

Survey of antibiotic resistance of *Pseudomonas* isolated from fresh cut red chicory (*Cichorium intybus* L., Asteraceae)

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SUMMARY

The present work was carried out to investigate the safety aspects of minimally processed red chicory (*Cichorium intybus* L., family Asteraceae) regarding the antibiotic resistance of *Pseudomonas* Migula, 1894 populations (Bacteria Pseudomonadaceae). The *Pseudomonas* strains tested for their characteristics were previously isolated from experimental monovarietal salads prepared with two different processes, a classic ready-to-eat (RTE) process and a process with entire leaves, and stored under refrigeration (4 °C) up to 15 days from production. Due to their dominance over the microbial community, *Pseudomonas* strains were better characterised for their undesirable features that might be exerted on the health of consumers. To this purpose, the 21 dominant *Pseudomonas* strains were subjected to the antimicrobial resistance test, applying the antibiotics commonly administered to combat intestinal bacteria responsible for human infections. The results revealed a certain sensibility to almost all the antibiotics diffused among these bacteria, with the exception of ampicillin, amoxicillin, and chloramphenicol. If, on one hand, this study has found that, generally, *Pseudomonas* are not related to the dissemination of the majority of antibiotics used for hospitalized patients, on the other hand, it poses the attention to this microbial group regarding ampicillin and amoxicillin.

INTRODUCTION

In the last decades, there has been the affirmation of a healthy life style of consumers in several countries. For this reason, vegetables became essential components of a healthy diet, thanks to their nutrients and micronutrients. They are a source of a wide variety of minerals, vitamins, and phytochemicals with antioxidant activity (SIRIAMORNUN ET AL., 2012; TANGO ET AL., 2018).

The high attention of consumers towards the composition of foods and the awareness of the risk factors represented by several foods of animal origin, determined an increasing demand for vegetables, especially those available in a ready-to-eat (RTE) form that require no further treatment before consumption (MAFFEI ET AL., 2016). This phenomenon determines a continuous increase of the request of vegetables (CHAKRABORTY & CHATTOPADHYAY, 2018).

In order to obtain RTE products, leafy vegetables are minimally processed through selection, prewashing, cutting, washing, drying, packaging, and cold storage (CODEX ALIMENTARIUS COMMISSION, 2003; KENNEDY & WALL, 2007). However, due to the large consumption, RTE vegetables may represent a public health issue because they can be a vehicle for the transmission of bacterial, parasitic, and viral pathogens (ABADIAS ET AL., 2008). The contamination of fresh cut products can take place during the pre- and post-harvest operations due to the contact with contaminated irrigation water, soil, manure of wild and domestic animals, following a cross-contamina-

tion in any point of the production process (BERGER ET AL., 2010; HARRIS ET AL., 2003; PARK ET AL., 2012; VERHOEFF-BAKKENES ET AL., 2011), and during the administration to the final consumers. Furthermore, the refrigerated temperature storage does not stop the growth of pathogenic and/or spoilage micro-organisms (AGUADO ET AL., 2004; GLEESON & O'BEIRNE, 2005).

Pseudomonads (Bacteria Pseudomonadaceae) are commonly associated to the spoilage of RTE salads (Nguyen-the and Prunier, 1989) and may represent up to 90% of the microbial community of fresh vegetables (GARG ET AL., 1990; ZAGORY, 1999; WULFKUEHLER ET AL., 2015). In a previous work (ALFONZO ET AL., 2018), *Pseudomonas* Migula, 1894 represented the dominant group of red chicory subjected to cut and uncut processes, with *P. fluorescens*, *P. endophytica*, *P. grimontii*, and *P. poae* as being the most frequently isolated species.

Antimicrobial resistant bacteria represent a public health issue, since their emergence, due to the antimicrobial agents used for clinical therapeutic use occurred worldwide (ECDC/EFSA/EMA 2015). In this work, the effect of the antibiotic commonly used against human pathogens was tested on *Pseudomonas* spp. from fresh cut *Cichorium intybus* L.

MATERIAL AND METHODS

The *Pseudomonas* strains used in this study belonged to the culture collection of the Agricultural Laboratory - Department of Agricultural, Food and Forest Science (University of Palermo, Italy). The strains were: *Pseudomonas azotoformans* (Ac. No. KY939740); *Pseudomonas beatica* (Ac. No. KY939741); *Pseudomonas brenneri* (Ac. No. KY939742); *Pseudomonas endophytica* (Ac. No. KY939743 – KY939744); *Pseudomonas extremaustralis* (Ac. No. KY939745); *Pseudomonas fluorescens* (Ac. No. KY939746 – KY939748); *Pseudomonas grimontii* (Ac. No. KY939749 – KY939750); *Pseudomonas helleri* (Ac. No. KY939751); *Pseudomonas libanensis* (Ac. No. KY939752); *Pseudomonas marginalis* (Ac. No. KY939753); *Pseudomonas poae* (Ac. No. KY939754 – KY939755); *Pseudomonas psychrophila* (Ac. No. KY939756); *Pseudomonas simiae* (Ac. No. KY939757); *Pseudomonas trivialis* (Ac. No. KY939758); *Pseudomonas viridiflava* (Ac. No. KY939759); *Pseudomonas yamanorum* (Ac. No. KY939760). All strains were reactivated in Luria Bertani broth (Oxoid) incubated overnight at 25 °C.

The assay for the antimicrobial susceptibility test was carried out according to the Clinical and Laboratory Standards Institute guidelines (CLSI, 2015) that, except *Pseudomonas aeruginosa*, are described for species other than *Pseudomonas*. The tests were performed applying the disk diffusion method. Bacterial inoculums were obtained from colonies developed overnight on PAB, suspended in Ringer's solution (5 ml) to a cell density of 0.5 McFarland standard and swabbed for confluent growth onto Mueller Hinton agar (Oxoid).

The following antibiotics at the concentrations generally used for the treatment of human and animal infections were tested: high-level gentamicin (GN—120 µg); high-level streptomycin (STR—300 µg); amoxicillin/clavulanic acid (AMC—30 µg); ampicillin (AMP—10 µg); chloramphenicol (C—30 µg); ciprofloxacin (CIP—5 µg); levofloxacin (LEV—5 µg); tetracycline (TE—30 µg). All antimicrobial compounds were purchased from Oxoid.

The effects of the antibiotics were evaluated after 18 hours of incubation at 37 °C, by measuring the inhibition halos according to the Clinical and Laboratory Standards Institute guidelines (CLSI, 2015). The antibiotics quinupristin/dalfopristin (QD—15 µg) was also included in the assay since Gram negative bacteria, such as *Pseudomonas* spp., are inherently resistant to this antibiotics because of cell wall impermeability (NADLER ET AL., 1999).

RESULTS AND CONCLUSIONS

Due to their dominance over prokaryotic and eukaryotic microorganisms found in red chicory samples by ALFONZO ET AL. (2018), *Pseudomonas* were deeply characterized for their resistance to the common antibiotics used for the treatment of infections.

The results of the resistance of the 21 *Pseudomonas* strains to the antibiotics tested is reported in Table 1. Only *P. viridiflava* 4G477 was susceptible to ampicillin. A low resistance was registered towards chloramphenicol and amoxicillin that inhibited 14 and five strains, respectively. Some strain-dependent behaviours were also observed, since only the strain 4G558 was inhibited by amoxicillin within the species *P. poae*. All strains were highly susceptible to gentamicin, streptomycin, ciprofloxacin, levofloxacin, and tetracycline with the diameter of the inhibition areas ranging between 16.0 to 36.0 mm. The most sensitive strain was *P. viridiflava* 4G477, inhibited by all antibiotics.

The epiphytic and endophytic microbial communities of plants play different roles. Basically, within

bacteria, some populations are plant pathogens, some can act as plant growth promoters and some others are responsible for human diseases (JACKSON ET AL., 2013). The operation of cutting leafy vegetables stimulates the growth of several microorganisms (STEELE, 2004) as a consequence of the higher and direct availability of nutrients generated by the damaged cells.

The involvement of pseudomonads in the deterioration of RTE salads is well known (NGUYEN-THE & PRUNIER, 1989). However, to our knowledge, no previous work characterised this dominating population of minimally processed red chicory for their antibiotic resistance. In the recent years, the rapid emergence of resistant bacteria is occurring across the globe, endangering the efficacy of antibiotics (VENTOLA, 2015). These bacteria present in foods can be a possible intermediate vehicle for the transmission of multidrug resistances to the human intestinal tract (GAGLIO ET AL., 2016; JAMET ET AL., 2012). Several initiatives have been launched worldwide to address

the biosafety concerns of starter cultures and probiotics (MATHUR & SINGH, 2005), but very little is known about the role of saprophytic microorganisms in the dissemination of antibiotic resistance characters. With this in mind, the dominant strains isolated from the red chicory processed in this study were evaluated for their resistance to the antibiotics commonly administered to combat intestinal bacteria responsible for human infections. In particular, eight antibiotics generally used to test the resistance of enterococci (CLSI, 2015), ubiquitous in several foods and human gut (LEBRETON ET AL., 2014), were used. Basically, the majority of the dominant *Pseudomonas* strains were susceptible to the action of the antibiotics.

In conclusion, *Pseudomonas* do not represent a risk of antibiotic resistance genes related to the consumption of fresh cut red chicory. However, their resistance to ampicillin and amoxicillin should be better considered in RTE salads.

Strains	Antibiotics ^a									Quality control
	Aminoglycosides		Penicillins		Phenicol	Fluoroquinolones		Tetra-cyclines	Strepto-gramins	
	GN	STR	AMP	AMC	C	CIP	LEV	T	QD	
<i>P. azotoformans</i> 4G504	18.5 ± 0.5	22.5 ± 0.5	0	0	10.0 ± 0	24.0 ± 0	30.5 ± 0.5	29.0 ± 1.0	0	
<i>P. baetica</i> 4G537	18.5 ± 0.5	19.0 ± 0	0	0	0	25.5 ± 0.5	24.5 ± 0.5	20.5 ± 0.5	0	
<i>P. brenneri</i> 4G513	23.0 ± 1.0	27.5 ± 0.5	0	0	20.5 ± 0.5	30.0 ± 0	34.0 ± 1.0	28.5 ± 1.5	0	
<i>P. endophytica</i> 4G414	21.0 ± 1.0	25.5 ± 1.5	0	16.0 ± 0	14.0 ± 0	24.0 ± 0	22.5 ± 0.5	30.0 ± 1.0	0	
<i>P. endophytica</i> 4G764	18.0 ± 0	21.0 ± 0	0	17.0 ± 1.0	12.5 ± 0.5	24.0 ± 1.0	23.5 ± 0.5	28.5 ± 1.5	0	
<i>P. extremaustralis</i> 4G619	20.0 ± 1.0	24.0 ± 1.0	0	0	9.5 ± 0.5	22.0 ± 0	23.0 ± 0	24.0 ± 1.0	0	
<i>P. fluorescens</i> 4G628	18.0 ± 0	20.0 ± 1.0	0	0	0	24.0 ± 1.0	26.0 ± 1.0	25.0 ± 0	0	
<i>P. fluorescens</i> 4G1034	18.0 ± 1.0	20.0 ± 0	0	0	0	25.0 ± 0	22.5 ± 0.5	23.0 ± 0	0	
<i>P. fluorescens</i> 4G1237	22.5 ± 0.5	27.0 ± 1.0	0	0	0	30.0 ± 1.0	30.0 ± 1.0	26.5 ± 0.5	0	
<i>P. grimontii</i> 4G769	18.5 ± 1.5	26.5 ± 0.5	0	0	12.5 ± 0.5	30.0 ± 1.0	30.5 ± 0.5	28.5 ± 0.5	0	
<i>P. grimontii</i> 4G483	18.0 ± 1.0	27.0 ± 1.0	0	0	11.5 ± 0.5	30.0 ± 1.0	30.0 ± 1.0	30.0 ± 1.0	0	
<i>P. helleri</i> 4G893	10.5 ± 0.5	16.0 ± 0	0	11.5 ± 0.5	10.5 ± 0.5	23.0 ± 0	20.0 ± 0	23.0 ± 1.0	0	
<i>P. libanensis</i> 4G787	20.0 ± 1.0	18.0 ± 0	0	0	0	27.5 ± 0.5	29.5 ± 0.5	25.5 ± 0.5	0	
<i>P. marginalis</i> 4G518	20.5 ± 1.5	2.65 ± 0.05	0	0	14.0 ± 2.0	28.5 ± 1.5	32.0 ± 1.0	31.5 ± 1.5	0	
<i>P. poae</i> 4G558	20.0 ± 0	24.0 ± 0	0	8.0 ± 0	0	30.5 ± 0.5	34.0 ± 1.0	30.0 ± 1.0	0	
<i>P. poae</i> 4G1030	18.0 ± 1.0	23.5 ± 0.5	0	0	8.0 ± 0	28.5 ± 0.5	28.0 ± 0	27.5 ± 0.5	0	
<i>P. psychrophila</i> 4G793	16.5 ± 0.5	19.0 ± 1.0	0	12.5 ± 0.5	18.5 ± 0.5	22.5 ± 0.5	22.5 ± 0.5	36.0 ± 2.0	0	
<i>P. simiae</i> 4G1010	18.0 ± 1.0	24.5 ± 0.5	0	0	0	26.0 ± 2.0	28.5 ± 1.5	24.5 ± 1.5	0	
<i>P. trivialis</i> 4G921	20.0 ± 1.0	25.5 ± 0.5	0	0	12.5 ± 0.5	23.0 ± 1.0	26.0 ± 0	25.5 ± 0.5	0	
<i>P. viridiflava</i> 4G477	24.0 ± 0	30.5 ± 0.5	7.5 ± 0.5	21.0 ± 0	15.5 ± 1.5	35.5 ± 1.5	33.0 ± 2.0	28.5 ± 1.5	0	
<i>P. yamanorum</i> 4G531	19.0 ± 1.0	23.0 ± 0	0	0	10.0 ± 0	27.5 ± 0.5	26.0 ± 1.0	25.5 ± 0.5	0	

Table 1. Antimicrobial resistance of *Pseudomonas* strains collected from cut and uncut red chicory samples during storage. GN: gentamicin; STR: streptomycin; AMP: ampicillin; AMC: amoxicillin; C: chloramphenicol; CIP: ciprofloxacin; LEV: levofloxacin; T: tetracycline; QD: quinupristin-dalfopristin.

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