1	Occurrence of Hepatitis A and E and Norovirus GI and GII in ready-to-eat
2	vegetables in Italy
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### 18 Abstract

Fresh vegetables and their ready-to-eat (RTE) salads have become increasingly recognized as potential vehicles for foodborne diseases. The EU Reg. 1441/2007 establishes microbiological criteria for bacterial pathogens for products placed on the market during their shelf-life (i.e. *Salmonella* spp., *Listeria monocytogenes*) for pre-cut fruits and vegetables (RTE) whilst it does not address the problem of contamination by enteric viruses.

24 In this study we investigated the contamination by hepatitis A virus (HAV), hepatitis E virus (HEV)

and norovirus (NoV) in 911 ready-to-eat vegetable samples taken from products at retail in Apulia
and in Lombardia.

The vegetable samples were tested using validated real-time PCR (RT-qPCR) assays, ISO
standardized virological methods and ISO culturing methods for bacteriological analysis.

The total prevalence of HAV and HEV was 1.9% (18/911) and 0.6% (6/911), respectively. None of the samples analyzed in this study was positive for NoV, *Salmonella* spp. or *Listeria monocytogenes*. The detection of HAV and HEV in RTE salads highlights a risk to consumers and the need to improve production hygiene.

33 Appropriate implementation of hygiene procedures is required at all the steps of the RTE vegetable

34 production chain and this should include monitoring of emerging viral pathogens.

**Keywords:** green vegetables, viruses, molecular methods

### 36 **1. Introduction**

37 Leafy green vegetables and their ready-to-eat (RTE) salads are important components of the current 38 human diets but are accompanied by new food safety threats since they are eaten raw and usually 39 without any further washing/decontamination procedures (Little and Gillespie, 2008). Apart from 40 psychrotrophic pathogens and spoilage microorganisms, RTE vegetables can be contaminated with 41 a number of human pathogens, including parasites, bacteria and viruses.

In Italy, the prevalence of bacterial pathogens in leafy green vegetables has been estimated between 3.7 for fresh and 1.8% for RTE products (Losio et al., 2015). Numerous foodborne virus outbreaks have been linked to the consumption of fresh produce, mostly attributable to enteric viruses such as norovirus (NoV), hepatitis A virus (HAV), hepatitis E virus (HEV), rotavirus (RV) and astrovirus (AstV) (European Food Safety Authority & European Centre for Disease Prevention and Control, 2013; Chiapponi et al., 2014; Collier et al., 2014; European Food Safety Authority, 2014; Kokkinos et al., 2012; Terio et al., 2015; Wheeler et al., 2005).

49 Enteric viruses may contaminate vegetables, during cultivation before harvest or post-harvest. 50 During pre-harvest cultivation, there are various routes of contamination, which usually include 51 application of organic wastes as fertilizer, contamination of water used for irrigation with faecal 52 material, contact with inadequately-treated sewage or sewage-polluted water. In addition, direct contamination by livestock, wild animals and birds should be considered (Heaton and Jones, 2008). 53 54 Water is the main critical vehicle of contamination in the farm-to-fork continuum. Spraying, washing or immersion of fruits and vegetables in water are common practices during post-harvest 55 56 processing (Gandhi et al., 2010). Moreover, minimal processing may induce cross-contamination of 57 clean produce during cutting, washing and packaging (López-Velasco et al., 2010). In addition, much emphasis has also been placed on the role of workers during collection, processing, storage, 58 59 distribution or final preparation (European Food Safety Authority BIOHAZ Panel, 2014; Koopmans and Duizer, 2004; Rzezutka and Cook, 2004). 60

61 Currently, the European legislation on the safety of pre-cut (ready-to-eat) fruit and vegetables requires the assessment of *Salmonella* spp. and *Listeria monocytogenes* contamination at the end of 62 63 the production chain (at retail). In addition, the Escherichia coli count is required during the 64 manufacturing processes (Commission Regulation (EC) No. 1441/2007). The regulations do not take into consideration the risks deriving from food contamination by enteric or emerging viral 65 pathogens, although there is a growing attention and interest for this important risk, as evidenced by 66 67 recent NoV and HAV outbreaks in Europe (European Centre for Disease Prevention and Control 68 and the European Food Safety Authority, 2014; Müller et al., 2016). Limited data exist in the 69 literature on the prevalence of foodborne viruses in RTE vegetables. In Italy, a study on RTE at 70 retail has been reported recently, revealing a very low rate (< 0.1%) of NoV contamination (Losio 71 et al., 2015). The purpose of this study was to investigate the presence of enteric viruses (HAV, 72 HEV and NoV) in RTE vegetables available for sale in Italy.

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#### 74 **2. Materials and Methods**

#### 75 2.1 Sampling

76 A total of 911 samples of bagged ready-to-eat vegetables belonging to different brands and 77 purchased from supermarkets in Apulia and Lombardy regions, Italy, were collected during 2014-78 2015. They included 619 mixed salads, 53 carrot (Daucus carota subsp. sativus), 40 valerian 79 (Valeriana officinalis), 104 rocket (Eruca vesicaria), 10 spinach (Spinacia oleracea), 18 Iceberg (Lactuca sativa) and 67 Romaine lettuce (Lactuca sativa L var. longifolia). All samples were 80 81 labelled as "Pre-washed and ready-to-eat". All samples were obtained in their original packaging 82 and analysed before the expiration date (up to 8 days). Samples were transported to the laboratory 83 in refrigerated boxes ( $< 8^{\circ}$ C) and analysed on the day of purchase.

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#### 85 2.2 Isolation of Salmonella spp. and Listeria monocytogenes

- 86 Vegetable samples were subjected to isolation of the pathogenic foodborne microorganisms
  87 Salmonella spp. and Listeria monocytogenes, using EN/ISO 6579 and EN/ISO 11290-1,
  88 respectively.
- 89
- 90 2.3 Virus concentration and nucleic acid extraction

Viral RNA was extracted following the ISO/TS 15216-2:2013 method for NoV and HAV detection
in foodstuffs. In brief, 25g of each sample were cut into small pieces and homogenized with TGBE
buffer pH 9.5 (100 mM Tris–HCl, 50 mM glycine, 1% beef extract) and 10 µl of process control
virus material (Mengovirus). The eluate was concentrated with 5X PEG/NaCl solution (50% (w/v)
PEG 8000, 1.5 M NaCl) and the viral nucleic acids were extracted and purified using commercial
kits (NucliSENS miniMAG kit, bioMérieux) according to the manufacturer's instructions.

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#### 98 2.4 Reverse transcriptase-polymerase chain reaction

99 Reverse transcription of viral RNA was performed using High-Capacity cDNA Reverse 100 transcription Kit (Applied Biosystems, Italy), containing 10X RT Buffer II, 5 mM MgCl<sub>2</sub>, 1 mM 101 dNTPs, 2.5 μM random hexamers, 20 U RNAse inhibitor and 50 U Reverse transcriptase according 102 to the manufacturer's instructions. The reaction conditions were 42°C for 30 min and 99°C for 5 103 min. The obtained cDNA was used for specific real-time PCR (RT-qPCR) for each target virus: 104 HAV, HEV, NoV GI and NoV GII.

105 The RT-qPCR reactions were performed in duplicate. All tests included negative controls for virus106 and for an internal amplification control (IAC).

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108 2.5 Hepatitis A virus qPCR

109 This assay was performed using the primers and conditions described in the ISO/TS 15216-2:2013 110 method with the inclusion of an internal amplification control (IAC). The reaction contained 1X 111 iTaq<sup>TM</sup> Universal Probes Supermix (Bio-Rad), 0.5  $\mu$ M primer HAV68, 0.9  $\mu$ M primer HAV240; 112 0.25  $\mu$ M probe HAV150 (-) (FAM labelled) (Costafreda et al., 2006), 50 nM IAC probe (VIC 113 labelled) and 300 copies of IAC (Martinez-Martinez et al., 2011). 20  $\mu$ l of cDNA was added to 114 make a final reaction volume of 50  $\mu$ l. The thermocycling conditions were 5 min at 95°C, followed 115 by 45 cycles of 15 s at 95°C and 1 min at 60°C and 1 min at 65°C.

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117 2.6 Hepatitis E virus qPCR

This assay used the primers and conditions described by Jothikumar et al. (2006) with the inclusion of an internal amplification control (IAC). The reaction contained 1X iTaq<sup>TM</sup> Universal Probes Supermix (Bio-Rad), 0.25  $\mu$ M for each primer (JVHEVF and JVHEVR), 0.1  $\mu$ M probe JVHEV-P (labelled with FAM), 50 nM IAC probe (labelled with VIC) and 300 copies of IAC (Martinez-Martinez et al., 2011). 20  $\mu$ l of cDNA was added to make a final reaction volume of 50  $\mu$ l. The thermocycling conditions were 10 min at 95°C, followed by 40 cycles of 15 s at 95°C and 1 min at 60°C.

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## 126 2.7 Norovirus GI and GII qPCR

This assay was performed using the primers and conditions described in the ISO/TS 15216-2:2013
method with the inclusion of an internal amplification control (IAC). The reaction was carried out
using 1X iTaq<sup>TM</sup> Universal Probes Supermix (Bio-Rad), 250nM probe (labelled with FAM), 500nM
forward primer and 900nM reverse primer, 50nM IAC probe (labelled with VIC) and 300 copies of
IAC (Martinez-Martinez et al., 2011).

Primers targeted the ORF2 region; for NoV GI: forward primer QNIF4, reverse primer NV1LCR
and probe NV1LCpr were employed; for NoV GII, forward primer QNIF2d, reverse primer
COG2R and probe QNIFS were used (da Silva et al., 2007; Kageyama et al., 2003; Loisy et al.,
2005; Svraka et al., 2007).

136 20  $\mu$ l of cDNA was added to make a final reaction volume of 50  $\mu$ l. The thermocycling conditions

137 were 5 min at 95°C, followed by 45 cycles of 15 s at 95°C and 1 min at 60°C and 1 min at 65°C.

#### 139 2.8 Extraction efficiency

140 For each sample, two aliquots of 5µl nucleic acid (NA) extract were added to adjacent wells of a 141 96-well optical reaction plate and made up to 25µl with Mengovirus-specific TaqMan reaction mix. 142 A dilution series prepared from the Mengovirus process control material was also tested. The 143 percentage extraction efficiency for each sample was determined by comparing the Ct values for the 144 sample NA extract with those for the Mengovirus dilution series. Any sample with an extraction 145 efficiency of <1% was subjected to retesting, in a first instance by re-extracting the viral RNA from 146 stored homogenate, then by testing the stored RTE salads. Results for any sample providing three 147 extraction efficiency results of <1% were considered invalid.

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## 149 **3. RESULTS**

- 150
- 151 3.1. Isolation of Salmonella spp. and Listeria monocytogenes
- 152 The pathogenic foodborne microorganisms *Salmonella* spp. and *L. monocytogenes*, were not 153 detected in any RTE salads.
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- 155 3.2 Detection of foodborne viruses by RT-qPCR
- 156 The mean viral extraction efficiency of the process was > 1% for each sample.
- 157 The results obtained from virological analysis of 911 RTE vegetable samples are summarized in158 Table 1.
- 159 HAV and HEV were detected in 2.6% of the tested samples (24/911): HAV was the most prevalent
- pathogenic enteric virus, detected in 1.9% of samples (18/911), and followed by HEV in 0.6% ofsamples (6/911).
- 162 In detail, 14 of the HAV positive samples were mixed salad samples (2.3% of mixed salad 163 samples), 1 valerian (2.5% of valerian samples) and 3 rocket samples (1.9% of rocket samples).

164 Two mixed salad samples resulted HEV positive (0.3 % of mixed salad samples), 2 rocket samples
165 (1.9%) and 2 spinach samples (20%). None of the samples analysed in this study tested positive for
166 NoV.

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#### 168 **4. DISCUSSION**

Over the years, foodborne NoV and HAV outbreaks have mainly been linked to shellfish consumption in the EU although more recently other matrices such as leafy vegetables and frozen red fruits (Chiapponi et al., 2014; Severi et al., 2015; Terio et al., 2015) have been associated with these infections. Foodborne transmission of HEV has been increasingly reported in sporadic cases and small outbreaks associated with the consumption of raw or uncooked boar or deer meat, of liver and liver sausages (Colson et al., 2010; Di Bartolo et al., 2015; Guillois et al., 2015), although HEV has also been detected in fresh produce (Kokkinos et al., 2012).

In this study, *Salmonella* spp. and *L. monocytogenes* were not detected in RTE salads. Our findings
are in general agreement with another investigation performed in Italy in which a low prevalence
(0.3-0.5%) of contamination by *Salmonella* spp. and *L. monocytogenes* was reported (Losio et al.,
2015).

180 On the opposite, in our study the virological investigations detected the presence of HAV and HEV 181 in 1.9% and 0.6% of the analysed samples, respectively. To our knowledge, this is the first report 182 documenting the presence of HAV and HEV in RTE salads in Italy. One limitation of this study was the inability to determine the viability of the viruses detected in the samples; indeed, although 183 184 the laboratory-adapted HAV strains can grow on foetal rhesus monkey kidney cells (Kingsley and 185 Richards, 2001), wild-type HAV strains are unable to replicate in vitro. Likewise, there is no 186 specific cell culture line for laboratory cultivation of HEV and cultivation of human NoV is 187 fastidious (Ettayebi et al., 2016; Jones et al., 2015). On the other hand, the fact that RT-PCR methods can detect inactivated viruses may not be a critical factor for RTE vegetables risk 188

assessment since a small number of viable virus particles are sufficient to cause illness (Fiore, 2004;

190 US Food and Drug Administration, 2012).

Whether the virus detected in RTE salads is viable or not, the presence of viral acid nucleic revealed by the molecular assays represents a clear index of contamination and potential risk for consumers considering that RTE salads do not undergo any treatment able to guarantee the virus inactivation. Moreover, this contamination represents an unacceptable risk that food handlers have to evaluate in the risk assessment plan (EC regulation N° 178/2002).

This study indicates that attention should be paid to enteric viruses and emerging viral pathogens in order to guarantee safety of RTE vegetables. Indeed, screening for enteric viruses, which are not correlated to the presence of *E. coli* and *Salmonella* spp. (Ethelberg et al., 2010; Fong and Lipp, 2005; Krog et al., 2014; Westrell et al., 2010), is crucial when assessing the health risks related with RTE salads, as already pointed out by the EU Regulation 1235/2012 for frozen strawberry.

For this reason, with a few exceptions in HACCP plans food businesses usually establish bacteria as a food safety hazard, whilst viruses are not considered as hazards. In addition, the measures taken to reduce the growth of, or eliminate bacteria, do not necessarily lead to a decrease in virus prevalence. On some occasions, these measures may even preserve viral particles, as is the case of refrigeration (Croci et al., 2002; Fiore, 2004).

Most of the available literature regarding the use of sanitizers during RTE production, such as chlorine, has concluded that washing with water or with disinfectant solutions reduces the natural microbial populations on the surface of the product by only 2 to 3 log units (Allende et al., 2007; Beuchat et al., 2004; Gómez-López et al., 2007; Gonzalez et al., 2004; Inatsu et al., 2005; Selma et al., 2008; Ukuku et al., 2005) and it has been reported that current industrial sanitizing treatments do not guarantee the total elimination of pathogens when present (Abadias et al., 2008; Beuchat, 1996; Parish et al., 2003; Pérez-Rodríguez et al., 2011; Posada-Izquierdo et al., 2014).

213 Studies on the efficacy of disinfectants for inactivating viruses show that viruses are relatively 214 resistant to chlorine decontamination (Seymour and Appleton, 2001), that viruses are generally more resistant than bacteria (Allwood et al., 2004), that virus sensitivity to disinfectants may vary widely between different virus species (Fraisse et al., 2011) and that sanitizing agents or disinfection techniques must be evaluated, usually on a case-by-case basis (Gil et al., 2009).

218 In our study, NoV was not detected in any of the 911 RTE samples analysed. The results of our 219 investigation are comparable with data reported in a previous Italian study in which NoV was detected in 1/1372 (< 0.1%) fresh leafy samples but not in 1160 RTE vegetable samples (Losio et 220 221 al., 2015). Although in these relatively small-scale studies the rate of NoV contamination was low, 222 the impact of NoV contamination in fresh and RTE vegetables has been clearly outlined by the 223 2013 report of the European Food Safety Authority (European Food Safety Authority & European 224 Centre for Disease Prevention and Control, 2013) that noted an increase in the number of NoV outbreaks caused by consumption of vegetables in Europe in 2011. The settings that were most 225 often reported were restaurants, households, schools and kindergartens and canteen or workplace 226 227 catering and residential institutions, and many NoV outbreaks were traced to food that was handled by one infected worker during food preparation (European Food Safety Authority & European 228 229 Centre for Disease Prevention and Control, 2011).

Viral contamination may occur in several times and points of the production chain. As viruses are not able to replicate extracellulary, their presence can only be accounted for by pre-harvesting contaminations of vegetables or by post-harvesting re-contaminations by food handlers (Berger et al., 2010; Heaton and Jones, 2008; Koopmans et al., 2002; Wu et al., 2005). Several factors, chiefly, the epidemiological patterns of viral infections (i.e. seasonal fluctuations in their prevalence, incubation times, duration of virus shedding after the clinical signs) should also be considered and could explain, at least in part, our findings.

The samples that tested positive for HAV and HEV were mixed salads, valerian, rocket and spinach. Mixed salads, valerian and rocket are characterized by a production process that requires strong manipulation during collection, whilst spinaches pose difficulties for washing of their leaves.

Water quality used for agricultural irrigation and the hands of the workers in the field affects the microbiological characteristics of products and therefore may pose a risk to consumer health.

Appropriate implementation of food-safety management systems including Good Agricultural Practices (GAP), Good Hygiene Practices (GHP) and Good Manufacturing Practices (GMP) should be the primary objective of operators producing RTE salads. Compliance with hygiene requirements, in particular hand hygiene, is an absolute necessity for food handlers at all stages of the vegetable production and supply chain to reduce the risks of foodborne virus contamination.

247 It is possible that the results of our and of other studies were biased by the sampling procedure. The 248 EU regulations do not provide indications for the virological risks and the sampling plan adopted in 249 our study was based on similar literature (Losio et al., 2015; Serracca et al., 2012). Indeed, the need 250 for adequate sampling plans for virological investigations in food matrices has arisen in recent years along with the understanding of the risks posed by viruses in food products (Perez-Rodriguez et al., 251 252 2014). Evaluation of the microbiological contamination status can be useful to define risk-based planning for official controls, while implementation of the management practices and knowledge of 253 254 the prevalence of foodborne viruses in RTE vegetables represent a contribution for assessing 255 consumer exposure.

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## Table

# 1 **Table 1**

2 Summary of virological results (no. of positive samples) obtained, for Ready-to-Eat vegetables,

3 using the reference ISO method, in Italy.

Type of samples		Total no.	Positive for	Positive for	Positive for
		911	HAV	HEV	NoV
Mixed salads		619	14	2	0
Carrot	Daucus carota subsp. sativus	53	0	0	0
Valerian	Valeriana officinalis	40	1	0	0
Rocket	Eruca vesicaria	104	3	2	0
Spinach	Spinacia oleracea	10	0	2	0
Iceberg lettuce	Lactuca sativa	18	0	0	0
Romaine Lettuce	Lactuca sativa L var. longifolia	67	0	0	0

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