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Evaluation of biological degradation of polyurethanes

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1. Abstract

Polyurethanes (PU) are a family of versatile synthetic polymers intended for diverse applications. Biological degradation of PU is a blooming research domain as it contributes to the design of eco-friendly materials sensitive to biodegradation phenomena and the development of green recycling processes. In this field, an increasing number of studies deal with the discovery and characterization of enzymes and microorganisms able to degrade PU chains. The synthesis of short lifespan PU material sensitive to biological degradation is also of growing interest. Measurement of PU degradation can be performed by a wide range of analytical tools depending on the architecture of the materials and the biological entities. Recent developments of these analytical techniques allowed for a better understanding of the mechanisms involved in PU biodegradation. Here, we reviewed the evaluation of biological PU degradation, including the required analytics. Advantages, drawbacks, specific uses, and results of these analytics are largely discussed to provide a critical overview and support future studies.

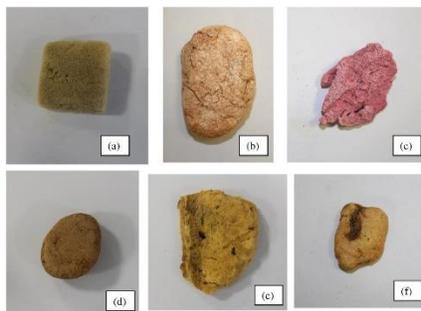
Keywords: Microbial degradation, enzymatic degradation, polyurethanes, technical review

2. Introduction

In 2019, the International Union of Pure and Applied Chemistry (IUPAC) published a list of ten emerging technologies in Chemistry with the potential to make our planet more sustainable. Among them, technologies permitting the transformation of plastic material into monomers are highlighted. These recycling technologies will help reduce plastic waste and save fossil resources. Despite comfort and incomparable uses to our everyday life, plastic materials are a cause of global and increasing pollution resulting from inadequate behaviors of both producers and users. The massive production involves a polluting exploitation of fossil resources and their poor waste management induces uncontrolled dissemination in the environment. In 2010, more than 275 million metric tons of plastic waste were generated in almost 200 coastal countries, out of which an estimated 5 to 13 million metric tons reached the oceans where plastic waste accumulate (Jambeck et al., 2015). With the problematic of the nano- or microplastic debris, ocean garbage patches are one of the major environmental concerns of this century (Cozar et al., 2014; Eriksen et al., 2013; Law et al., 2010). Even if some very minor studies still question the actual impact of plastic waste (Duis and Coors, 2016), most are warnings of irrevocable environmental damages (Clukey et al., 2018; Darmon et al., 2017; Galloway and Lewis, 2016). Furthermore, by entering

38 the food chain, plastic materials finally attain human beings, thereby causing health concerns
39 (Barboza et al., 2018; Bouwmeester et al., 2015; Chae and An, 2017).

40 Among the vast families of resistant plastic materials, we can find the polyurethanes (PUs). Low-
41 density and easily dispersible foams (soft to rigid) represent around 70% of the PU production.
42 The presence of PUs as pollutants in marine ecosystems has been largely attested (Frère et al.,
43 2016; Reddy et al., 2006). In 2016, Turner et al. revealed that over the 70 foamed plastics
44 fragments collected on a Britain beach, 39 were identified as PU (Figure 1), thus pointing out the
45 significant role of PU in plastic pollution (Turner and Lau, 2016).

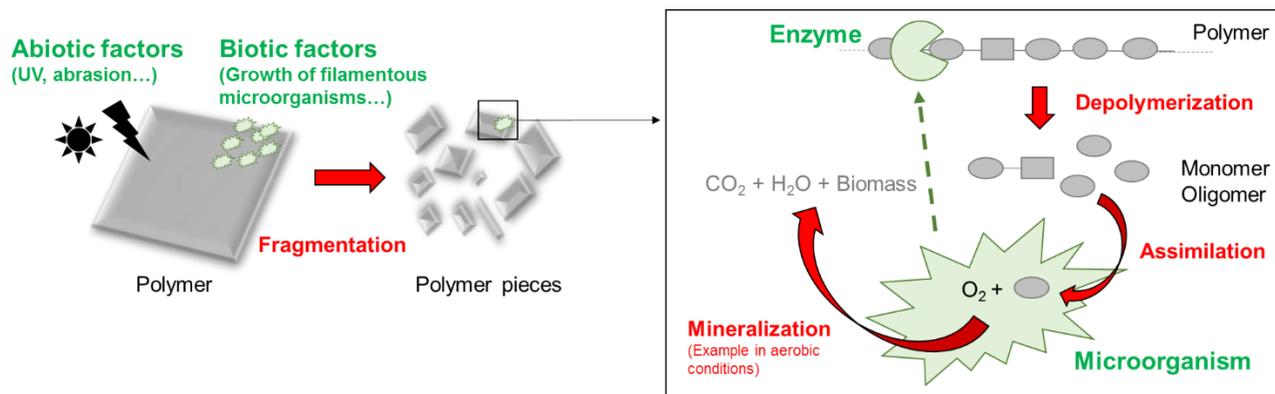


46
47 *Figure 1 –Foamed plastic debris collected on a Britain beach, pieces a, b d, e and f are PU (Turner and Lau, 2016)*

48
49 First synthesized in the 1930s by the German chemist Otto Bayer, PU products were
50 commercialized about 10 years later (Bayer, 1948). The use of PU spread during World War II,
51 where it replaced natural rubber for elastomer production. Rapidly, other applications emerged
52 in aviation and textile, which were flourishing markets at the time. Diversification of PU properties
53 allowed to reach other numerous markets. In 2016, 18 million tons of PU were produced
54 worldwide, representing 5.3% of global plastic production (Furtwengler et al., 2018a;
55 PlasticsEurope, 2017). PU rank at the 6th most produced synthetic polymers. About 22 million tons
56 are expected to be produced in 2020 (Akindoyo et al., 2016).

57 In 2014, the PU foam market was valued at \$46.8 billion and is expected to reach \$72.2 billion by
58 2020 (Pillai et al., 2016). Flexible and rigid foams represent respectively 32% and 36% of the global
59 PU production. Flexible foams, renowned for the comfort they provide, are used for the
60 cushioning of furniture, bedding or automotive seats. In the construction area, rigid foams are
61 preferred for thermal insulation and their use increases in agreement with a growing demand for
62 energetically efficient buildings. PU are also widely used as coatings, adhesives, sealants and
63 elastomers (CASE). PU coatings provide a protection layer against weather, abrasion and
64 corrosion. Elastomers are both elastic and flexible and can adopt any desired shape, such as
65 wheels for rollerblades. Biocompatibility of certain types of polyurethanes make them polymers
66 of choice for medical application, for instance, cardiovascular devices or orthopedic prosthesis
67 (Gunatillake et al., 2011; Zhou et al., 2012). The common thread between most of these PU
68 materials is that they are mostly intended for long-term applications. They are mainly designed
69 to resist environmental factors such as microbial degradation, abrasion, hydrolytic (moisture) or
70 UV degradation.

71 Biodegradation is generally defined as the decomposition/degradation of materials by the means
 72 of biological entities such as microorganisms or enzymes. This process is used for numerous
 73 industrial applications such as waste water treatment (Watanabe, 2001) or depollution of
 74 contaminated site by, for instance, polycyclic aromatic hydrocarbons (Shuttleworth and Cerniglia,
 75 1995). Polymer degradation by microorganisms is performed through several steps (Figure 2).
 76 First, materials are fragmented into pieces thanks to abiotic and biotic factors such as UV,
 77 hydrolysis, abrasion or pressure exerted by filamentous microorganisms. Growth of filaments into
 78 polymer pores provokes cracks. Then, macromolecules are cleaved by enzymatic hydrolysis
 79 and/or oxidation, leading to the release of low molar mass molecules such as oligomers and
 80 monomers. These molecules are finally assimilated and mineralized by microorganisms to
 81 promote microbial growth (Lucas et al., 2008; Shah et al., 2008b). A countless number of
 82 mineralization paths exists in nature. The step of enzymatic depolymerization or enzymatic
 83 degradation can be reproduced or mimicked *in vitro*, independently of the microbial degradation.
 84 Interestingly, the released molecules resulting from enzymatic depolymerization of polymers
 85 could be turned into building blocks and high value products for the chemistry market (Wierckx
 86 et al., 2015).



87
 88 *Figure 2 – Polymer biodegradation process*

89
 90 PU are not biodegradable polymers (Wierckx et al., 2018). Even if some PU are partly sensitive to
 91 biological degradation, they do not answer, for instance, the requirements of the European norm
 92 EN 13432 defining biodegradable and compostable materials (Avérous and Pollet, 2012; Bastioli,
 93 2005). This norm considers a material as biodegradable if the degradation reaches 90% after 6
 94 months, under composting conditions. The non-toxicity of the degradation products is also a
 95 requirement of this norm to declare a material as biodegradable or compostable. Biodegradation
 96 of PU has been studied since the 1960s. The first scientific publications on this topic aimed to
 97 evaluate the microbial degradation susceptibility of PU formulations to promote the development
 98 of highly resistant materials (Cooney, 1969; Darby and Kaplan, 1968; Edmonds and Cooney, 1968;
 99 Kanavel et al., 1966; Kaplan et al., 1968). Today, this approach is reversed to address the PU
 100 materials end-of-life issues. Due to increased environmental concerns, sensitivity to microbial
 101 degradation has become a desired feature to reduce the environmental footprint of PU materials,
 102 mainly at their ends of life (Prieto, 2016). Meanwhile, the development of bioresorbable PU
 103 materials for the biomedical industry raised interest (Pavlova and Draganova, 1993; Špírková et

104 al., 2017). Currently, PU biodegradation assessments are focused on two main purposes: (i)
105 biodegradation susceptibility of new eco-materials or materials intended for biomedical
106 purposes, thus focusing on the polymer synthesis and (ii) bioremediation or biological recycling
107 of PU, thus focusing on the biological entities capable of degradation.

108 Techniques used to evaluate the biodegradation of PU are diverse, depending on the type of PU
109 and the degrading entity. Furthermore, conflicting results and conclusions are found in the
110 scientific literature, often due to the difficulty in interpreting the analytical results. Nevertheless,
111 significant advances have recently been made on the evaluation of PU biodegradation and
112 consequently on the understanding of degradation mechanisms.

113 Review articles recently published on PU biodegradation have mainly focused on the degradation
114 by microorganisms (Mahajan and Gupta, 2015), enzymatic degradation (Loredo-Treviño et al.,
115 2011) or PU biodegradation for recycling (Cregut et al., 2013). Here, we reviewed
116 comprehensively PU biodegradation including used analytics. For a full understanding of the
117 different approaches, a first part is dedicated to the diversity of PU in term of compositions,
118 architectures and corresponding waste management. Secondly, biological entities
119 (microorganisms, enzymes) able to degrade PU and their degradation mechanisms, if known, are
120 reviewed. Finally, analytical techniques used to assess PU biodegradation are gathered. Their
121 applications, advantages and drawbacks are fully discussed.

122

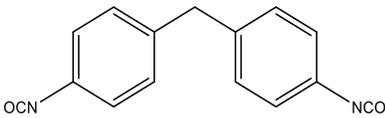
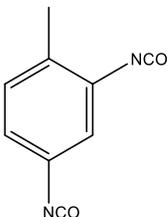
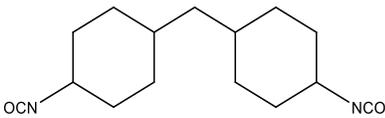
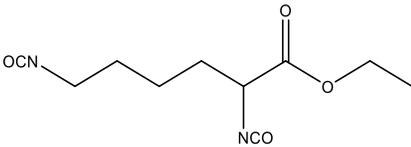
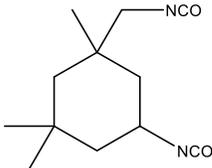
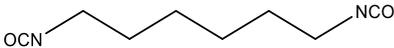
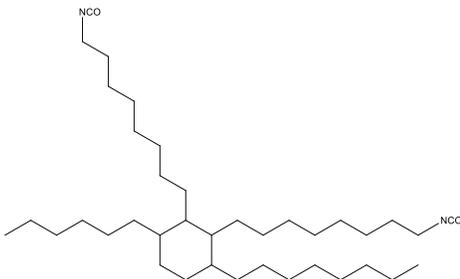
123 3. Polyurethane structure and biological degradation susceptibility 124 relationships

125 3.1. Diversity of the compositions and architectures of PU

126 Chemical composition and macromolecular architectures are of prime importance for the
127 biodegradation of polymers (Kim and Kim, 1998). The nature of chemical bonds, crystallinity and
128 molar mass are key parameters influencing the polymer susceptibility to biological attacks (Zeng
129 et al., 2016). PUs are characterized by the urethane or carbamate linkage, generally obtained by
130 addition of an isocyanate to a hydroxyl group (Figure 3). For instance, in thermoplastic PU (TPU)
131 synthesis, an excess of polyisocyanate reacts with a polyol, principally long polyester- or
132 polyether-based polyols, with a controlled functionality close to 2. A linear prepolymer with
133 isocyanate end groups is formed, followed by addition of a chain extender, usually a short diol,
134 obtain high molar mass polymers. Molar mass is a parameter influencing the biological
135 degradation susceptibility of polymers. It has been shown that for polymers with the same
136 chemical structures and different molar mass, the higher the molar mass, the lower the biological
137 degradation susceptibility (Philip et al., 2007; Zheng et al., 2005).

138

Table 1 – Structure of the some common biobased and fossil-based isocyanates

IUPAC name	Abbrev.	Type	Structure	Potentially biobased	Reference
1-isocyanato-4-[(4-isocyanatophenyl)methyl]benzene	4, 4'-MDI	Aromatic		No	(Shah et al., 2016)
2,4-diisocyanato-1-methylbenzene	2, 4-TDI	Aromatic		No	(Spontón et al., 2013)
1-Isocyanato-4-[(4-isocyanatocyclohexyl)methyl]cyclohexane	H ₁₂ MDI	Aliphatic cyclic		No	(Brzeska et al., 2015)
Ethyl Ester L-Lysine Diisocyanate	LDI	Aliphatic linear		Yes	(Zhou et al., 2012)
5-isocyanato-1-(isocyanatomethyl)-1,3,3-trimethyl-cyclohexane	IPDI	Aliphatic cyclic		No	(Pereira et al., 2012)
1,6-diisocyanatohexane	HDI	Aliphatic linear		No	(Tang et al., 2001a)
Dimer fatty acid-based diisocyanate	DDI	Aliphatic cyclic		Yes	(Charlon et al., 2014)

163

164 Common polyols are polyether, polyester or, more rarely, polycarbonates. Higher flexibility of
 165 polyether polyols makes them more convenient for polyurethane production (Krasowska et al.,
 166 2012). A non-exhaustive list of polyols with their structures is available in

167 Table 2. Polyols ordinarily used are fossil-based molecules, but an increasing number of studies
168 deal with bio-based polyols, on agreement with green chemistry principles. Polyols from vegetal
169 sources such as castor oil (Hablott et al., 2008; Trovati et al., 2010), starch (Duarah et al., 2016) or
170 aromatic biopolymers such as tannins or lignin (Ignat et al., 2011; Laurichesse et al., 2014) are
171 increasingly incorporated in PU formulations. Based on short diols such as 1,4-butanediol,
172 ethylene glycol or 1,6-hexanediol, (Akindoyo et al., 2016) chain extenders are used to obtain high-
173 molar mass polymers. Low molar mass diamines can also be used, such as ethylene diamine (Tang
174 et al., 1997), thus generating urea instead of urethane bonds.

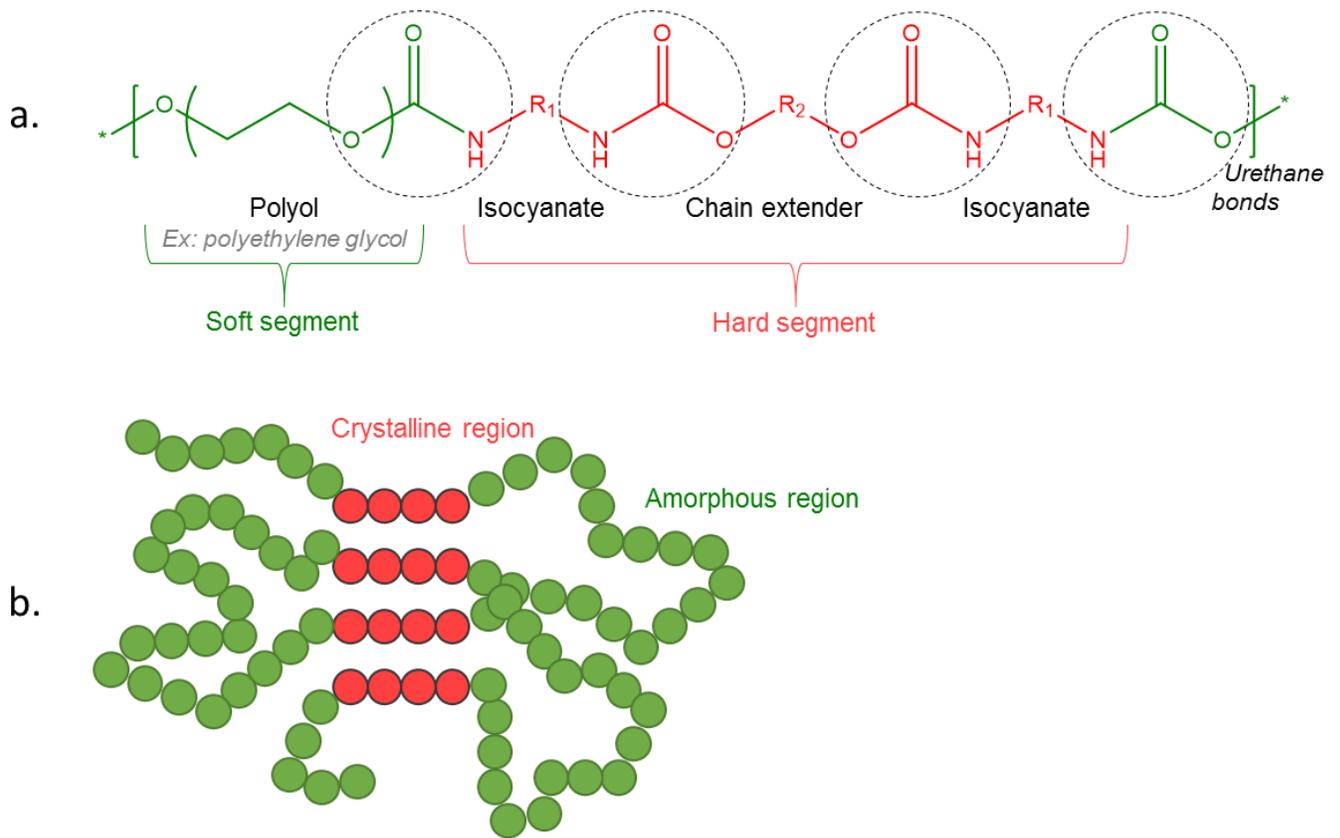
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176

Table 2 – Some biobased and fossil-based polyols used for PU synthesis

Polymer name	Abbrev.	Structure	Potentially biobased	Reference
Polyester				
Poly(caprolactone)	PCL		No	(Yeganeh and Hojati-Talemi, 2007)
Poly(lactic acid)	PLA		Yes	(Izadi-Vasafi et al., 2017)
Poly(hydroxyalkanoates)	PHA		Yes	(Debuissy et al., 2017)
Poly(butylene succinate)	PBS		Yes	(Li et al., 2015)
Poly(butylene adipate)	PBA		Yes	(Shah et al., 2013b)
Polyether				
Poly(ethylene glycol)	PEG		No	(Zhang et al., 2013)
Poly(propylene glycol)	PPG		No	(Chattopadhyay et al., 2008)
Poly(tetramethylene glycol)	PTMEG		No	(Wiggins et al., 2003)
Polycarbonate				
Poly(propylene carbonate)	PPC		No	(Chen et al., 2016)
Poly(1,6-hexyl 1,2-ethyl carbonate)	PHEC		No	(Christenson et al., 2004)

179 PUs can be thermoplastics or thermosets. Thermoplastics are linear or slightly cross-
 180 linked/branched structures. Isocyanates, polyols and chain extenders used for TPU synthesis have
 181 only two functional groups (diols or diisocyanates). Thermoplastics can be soluble in organic
 182 solvent and can be melted, or present a liquid-like behavior with increased temperature. TPUs
 183 are commonly described by two types of segments, hard and soft segments. The segments are
 184 generally organized with a specific micro-segregation which can lead to micro-crystalline phases.
 185 The hard segment (HS) is a block segment with low mobility mainly formed by the isocyanate and
 186 the short-chain extender. By contrast, the soft segment (SS) is mainly based on the long polyol
 187 part (Figure 4a). TPUs are often semi-crystalline structures (Figure 4b). HS content and chemistry
 188 influenced the biodegradation susceptibility of a polycarbonate PU (Tang et al., 2001a, b).
 189 Interactions between enzymes and mobile SS are higher than with the HS. Consequently, the
 190 higher the HS content, the lower the biological degradation susceptibility.

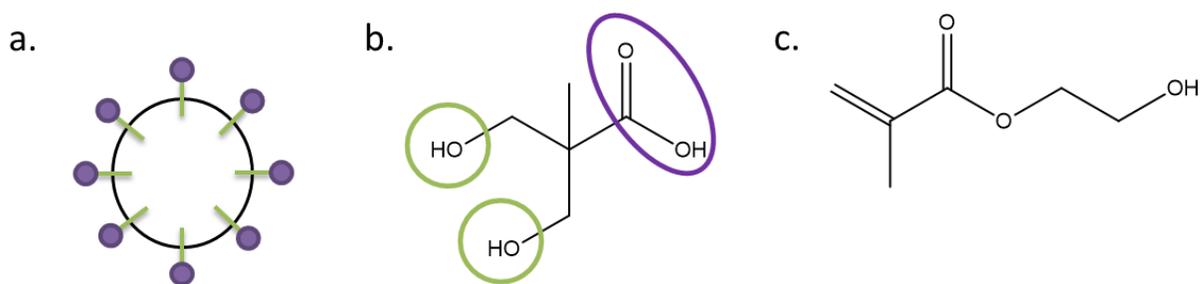


191
 192 *Figure 4 – (a) Schematic representation of a semi-crystalline polymer and (b) linear polyurethane structure*

193
 194 Thermosets are highly cross-linked polymers with 3D structures brought by molecules with a
 195 functionality higher than 2. Due to their architecture, thermosets are not soluble, do not melt and
 196 are denatured at high temperature, and thus cannot be reshaped by heating, unlike
 197 thermoplastics. PU foams are thermoset materials. The alveolar structures of these complex and
 198 multicomponent systems are obtained thanks to physical or chemical blowing with air, carbon
 199 dioxide, hydrocarbons such as isopentane or other gaseous substances. Foams are structured by

200 struts and walls defining cell cavities. Cells can be closed (closed-cell foams, mainly rigid e.g.
201 thermal insulation) or open (open-cell foams, mainly soft e.g. damping) (Gautam et al., 2007a).
202 Side reactions during foaming induce the formation of various reversible and irreversible bonds
203 such as urea or isocyanurate moieties (Furtwengler and Avérous, 2018). Isocyanurate linkage is
204 obtained by isocyanate trimerization. Polyisocyanurate foams are increasingly produced because
205 of their better properties such as fire resistance and thermal stability (Arbenz et al., 2016;
206 Furtwengler et al., 2018b). It is then important to consider the bond nature when dealing with
207 foam degradation. Because of the variety of bonds, it is difficult to anticipate and know if the
208 urethane bond was degraded. Also, commercial foams are supplemented with several additives
209 such as fire retardants, antioxidant, processing aids, and anti-microbial compounds (mainly
210 fungicides) that can prevent biodegradation.

211 Due to the nanometric size of the particles and their hydrophilicity, waterborne PU dispersions
212 (WPUDs) are particularly suitable for biological assays (Figure 5a). To provide hydrophilicity,
213 WPUDs are synthesized using an emulsifier, often the 2,2-dimethylol propionic acid. This
214 molecule contains two hydroxyl groups which react with the isocyanates to form urethane linkage
215 (Figure 5b). The hydroxyl group of the carboxylic acid does not react with isocyanate because of
216 steric hindrance and the lower reactivity (Coutinho et al., 2001). Hydrophilic carboxylic acid then
217 forms a stabilizing top-layer around the hydrophobic polymer (Zhang et al., 2011). Systems
218 containing acrylic polyols are readily dispersed in water (Ionescu, 2005). Acrylic polyols are thus
219 widely used in WPUD formulations. These polyols provide strength and resistance to coatings
220 (Akindoyo et al., 2016). Acrylic polyols are generally based on hydroxyethyl methacrylate or
221 hydroxyethyl acrylate (Figure 5c). The hydroxyl groups of the lateral chains then react with
222 isocyanate for urethane formation.



223
224 *Figure 5 – (a) Schematic structure of a polymer particle, (b) structure of the 2,2-dimethylol propionic acid and (c)*
225 *structure of the hydroxyethyl methacrylate*

226

227 3.2. PU waste disposal

228 The diversity of PUs macromolecular structures and chemistry is a clear obstacle for efficient
229 waste management. To appreciate the importance of PU biodegradation, it is necessary to
230 analyze the current PU waste disposal. PU waste is made of post-consumer products as well as
231 PU production waste, mostly from foam. Indeed, scrap from slabstock foam can reach up to 10%
232 of the production (Simon et al., 2018). In France, scrap PUs were estimated at 13 kTons in 2011
233 while end-of-life PU volume is about 198 kTons (Boujard et al., 2014). This source of PU waste

234 mainly arises from construction, furniture, bedding, automotive, shoes and home appliances.
235 Efficient collection and product dismantling are required to recycle these materials, thus limiting
236 their valorization.

237 PU wastes are mainly treated by three different methods: landfilling, incineration (which can also
238 be considered as quaternary recycling (Ignatyev et al., 2014)) and conventional recycling. Landfill
239 discharge is often the main option but is gradually decreasing, especially in Europe, since it
240 requires large land areas and no value is brought from the waste. European Union aims at
241 reducing municipal waste landfilling to a maximum of 10% by 2030 (Castillo-Gimenez et al., 2019;
242 Makarichi et al., 2018). Landfilling and the absence of waste management lead generally to
243 pollution (Jambeck et al., 2015).

244 Incineration presents the advantage of being a mature technique, practiced for several decades.
245 Energy is recovered by burning waste and can totally or partly offset the energy spent in the
246 heating process. Even if some CO₂ emission exists and may cause greenhouse effect and
247 contribute to global warming, new generations of plants are equipped with dry and wet air
248 pollution control system to make this process as clean as possible (Brunner and Rechberger, 2015;
249 Makarichi et al., 2018). Because of the low value recovered it is hardly considered as recycling. It
250 is a proper solution to reduce landfill volumes, yet incineration is not a satisfying strategy since
251 the richness from the chemical architecture is fully lost.

252 Depending on the nature (thermoplastics vs. thermosets), recycling processes differ. TPUs can be
253 heated and remolded therefore making the physical recycling process easier. However, it is
254 estimated that only 1% of PU are recycled thanks to physical methods (Behrendt and Naber,
255 2009). The recycling of PU foams is more challenging since foams cannot be remolded. The main
256 path for foam recycling is regrinding. In 2002, more than 380 kTons were used for carpet underlay
257 (Zia et al., 2007).

258 Chemical recycling can address both thermoplastic and thermoset architectures (Simón et al.,
259 2016; Wang et al., 2011). Glycolysis appears as the most promising technique (Simon et al., 2018).
260 Glycolysis is a transesterification reaction. The ester group of the urethane bond is interchanged
261 by the hydroxyl group of a diol (glycol) added in large excess (Simón et al., 2013). Simón et al.,
262 developed a glycolysis process allowing polyether polyol recovery from high resilience PU foams
263 (Simón et al., 2016). These polyols can then serve as building blocks for the synthesis of second-
264 generation polymers. The major limits of chemical recycling are the processing temperature that
265 leads to high energy consumption (in the example mentioned above, the glycolysis temperature
266 is 190°C) and the side chemical reactions occurring on the urethane bond during the chemical
267 reactions (Gadhve et al., 2019).

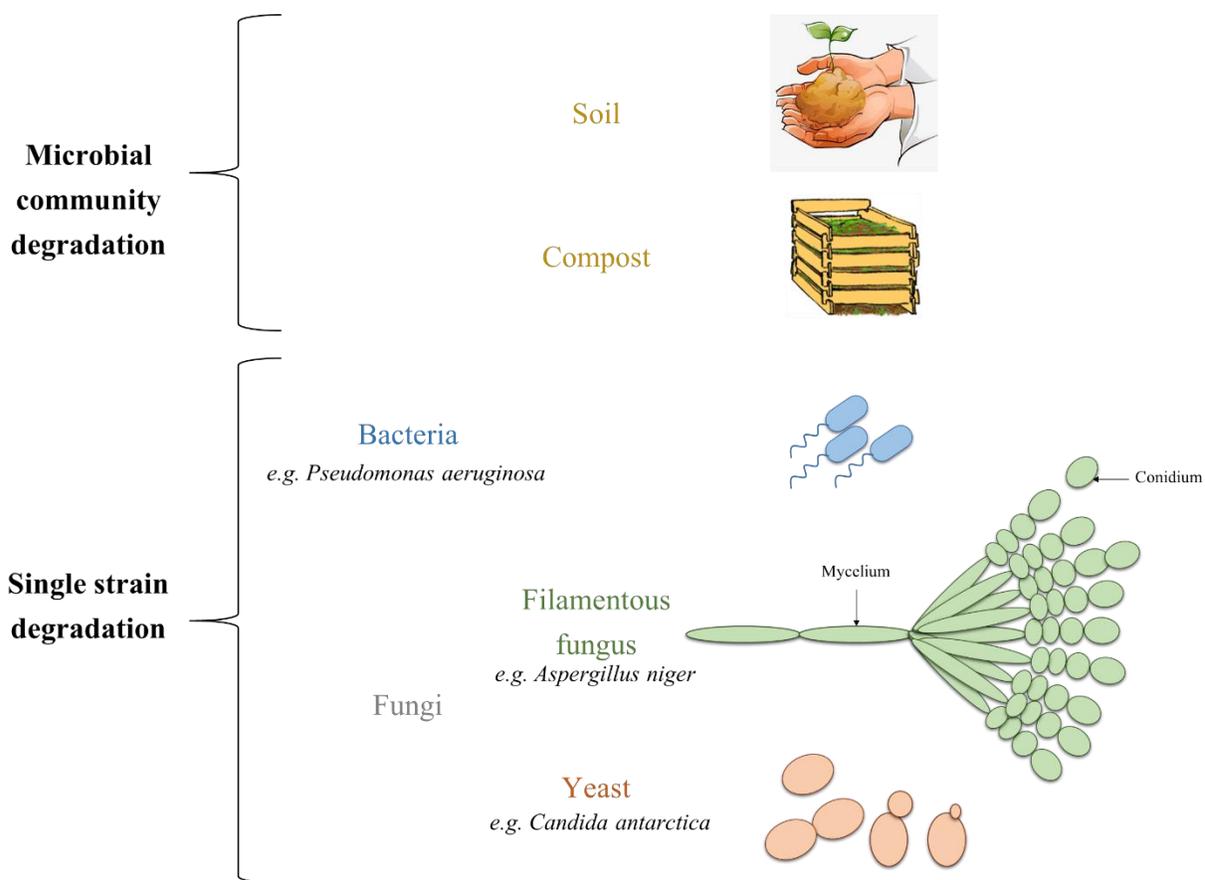
268 Biological recycling is a growing route with high potential that might answer the need for PU
269 recycling in the coming years. This is a soft process that can be implemented at low temperatures
270 (less than 70°C) (Mueller, 2006; Valerio, 2010). This process is catalyzed by biological entities,
271 namely enzymes. The resulting degradation products may then be valorized. Efficient enzymatic
272 depolymerization of poly(ethylene terephthalate) (PET) has been demonstrated at 60°C leading
273 to the release of valuable building blocks such as terephthalic acid and mono(2-hydroxyethyl)-
274 terephthalate (Gamerith et al., 2017). This result, based on PET, can be considered as the first

275 benchmark for PU. For instance, the company Carbios (France) is starting the biological recycling
276 of PET for the synthesis of second-generation polymers.

277
278 4. Actors of the PU biodegradation: Biological entities and associated
279 mechanisms

280 4.1. Microorganisms

281 Biodegradation involving microorganisms can be performed by a microbial community or a single
282 strain (Figure 6). Microorganisms can form biofilms on the polymer surface by adhesion (Sivan,
283 2011). Once colonized, the material constitutes a source of carbon and nitrogen thus promoting
284 microbial growth.



285
286 Figure 6 – Different paths for microbial PU degradation

287
288 4.1.1. PU degradation by microbial communities

289 A microbial community is a group of microorganisms sharing a common living place. These
290 microorganisms interact in different ways such as mutualism, predation or competition (Faust
291 and Raes, 2012). Mutualism, also called symbiosis, may occur during the microbial degradation of

292 xenobiotic, such as synthetic polymers, leading to an improved degradation (Tsoi et al., 2019;
 293 Vaclavkova et al., 2007). Comparison of degradation skills of a strain of fungi (*Aspergillus niger*)
 294 and a strain of bacteria (*Pseudomonas aeruginosa*) revealed a slightly higher TPU degradation
 295 with the bacteria but, above all, an impressive synergistic effect was observed when the polymer
 296 was incubated with both strains (Fernandes et al., 2016). Weight losses were approximately
 297 doubled for polyester TPU incubated with both microorganisms compared to single strain
 298 incubation. PU substrates degraded by communities range from simple structures such as
 299 thermoplastic polyester PU (Genovese et al., 2016) to recalcitrant material such as polyether PU
 300 foam (Ge et al., 2000). As PU are recalcitrant material, degradation experiments by microbial
 301 communities are often performed on a long time scale going from 28 days (Bentham et al., 1987)
 302 to two years of incubation (Seal and Pantke, 1988). The predominant systems for studying PU
 303 degradation have been composting and soil burial (Table 3 – PU degradation by microbial
 304 communities and associated PU substrates. These ecosystems are rich in degrading
 305 microorganisms. Other strategies to obtain communities that are more acclimated to PU are also
 306 developed. For instance, Cregut et al., selected microbial communities from the soil of a PU foam
 307 industrial plant (Cregut et al., 2014).

308

309

Table 3 – PU degradation by microbial communities and associated PU substrates

Microbial communities	PU substrates	Time of incubation	Reference
Composting	Polyester PU foam	50 days	(Gómez et al., 2014)
	Thermoplastic polyester PU	90 days	(Genovese et al., 2016; Kucharczyk et al., 2016)
		12 weeks	(Zafar et al., 2013)
		24 months	(Krasowska et al., 2012)
	Thermoset polyester PU	90 days	(Das et al., 2017)
Microbial communities from Garbage Landfill Leakage Water	Polyester PU foam	3 months	(Filip, 1978)
	Polyether PU foam	3 months	(Filip, 1978)
Microbial communities from sewage water of a latex rubber factory	Polyester PU foam	60 days	(Rattanapan et al., 2016)
Microbial communities from soil of a PU foam industrial plant	Polyether PU foam	28 days	(Cregut et al., 2014)
Mixed culture of <i>Aspergillus niger</i> and <i>Pseudomonas aeruginosa</i>	Thermoplastic polyester PU	30 days	(Fernandes et al., 2016)
Soil burial	Polyester PU	383 days	(Aranguren et al., 2012)
		24 months	(Seal and Pantke, 1988)
	Polyester PU foam	28 days	(Bentham et al., 1987)
		320 days	(Gómez et al., 2014)
	Polyether PU foam	6 months	(Ge et al., 2000)

	Polyether PU foam	12 months	(Zhang et al., 2013)
	PU coating	12 months	(Lu et al., 2016)
	Thermoplastic polyester PU	44 days	(Barratt et al., 2003)
		12 weeks	(Zafar et al., 2013)
		16 weeks	(Huang et al., 2016)
		140 days	(Umare and Chandure, 2008)
		5 months	(Cosgrove et al., 2007)
		6 months	(Fernandes et al., 2016)
	Thermoplastic polyester PU	12 months	(Oprea et al., 2016; Tajau et al., 2016)
	Thermoplastic polyether PU	12 months	(Oprea et al., 2016)
Soil microbial communities, bioaugmentation with PU-degrading fungi	Thermoplastic polyester PU	4 weeks	(Cosgrove et al., 2010)
Vermiculite inoculated with degrading microorganisms, notably <i>Pseudomonas aeruginosa</i> and <i>Achromobacter marplatensis</i>	Polyester PU	383 days	(Aranguren et al., 2012)

310

311

4.1.2. Single-strain degradation

312 Single species of bacteria and fungi can be isolated and identified using molecular tools. These
313 isolated strains are then used alone for PU degradation assays (Khan et al., 2017; Nair and Kumar,
314 2007). Another approach is to establish a collection of microorganisms and to screen it for the
315 identification of PU-degrading strain. For instance, Russell *et al.* collected endophytic fungi from
316 wood of the Ecuadorian Amazonian rainforest (Russell et al., 2011). Equatorial are hot and humid
317 environments, and above-all, well-known for the tremendous richness of their biodiversity. PU
318 debris were sampled in e.g., dump-site for isolation of already adapted microorganisms (Álvarez-
319 Barragán et al., 2016; Oceguera-Cervantes et al., 2007).

320

Bacteria

321 Bacteria are mainly studied for the degradation of TPU and coatings (Table 4). Only a few
322 publications described the bacterial degradation of polyester-based PU foams, notably by
323 *Pseudomonas aeruginosa* (Cooney, 1969; Gautam et al., 2007c; Hedrick and Crum, 1968; Kay et
324 al., 1991) or by a strain of *Corynebacterium* (Kay et al., 1991). *Pseudomonas* is the most studied
325 genus. The strain of *P. aeruginosa* ATCC 13388 is the only recommended strain by the ASTM for
326 testing material resistance to bacterial degradation (ASTM code G22-76: Standard Practice for
327 Determining Resistance of Plastics to Bacteria) (Gu and Gu, 2005; Kay et al., 1991). Other
328 *Pseudomonas* species such as *P. chlorographis* or *P. putida* were described as PU-degrading
329 entities (Gautam et al., 2007c; Peng et al., 2014). Furthermore, *Pseudomonas* strains are known
330 to be of high interest in white biotechnology (Wierckx et al., 2015).

331

Table 4 – PU-degrading bacteria and associated PU substrates

Species and/or strains	PU substrates	References
<i>Acinetobacter gerveri</i> P7	Polyester PU coating (Impranil)	(Howard et al., 2012)

<i>Alicyclophilus sp.</i> BQ1	Polyester PU coating	(Oceguera-Cervantes et al., 2007)
<i>Alicyclophilus sp.</i> BQ8	Polyester PU coating	(Oceguera-Cervantes et al., 2007)
	Polyester PU foam	(Pérez-Lara et al., 2016)
<i>Arthrobacter sp.</i>	Thermoplastic polyester PU	(Shah et al., 2008a)
<i>Arthrobacter calcoaceticus</i> ATCC 31012	Polyester PU coating	(El-Sayed et al., 1996)
<i>Arthrobacter calcoaceticus</i> NAV-2	Polyester PU coating	(El-Sayed et al., 1996)
<i>Arthrobacter globiformis</i>	Polyester PU coating	(El-Sayed et al., 1996)
<i>Bacillus sp.</i>	Polyester PU coating (Impranil)	(Ii et al., 1998)
	Thermoplastic polyester PU	(Shah et al., 2008a)
<i>Bacillus amyloliquefaciens</i>	Thermoplastic poly(ether urea) PU	(Rafiemanzelat et al., 2015)
<i>Bacillus pumilus</i> NMSN-1d	Polyester PU coating (Impranil)	(Nair and Kumar, 2007)
<i>Bacillus subtilis</i> MZA-75	Thermoplastic polyester PU	(Shah et al., 2016)
		(Shah et al., 2013b)
<i>Chryseobacterium meningosepticum</i>	Polyester PU foam	(Cangemi et al., 2008)
<i>Comamonas acidovorans</i> TB-35	Thermoplastic polyester PU	(Akutsu et al., 1998; Nakajima-Kambe et al., 1997; Nakajima-Kambe et al., 1995)
	Thermoplastic polyether PU	(Nakajima-Kambe et al., 1995)
<i>Corynebacterium sp.</i>	Thermoplastic polyester PU	(Kay et al., 1993)
		(Shah et al., 2008a)
	Polyester PU foam	(Kay et al., 1991)
<i>Escherichia coli</i>	Thermoplastic poly(ether urea) PU	(Rafiemanzelat et al., 2013)
<i>Micrococcus sp.</i>	Thermoplastic polyester PU	(Shah et al., 2008a)
	Thermoplastic poly(ether urea) PU	(Rafiemanzelat et al., 2013)
<i>Pseudomonas sp.</i>	Polyester PU foam	(Spontón et al., 2013)
	Thermoplastic polyester PU	(Shah et al., 2008a)
<i>Pseudomonas aeruginosa</i>	Thermoset poly(amido amine) PU	(Gogoi and Karak, 2015)
	Thermoset polyester PU	(Duarah et al., 2016)
	Polyester PU coating (Impranil)	(Mukherjee et al., 2011)
	Thermoset poly(ester amide) PU	(Gogoi and Karak, 2017)
	Polyester PU foam	(Cooney, 1969; Edmonds and Cooney, 1968)
<i>Pseudomonas aeruginosa</i> ATCC 13388	Polyester PU foam	(Kay et al., 1991)
<i>Pseudomonas aeruginosa</i> ATCC 9027	Thermoplastic polyester PU	(Fernandes et al., 2016)
<i>Pseudomonas aeruginosa</i> MTCC 7814	Thermoset polyester PU	(Bayan and Karak, 2017; Gogoi and Karak, 2014)
<i>Pseudomonas aeruginosa</i> MZA-85	Thermoplastic polyester PU	(Shah et al., 2016; Shah et al., 2013a)
<i>Pseudomonas aeruginosa</i> NAV-6	Polyester PU coating	(El-Sayed et al., 1996)
<i>Pseudomonas cepacia</i>	Polyester PU coating	(El-Sayed et al., 1996)
<i>Pseudomonas chlororaphis</i> ATCC 55729	Polyester PU foam	(Gautam et al., 2007c)
<i>Pseudomonas fluorescens</i>	Polyether PU coating	(Crookes-Goodson et al., 2013)
	Polyester PU coating (Impranil)	(Howard and Blake, 1998; Vega et al., 1999)
<i>Pseudomonas protegens</i> Pf-5	Polyester PU coating (Impranil)	(Biffinger et al., 2014)
	Thermoplastic polyether PU	(Barlow et al., 2016)
<i>Pseudomonas putida</i>	Polyester PU coating (Impranil)	(Peng et al., 2014)
<i>Pseudomonas putida</i> ATCC 17484	Polyester PU coating	(El-Sayed et al., 1996)

<i>Rhodococcus equi</i> strain TB-60	Model urethane molecule (toluene-2,4-dicarbamic acid dibutyl ester)	(Akutsu-Shigeno et al., 2006)
<i>Staphylococcus aureus</i>	Poly(ether urea) PU Undefined PU	(Rafiemanzelat et al., 2013) (Curia et al., 2014)
<i>Staphylococcus epidermidis</i> strain KH 11	Thermoplastic polyether PU	(Jansen et al., 1991)

332

333 *Fungi*

334 It is possible to appraise only the fungal activity of a consortium or isolate only fungal strain by
335 adding antibacterial molecules which prevent bacterial growth. For instance, 50 µg/mL of
336 chloramphenicol can be added to the medium to prevent bacterial growth and therefore perform
337 analysis only on fungal strains (Zafar et al., 2013). Microorganisms from the fungi kingdom
338 described as PU degrading entities are almost only filamentous fungi. Strains belonging to the
339 genus of *Alternaria* (Magnin et al., 2018; Matsumiya et al., 2010; Oprea et al., 2018), *Aspergillus*
340 (Khan et al., 2017; Magnin et al., 2018; Mathur and Prasad, 2012; Osman et al., 2018) and
341 *Cladosporium* (Álvarez-Barragán et al., 2016) are frequently isolated for PU degradation (Table 5).
342 Only one study on yeast was found, describing the growth of *Cryptococcus sp.* MTCC 5455 on fish
343 waste to produce a lipase with activity on PU (Thirunavukarasu et al., 2015). Five strains are
344 recommended by the American Society for Testing and Materials (ASTM) to evaluate the fungal
345 resistance of a material: *Aspergillus niger* ATCC 9642, *Aureobasidium pullulans* ATCC15233,
346 *Chaetomium globosum* ATCC6205, *Gliocladium virens* ATCC9645, *Penicillium pinophilum*
347 ATCC11797 (ASTM code: G21 - 90 Standard Practice for Determining Resistance of Synthetic
348 Polymetric Materials to Fungi) (Gu and Gu, 2005). Oprea et al. who developed new bio-based
349 materials and tested their fungal degradation susceptibility with the strain of *Chaetomium*
350 *globosum* (Oprea, 2010; Oprea and Doroftei, 2011; Oprea et al., 2016). Only a few studies are
351 based on this norm and mentioned these strains. However, using such reference fungal strains
352 would allow better reproducibility and comparison of results in-between studies.

353 A review on the biodegradation of fossil-based polymers interestingly shows the significant
354 importance of the abiotic effect of fungal biodegradation (Lucas et al., 2008). The formation of
355 filaments exerts physical pressure leading to polymer breaking. Filament apices penetrate in the
356 material increasing the size of pores and provoking cracks. For instance, rifts under the form of a
357 fungal filament network were observed by microscopy on the surface of a TPU incubated two
358 months with a strain of *Penicillium brasilianum* (Magnin et al., 2018). Fungal degradation has been
359 demonstrated on both polyester- and polyether-based PU, on TPU, foams and coatings. However,
360 mechanisms of degradation have not been fully elucidated. The importance of each biotic and
361 abiotic steps in fungal degradation still needs to be clarified.

362

Table 5 – PU-degrading fungi and associated PU substrates

Species and/or strain	PU substrates	Reference
<i>Alternaria sp.</i>	Thermoplastic polyester PU	(Magnin et al., 2018)
<i>Alternaria Solani</i> Number Ss.1-3	Thermoplastic polyester PU	(Ibrahim N. Ibrahim, 2009)
<i>Alternaria sp.</i> strain PURDK2	Polyether PU foam	(Matsumiya et al., 2010)
<i>Alternaria tenuissima</i>	Thermoplastic polyether PU	(Oprea et al., 2018)
<i>Aspergillus sp.</i>	Polyester PU foam	(Cangemi et al., 2006; Cangemi et al., 2008)
<i>Aspergillus flavus</i>	Thermoplastic polyester PU	(Mathur and Prasad, 2012)
<i>Aspergillus fumigatus</i>	Polyester PU coating (Impranil)	(Álvarez-Barragán et al., 2016)
	Polyether PU foam	(Álvarez-Barragán et al., 2016)
	Thermoplastic polyester PU	(Osman et al., 2018)
<i>Aspergillus niger</i>	Polyether PU foam	(Filip, 1979)
<i>Aspergillus niger</i> ATCC 9642	Thermoplastic polyester PU	(Kanel et al., 1966)
<i>Aspergillus section flavi</i>	Thermoplastic polyester PU	(Magnin et al., 2018)
<i>Aspergillus tubingensis</i>	Thermoplastic polyester PU	(Khan et al., 2017)
<i>Aureobasidium pullulans</i>	Polyester PU coating (Impranil)	(Crabbe et al., 1994)
<i>Chaetomium globosum</i>	Thermoset polyester PU	(Oprea and Doroftei, 2011)
	Thermoplastic polyester PU	(Oprea et al., 2016)
<i>Cladosporium sp.</i>	Polyester PU coating (Impranil)	(Crabbe et al., 1994)
	Polyether PU foam	(Cooney, 1969; Edmonds and Cooney, 1968)
<i>Cladosporium tenuissimum</i>	Polyester PU coating (Impranil)	(Álvarez-Barragán et al., 2016)
	Polyether PU foam	(Álvarez-Barragán et al., 2016)
<i>Cladosporium asperulatum</i>	Polyester PU coating (Impranil)	(Álvarez-Barragán et al., 2016)
	Polyether PU foam	(Álvarez-Barragán et al., 2016)
<i>Cladosporium herbarum</i>	Polyether PU foam	(Filip, 1979)
<i>Cladosporium montecillanum</i>	Polyester PU coating (Impranil)	(Álvarez-Barragán et al., 2016)
	Polyether PU foam	(Álvarez-Barragán et al., 2016)
<i>Cladosporium pseudocladosporioides</i>	Polyester PU coating (Impranil)	(Álvarez-Barragán et al., 2016)
	Polyether PU foam	(Álvarez-Barragán et al., 2016)
<i>Cryptococcus laurentii</i>	Polyester PU coating (Impranil)	(Zicht, 2017)
<i>Curvularia senegalensis</i>	Polyester PU coating (Impranil)	(Crabbe et al., 1994)
<i>Exophiala jeanselmei</i>	Model urethane molecule (N-tolylcarbamate)	(Owen et al., 1996)
<i>Fusarium solani</i>	Polyester PU coating (Impranil)	(Crabbe et al., 1994)
<i>Gliocladium roseum</i>	Thermoplastic polyester PU	(Shuttleworth and Seal, 1986)
<i>Penicillium chrysogenum</i>	Polyester PU coating (Impranil)	(Álvarez-Barragán et al., 2016)
	Polyether PU foam	(Álvarez-Barragán et al., 2016)
<i>Penicillium section lanata-divaricata</i>	Thermoplastic polyester PU	(Magnin et al., 2018)
<i>Pestalotiopsis microspora</i>	Polyester PU coating (Impranil)	(Russell et al., 2011)

365

4.2. Enzymes

366 Enzymes are biological catalysts (biocatalysts). Enzymes identified as PU degrading entities
 367 originate from microorganisms but also mammalian cells such as lipase porcine pancreas (Ng et
 368 al., 2017) or from plants such as papain from *Carica papaya* (Ferris et al., 2010). Enzymes used for
 369 PU degradation assays are either commercial enzymes or enzymes over-expressed in
 370 heterologous microorganisms. In this latter case, they correspond to enzymes identified in PU-
 371 degrading microorganisms for which the encoding genes were cloned into model organism.
 372 Enzymes are then over-expressed and even purified in some cases. Enzymes are mainly described
 373 for the depolymerization of TPU or coatings (Table 6 – PU-degrading enzymes and associated PU
 374 substrates). As far as we know, only one publication has addressed the enzymatic degradation of
 375 foams (Ng et al., 2017) all other studies involved degradation by microorganisms. A set of
 376 different poly(ester ether) PU foam containing PCL, PEG and polyester from palm oil was studied
 377 by enzymatic degradation. A maximal weight loss of 70% was measured after 28 days of
 378 incubation with lipase from porcine pancreas for a polymer with 7.7% of PCL, 34.8% of PEG and
 379 7.5% of polyester from palm oil.

380

Table 6 – PU-degrading enzymes and associated PU substrates

EC number	Enzyme name in the publication	Type of PU	Reference
EC 1.10.3.2	Laccase	Thermoplastic polyester PU	(Ignat et al., 2011)
EC 1.11.1.7	Fungal peroxidase	Thermoplastic polyester PU	(Ignat et al., 2011)
EC 3	Tcur0390 (<i>Thermomonospora curvata</i> DSM43183 hydrolase)	Thermoplastic polyester PU	(Schmidt, J. et al., 2017)
	Tcur1278 (<i>Thermomonospora curvata</i> DSM43183 hydrolase)	Thermoplastic polyester PU	(Schmidt, J. et al., 2017)
EC 3.1	<i>Bacillus subtilis</i> esterase	Polyester PU (Impranil)	(Rowe and Howard, 2002)
	<i>Comamonas acidovorans</i> TB-35 esterase	Polyester PU (Impranil)	(Allen et al., 1999)
	<i>Curvularia senegalensis</i> esterase	Polyester PU (Impranil)	(Crabbe et al., 1994)
	E3576 (esterase)	Polyester PU Impranil	(Magnin et al., 2019)
		Thermoplastic polyester PU	(Magnin et al., 2019)
	<i>Pseudomonas fluorescens</i> esterase	Polyester PU (Impranil)	(Biffinger et al., 2015)
	PudA (<i>Comamonas acidovorans</i> TB-35 esterase)	Thermoplastic polyester PU	(Akutsu et al., 1998; Nomura et al., 1998)
	PulA (<i>Pseudomonas fluorescens</i> esterase)	Polyester PU (Impranil)	(Ruiz and Howard, 1999)
EC 3.1.1	<i>Cryptococcus</i> sp. MTCC 5455 lipase	Thermoplastic polyester PU	(Thirunavukarasu et al., 2015)
	Lipase	Thermoplastic poly(ester ether) PU	(Feng et al., 2017)
	Lipase AK	Thermoplastic polyester PU	(Zhou et al., 2012)
		Thermoset poly(ester ether) PU	(Jiang et al., 2007)
	Lipase PS	Thermoplastic polycarbonate PU	(Chen et al., 2016)
		Thermoplastic polyester PU	(Xu et al., 2014)
	Lipolase 100L	Polyester PU coating	(Pilch-Pitera, 2012)
	Novozym 51,032	Polyester PU coating	(Pilch-Pitera, 2012)

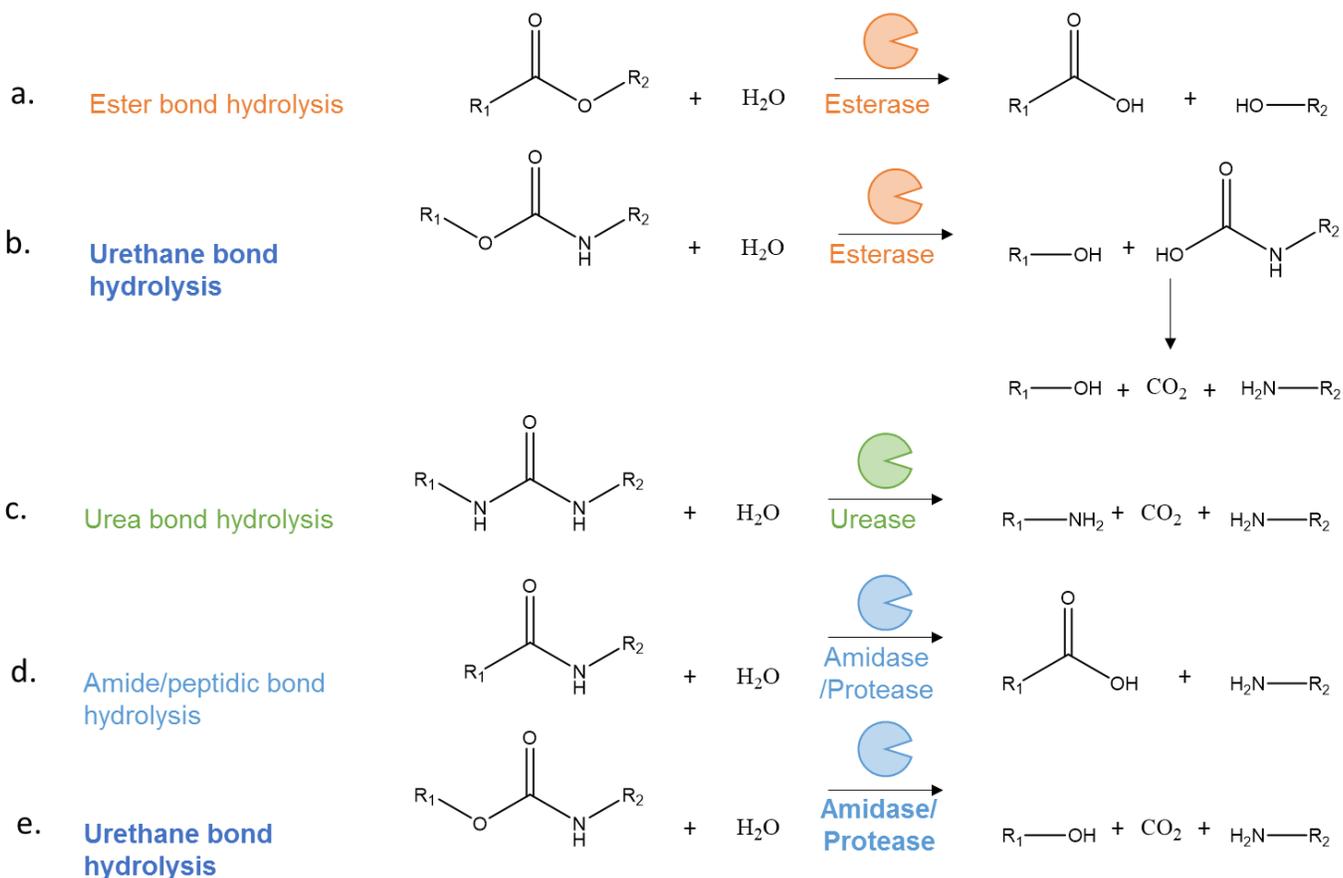
	Novozym 735	Polyester PU coating	(Pilch-Pitera, 2012)
	Palatase 20,000	Polyester PU coating	(Pilch-Pitera, 2012)
	<i>Pseudomonas cepacia</i> lipase	Thermoplastic poly(ester ether) PU	(Zhou and Xie, 2017)
		Thermoset polyester PU	(Schöne et al., 2016)
	<i>Pseudomonas sp.</i> lipase	Polyester PU (Impranil)	(Biffinger et al., 2015)
		Thermoplastic polyester PU	(Daemi et al., 2016)
	PueA (<i>Ps chlororaphis</i> lipase)	Polyester PU (Impranil)	(Howard et al., 2007; Hung et al., 2016; Langlois and Howard, 2002; Ruiz et al., 1999; Stern and Howard, 2000)
	PueB (<i>Pseudomonas chlororaphis</i> lipase)	Polyester PU (Impranil)	(Howard et al., 2007; Howard et al., 2001; Hung et al., 2016; Ruiz et al., 1999)
	<i>Rhizopus arrhizus</i> lipase	Thermoplastic polyester PU	Tokiwa (Tokiwa et al., 1988)
	<i>Rhizopus delemar</i> lipase	Thermoplastic polyester PU	Tokiwa (Tokiwa et al., 1988)
	<i>Thermomyces lanuginosus</i> lipase	Thermoplastic poly(ester urea) PU	(Fang et al., 2014)
		Thermoset polyester PU	(Wu et al., 2016)
EC 3.1.1.1	Esterase	Thermoplastic polyether PU	(Smith et al., 1987)
	Porcine liver esterase	Thermoplastic polyester PU	(Li and Yang, 2006)
EC 3.1.1.3	<i>Candida antarctica</i> lipase	Thermoplastic polyester PU	(Takamoto et al., 2001)
		Polyester PU coating	(Liu et al., 2016)
	<i>Candida cylindracea</i> lipase	Thermoplastic polyester PU	(Kim and Kim, 1998)
	<i>Candida rugosa</i> lipase	Thermoplastic polyester PU	(Li et al., 2015)
		Polyester PU (Impranil)	(Gautam et al., 2007b)
	Porcine pancreas lipase	Polyester PU foam	(Ng et al., 2017)
		Thermoplastic poly(ester ether) PU	(Brzeska et al., 2015)
		Thermoplastic polyester PU	(Brzeska et al., 2015)
		Thermoplastic polyether PU	(Ferris et al., 2010)
EC 3.1.1.13	Cholesterol esterase	Thermoplastic poly(ester urea) PU	(Santerre et al., 1993; Santerre et al., 1994; Wang et al., 1997a)
		Thermoplastic poly(ether urea) PU	(Santerre et al., 1994)
		Thermoplastic polycarbonate PU	(Christenson et al., 2006; Tang et al., 2001a, b; Tang et al., 2003)
		Thermoplastic polyester PU	(Woo et al., 2000)
		Thermoplastic polyether PU	(Christenson et al., 2006)
EC 3.1.1.74	LC cutinase (LCC)	Thermoplastic polyester PU	(Schmidt, J. et al., 2017)
	TfCut 2 (<i>Thermobifida fusca</i> KW3 cutinase)	Thermoplastic polyester PU	(Schmidt, J. et al., 2017)
		Thermoset polyester PU	(Wu et al., 2016)
EC 3.4	<i>Bacillus sp.</i> Protease	Polyester PU (Impranil)	(Biffinger et al., 2015)
EC 3.4.11.1	Leucine aminopeptidase	Thermoplastic polyester PU	(Ratner et al., 1988)
		Thermoplastic polyether PU	(Ratner et al., 1988)
EC 3.4.14.1	Cathepsin C	Thermoplastic polyether PU	(Smith et al., 1987)
EC 3.4.21.1	Chymotrypsin	Thermoplastic poly(ester ether) PU	(Ciardelli et al., 2004)
		Thermoplastic poly(ester urea) PU	(Elliott et al., 2002)

		Thermoplastic polyester PU	(Ratner et al., 1988; Yamamoto et al., 2007)
		Thermoplastic polyether PU	(Campinez et al., 2013; Ferris et al., 2010; Ratner et al., 1988; Smith et al., 1987)
EC 3.4.21.36	Porcine pancreatic elastase	Thermoplastic poly(ester urea) PU	(Guan et al., 2008; Labow et al., 1996)
		Thermoplastic poly(ether urea) PU	(Labow et al., 1996)
EC 3.4.21.62	Subtilisin	Thermoplastic polyamide PU	(Huang et al., 2016)
EC 3.4.21.64	Protease K	Thermoplastic polyester PU	(Dogan et al., 2017; Yamamoto et al., 2007)
EC 3.4.22.2	Papain	Thermoplastic poly(ether urea) PU	(Zhao et al., 1987)
		Thermoplastic polyester PU	(Ratner et al., 1988; Yamamoto et al., 2007)
		Thermoplastic polyether PU	(Campinez et al., 2013; Ferris et al., 2010; Phua et al., 1987; Ratner et al., 1988; Smith et al., 1987)
		poly(ether urethane urea) elastomer	(Marchant et al., 1987)
EC 3.4.22.3	Ficin	Thermoplastic polyester PU	(Yamamoto et al., 2007)
		Thermoplastic polyether PU	(Smith et al., 1987)
EC 3.4.4.24	Bromelain	Thermoplastic polyester PU	(Smith et al., 1987; Yamamoto et al., 2007)
EC 3.4.21.37	Human neutrophil elastase	Thermoplastic poly(ester urea) PU	(Labow et al., 1996)
		Thermoplastic poly(ether urea) PU	(Labow et al., 1996)
EC 3.4.24	Collagenase	Thermoplastic polyester PU	(Zhang et al., 1994)
		Thermoplastic polyether PU	(Mendoza-Novelo et al., 2013)
EC 3.5.1.4	E4143 (amidase)	Thermoplastic polyester PU	(Magnin et al., 2019)
	<i>Nocardia farcinica</i> polyamidase	Thermoplastic polyester PU	(Gamerith et al., 2016)
EC 3.5.1.5	Urease	Thermoplastic polyether PU	(Phua et al., 1987)
No EC number	Pancreatine (enzyme mixture)	Thermoplastic polyester PU	(Zhang et al., 1994)

381

382 Enzymatic degradation assays on PU using oxidase enzymes (Enzyme Classification 1, EC 1) e.g.,
383 fungal peroxidase (EC 1.11.1.7) and laccase (E.C. 1.10.3.2) have been performed with success on
384 a polyester PU (Ignat et al., 2011). All the other assays dealing with enzymatic degradation of PU
385 relies on hydrolytic enzymes (EC 3). Several hydrolytic mechanisms have indeed been highlighted
386 for PU degradation. The most common is the hydrolysis of the polyester moieties of polyester-
387 based PU by esterases (EC 3.1). Ester hydrolysis leads to the release of a carboxylic acid and an
388 alcohol (Figure 7a). Other esterases such as lipases (EC 3.1.1) (Fang et al., 2014; Schöne et al.,
389 2016), cutinases (EC 3.1.1.74) (Schmidt, J. et al., 2017; Yang et al., 2013) or unspecific esterases
390 (EC 3.1) (Kang et al., 2011) have been described as polyester PU degrading enzymes. Esterase has
391 also been described as hydrolyzing the urethane linkage. Some of these studies refer to a
392 mechanism resulting in carbamic acid and alcohol chain-ends after hydrolysis (Mahajan and
393 Gupta, 2015; Wei and Zimmermann, 2017). However, this mechanism does not seem conceivable
394 because of the instability of the carbamic acid which immediately breaks down into an amine with
395 the release of a molecule of carbon dioxide (Ionescu, 2005; Ozaki, 1972) (Figure 7b). Moreover,
396 most of the assays involving esterases concern polyester-based PU and do not allow

397 differentiation between ester and urethane bond hydrolysis. To evaluate the urethane bond
 398 hydrolysis by an esterase, assays must be performed on substrates that do not contain ester
 399 bonds. Publications describing slight esterase activity on polyether PU and showing the potential
 400 ability of esterase to hydrolyze the urethane bond are scarce (Santerre et al., 1994; Smith et al.,
 401 1987). A cholesterol lipase was reported to display activity on a PU based on triethylene glycol
 402 and 1,4-di-S-benzyl-D,L-dithiothreitol (Ferris et al., 2010). Urease (EC 3.5.1.5) also showed activity
 403 on poly(ether urea) PU (Phua et al., 1987) but the degradation is mainly attributed to the urea
 404 bond hydrolysis (Figure 7c). Amidases (EC 3.5.1.4) and proteases hydrolyze amide or peptidic
 405 bonds leading to the release of a carboxylic acid and an amine (Figure 7d). These enzymes
 406 appeared to be also efficient for the hydrolysis of the urethane bond leading to the release of an
 407 amine, an alcohol and a carbon dioxide molecule (Figure 7e). Proteases such as papain (EC
 408 3.4.22.2) (Campinez et al., 2013; Ferris et al., 2010; Marchant et al., 1987; Yamamoto et al., 2007),
 409 bromelain (EC 3.4.22.32/33) (Yamamoto et al., 2007), ficain (EC 3.4.22.3) (Yamamoto et al., 2007)
 410 and chymotrypsin EC 3.4.21.1 (Ciardelli et al., 2004; Elliott et al., 2002; Ferris et al., 2010) are also
 411 described for the degradation of PU. Recently, an amidase drew attention. This enzyme was
 412 isolated from *Nocardia farcinica* with the specificity of being able to hydrolyze polyamides (Guo
 413 et al., 2014) but also both the ester and amide bonds of non-water soluble model substrates
 414 (Heumann et al., 2009) and polyester-based PUs (Gamerith et al., 2016). Even if esterase is the
 415 main class of enzyme describes for PU degradation, others such as amidase or oxidase are rising
 416 interest for the full degradation of PU material.



417

418

Figure 7 – Main mechanisms of enzymatic degradation of PU

419

420 An alternative strategy aiming at discovering efficient PU-degrading enzymes was recently
421 published. Metagenomics tools allowing to screen bovine rumen microbiota were developed to
422 select enzymes with activity towards carbamate insecticides and PUs (Ufarte et al., 2017). The
423 main advantage of this strategy is the possibility of studying enzymes from uncultivable
424 microorganisms that are predominant in microbial communities, and thus having potential access
425 to new degrading enzymes.

426 To date, PU depolymerization using enzymes is not efficient enough for the development of
427 recycling processes. Recent publications on this topic aimed to improve the efficiency of the
428 depolymerization reactions. Complementary activities of enzymes presented above justify testing
429 cocktail of enzymes with different activities. A mixture of an esterase and an amidase revealed a
430 synergistic effect between these two enzymes for the degradation of a polyester PU.
431 Corresponding product analysis showed that hydrolysis of ester moieties from the SS by the
432 esterase released low molar mass molecules containing urethane bonds that are then hydrolyzed
433 by the amidase (Magnin et al., 2019). Another strategy for enzymatic depolymerization
434 improvement is to use protein engineering. This approach already proved itself on PET-degrading
435 enzymes. As an example, decreased inhibitor sensitivity of a PET-degrading cutinase has been
436 successfully achieved by amino-acid modification into the catalytic site (Wei et al., 2016).
437 Thermostability is also a key parameter for polymer degradation (Kawai et al., 2014). Ribitsch et
438 al., fused a PET-degrading enzyme (a cutinase from *Thermomyces cellulosylitica*) with a binding
439 module of a PHA depolymerase from *Alcaligenes faecalis* (Ribitsch et al., 2013) to improve
440 enzyme/polymer interactions. This binding domain was recently added to the amidase from
441 *Nocardia farcinica* to improve the degradation of polyester PU pellets (Gamerith et al., 2016).
442 These improvements will help reaching efficient depolymerization processes for PU biorecycling.

443 5. Analytical solutions for the measurement of polyurethane 444 biodegradation

445 The great diversity of PU structures and biological entities gives rise to numerous analytical
446 approaches to evaluate PU biological degradation. The methods have evolved lately with
447 powerful tools for a better understanding of the mechanisms of PU biodegradation. This chapter
448 offers an overview of the different techniques. The implementation of tools using urethane-based
449 model molecules is first addressed. Then, the degradation of more complex substrates, such as
450 TPU and PU foams, is presented.

451 5.1. Biological degradation of urethane-based model molecules

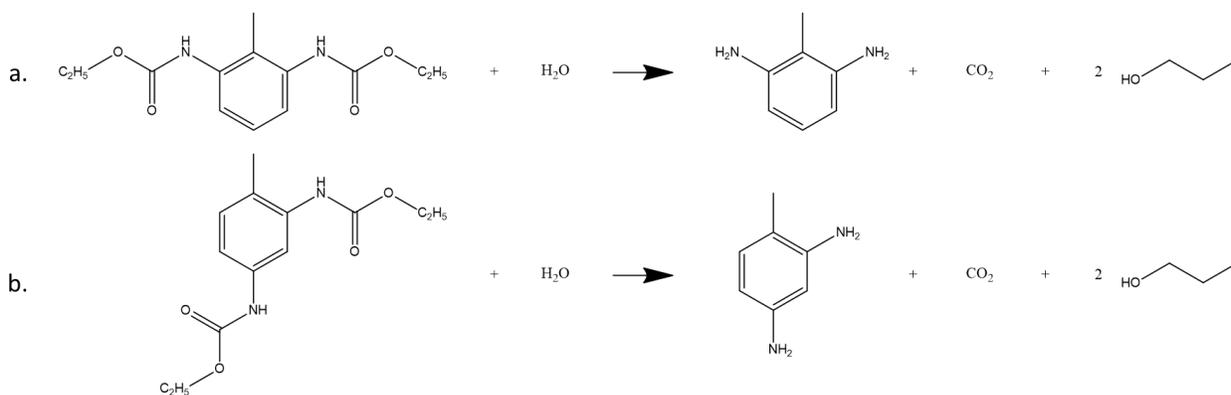
452 For the development of bioremediation or biological recycling processes, degradation of model
453 molecules is generally the first step as it allows easy identification of efficient degrading entities.
454 Urethane-based model molecules are readily hydrolysable and generally propose a simplified
455 hydrolysis detection. These model molecules can be either low molar mass molecules or
456 hydrophilic PU dispersion.

457

5.1.1. Analysis of low molar mass molecules

458 To cover the degradation of different PU structures, tracking the urethane bond hydrolysis
459 appears as a relevant solution. Low molar mass molecules containing a single urethane linkage
460 can be designed for this purpose. Urethane-based molecules are not soluble in water. Pre-dilution
461 in organic solvents such as ethanol (Akutsu-Shigeno et al., 2006) or DMSO (Gamerith et al., 2016)
462 is thus required.

463 Low molar mass N-tolylcarbamate molecules correspond to toluene with urethane linkage on one
464 or two carbons of the aromatic ring bound to ethanol moieties (Owen et al., 1996). In Owen et
465 al., aromatic amines resulting from the N-tolylcarbamate hydrolysis were extracted in chloroform
466 and quantified by Gas Chromatography coupled with Mass Spectrometry (GC/MS). This assay
467 revealed that the degrading activity of the *Exophiala jeanselmei* strain REN-11A depends on the
468 position of the urethane(s) around the aromatic ring. Toluene-2,4- and -2,6-dicarbamic acid
469 diethyl ester were the most readily biodegradable molecules (Figure 8).



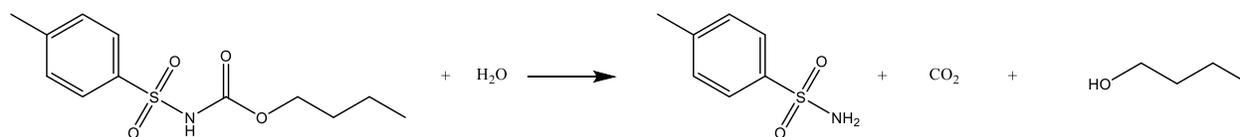
470

471 *Figure 8 – Hydrolysis of (a) the toluene-2,6-dicarbamic acid diethyl ester into 2,6-toluene diamine and propanol and*
472 *(b) the toluene-2,4-dicarbamic acid diethyl ester into 2,4-toluene diamine and propanol*

473

474 Akutsu-Shigeno et al., also described a set of molecules bearing a single urethane bond formed
475 by reacting a di-isocyanate (2,4-TDI, 4,4'-MDI and HDI) with butanol (Akutsu-Shigeno et al., 2006).
476 These compounds were degraded by both *Rhodococcus equi* strain TB-60 and a purified urethane-
477 degrading enzyme secreted by this bacterium. Degradation products were extracted with ethyl
478 acetate and analyzed by GC/MS except for the HDI-based model molecules which degradation
479 products were extracted with toluene under alkaline conditions. Because of the difficulties to
480 detect aliphatic amines in GC/MS, amines coming from HDI-based molecules hydrolysis were
481 derivatized using heptafluorobutyric acid anhydride following the method of Skarping et al.
482 (Skarping et al., 1988).

483 Coupling a 96-wells microplate assay with HPLC analysis was proposed for the development of a
484 medium-throughput screening (Magnin et al., 2019). A model molecule based on p-
485 Toluenesulfonyl isocyanate was synthesized for this microplate assay (Figure 9). Both the
486 substrate and the degradation products were analyzed by HPLC. Finally, 55 enzymes were
487 screened resulting in the identification of two amidases able to hydrolyze the urethane bond.



489 *Figure 9 – Hydrolysis of p-Toluenesulfonyl isocyanate model substrate (Magnin et al., 2019)*

490

491 To avoid the setup of complex analytical procedures such as GC/MS, Gamerith et al., proposed

492 the synthesis of a model molecule based on 4-nitrophenol (Gamerith et al., 2016). Molecules

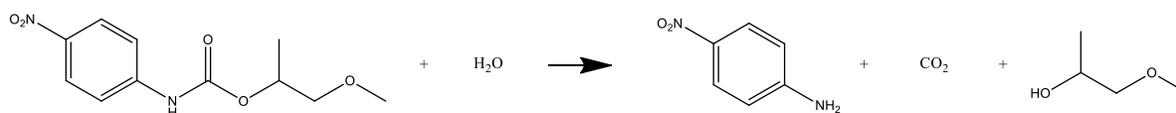
493 based on this aromatic compound are well known as model substrates for enzymes such as

494 esterase (4-nitrophenyl acetate) or amidase (4-nitroacetanilide). 1-methoxypropan-2-yl (4-

495 nitrophenyl) carbamate was synthesized (Figure 10) and the subsequent hydrolysis leads to the

496 release of 4-nitroaniline that can be tracked and quantified by UV-vis absorbance measurements

497 at 405 nm.



499 *Figure 10 - Hydrolysis of the 1-methoxypropan-2-yl (4-nitrophenyl) carbamate leading to 4-nitroaniline and 1-*

500 *methoxy-2-propanol (Gamerith et al., 2016)*

501

502 The use of low molar mass urethane substrates is a good way to identify efficient degrading

503 entities. However, the low steric hindrance of these molecules makes them far from being

504 representative of actual PU materials which often present organized and crystalline structures

505 and are much more hydrophobic. Activity assays on real and complex polymers must then be

506 performed with the identified degrading entities.

507 *5.1.2. Waterborne polyurethane dispersions as a PU-based model*

508 WPUD are particularly suitable to assess the degradation ability of enzymes and microorganisms

509 thanks to the polymer particles nanometric size and homogeneity in water. Indeed, their specific

510 surface is higher than for previously described polymers, thus maximizing interactions between

511 the degrading entity and the polymer. Moreover, all biodegradation reactions occur in aqueous

512 media. The most famous WPUD is the Impranil-DLN[®], commercialized by Covestro (Germany) for

513 textile coating applications. Particle size is estimated to range between 0.1 to 0.2 μm (Biffinger et

514 al., 2015). This dispersion appears as a white, milky suspension containing 40% of polymer. The

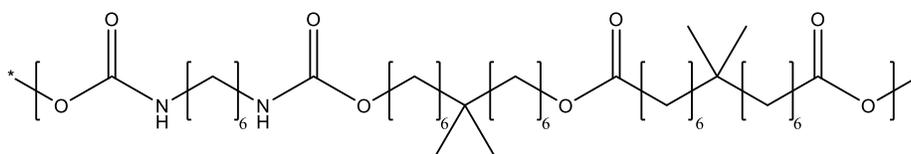
515 exact composition and structure of Impranil-DLN[®] are not precisely known. A tentative structure

516 has been proposed by Biffinger et al. based on polyhexane/neopentyl adipate polyester and HDI

517 (Figure 11). Diethylene glycol is also a component of Impranil-DLN[®] (Gautam et al., 2007b).

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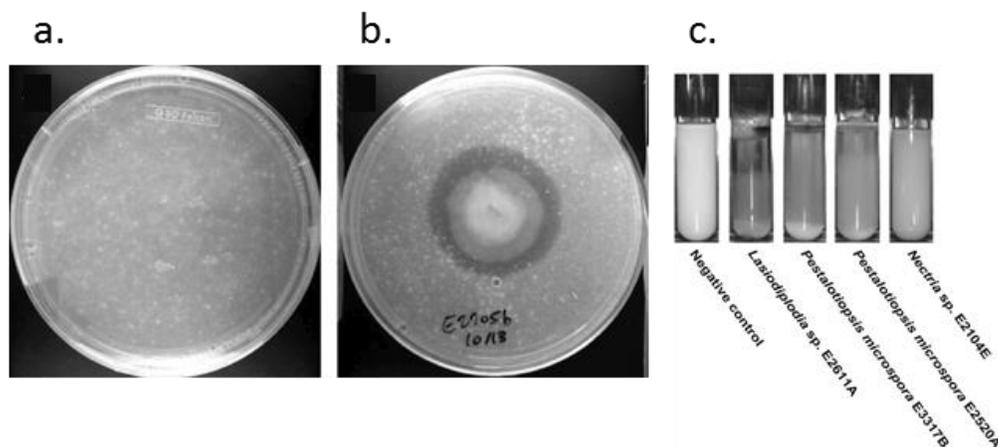
Figure 11 – Impranil-DLN® tentative structure adapted from Biffinger et al., 2015 (Biffinger et al., 2015)

521

522

Impranil-DLN® has been widely studied as a PU-based model since it presents the particularity to become translucent when hydrolyzed as a result of water-soluble molecules being released in the medium. Impranil-DLN® was used as “polyurethane biodegradation benchmark” for the first time in 1994 (Crabbe et al., 1994). Screening of soil fungi was performed on a plate where Impranil-DLN® was mixed with an agar medium. Fungi were allowed to grow on it and a transparent halo appeared after a few days when the microorganism was producing degrading enzymes. This agar plate technique has subsequently been used intensively (Howard et al., 2001; Peng et al., 2014; Rowe and Howard, 2002; Vega et al., 1999) (Figure 12a & b). Impranil-DLN® is also suitable for assays in liquid media for both microorganisms (Álvarez-Barragán et al., 2016; Russell et al., 2011) (Figure 12c) and enzymes (Gautam et al., 2007b; Schmidt, J. et al., 2017).

531



532

533

Figure 12 – Hydrolysis of Impranil-DLN®. Degradation assay in agar plate (a) negative control and (b) Pleosporales sp. strain E2705B after two weeks of incubation. (c) Assay in liquid media. Adapted from Russel et al. (Russell et al., 2011)

535

536

537

As the exact polymer structure is unknown, it is difficult to appraise the mechanism of degradation. Biffinger et al., used NMR and FT-IR to offer quantitative analysis of alcohol and carboxylic acid release after the ester bond cleavage (Biffinger et al., 2015). They also highlighted that polymer aggregation can occur when incubated with enzymes without any measurable degradation. Observation of the polymer is thus not sufficient to conclude on enzymatic degradation activity. Ufarté et al., proposed to use Matrix Assisted Laser Desorption Ionisation - Time of Flight/Mass Spectrometry (MALDI-TOF/MS) to identify degradation by-products. Three peaks at m/z 682, 683 and 782 were specific of the bacterial degradation of Impranil-DLN®. All

544

545 seemed to correspond to the formula $C_{36}H_{68}O_8N_2$, ionized with either Na^+ or I^- . The peak at m/z
546 683 could correspond to the isotope form of the molecule at m/z 682. However, no specific
547 chemical structure has been suggested. Alvarez-Barragan et al., presented the analysis of
548 degradation by-products by GC/MS (Álvarez-Barragán et al., 2016). Almost none of the products
549 identified correspond to the Impranil putative structure shown above (Figure 11) suggesting that
550 the structure is much more complex than expected. The only corresponding molecule is HDI.
551 However, HDI cannot be a degradation product as isocyanates are not stable in water. A possible
552 explanation would be the detection of either hexane diamine (HDA) or HDI-derivatives.

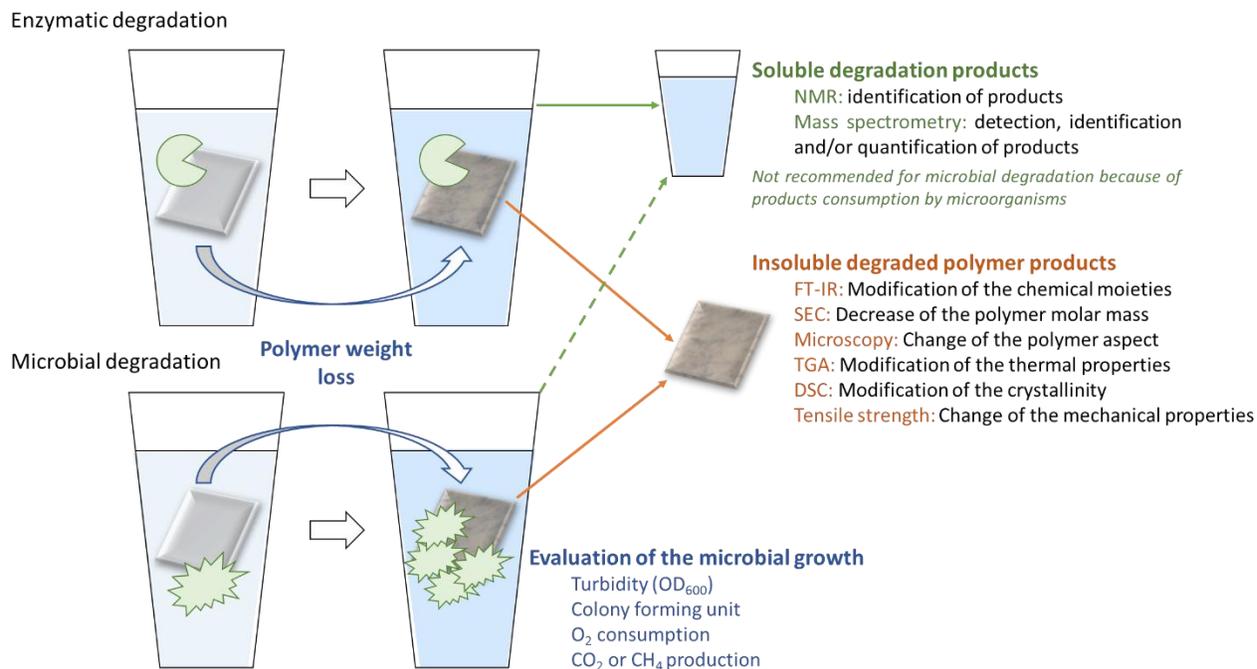
553 Despite some important limitations, Impranil-DLN[®] is thus a good model substrate to identify
554 degrading entities that have a great chance to be efficient afterwards on polyester PUs. It provides
555 a first approach that must then be confirmed. Hung et al. (2016) did not only present Impranil-
556 DLN[®] as a model but also as a common coating which integrity can be affected by microbial
557 degradation (Hung et al., 2016). Esterase and lipase are mainly involved in the enzymatic
558 degradation of Impranil-DLN[®]. However, up to now, there is no clear evidence that these
559 enzymes, improperly called “Polyurethanase” (Ruiz et al., 1999; Stern and Howard, 2000),
560 effectively hydrolyze the urethane bond.

561 Few other WPUDs have been tested for degradation activities. Bayhydrol 110 (Covestro,
562 Germany) is a polyester PU dispersion presenting the same clarification properties as Impranil-
563 DLN[®]: translucent halo appeared when incubated with a strain of *Pseudomonas chlorographis* on
564 an agar plate containing this WPUD (Howard et al., 2001). Poly Lack (Sayer Lack Mexicana,
565 Mexico), a polyether PU, was also tested on agar plate containing the polymer and minimal media
566 (Álvarez-Barragán et al., 2016). Strains able to grow on Poly Lack as sole carbon source were
567 isolated, yet, no clarification zone could have been observed on this polymer.

568

569 5.2. Biological degradation of TPU and PU foams

570 The most mainstream PU systems, i.e. TPU and PU foams, have been tested in biodegradation
571 assays. These products present a wide and varied range of chemical structures and groups. When
572 the purpose is to evaluate the susceptibility to biological degradation of a material, the structure
573 and formulation are precisely described in the study. In contrast, bioremediation and biological
574 recycling studies on PU mainly involve commercial products of complex and often unknown
575 chemical structure, additives and composition. The biological degradation experiments described
576 in the literature are mainly partial, leading to the recovery of the degraded polymer and, possibly,
577 soluble degradation products released in the aqueous media (Figure 13). Techniques developed
578 to evaluate the biological degradation of polymers can thus be oriented towards the efficiency of
579 the entire degradation system with the analysis of the degraded polymer products (soluble and
580 insoluble).



581
 582 *Figure 13 – Strategies and diversity of the analytical methods for TPU and PU foam biodegradation monitoring*

583
 584 *Samples structures and preparation*

585 Evaluation of PU foam degradation is more challenging than that of TPU, since foams are highly
 586 complex systems based on crosslinked architectures with different components and additives.
 587 Side reactions during synthesis and foaming steps can also lead to the formation of isocyanurate
 588 and other bonds. Foam analysis is limited by their insolubility in solvent.

589 PUs shape drives the specific surface and thus the bioavailability which is of importance for the
 590 biological degradation. Thanks to their alveolar structure, bioavailability is greater for foams,
 591 especially for open-cell foams, as microorganisms can easily circulate inside the material. Foams
 592 can also be ground to increase bioavailability (Cregut et al., 2014). TPUs for degradation assays
 593 can be used as thin films obtained by coating after solubilizing in an appropriate solvent, pouring
 594 on glassware and solvent evaporation (Chen et al., 2016; Woo et al., 2000). Thin films obtained
 595 can reach a few dozen micrometers (Phua et al., 1987). TPU films can also be obtained by
 596 thermoforming (Zhou and Xie, 2017). TPU cubes (Nakajima-Kambe et al., 1995), pellets (Cosgrove
 597 et al., 2007) or sheets (Ibrahim N. Ibrahim, 2009) are also used for PU degradation assays.

598 *Preparation of samples*

599 It is not recommended to sterilize TPUs by autoclaving for microbial experiments as most of the
 600 TPU becomes liquid-like or very soft at autoclaving temperatures (121°C). Degradation can occur
 601 e.g., a study comparing autoclaved and non-autoclaved poly(ether urea) PU material revealed
 602 that no weight loss was observed after autoclaving but a surface alteration appeared, leading to
 603 bias in degradation measurement (Rafiemanzelat et al., 2015). Alternatives such as rinsing with
 604 ethanol (Cosgrove et al., 2010; Mathur and Prasad, 2012), UV exposure (Gogoi and Karak, 2014)

605 or both (Osman et al., 2018) are thus frequently employed to sterilize the samples. Because of
606 better thermal resistance of thermosetting materials, foams can be tested after autoclaving
607 (Álvarez-Barragán et al., 2016).

608 *5.2.1. Evaluation of the global biodegradation efficiency*

609 The most widespread method to evaluate global biodegradation is the weight loss of the PU.
610 Samples are weighted before and after degradation assay to evaluate the weight of degradation
611 products released in the liquid media. It is a straightforward and easy method to implement,
612 although some bias can be noticed. For instance, if the degradation is too superficial and too low
613 to lead to detectable product release, the test will be considered as inefficient as no weight loss
614 will be measured. Therefore, weight loss measurement must be associated with a surface analysis
615 of the sample (Rafiemanzelat et al., 2015) in case of a low degradation extent, or to analyze the
616 early steps of the degradation. It is necessary to remove all the biological materials that can
617 remain on the polymer surface. Hard washing with ethanol is generally performed (Mathur and
618 Prasad, 2012; Urgun-Demirtas et al., 2007). Non-ionic surfactants such as TritonX-100
619 (Polyethylene glycol tert-octylphenyl ether) (1% v/v) have also been suggested to remove
620 reversibly bounded enzymes and cells (Ciardelli et al., 2004). This cleaning step is particularly
621 challenging for the fungal degradation of foams as filaments could be deeply trapped into the
622 bulk structure of the samples. It was recently suggested to use 0.88% (wt/vol) sodium
623 hypochlorite for 18 h to destroy and remove the remaining mycelium (Álvarez-Barragán et al.,
624 2016). When enzymatic degradations are performed, kinetic weight loss is generally set up
625 (Dogan et al., 2017; Zhou and Xie, 2017). In such experiments, a loss of enzymatic activity is often
626 observed with time. This loss of activity may be due to the thermal denaturation of enzymes
627 (Pastorino et al., 2004) or to the release of inhibiting compounds (Barth et al., 2015). To cope with
628 this phenomenon, enzymatic solutions are frequently renewed at regular time intervals (Phua et
629 al., 1987). Between the removal step and the renewal step, polymer pieces are usually washed,
630 dried and weighed.

631 As a parameter of global assay efficiency, the evaluation of the ability of a microorganism to grow
632 on PUs when the polymer is used as the sole source of carbon (or carbon and nitrogen) is common
633 practice (Cooney, 1969). Indeed, microbial development means that microorganisms can
634 depolymerize PUs and metabolize degradation products for growth. It is possible to quantitatively
635 follow the bacterial growth through the McFarland method which estimates the number of
636 bacteria thanks to turbidity measurement using UV-vis at 600 nm (Bayan and Karak, 2017;
637 Fernandes et al., 2016; Gogoi and Karak, 2014). Colony forming unit (CFU) is another method to
638 count bacteria: after being incubated with polymers, bacteria are sampled, diluted and poured
639 on an agar plate containing a rich medium (Crookes-Goodson et al., 2013; Urgun-Demirtas et al.,
640 2007). Colonies are counted after overnight incubation. The dry or wet weight of the biomass
641 corresponding to microorganisms growth can be measured (Oceguera-Cervantes et al., 2007).
642 This technique could be particularly suitable for bacteria that form aggregates or for filamentous
643 fungi.

644 Mineralization of polymers by microorganisms in aerobic conditions leads to the production of
645 CO₂ with O₂ consumption. Online sensors are used to measure both evolutions. These variations
646 must be compared to a negative control made without polymers (Cregut et al., 2014) or with an

647 inert polymer such as low-density polyethylene (Rattanapan et al., 2016), and to a conventional
648 biodegradable positive control such as cellulose (Gómez et al., 2014) or sodium benzoate
649 (Rattanapan et al., 2016) incubated with the same inoculum. For instance, low O₂ consumption
650 and CO₂ release were measured during the 28 days of degradation of a ground polyether PU foam
651 revealing low degradation by an acclimated microbial consortium (Cregut et al., 2014). CO₂
652 release during PU mineralization is associated with pressure increase. The pressure can be
653 measured by a Sturm test (Standard OECD 301 B) (Shah et al., 2016). This test is used for readily
654 biodegradable materials and usually lasts 28 days. Rattanapan et al., used this assay to measure
655 the biodegradation of a biobased polyester PU foam with long incubation time (60 days)
656 (Rattanapan et al., 2016). In fact, after 30 days, 7 to 11% of degradation was measured while a
657 higher degradation rate occurred during the last 30 days leading to a maximal degradation yield
658 of 46 wt%. Under anaerobic digestion, CH₄ is produced proportionally to polymer consumption.
659 Gomez et al., proposed to compare the biological degradation susceptibility of polyether PU
660 foams under composting, soil burial and anaerobic digestion according to three ASTM standard
661 methods based on CO₂ or CH₄ measurement (Gómez et al., 2014). These methods are ASTM
662 D5988-03 (Standard Test Method for Determining Aerobic Biodegradation in Soil of Plastic
663 Materials or Residual Plastic Materials After Composting) (International, 2003b), the ASTM
664 D5338-98 (Standard Test Method for Determining Aerobic Biodegradation of Plastic Materials
665 Under Controlled Composting Conditions) (International, 2003a) and the ASTM D5511-02
666 (Standard test method for determining anaerobic biodegradation of plastic materials under high-
667 solid anaerobic-digestion conditions) (International, 2002). The most pronounced degradation
668 was observed for a bio-based PU foam after 320 days of soil burial. Methods involving CO₂, O₂ or
669 CH₄ measurement present a low throughput. Incubation time is superior to 28 days. Moreover, a
670 pressure or gas monitor is required for each reaction which is generally performed in flasks from
671 2 to 5 liters (recommendation for OECD 301 series). These methods are mainly oriented towards
672 the evaluation of the biological sensitivity or resistance of newly synthesized or commercial PUs,
673 especially foams. These techniques are not suitable for screening of PU degrading entities.

674 *5.2.2. Study of the degraded polymers samples and insoluble products*

675 The first assessment of polymer degradation is a naked eye observation, sometimes sufficient to
676 evaluate the degradation onset. Change in color, roughness or shape can be noticed (Pilch-Pitera,
677 2012). These observations can be completed by surface analysis such as spectroscopy and/or
678 microscopy, polymer molar mass evolution by SEC or modifications of physical properties.

679 *Spectroscopy techniques*

680 FT-IR (Fourier-transform infrared spectroscopy) analysis relies on the fact that most molecules
681 absorb in the infrared region. This absorption corresponds specifically to the vibration modes of
682 the different bonds present in the analytes. Absorption spectra thus provide information on the
683 chemical structure of the polymer as a fingerprint. FT-IR is particularly popular for PU degradation
684 analysis because of its accuracy and rapidity and because FT-IR is a non-destructive method: the
685 sample is recovered without damage after the analysis. Surface analysis can be easily performed
686 to measure superficial biological degradation. Different issues can be encountered in the analysis
687 and interpretation of a spectrum. For instance, partial similarities between bonds e.g., urethane
688 and ester groups complicate the spectra interpretation for polyester PUs. Moreover, both

689 hydroxyl (OH) moieties, resulting from ester and urethane degradation, and amine (NH) moieties,
690 resulting from urethane degradation, absorb around 3400 cm^{-1} . For polyester PUs, the increase
691 of this large band is generally attributed to ester- or both ester and urethane hydrolysis (Oprea,
692 2010; Spontón et al., 2013), but it has also already been interpreted exclusively as the cleavage
693 of the urethane bond (Umare and Chandure, 2008).

694 Similarly, it is difficult to interpret changes in spectra presenting carbonyl bonds (C=O) that appear
695 in ester ($1750\text{-}1725\text{ cm}^{-1}$), urethane (1700 cm^{-1}) and urea bonds (1630 cm^{-1}). In polyester PUs,
696 there is often a unique broad signal representing both urethane and ester carbonyl bonds. Its
697 decrease is generally attributed to ester bond hydrolysis (Schmidt, J. et al., 2017; Shah et al., 2016)
698 but sometimes has been attributed only to urethane bond hydrolysis in polyester PUs (Gómez et
699 al., 2014; Ozsagiroglu et al., 2012). The signal at 1530 cm^{-1} is generally attributed to the nitrogen
700 of the urethane moieties. A comparison of the polyester PU based on PCL and the constitutive
701 PCL polyester showed that the signal at 1530 cm^{-1} only appeared on the PU spectrum thus
702 confirming that this signal is attributed to the urethane (Magnin et al., 2019). Oprea *et al.*
703 suggested that an increase of this signal is related to urethane bond hydrolysis (Oprea, 2010)
704 while others suggested that a decrease of this signal attests to urethane bond degradation
705 (Oceguera-Cervantes et al., 2007; Sarkar and Lopina, 2007). It is also conceivable that an increase
706 of this signal is correlated to the increase of the urethane proportion in the polymer after the
707 biological hydrolysis of the soft segment. Concluding on the variation of this signal upon
708 degradation appeared therefore complicated. Other peaks are sometimes considered as proof of
709 PU degradation. For instance, the emergence of a peak at 2250 cm^{-1} after degradation has been
710 attributed to isocyanate (NCO) release (Shah et al., 2016) (Figure 14b). However, isocyanates are
711 unstable in water and cannot be released through biological degradation in aqueous media. This
712 signal could correspond to atmospheric CO_2 resulting from improperly done FT-IR background
713 spectra (Gerakines et al., 1994). Since high variation on FT-IR spectra interpretations are observed
714 through the literature, especially concerning polyester PUs, additional methods must be used to
715 confirm urethane bond cleavage for PU degradation.

716 Kay *et al.*, suggested to consider the decrease of the ratio ester (C=O) /ether ($1720\text{ cm}^{-1}/1125\text{ cm}^{-1}$),
717 the ratio urethane (NH)/ether ($1630\text{ cm}^{-1}/1125\text{ cm}^{-1}$) and the ratio aryl (C=C)/ether in order to
718 provide a semi-quantitative analysis of the degradation of a polyether PU (Kay et al., 1993). The
719 ratio ester/ether decreases after degradation with a strain of *Corynebacterium* while the ratio
720 urethane/ether and aryl/ether remain stable meaning that the ester bonds are affected by the
721 bacterial degradation. Zhang et al., have used the band at 1463 cm^{-1} corresponding to CH_2
722 moieties to normalize their results (Zhang et al., 1994). The decrease of the normalized signals of
723 $1239\text{ cm}^{-1}/1463\text{ cm}^{-1}$ revealed an alteration of the polyester part of an arterial prosthesis made in
724 polyester PU after 100 days of incubation with pancreatin and collagenase. However, no change
725 of the normalized signal of $1695\text{ cm}^{-1}/1463\text{ cm}^{-1}$ showed the stability of the urethane moieties.

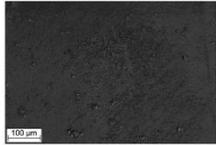
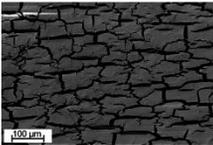
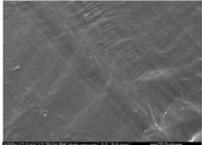
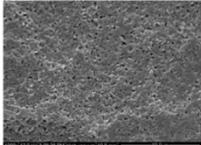
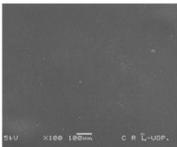
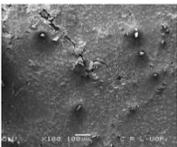
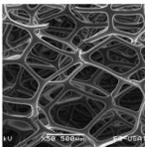
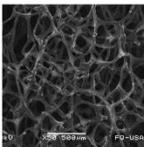
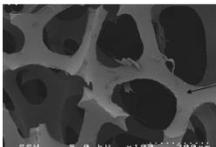
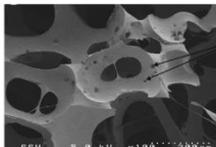
726 Recently, a Raman spectroscopy has been applied to monitor the biodegradation of a polyether
727 polyurethane foam which is among the most recalcitrant PU (Cregut et al., 2013). This technique
728 allowed conclusions to be drawn on the amorphous region degradation of the foam by a microbial
729 consortium while the crystalline region remained unaffected.

730 *Microscopy*

731 For the assessment of morphological surface modification of PUs, microscopy, particularly
732 scanning electron microscopy (SEM) is employed. SEM allows for a qualitative evaluation of the
733 degradation on the surface after biological treatment by observation of cracks or holes on the
734 degraded polymers. Enzymatic degradation generally leads to cracks (Figure 14a) or holes (Figure
735 14b) homogeneously spread at the TPU surface (Ozsagiroglu et al., 2012; Schmidt, J. et al., 2017)
736 while degradation with a microbial consortium leads to irregularities (Figure 14c) (Das et al., 2017;
737 Thirunavukarasu et al., 2015; Zafar et al., 2013). For instance, Das et al., showed the appearance
738 of cracks at the surface of a polyester TPU degraded under composting conditions. Depth of the
739 cracks, corresponding to fungal mycelium development, increased until the formation of holes
740 (Das et al., 2017).

741 SEM observations of the fungal mycelium propagation inside a PU foam highlights the higher
742 biodegradability of open-cell foams compared to closed-cell foams. The strut of cells appeared
743 distended, leading to the collapse of the alveolar structure (Figure 14d) (Álvarez-Barragán et al.,
744 2016). Degradation is efficient in a PDB medium (Potatoes Dextrose Broth, rich medium) but it is
745 specified that no degradation was observed by either weight loss or microscopy in minimal media.
746 Small holes appeared when foams were incubated with one of the three tested strains, confirming
747 enzymatic action. Holes in the walls and struts of the foam structure were already described
748 previously (Figure 14e) (Gautam et al., 2007c).

749

Observation	Abiotic control	Biological degradation	Substrate	Time of degradation/degraders
a. Regular enzymatic cracks on TPU surface			Polyester TPU	200 hours with LC cutinase
b. Regular enzymatic holes on TPU surface			Polyester TPU	24 hours with lipase from <i>Cryptococcus sp.</i>
c. Irregular microbial cracks on TPU surface			Polyester TPU	4 months soil burial
d. Collapse of the foam alveolar structure			Polyether PU foam	21 days degradation with <i>Cladosporium tenuissimum</i> (Fungi)
e. Holes in foams			Polyester PU foam	6 days degradation with <i>Pseudomonas chlororaphis</i> (Bacteria)

750
751 *Figure 14 – SEM images showing the morphological modifications of (a, b and c) thermoplastics PU and (d and e) PU*
752 *foams degraded by (a and b) enzymes, (c) microbial communities or (d and e) single strains. Photographs are*
753 *adapted from (a) Schmidt et al., 2017, (b) Thirunavukarasu et al., 2015, (c) Khan et al., 2017, (d) Alvarez-Barragan*
754 *et al., 2016 and (e) Gautam et al., 2007*

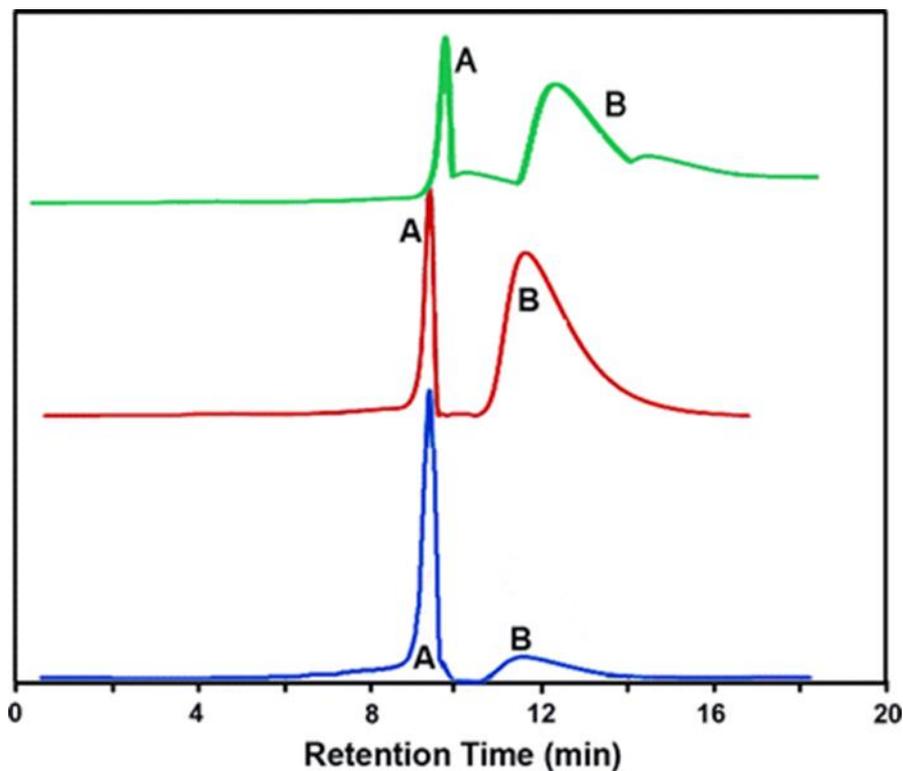
755 SEM is also used to evaluate microbial growth. For instance, *Micrococcus* biofilm formation on
756 the surface of a poly(ether urea) PU was shown by microscopy (Rafiemanzelat et al., 2013). SEM
757 can cope with the limitation of available assays to evaluate fungal growth on carbon depleted
758 media containing polymers. Huang et al. indeed highlighted a higher fungal growth on the surface
759 of an adhesive containing 70% of a polyester PU than the one containing 40% (Huang et al., 2016).

760 *Size Exclusion Chromatography (SEC)*

761 SEC analysis allows the determination of the molar mass distribution of polymers. This
762 quantitative analysis is more powerful than weight loss because it can appraise the change of
763 polymer mass distribution from the beginning of chains cut off even if soluble products are not
764 released. This measurement relies on the separation of the polymer chains in a column according
765 to their length. The polymer is solubilized in an organic solvent such as THF, chloroform,
766 dimethylformamide (DMF), then the solution runs through a fixed column packed with porous
767 beads (gel) with different sizes pores. Short chains pass through the pores while longer chains
768 cannot enter and are eluted more rapidly, then the higher the retention time, the lower the molar
769 mass is. Detection can be performed with a UV diode array detector and/or refractive index (RI)
770 detector. UV-vis analysis is efficient for PU containing aromatic rings such as 4,4'-MDI- or TDI-

771 based PUs (generally at 254 nm). Since the sample must be soluble in an organic solvent, this
772 method is not suitable for PU foams. Analysis and comparison of chromatograms can be
773 performed to evaluate PU degradation (Christenson et al., 2006; Rafiemanzelat et al., 2013).
774 Molar masses are usually determined with polystyrene standards. Because of its aromatic rings,
775 this standard is adapted to both UV and RI detection. Three main parameters are usually
776 considered with the number average molar mass (M_n), the weight average molar mass (M_w) and
777 the dispersity (\mathcal{D}) (ratio of M_w over M_n).

778 Polymer degradation leads to changes in the molar mass distribution. The most common
779 observed variation on PU biodegradation studies is a decrease of the M_w whereas M_n remains
780 unchanged, leading to a decreasing \mathcal{D} value (Schmidt, J. et al., 2017). M_w being more sensitive to
781 long polymer chains contribution, this is consistent with the cleavage of the long chains into lower
782 molar mass molecules. In Ferris et al., only the M_w was found to decrease (Ferris et al., 2010).
783 Changes in molar mass distribution reveal global degradation in the bulk material and not only
784 what is occurring at the surface of the polymer (Shah et al., 2013a; Shah et al., 2013b).
785 Rafiemanzelat et al. described a bi-modal SEC profile with a high and a low molar mass
786 distribution after 4 months soil burial degradation of a poly(ether urea) PU (Rafiemanzelat et al.,
787 2013) resulting from the cleavage of the long polymer chains into shorter ones (Figure 15). The
788 decrease of the peak area corresponding to the main polymer chain in favor of lower molar mass
789 chains, after fungal degradation of two polyester PUs, has also been reported (Magnin et al.,
790 2019).



791
792 *Figure 15 – SEC chromatograms of a poly(ether urea) PU before (bottom), after 4 months (middle) and after 6*
793 *months (top) of soil burial (Rafiemanzelat et al., 2013). A et B correspond respectively to the long and the short*
794 *polymers chains.*

795
796 Throughput of SEC analysis is rather low since a run generally lasts around one hour. Only a few
797 publications on PU biodegradation offered robust SEC data with several repetitions of the analysis
798 (Kuang and Mather, 2018; Magnin et al., 2019; Schmidt, J. et al., 2017).

799 *Modifications of the physical and physico-chemical properties of PU degraded*
800 *materials*

801 Observation of modifications in the physical and physicochemical properties of a sample can be
802 an indirect method of assessing biodegradation since the polymer chain cleavage often affects
803 some of these properties. These assays are thus used as complementary methods. Loss of
804 mechanical properties such as uniaxial tensile strength is generally observed after significant
805 biological degradation of a material. Tensile test measurement allows, for example, to evaluate
806 material elasticity and the behavior at break (Phua et al., 1987). For instance, after 24 months of
807 composting, a decrease of the tensile strength from 20 to 10 MPa was measured for a poly(ether
808 urea) TPU. In similar degradation conditions, samples of polyester PU were already broken down
809 into pieces and cannot be evaluated by uniaxial tensile tests (Krasowska et al., 2012). This
810 technique was also adapted for flexible PU foams. The tensile strength increased and the
811 elongation at break decreased after 60 days of incubation with a strain of *Pseudomonas* for
812 polyester PU foams (Spontón et al., 2013) (Figure 17, PU-1 and PU-2). In contrast, no change in
813 static mechanical properties was observed for the polyether PU foam before and after
814 degradation (Figure 17, PU-3). The polyether PU is thus more stable than the polyester PU.

815 PUs thermal stability may also be affected by biological degradation. This property can be
816 measured by thermogravimetric analysis (TGA) where the weight evolution of a sample is
817 recorded while the temperature is increased in a furnace, under air (oxidative) or N₂ (non-
818 oxidative) environments. For most polymers, the analysis temperatures range from 0 to 600-
819 800°C. Weight loss variations correspond to specific structure degradation and/or distinct
820 mechanisms. The first derivative of the TGA curve (DTG) gives curve with different peaks, to
821 determine different specific temperatures for each maximum in the case of a multistep
822 degradation, for instance. The maximum thermal degradation temperature can be used to
823 compare thermal stability between samples. For polyester TPUs, the first window from 100 to
824 300°C corresponds to the release of volatile compounds such as additives. Although urethane
825 bonds present a reversibility at around 200°C (Delebecq et al., 2013), urethane bonds degradation
826 induces a weight loss between 300 and 400°C while ester bond cleavage results in a weight loss
827 between 400 and 500°C (Cangemi et al., 2006; Mathur and Prasad, 2012). A decrease in weight
828 loss occurring between 400 and 500°C was observed after biological degradation (Mathur and
829 Prasad, 2012). This revealed a decrease in the ester bond content per polymer and thus evidenced
830 the biological hydrolysis of these linkages. Beyond the type of linkage affected, TGA may,
831 therefore, provide information on the material part affected. For instance, a poly(ether urea) PU
832 was incubated for one month with a strain of *Bacillus*. The observed changes in the material
833 thermal stability attested for a higher proportion of hard segments and consequently a
834 degradation occurring preferentially at the SS domains containing ether bonds (Rafiamanzelat et
835 al., 2015). TGA is perfectly appropriate to analyze cross-linked foams. In their study, Gomez et al.,
836 compared the composting of a fossil-based and a biobased PU foam (Gómez et al., 2014). TGA

837 was combined with MS analysis to identify the gaseous products of thermal degradation. They
838 found that a bio-based, aliphatic polyester PU is more sensitive to biodegradation than the fossil-
839 based, polyester ether PU with aromatic rings.

840 Differential scanning calorimetry (DSC) is a thermal analysis which appraises the phase transitions
841 of material. Glass transition temperature (T_g) is the temperature over which the amorphous
842 region of a polymer transitions from a hard to a more mobile/viscous state. Working above the
843 glass transition temperature (T_g) is an advantage for polymer degradation as it promotes chain
844 mobility of the amorphous zone of semi-crystalline material. Temperature changes depending on
845 the biological degrading-entities. Temperature of composting process can reach up to 58°C
846 (Genovese et al., 2016) while degradation with single strain of bacteria or fungi are generally
847 performed between 25 and 37°C. Enzymatic reaction can be performed at 60 or 70°C (Schmidt, J.
848 et al., 2017). The melting temperature (T_m) is defined as the temperature of transition from a solid
849 to a liquid state, which corresponds to the fusion of the crystalline regions. In practice, the area
850 of this fusion peak allows determining the crystallinity of the material. The crystallinity of a
851 poly(ether-urea) PU was found to increase after 4 months of soil incubation because of the
852 decrease in the amorphous region preferentially degraded (Rafiemanzelat et al., 2013). This
853 modification was accompanied by an increase in the T_m (Figure 19). Similarly, Osman et al.,
854 observed a shift of T_m from 191 to 196°C after the fungal degradation of a polyester PU (Osman
855 et al., 2018). In addition to this change, a decrease of T_g was measured. Pilch-Pitera et al., also
856 measured a lower T_g after PU degradation with an enzyme, associated with lower rigidity of the
857 material (Pilch-Pitera, 2012).

858 Biological degradation can alter the hydrophobicity of a PU material. Degradation induces
859 exposure of hydrophilic moieties on the polymer surface such as alcohol and carboxylic acid.
860 Moreover, disruption (cracks, holes) observed on the material surface offers a higher surface area
861 of access for water and thus higher hydrophilicity. To evaluate the hydrophobic/hydrophilic
862 balance, a drop of known liquid, generally water, is deposited on the polymer surface and the
863 contact angle is measured. With a hydrophilic material, the drop of water will collapse and the
864 contact angle will be low. A shift from 90° to 63° has been measured after 320 days of soil burial
865 of a polyester PU (Aranguren et al., 2012). Therefore, this polyester PU is more hydrophilic after
866 biological degradation.

867 *5.2.3. Analysis of the soluble degradation products*

868 The identification of soluble degradation products found in the liquid phase is the best route to
869 understand PU degradation mechanisms. Enzymatic degradation is more relevant than microbial
870 degradation in this case. Indeed, with microbial degradation, the quantifications are biased by
871 the possible assimilation of some degradation products by the microorganisms. Consequently,
872 the recovered products are those that cannot be metabolized by the microorganisms.

873 Quantification of these products is possible with the measurement of the Total Organic Carbon
874 (TOC) of the soluble fraction (Yamamoto et al., 2007). This measurement can be performed only
875 for enzymatic reactions as the amount of carbon brought by the enzyme remains stable in time
876 contrary to microbial cultures which involve growth and thus an increase of the carbon content.
877 Yamamoto et al., used this method to evaluate the degradation of several PUs based on LDI with

878 various proteases. For instance, 518 ppm of carbon from degradation products were released
879 after the degradation of a PU based on LDI and ethylene glycol, representing 44% of the
880 theoretical carbon of the polymer. Another way to measure the release of soluble degradation
881 products is the use of radiolabeled polymers. They were synthesized with ^{14}C molecules such as
882 ^{14}C -TDI, ^{14}C ethylene diamine, ^{14}C 1,4-butanediol or ^{14}C 1,6-HDI (Santerre et al., 1994; Woo et al.,
883 2000). The increase of radiolabeled-based molecules in the supernatant is quantified in counts
884 per minute (CPM). This is a very sensitive method, but the cost and the hazardous exposure to
885 radioactivity for the experimenter limit its usage. Both TOC and radioactivity measurements allow
886 precise quantification of the degradation products release, but these techniques do not give
887 information about the chemical structure of the corresponding degradation products.

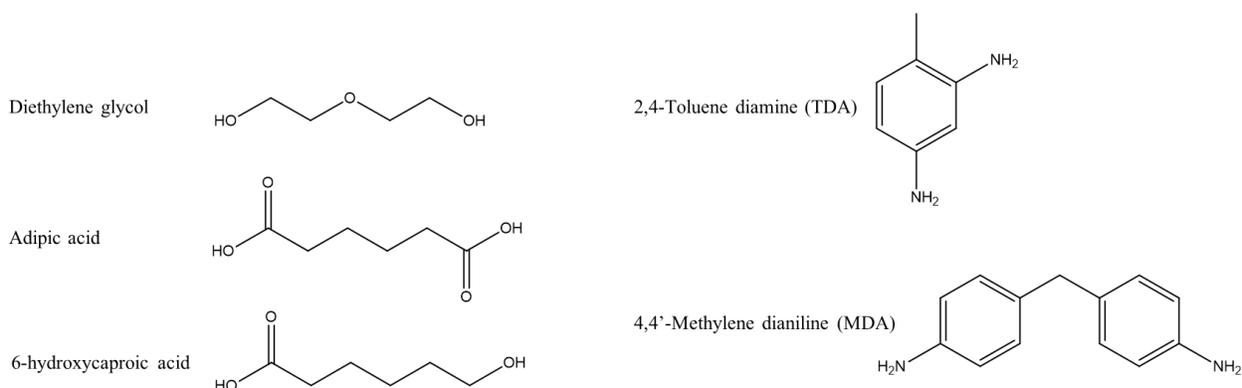
888 Another indirect way to assess PU biodegradation is to monitor the toxicity of the soluble fraction
889 resulting from the release of toxic degradation by-products. This is the case of 4,4'-methylene
890 dianiline (MDA), coming from the hydrolysis of MDI-based PU, which is known to be carcinogenic
891 (McQueen and Williams, 1990). The toxicity of the soluble fraction containing degradation
892 products can be evaluated using Microtox[®] bio-assay. This assay relies on a naturally luminescent
893 bacterium, *Photobacterium phosphoreum*. The parameter considered is the IC_{50} which is, in this
894 case, the volume of the soluble fraction that induces a decrease of the luminescence of 50%
895 (Spontón et al., 2013). A liquid medium of a polyester PU foam incubated for 60 days with
896 *Pseudomonas sp.*, presented an IC_{50} of 13.29% (V/V) thus attesting the release of toxic products
897 during the degradation, probably MDA or MDA-based molecules.

898 The efficient recovery of the degradation products for their analysis is an important issue. Indeed,
899 the liquid fraction of a degradation assay is a mixture containing salts, enzymes, and eventually
900 microbial cellular debris and degradation products which are molecules released from the
901 polymer. Several strategies were suggested to recover only the degradation products from PU
902 degradation assays. This recovery can be performed by solvent extraction using, for example,
903 ethyl acetate (Shah et al., 2016), acetonitrile (Tang et al., 2003) or ethyl ether (Spontón et al.,
904 2013). Instead of solvent extraction, selective recovery of degradation products can also be
905 achieved by removing enzymes using filtration (Wang et al., 1997a). Gamerith et al., added one
906 volume of methanol and acidified the supernatant to pH 3.5 so that proteins precipitated and
907 could be removed by centrifugation (Gamerith et al., 2016).

908 High Performance Liquid Chromatography (HPLC) with UV detection (Thirunavukarasu et al.,
909 2015) and mass spectrometry analyses associated with liquid chromatography (LC-MS) (Elliott et
910 al., 2002; Wang et al., 1997b) or with gas chromatography (GC-MS) (Pérez-Lara et al., 2016) are
911 methods of choice to identify the degradation products. These methods coupled a
912 chromatographic, for the separation of the mix of degradation products and an analytical method
913 such as the mass spectrometry. Detected degradation products highly depend on the initial
914 structure of polymers. Some chemical structures of identified degradation products are shown in
915 Figure 16. Thirunavukarasu et al., monitored the degradation of 50 mg of a poly(diethylene glycol
916 adipate)-based PU by quantifying diethylene glycol and adipic acid with HPLC analysis. After 4
917 days of PU incubation with a *Cryptococcus sp.* lipase, about 25 mg of adipic acid and about 8 mg
918 of diethylene glycol were released (Thirunavukarasu et al., 2015). The detection of specific amines
919 appears as the best way to confirm the cleavage of urethane bonds. It is interesting to notice that

920 the degradation of a PCL-based TPU (based on TDI as isocyanate starting material) with
921 cholesterol esterase leads to the release of low molar mass urethane molecules containing TDI-
922 based moieties but no TDA was detected (Wang et al., 1997a) (Figure 16). Then, the urethane
923 bond has not been cleaved. MDA was the unique aromatic amine released from polycarbonate-
924 based PUs synthesized with diverse diisocyanates (HDI, HMDI and MDI) after hydrolysis with
925 cholesterol esterase (Tang et al., 2003). This enzyme is thus able to cleave urethane linkages of
926 MDA based PU but not HDI- and HMDI-based PU. Gamerith et al., used liquid
927 chromatography/electrospray/time-of-flight mass spectrometry (LC/ESI/TOF-MS) and also
928 detected MDA as well as MDA derivatives after the hydrolysis of a polyester PU incubated with a
929 *Nocardia farcinica* polyamidase enzyme. Using the same procedure, Magnin et al., quantified
930 MDA at 0.3 mg/L after incubation of a PCL-based TPU for 50 days with an amidase (Magnin et al.,
931 2019).

932 NMR analysis was used for the identification of degradation products from the enzymatic
933 hydrolysis of a polyester PU based on PCL. 6-hydroxycaproic acid was predominantly identified
934 showing the efficient depolymerization of the ester linkage of the SS into the constitutive building
935 block by the esterase E3576 (Magnin et al., 2019).



936
937 *Figure 16 – Degradation products identified after PU biological degradation*

938

939 6. Conclusion: The challenge of PU biological degradation: the 940 urethane bond hydrolysis

941 PU are versatile polymers with high variability of structures, chemical compositions, formulations,
942 morphologies, shapes, with a direct impact on the biodegradation mechanisms and kinetics. From
943 the published literature, a large variety of potential or efficient biological degrading entities can
944 be identified among fungi, bacteria or enzymes. Because of the diversity of substrates and
945 analytical tools, a direct comparison of these degrading entities does not appear as an easy task.
946 Microbial degradation of PUs remains a complex and rather obscure process. For instance,
947 analysis of the set of enzymes produced by microorganisms during degradation often fails to
948 understand the mechanisms involved in PU degradation.

949 To tackle the degradation of the widest range of PU, the urethane bond cleavage appears as the
950 key parameter. However, only a few techniques provide undeniable proof of urethane bond

951 cleavage. Detection of amines derived from isocyanates as degradation products seems to be the
952 main direct way to prove urethane bond hydrolysis. For that purpose, the recent development of
953 new techniques such as LC/ESI/TOF-MS helps going forward in the resolution of this issue. Once
954 mastered, the efficient enzymatic hydrolysis of the urethane bond will undoubtedly pave the way
955 for a biological recycling of PU. Indeed, besides giving information on the degradation mechanism,
956 released molecules resulting from PU biological depolymerization can also be considered as
957 valuable products and used as building blocks for second-generation polymer synthesis. In the
958 review of Cregut *et al.*, the economic value of major building blocks was evaluated, showing the
959 interest of recovering molecules such as diethylene glycol, adipic acid or trimethylol propane,
960 which are products often identified after PU biological degradation (Figure 16) (Cregut *et al.*,
961 2013).

962 Applicability of the biological recycling on mainstream PU waste still needs to be attested. As far
963 as we know, only one study deals with the biodegradation of a real PU waste. Gautam *et al.*,
964 described the successful degradation of a waste polyester PU foam with a strain of *Pseudomonas*
965 *chlororaphis* (Gautam *et al.*, 2007c). Attempts to work with "real" PU waste (and no PU-based
966 models) have recently been performed by Alvarez-Barragan *et al.* by studying the degradation of
967 polyether-PU foam synthesized with and without the addition of a fire retardant tris(1,3-dichloro-
968 2-propyl)phosphate (TDCPP) (Álvarez-Barragán *et al.*, 2016). The TDCPP-containing foam was
969 found to be less sensitive to biodegradation thus highlighting the need for considering thoroughly
970 the presence of additives in real PU biological degradation assessment.

971 Polluting waste management such as landfilling will no longer be suitable solution. Limitation of
972 the pollution linked to PU waste is an outcome deeply needed and expected from recycling. An
973 efficient biological recycling path for PUs will support the economic value of PU waste towards
974 the development of a circular economy for plastic material.

975

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980

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