitative study based on concentration gradients
 in transient continuous cultures. J Biotechnol 128, 824–837.
 https://doi.org/10.1016/j.jbiotec.2006.12.024
- Jungo, C., Schenk, J., Pasquier, M., Marison, I.W., von Stockar, U., 2007b. A quantitative analysis of the
 benefits of mixed feeds of sorbitol and methanol for the production of recombinant avidin with
 Pichia pastoris. J Biotechnol 131, 57–66. https://doi.org/10.1016/J.JBIOTEC.2007.05.019
- Kar, T., Destain, J., Thonart, P., Delvigne, F., 2010. Impact of scaled-down on dissolved oxygen
 fluctuations at different levels of the lipase synthesis pathway of *Yarrowia lipolytica*.
 Biotechnology, Agronomy, Society and Environment 14, 523–529.
- Karim, A., Gerliani, N., Aïder, M., 2020. *Kluyveromyces marxianus*: An emerging yeast cell factory for
 applications in food and biotechnology. Int J Food Microbiol 333, Article 108818.
 https://doi.org/10.1016/j.ijfoodmicro.2020.108818
- Knoll, A., Bartsch, S., Husemann, B., Engel, P., Schroer, K., Ribeiro, B., Stöckmann, C., Seletzky, J., Büchs,
 J., 2007. High cell density cultivation of recombinant yeasts and bacteria under non-pressurized
 and pressurized conditions in stirred tank bioreactors. J Biotechnol 132, 167–179.
 https://doi.org/10.1016/J.JBIOTEC.2007.06.010
- Kong, N., Mu, X., Han, H., Yan, W., 2009. Pilot-scale fermentation, purification, and characterization of
 recombinant human Oncostatin M in *Pichia pastoris*. Protein Expr Purif 63, 134–139.
 https://doi.org/10.1016/j.pep.2008.10.002
- Krainer, F.W., Dietzsch, C., Hajek, T., Herwig, C., Spadiut, O., Glieder, A., 2012. Recombinant protein
 expression in *Pichia pastoris* strains with an engineered methanol utilization pathway. Microb
 Cell Fact 11, 1–14. https://doi.org/10.1186/1475-2859-11-22
- Labrou, N.E., 2014. Protein Purification: An Overview, in: Methods in Molecular Biology. Humana Press,
 Totowa, NJ, pp. 3–10. https://doi.org/10.1007/978-1-62703-977-2_1
- Lane, M.M., Morrissey, J.P., 2010. *Kluyveromyces marxianus*: A yeast emerging from its sister's
 shadow. Fungal Biol Rev 24, 17–26. https://doi.org/10.1016/J.FBR.2010.01.001
- Lara, A.R., Galindo, E., Ramírez, O.T., Palomares, L.A., 2006. Living with heterogeneities in bioreactors:
 Understanding the effects of environmental gradients on cells. Mol Biotechnol 34, 355–381.
 https://doi.org/10.1385/MB:34:3:355
- Larsson, G., Törnkvist, M., Ståhl Wernersson, E., Trägårdh, C., Noorman, H., Enfors, S.O., 1996.
 Substrate gradients in bioreactors: Origin and consequences. Bioprocess Engineering 14, 281–
 289. https://doi.org/10.1007/BF00369471
- Leblond, Y., Marty, A., Mouz, N., Uribelarrea, J.L., 2012. Method for producing lipase, trans-formed
 Yarrowia lipolytica cell capable of producing said lipase and their uses. US8834867B2.
- Lee, J., Liu, Z., Chen, W.H., Wei, J., Kundu, R., Adhikari, R., Rivera, J.A., Gillespie, P.M., Strych, U., Zhan,
 B., Hotez, P.J., Bottazzi, M.E., 2021. Process development and scale-up optimization of the SARS CoV-2 receptor binding domain-based vaccine candidate, RBD219-N1C1. Appl Microbiol
 Biotechnol 105, 4153–4165. https://doi.org/10.1007/s00253-021-11281-3

- Lehnen, M., Ebert, B.E., Blank, L.M., 2019. Elevated temperatures do not trigger a conserved metabolic
 network response among thermotolerant yeasts. BMC Microbiol 19, 1–11.
 https://doi.org/10.1186/s12866-019-1453-3
- Lejeune, A., Delvigne, F., Thonart, P., 2010. Influence of bioreactor hydraulic characteristics on a
 Saccharomyces cerevisiae fed-batch culture: Hydrodynamic modelling and scale-down
 investigations. J Ind Microbiol Biotechnol 37, 225–236. https://doi.org/10.1007/s10295-009 0564-5
- Li, P., Anumanthan, A., Gao, X.G., Ilangovan, K., Suzara, V. V., Düzgüneş, N., Renugopalakrishnan, V.,
 2007. Expression of recombinant proteins in *Pichia pastoris*. Appl Biochem Biotechnol 142, 105–
 124. https://doi.org/10.1007/s12010-007-0003-x
- Li, X., He, X., Li, Z., Wang, F., 2013. Combined strategies for improving the production of recombinant
 Rhizopus oryzae lipase in pichia pastoris. Bioresources 8, 2867–2880.
 https://doi.org/10.15376/biores.8.2.2867-2880
- Liu, W.C., Gong, T., Wang, Q.H., Liang, X., Chen, J.J., Zhu, P., 2016. Scaling-up Fermentation of *Pichia pastoris* to demonstration-scale using new methanol-feeding strategy and increased air pressure
 instead of pure oxygen supplement. Sci Rep 6, 1–12. https://doi.org/10.1038/srep18439
- Liu, Z., Hou, J., Martínez, J.L., Petranovic, D., Nielsen, J., 2013. Correlation of cell growth and
 heterologous protein production by *Saccharomyces cerevisiae*. Appl Microbiol Biotechnol 97,
 8955–8962. https://doi.org/10.1007/s00253-013-4715-2
- Looser, V., Bruhlmann, B., Bumbak, F., Stenger, C., Costa, M., Camattari, A., Fotiadis, D., Kovar, K., 2014.
 Cultivation strategies to enhance productivity of *Pichia pastoris*: A review. Biotechnol Adv 33,
 1177–1193. https://doi.org/10.1016/j.biotechadv.2015.05.008
- Lopes, M., Gomes, N., Gonçalves, C., Coelho, M.A.Z., Mota, M., Belo, I., 2008. Yarrowia lipolytica lipase
 production enhanced by increased air pressure. Lett Appl Microbiol 46, 255–260.
 https://doi.org/10.1111/j.1472-765X.2007.02299.x
- Lorantfy, B., Jazini, M., Herwig, C., 2013. Investigation of the physiological response to oxygen limited
 process conditions of *Pichia pastoris* Mut+ strain using a two-compartment scale-down system. J
 Biosci Bioeng 116, 371–379. https://doi.org/10.1016/J.JBIOSC.2013.03.021
- Ma, B., Guan, X., Li, Y., Shang, S., Li, J., Tan, Z., 2020. Protein Glycoengineering: An Approach for
 Improving Protein Properties. Front Chem 8, 622. https://doi.org/10.3389/fchem.2020.00622
- Madhavan, A., Arun, K.B., Sindhu, R., Krishnamoorthy, J., Reshmy, R., Sirohi, R., Pugazhendi, A.,
 Awasthi, M.K., Szakacs, G., Binod, P., 2021. Customized yeast cell factories for
 biopharmaceuticals: from cell engineering to process scale up. Microb Cell Fact.
 https://doi.org/10.1186/s12934-021-01617-z
- Madzak, C., Beckerich, J.-M., 2013. Heterologous Protein Expression and Secretion in *Yarrowia lipolytica*. Springer, Berlin, Heidelberg, pp. 1–76. https://doi.org/10.1007/978-3-642-38583-4_1
- Mattanovich, D., Branduardi, P., Dato, L., Gasser, B., Sauer, M., Porro, D., 2012. Recombinant protein
 production in yeasts. Methods in Molecular Biology 824, 329–358. https://doi.org/10.1007/978 1-61779-433-9_17
- Mears, L., Feldman, H., Falco, F.C., Bach, C., Wu, M., Nørregaard, A., 2017. Continuous fermentation
 for biopharmaceuticals?, in: Kleinebudde, P., Khinast, J., Rantanen, J. (Eds.), Continuous

- Manufacturing of Pharmaceuticals. John Wiley & Sons, pp. 227–246.
 https://doi.org/10.1002/9781119001348
- Mears, Lisa, Stocks, S.M., Sin, G., Gernaey, K. V., 2017. A review of control strategies for manipulating
 the feed rate in fed-batch fermentation processes. J Biotechnol 245, 34–46.
 https://doi.org/10.1016/j.jbiotec.2017.01.008
- Mordor Intelligence, 2021. Recombinant Protein Market | 2021 26 | Industry Share, Size, Growth Mordor Intelligence [WWW Document]. URL https://www.mordorintelligence.com/industry reports/recombinant-protein-market (accessed 2.8.22).
- Nadal-Rey, G., McClure, D.D., Kavanagh, J.M., Cornelissen, S., Fletcher, D.F., Gernaey, K. V., 2021.
 Understanding gradients in industrial bioreactors. Biotechnol Adv 46, 107660.
 https://doi.org/10.1016/j.biotechadv.2020.107660
- Nieto-Taype, M.A., Garcia-Ortega, X., Albiol, J., Montesinos-Seguí, J.L., Valero, F., 2020. Continuous
 Cultivation as a Tool Toward the Rational Bioprocess Development With *Pichia Pastoris* Cell
 Factory. Front Bioeng Biotechnol 8, 632. https://doi.org/10.3389/fbioe.2020.00632
- Niu, H., Jost, L., Pirlot, N., Sassi, H., Daukandt, M., Rodriguez, C., Fickers, P., 2013. A quantitative study
 of methanol/sorbitol co-feeding process of a *Pichia pastoris* Mut+/pAOX1-lacZ strain. Microb Cell
 Fact 12, 1–8. https://doi.org/10.1186/1475-2859-12-33
- Nurcholis, M., Lertwattanasakul, N., Rodrussamee, N., Kosaka, T., Murata, M., Yamada, M., 2020.
 Integration of comprehensive data and biotechnological tools for industrial applications of
 Kluyveromyces marxianus. Appl Microbiol Biotechnol 104, 475–488.
 https://doi.org/10.1007/S00253-019-10224-3/METRICS
- Olughu, W., Deepika, G., Hewitt, C., Rielly, C., 2019. Insight into the large-scale upstream fermentation
 environment using scaled-down models. Journal of Chemical Technology and Biotechnology 94,
 647–657. https://doi.org/10.1002/jctb.5804
- Pan, Y., Yang, J., Wu, J., Yang, L., Fang, H., 2022. Current advances of *Pichia pastoris* as cell factories for
 production of recombinant proteins. Front Microbiol 13, 4702.
 https://doi.org/10.3389/FMICB.2022.1059777/BIBTEX
- Park, Y.-K., Ledesma-Amaro, R., 2022. What makes *Yarrowia lipolytica* well suited for industry? Trends
 Biotechnol. https://doi.org/10.1016/J.TIBTECH.2022.07.006
- Partow, S., Siewers, V., Bjørn, S., Nielsen, J., Maury, J., 2010. Characterization of different promoters
 for designing a new expression vector in *Saccharomyces cerevisiae*. Yeast 27, 955–964.
 https://doi.org/10.1002/YEA.1806
- Patra, P., Das, M., Kundu, P., Ghosh, A., 2021. Recent advances in systems and synthetic biology
 approaches for developing novel cell-factories in non-conventional yeasts. Biotechnol Adv 47,
 Article 107695. https://doi.org/10.1016/j.biotechadv.2021.107695
- 901Peebo, K., Neubauer, P., 2018. Application of continuous culture methods to recombinant protein902productioninmicroorganisms.Microorganisms6,56.903https://doi.org/10.3390/microorganisms6030056
- Ponte, X., Barrigón, J.M., Maurer, M., Mattanovich, D., Valero, F., Montesinos-Seguí, J.L., 2018.
 Towards optimal substrate feeding for heterologous protein production in *Pichia pastoris* (Komagataella spp) fed-batch processes under PAOX1 control: a modeling aided approach.

- 907JournalofChemicalTechnologyandBiotechnology93,3208–3218.908https://doi.org/10.1002/jctb.5677
- Potvin, G., Ahmad, A., Zhang, Z., 2012. Bioprocess engineering aspects of heterologous protein
 production in *Pichia pastoris*: A review. Biochem Eng J. https://doi.org/10.1016/j.bej.2010.07.017
- Puetz, J., Wurm, F.M., 2019. Recombinant Proteins for Industrial versus Pharmaceutical Purposes: A
 Review of Process and Pricing. Processes 7, 476. https://doi.org/10.3390/pr7080476
- Rahimi, A., Hosseini, S.N., Karimi, A., Aghdasinia, H., Arabi Mianroodi, R., 2019. Enhancing the efficiency
 of recombinant hepatitis B surface antigen production in *Pichia pastoris* by employing continuous
 fermentation. Biochem Eng J 141, 112–119. https://doi.org/10.1016/j.bej.2018.10.019
- 916 Raschmanová, H., Weninger, A., Glieder, A., Kovar, K., Vogl, T., 2018. Implementing CRISPR-Cas
 917 technologies in conventional and non-conventional yeasts: Current state and future prospects.
 918 Biotechnol Adv 36, 641–665. https://doi.org/10.1016/J.BIOTECHADV.2018.01.006
- Raschmanová, H., Weninger, A., Knejzlík, Z., Melzoch, K., Kovar, K., 2021. Engineering of the unfolded
 protein response pathway in *Pichia pastoris*: enhancing production of secreted recombinant
 proteins. Applied Microbiology and Biotechnology 2021 105:11 105, 4397–4414.
 https://doi.org/10.1007/S00253-021-11336-5
- Rebnegger, C., Vos, T., Graf, A.B., Valli, M., Pronk, J.T., Daran-Lapujade, P., Mattanovich, D., 2016. *Pichia Pastoris* exhibits high viability and a low maintenance energy requirement at near-zero
 specific growth rates. Appl Environ Microbiol 82, 4570–4583.
 https://doi.org/10.1128/AEM.00638-16
- Reuss, M., Schmalzriedt, S., Jenne, M., 1994. Structured Modelling of Bioreactors, in: Advances in
 Bioprocess Engineering. Springer, Dordrecht, pp. 207–215. https://doi.org/10.1007/978-94-017 0641-4_29
- Risager Wright, N., Rønnest, N.P., Thykaer, J., 2016. Scale-down of continuous protein producing
 Saccharomyces cerevisiae cultivations using a two-compartment system. Biotechnol Prog 32,
 152–159. https://doi.org/10.1002/btpr.2184
- Ritala, A., Häkkinen, S.T., Toivari, M., Wiebe, M.G., 2017. Single cell protein-state-of-the-art, industrial
 landscape and patents 2001-2016. Front Microbiol. https://doi.org/10.3389/fmicb.2017.02009
- Roe, S., Streck, C., Obersteiner, M., Frank, S., Griscom, B., Drouet, L., Fricko, O., Gusti, M., Harris, N.,
 Hasegawa, T., Hausfather, Z., Havlík, P., House, J., Nabuurs, G.J., Popp, A., Sánchez, M.J.S.,
 Sanderman, J., Smith, P., Stehfest, E., Lawrence, D., 2019. Contribution of the land sector to a 1.5
 °C world. Nat Clim Chang 9, 817–828. https://doi.org/10.1038/s41558-019-0591-9
- Shenoy, A., Yalamanchili, S., Davis, A.R., Barb, A.W., 2021. Expression and display of glycoengineered
 antibodies and antibody fragments with an engineered yeast strain. Antibodies 10, 38.
 https://doi.org/10.3390/antib10040038
- Signori, L., Passolunghi, S., Ruohonen, L., Porro, D., Branduardi, P., 2014. Effect of oxygenation and
 temperature on glucose-xylose fermentation in *Kluyveromyces marxianus* CBS712 strain. Microb
 Cell Fact 13, 1–13. https://doi.org/10.1186/1475-2859-13-51
- Soyaslan, E.Ş., Çalik, P., 2011. Enhanced recombinant human erythropoietin production by Pichia
 pastoris in methanol fed-batch/sorbitol batch fermentation through pH optimization. Biochem
 Eng J 55, 59–65. https://doi.org/10.1016/j.bej.2011.03.007

- Spohner, S.C., Schaum, V., Quitmann, H., Czermak, P., 2016. *Kluyveromyces lactis*: An emerging tool in
 biotechnology. J Biotechnol 222, 104–116. https://doi.org/10.1016/j.jbiotec.2016.02.023
- Stehfest, E., van Zeist, W.J., Valin, H., Havlik, P., Popp, A., Kyle, P., Tabeau, A., Mason-D'Croz, D.,
 Hasegawa, T., Bodirsky, B.L., Calvin, K., Doelman, J.C., Fujimori, S., Humpenöder, F., LotzeCampen, H., van Meijl, H., Wiebe, K., 2019. Key determinants of global land-use projections. Nat
 Commun 10, 1–10. https://doi.org/10.1038/s41467-019-09945-w
- Stratton, J., Chiruvolu, V., Meagher, M., 1998. High cell-density fermentation. Methods Mol Biol 103,
 107–120. https://doi.org/10.1385/0-89603-421-6:107
- Sun, M.L., Shi, T.Q., Lin, L., Ledesma-Amaro, R., Ji, X.J., 2022. Advancing *Yarrowia lipolytica* as a superior
 biomanufacturing platform by tuning gene expression using promoter engineering. Bioresour
 Technol 347, 126717. https://doi.org/10.1016/J.BIORTECH.2022.126717
- 959 Tavasoli, T., Arjmand, S., Ranaei Siadat, S.O., Shojaosadati, S.A., Sahebghadam Lotfi, A., 2019. A robust 960 feeding control strategy adjusted and optimized by a neural network for enhancing of alpha 1-144, 961 antitrypsin production Pichia pastoris. Biochem Eng 18-27. in J https://doi.org/10.1016/j.bej.2019.01.005 962
- Timoumi, A., Cléret, M., Bideaux, C., Guillouet, S.E., Allouche, Y., Molina-Jouve, C., Fillaudeau, L.,
 Gorret, N., 2017. Dynamic behavior of *Yarrowia lipolytica* in response to pH perturbations:
 dependence of the stress response on the culture mode. Appl Microbiol Biotechnol 101, 351–
 366. https://doi.org/10.1007/s00253-016-7856-2
- Tolner, B., Smith, L., Begent, R.H.J., Chester, K.A., 2006. Production of recombinant protein in *Pichia pastoris* by fermentation. Nat Protoc 1, 1006–1021. https://doi.org/10.1038/nprot.2006.126
- Trentmann, O., Khatri, N.K., Hoffmann, F., 2004. Reduced oxygen supply increases process stability and
 product yield with recombinant *Pichia pastoris*. Biotechnol Prog 20, 1766–1775.
 https://doi.org/10.1021/bp049711h
- Uribelarrea, J. -L, Winter, J., Goma, G., Pareilleux, A., 1990. Determination of maintenance coefficients
 of *Saccharomyces cerevisiae* cultures with cell recycle by cross-flow membrane filtration.
 Biotechnol Bioeng 35, 201–206. https://doi.org/10.1002/BIT.260350211
- 975 Vandermies, M., Fickers, P., 2019. Bioreactor-scale strategies for the production of recombinant
 976 protein in the yeast *Yarrowia lipolytica*. Microorganisms 7, 40.
 977 https://doi.org/10.3390/microorganisms7020040
- 978 Vantage Market Research, 2022. Alternative Protein Market | \$126.84 Mn by 2028 | Growth [WWW
 979 Document]. URL https://www.globenewswire.com/news 980 release/2022/03/16/2404160/0/en/Alternative-Protein-Market-126-84-Mn-by-2028-Growth-
- 981 Forecast-at-16-90-CAGR-During-2022-to-2028-COVID19-Impact-and-Global-Analysis-by-
- 982 Vantage-Market-Research.html (accessed 6.9.22).
- Verfied Market Research, 2021. Industrial Enzymes Market Size, Share, Trends, Opportunities &
 Forecast [WWW Document]. URL https://www.verifiedmarketresearch.com/product/industrial enzymes-market/ (accessed 2.8.22).
- Vogl, T., Glieder, A., 2013. Regulation of *Pichia pastoris* promoters and its consequences for protein
 production. N Biotechnol 30, 385–404. https://doi.org/10.1016/j.nbt.2012.11.010

- Vuree, S., 2020. Pichia pastoris expression system: An impending candidate to express protein in industrial and biopharmaceutical domains. New and Future Developments in Microbial
 Biotechnology and Bioengineering 223–234. https://doi.org/10.1016/B978-0-12-821007 9.00017-6
- Waegeman, H., Soetaert, W., 2011. Increasing recombinant protein production in *Escherichia coli* through metabolic and genetic engineering. J Ind Microbiol Biotechnol 38, 1891–1910.
 https://doi.org/10.1007/s10295-011-1034-4
- Walsh, G., Walsh, E., 2022. Biopharmaceutical benchmarks 2022. Nature Biotechnology 2022 40:12 40,
 1722–1760. https://doi.org/10.1038/s41587-022-01582-x
- Walther, J., Godawat, R., Hwang, C., Abe, Y., Sinclair, A., Konstantinov, K., 2015. The business impact
 of an integrated continuous biomanufacturing platform for recombinant protein production. J
 Biotechnol 213, 3–12. https://doi.org/10.1016/j.jbiotec.2015.05.010
- 1000 Wang, G., Huang, M., Nielsen, J., 2017. Exploring the potential of *Saccharomyces cerevisiae* for
 1001 biopharmaceutical protein production. Curr Opin Biotechnol.
 1002 https://doi.org/10.1016/j.copbio.2017.03.017
- Wang, J., Wang, X., Shi, L., Qi, F., Zhang, P., Zhang, Y., Zhou, X., Song, Z., Cai, M., 2017. Methanol Independent Protein Expression by AOX1 Promoter with trans-Acting Elements Engineering and
 Glucose-Glycerol-Shift Induction in *Pichia pastoris*. Scientific Reports 2017 7:1 7, 1–12.
 https://doi.org/10.1038/srep41850
- 1007 Wang, X., Sun, Y., Ke, F., Zhao, H., Liu, T., Xu, L., Liu, Y., Yan, Y., 2012a. Constitutive expression of
 1008 *Yarrowia lipolytica* lipase LIP2 in *Pichia pastoris* using GAP as promoter. Appl Biochem Biotechnol
 1009 166, 1355–1367. https://doi.org/10.1007/s12010-011-9524-4
- Wang, X., Sun, Y., Shen, X., Ke, F., Zhao, H., Liu, Y., Xu, L., Yan, Y., 2012b. Intracellular expression of
 Vitreoscilla hemoglobin improves production of *Yarrowia lipolytica* lipase LIP2 in a recombinant
 Pichia pastoris. Enzyme Microb Technol 50, 22–28.
 https://doi.org/10.1016/j.enzmictec.2011.09.003
- Xiong, Z.Q., Guo, M.J., Chu, J., Zhuang, Y.P., Zhang, S.L., 2015. On-line specific growth rate control for
 improving reduced glutathione production in *Saccharomyces cerevisiae*. Biotechnology and
 Bioprocess Engineering 20, 887–893. https://doi.org/10.1007/s12257-015-0018-z
- 1017Yang, X., 2010. Scale-Up of Microbial Fermentation Process, in: Manual of Industrial Microbiology and1018Biotechnology.JohnWiley& Sons,Ltd,pp.669–675.1019https://doi.org/10.1128/9781555816827.CH47
- Yang, Z., Zhang, Z., 2018. Engineering strategies for enhanced production of protein and bio-products
 in *Pichia pastoris*: A review. Biotechnol Adv 36, 182–195.
 https://doi.org/10.1016/j.biotechadv.2017.11.002
- Ye, J., Ly, J., Watts, K., Hsu, A., Walker, A., Mclaughlin, K., Berdichevsky, M., Prinz, B., Sean Kersey, D.,
 D'Anjou, M., Pollard, D., Potgieter, T., 2011. Optimization of a glycoengineered *Pichia pastoris* cultivation process for commercial antibody production. Biotechnol Prog 27, 1744–1750.
 https://doi.org/10.1002/btpr.695

- Yu, X.W., Sun, W.H., Wang, Y.Z., Xu, Y., 2017. Identification of novel factors enhancing recombinant
 protein production in multi-copy *Komagataella phaffii* based on transcriptomic analysis of
 overexpression effects. Sci Rep 7, Article 16249. https://doi.org/10.1038/S41598-017-16577-X
- Zahrl, R.J., Peña, D.A., Mattanovich, D., Gasser, B., 2017. Systems biotechnology for protein production
 in Pichia pastoris. FEMS Yeast Res 17, 68. https://doi.org/10.1093/femsyr/fox068
- Zalai, D., Dietzsch, C., Herwig, C., Spadiut, O., 2012. A dynamic fed batch strategy for a *Pichia pastoris*mixed feed system to increase process understanding. Biotechnol Prog 28, 878–886.
 https://doi.org/10.1002/btpr.1551
- Zavec, D., Gasser, B., Mattanovich, D., 2020. Characterization of methanol utilization negative *Pichia pastoris* for secreted protein production: New cultivation strategies for current and future
 applications. Biotechnol Bioeng 117, 1394–1405. https://doi.org/10.1002/bit.27303
- Zhao, W., Wang, J., Deng, R., Wang, X., 2008. Scale-up fermentation of recombinant *Candida rugosa* lipase expressed in *Pichia pastoris* using the GAP promoter. J Ind Microbiol Biotechnol 35, 189–
 195. https://doi.org/10.1007/s10295-007-0283-8
- Zheng, J.Y., Janis, L.J., 2006. Influence of pH, buffer species, and storage temperature on
 physicochemical stability of a humanized monoclonal antibody LA298. Int J Pharm 308, 46–51.
 https://doi.org/10.1016/J.IJPHARM.2005.10.024
- Zhong, Y., Yang, L., Guo, Y., Fang, F., Wang, D., Li, R., Jiang, M., Kang, W., Ma, J., Sun, J., Xiao, W., 2014.
 High-temperature cultivation of recombinant *Pichia pastoris* increases endoplasmic reticulum
 stress and decreases production of human interleukin-10. Microb Cell Fact 13, 1–10.
 https://doi.org/10.1186/s12934-014-0163-7
- Zhou, J., Zhu, P., Hu, X., Lu, H., Yu, Y., 2018. Improved secretory expression of lignocellulolytic enzymes
 in *Kluyveromyces marxianus* by promoter and signal sequence engineering. Biotechnol Biofuels
 11, 1–14. https://doi.org/10.1186/S13068-018-1232-7/FIGURES/6
- 1051 Zhu, T., Sun, H., Li, P., Xue, Y., Li, Y., Ma, Y., 2014. Constitutive expression of alkaline β-mannanase in
 1052 recombinant *Pichia pastoris*. Process Biochemistry 49, 2025–2029.
 1053 https://doi.org/10.1016/J.PROCBIO.2014.08.014

1054