

Meta-Analysis of the Effects of Sanitizing Treatments on *Salmonella* spp., *Escherichia coli* O157:H7, and *Listeria monocytogenes* Inactivation in Fresh Produce

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The aim of this study was to perform a meta-analysis of the effects of sanitizing treatments of fresh produce on *Salmonella* spp., *Escherichia coli* O157:H7, and *Listeria monocytogenes*. From 55 primary studies found to report on such effects, 40 were selected based on specific criteria, leading to more than 1,000 data on mean log reductions of these three bacterial pathogens impairing the safety of fresh produce. Data were partitioned to build three meta-analytical models that could allow the assessment of differences in mean log reductions among pathogens, fresh produce, and sanitizers. Moderating variables assessed in the meta-analytical models included type of fresh produce, type of sanitizer, concentration, and treatment time and temperature. Further, a proposal was done to classify the sanitizers according to bactericidal efficacy by means of a meta-analytical dendrogram. The results indicated that both time and temperature significantly affected the mean log reductions of the sanitizing treatment ($P < 0.0001$). In general, sanitizer treatments led to lower mean log reductions when applied to leafy greens (for example, 0.68 log reductions [0.00 to 1.37] achieved in lettuce) compared to other, nonleafy vegetables (for example, 3.04 mean log reductions [2.32 to 3.76] obtained for carrots). Among the pathogens, *E. coli* O157:H7 was more resistant to ozone (1.6 mean log reductions), while *L. monocytogenes* and *Salmonella* presented high resistance to organic acids, such as citric acid, acetic acid, and lactic acid (~3.0 mean log reductions). With regard to the sanitizers, it has been found that slightly acidic electrolyzed water, acidified sodium chlorite, and the gaseous chlorine dioxide clustered together, indicating that they possessed the strongest bactericidal effect. The results reported seem to be an important achievement for advancing the global understanding of the effectiveness of sanitizers for microbial safety of fresh produce.

The consumption of fresh fruits and vegetables comprises an essential element of a healthy diet and a protective factor against several chronic diseases (1, 2). Even though the ingestion of these products is highly recommended by health authorities, guaranteeing fresh, safe, and high-quality fruits and vegetables remains an enormous challenge for fresh-produce industries.

In order to deliver the health benefits (2), fruits and vegetables must be safe. One of the chief concerns related to the safety of these products is their recurrent and increased association with disease outbreaks (3–6). Epidemiological investigations indicate that *Salmonella*, pathogenic *Escherichia coli* and *Listeria monocytogenes* stand out as the most important bacterial agents linked to fresh-produce disease outbreaks (3, 5–7). Recent studies have reported the occurrence and high diversity of these microorganisms in the environment or in close areas of produce farming areas (8–13).

Given the above, fresh-produce industries have been implementing measures at pre- and postharvest steps to reduce or avoid the contamination of these products and, consequently, to diminish the burden of disease outbreaks. At postharvest steps, disinfection is the critical step for reduction of microbial contamination (14, 15). During disinfection, fresh produce is allowed to stay in contact with sanitizers added to the washing tanks aiming to reduce their microbial load. Finally, yet importantly, during disinfection, washing water should not become a point of cross-contamination (15, 16). The phenomenon of cross-contamination during produce washing has been indicated as the potential cause of the spinach *E. coli* O157:H7 outbreak that resulted in 205 illnesses and three deaths in the fall of 2006 in the United States (17,

18). As disinfection of fresh produce constitutes a critical control point, several reports quantify the pathogens' concentrations in these foods before and after disinfection with different sanitizers (19–26). However, because the log reductions of pathogens attained by the sanitizers are affected by the particular conditions or study characteristics (protocols for washing, type of fruit and vegetable, whole or cut fresh produce, type of sanitizer and concentration, washing time and temperature, pathogenic strains, microbiological essays, etc.) under which the measurements were obtained, variability in the effect size reported in the primary studies is expected to occur. This variability can be observed even among studies investigating the same type of fresh produce and sanitizer. Nonetheless, by means of a *posteriori* analysis and identification of the sources of variability likely to affect the log reduction of the pathogenic flora due to disinfection, it may be possible

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to explain, at least to some extent, the differences found among the study outcomes. Most importantly, it may be realistic to build a model that can be generalized to different types of fresh produce.

Meta-analysis has been defined as a statistical analysis of a collection of analytic results for the purpose of integrating the findings from a large amount of primary studies (27). Meta-analysis allows the explanation of the divergences in the study outcomes by the codification of moderating variables representing study characteristics related to research design features, data collection procedures, type of samples, etc., aiming to reduce the between-study heterogeneity or variability (28). Through meta-analysis, it may also be possible to accurately estimate the overall outcome measure, with increased statistical power, than is possible using only a single study (29).

Considering the capabilities of the meta-analysis and the significance of *Salmonella*, pathogenic *E. coli*, and *L. monocytogenes* for the microbial safety of fresh produce and for the public health (3, 5–7), the objective of this study is 4-fold: (i) first, to compile all publicly accessible published findings on the effects of sanitizers on the mean reduction in the log counts of *Salmonella* spp., *E. coli* O157:H7, and *L. monocytogenes* on fresh produce and quantitatively summarize the outcomes by means of meta-analytical models based on mixed-effects linear regressions; (ii) second, to explain a proportion of the total between-study heterogeneity in the reduction of the pathogens' populations by incorporating available study characteristics, such as type of fresh produce, type of sanitizer, concentration, and treatment time and temperature, into the basic meta-analysis model; (iii) third, to assess possible differences in the susceptibilities of *Salmonella* spp., *E. coli* O157:H7, and *L. monocytogenes* to selected sanitizers, as well as differences in the overall microbial log reduction among fresh produce for a given disinfectant treatment; and (iv) finally, to evaluate the effectiveness of the sanitizers to reduce each of the pathogen's populations using a common disinfectant treatment and to propose a classification of sanitizers according to their bactericidal efficacies by means of a meta-analytical dendrogram. The resulting meta-analysis models have the capability to provide overall log reduction estimates for a particular pathogen when using a given sanitizer and sanitizing treatment.

MATERIALS AND METHODS

Data collection and effect size parameterization. Before commencing any meta-analysis study, the research problem must be stated and three important facets should be defined: population, intervention or treatment, and measured outcome. In this meta-analysis, the problem statement was the estimation of the overall effect of disinfecting fresh and minimally processed fruits and vegetables with aqueous and gaseous chemicals on the final microbial concentration (number of log reductions) of three pathogens (i.e., *Salmonella* spp., *L. monocytogenes*, and *E. coli* O157:H7). The population was specified as fresh produce, fruits and vegetables, prior to the sanitizing treatment, while the intervention or treatment was represented by the disinfection unit operation using aqueous or gaseous sanitizers. The measured outcome is derived from the pathogen's concentration on the fruits or vegetables before and after treatment, giving the number of log reductions.

The next step of literature identification was conducted using electronic search through Google with key terms, both in English and in Portuguese, encompassing the following: "*Salmonella*," "*Escherichia coli* O157:H7," "*Listeria monocytogenes*," "pathogens," "sanitizers," "chemicals," "solutions," "organic acids," "detergents," "washing," "inactivation," "antibacterial effect," "reduction," "fruits," "vegetables," and "produce." Electronic searches were conducted using sensible combinations of

at least three of the above-listed terms. Also, literature for inclusion in the study was identified from bibliographic databases such as PubMed, Science Direct, and Scopus using the same key words. Data included considered studies available in scientific journals and electronically from 1990 to 2014. After an initial screening for research quality, a total of 55 studies were found to report on the effect of sanitizers on the concentration of pathogens in fresh produce. However, these encompassed also studies using ultrasound and irradiation treatments, which were disregarded, as only conventional washing with sanitizers was considered for inclusion in the meta-analysis. A second criterion used in the screening was the need for the primary study to report the concentration of either of the pathogens (*Salmonella* spp., *E. coli* O157:H7, and *L. monocytogenes*) in fresh produce both before (i.e., control) and after the disinfection treatment or, alternatively, the microbial log reduction attained by the disinfection treatment. As a third criterion for inclusion, the primary study had to clearly specify the type of sanitizer and its concentration, washing or exposure time and temperature, and sample size and/or standard deviations. As a fourth criterion, an approved microbiological method for pathogen enumeration had to be employed. Considering all of those requirements, 40 primary studies were regarded as appropriate for inclusion in the present meta-analysis study (19–26, 30–62).

As a next step, a parameterization of the intervention's effect size should be determined. The effect size (θ) refers to the degree to which the hypothetical phenomenon (i.e., reduction in the concentration of pathogens on fresh produce due to disinfection treatment) is present in the population. For the studies' outcomes to be compatible and meaningful for analysis, such effect size should be converted to a common scale that permits direct comparison and summation of the primary studies. Because the data generated by the primary studies are the means of a continuous variable (i.e., microbial concentration in or on fresh produce), the possible parameters to measure effect size or treatment difference are raw (unstandardized) mean difference, standardized mean difference, and response ratios (63). The most suitable parameter to measure effect size was the raw mean difference between control and treatment means, because all the primary studies reported in the same log CFU scale and it is an intuitively meaningful parameter. That is, referring in terms of, say, a 2- or 3-log microbial reduction is a widespread use among food microbiologists.

Consider a primary study j that reports means for two groups (control, or before disinfection treatment, and treated, or after disinfection treatment). Let \bar{x}_C and \bar{x}_T be the sample means of the two independent groups; hence, the effect size estimate $\hat{\theta}$, which, in our case, is the difference in sample means or mean log reduction R , is defined as

$$\hat{\theta}_j = R_j = \bar{x}_{Cj} - \bar{x}_{Tj} \quad (1)$$

Now, let s_C and s_T be the sample standard deviations of the two groups and n_C and n_T be the sample sizes in the two groups, control and treated, respectively. If we assume that the two population standard deviations are different, then the SE of the mean log reduction R can be estimated as

$$SE(R_j) = \sqrt{\frac{s_C^2}{n_C} + \frac{s_T^2}{n_T}} \quad (2)$$

Mean log reductions (R) and their standard errors (SE) for the three pathogens were estimated from the primary studies whose results were reported separately for the control and treated groups. Nonetheless, in some primary studies, mean log reductions and their standard errors were provided as such, so none of the above formulas needed to be applied, and their values were extracted directly from tables or charts.

Description of meta-analytical data set. The microbial mean log reduction values for the three pathogens, whether estimated using equations 1 and 2 or directly extracted from the primary studies, were the outcomes of experiments carried out with a specific fresh produce under a certain disinfection treatment. Hence, all this additional information was also annotated from the primary studies in the form of study characteristics or moderating variables. The study characteristics considered

were bacteria (a categorical variable), type of sanitizer (a categorical variable), sanitizer concentration (a continuous variable), type of produce (a categorical variable), treatment or washing time (a continuous variable), and treatment or washing temperature (a continuous variable). As explained above, sample sizes (n_C and n_T) and standard deviations (s_C and s_T) of the control (preinfection) and treated (postinfection) groups were also extracted. Depending upon the sanitizer, a specific concentration unit was used in a primary study. For instance, for gaseous chlorine dioxide, the concentration was often expressed in parts per million, while for sodium chlorite, it was in grams per liter. In order to facilitate comparisons among sanitizers, all concentrations were converted to grams per 100 ml. For the 27 sanitizers recovered (namely, acetic acid [AA], acidified sodium chlorite [ASC], benzalkonium chloride [Bzc], citric acid [CA], calcined oyster shell [Ca-Oy], calcined Sakhalin surf clam [Ca-SS], calcium hypochlorite [CH], Citrox, chlorine dioxide gas [CD], dodecyl-benzenesulfonic acid [DA], sodium 2-ethylhexyl-sulfate [EHS], hydrogen peroxide [HP], lactic acid [LA], malic acid [MA], nisin, ozonized water [OW], ozone gas [Oz], pediocin, peroxyacetic acid [PAA], phytic acid [Phy], slightly acidic electrolyzed water [SAEW], sodium chlorite [SC], sodium-dodecyl sulfate [SDS], sodium hypochlorite [SH], tartaric acid [TA], trisodium phosphate [TSP], and Tsunami 100), the concentrations ranged from 0.0001 to 4.8 g/100 ml, although it is important to bear in mind that every sanitizer is associated with a specific concentration range. For instance, in the primary studies, fresh fruits and vegetables are treated with chlorine dioxide gas in concentrations from 0.00015 to 0.00030 g/100 ml, while they are washed with lactic acid in higher concentrations, from 0.003 to 2.0 g/100 ml. In addition to the 27 sanitizers identified, primary studies also provided microbial mean log reductions from washing using only water (i.e., a blank treatment). The water types were tap water (W), distilled water (DisW), and deionized water (DioW), which were categorized within sanitizers, although a solute concentration of 0 g/100 ml was assigned to all water types. Washing times were in the range of 0.15 to 180 min, yet the longer times belonged to chlorine dioxide gas treatments. Temperatures for washing were mostly ambient, although overall they were in the interval from 4 to 55°C.

To get some insight into the spread of the microbial log reduction data among the categorical study characteristics, Table 1 compiles the number of mean log reduction observations partitioned by sanitizer, pathogen, and fresh produce. It should be noticed that the meta-analytical data are highly sparse, meaning that for some sanitizers, fewer data are available. For instance, for SAEW, microbial mean log reduction observations were reported for the three pathogens, while for ozone gas, data were limited to *E. coli* O17:H7 only. Moreover, the heterogeneity in the distribution of fresh produce across pathogens and sanitizers caused further sparseness. Said otherwise, for a given sanitizer, the types of produce studied did not coincide for all pathogens. From the 27 sanitizers whose microbial mean log reduction information was available in the literature, 9 were excluded from the meta-analyses (i.e., Bzc, Ca-Oy, Ca-SS, DA, EHS, nisin, pediocin, Phy, and SC) because too few observations were presented (the threshold was set as equal to or less than four observations per sanitizer) (Table 1).

Thus, for the 18 sanitizers remaining for the meta-analyses plus the three water types (i.e., 21 sanitizers), a total of 1,025 microbial mean log reduction values were brought together. In the primary studies, those values were obtained by measuring the effects of disinfectant treatment in 30 types of fresh produce: apple, baby spinach, blueberry, broccoli, buckwheat, cabbage, cantaloupe, carrot, cherry tomato, Chinese cabbage, Chinese celery, cilantro, cucumber, daikon, green onion, honeydew, lettuce, mung bean, mung bean sprouts, onion, peach, pepper, rocket leaves, romaine lettuce, sesame leaf, spinach, spring onion, strawberry, tatsoi, and tomato. For space reasons, the meta-analytical raw data are not presented here, but they can be found in the supplemental material in Tables S1, S2, and S3, separately for *E. coli* O157:H7, *L. monocytogenes*, and *Salmonella* spp., respectively. Such tables compile the microbial mean log reductions and standard errors when available, sanitizer agents, sanitizer concentra-

tions, treatment times and temperatures, types of fresh produce, sample size, and primary studies.

The sparseness of the data has some implications in the choice and the design of the meta-analysis mixed-effect models. Because of the considerable dispersion in the number of microbial mean log reduction values among the sanitizer-pathogen combinations (Table 1), a general meta-analysis model encompassing all data could not be adjusted. Hence, separate meta-analysis studies were conducted, first on data partitioned by sanitizer—in order to make comparisons among pathogens and fresh produce—and, subsequently, on data partitioned by pathogen—in order to make comparisons among the bactericidal efficacies of sanitizers. This is explained in detail in the following subsections.

Meta-analysis models by sanitizer. When constructing separate meta-analysis models by sanitizer, it is possible to assess both whether there are differences in the resistances to the sanitizer agent among the three pathogens and whether there are differences among produce in the microbial log-reduction attained by a disinfection treatment.

Assessing differences among pathogens. To assess the bactericidal efficacy of fresh produce disinfection among pathogens, nine sanitizing agents with the least sparseness in the number of observations across pathogens were selected. These were acetic acid, acidified sodium chlorite, chlorine dioxide gas, citric acid, hydrogen peroxide, malic acid, peroxyacetic acid, slightly acidic electrolyzed water, and sodium hypochlorite. Tap water was also selected for comparison, as it can be regarded as a blank treatment for washing (i.e., washing without sanitizer).

A meta-analysis model can be considered a special case of a multilevel analysis using hierarchical linear models, with subjects between studies at the first level and studies at the second level. In a multilevel meta-analysis, one usually starts from the random-effects model, and if the between-study variance is shown to be noteworthy, study characteristic can be added to the model to account for at least part of the heterogeneity in the true effect size θ (in our case, the mean log reduction R). Thus, for each of the 10 selected sanitizers (including tap water), the microbial mean log reduction R was modeled as

$$R_{ijk} = \beta_{0i} + \beta_1 \log C + \beta_2 T + \beta_3 t + u_{jk} + \varepsilon_{ijk} \quad (3)$$

where β_0 is the fixed effect of the pathogen i , β_1 the mean effect of the increment in the logarithm of the sanitizer concentration C , β_2 the mean effect of a 1°C increment in disinfection temperature T , and β_3 the mean effect of a 1-min increment in disinfection time t . The wide range (0.00005 to 2.0 g/100 ml) of the sanitizer concentration was reduced by log transformation in order to stabilize the parameter estimation of the regression models.

Because of the sparse nature of the data structure, whereby in many cases one primary study reported results for only one or two types of fresh produce, for this meta-analysis it was not feasible either to separate the between-produce variability from the between-study variability or to build a nested covariance of primary studies within a type of fresh produce. To overcome this problem and still be able to account for the evident variability due to the different primary studies j and the different fresh produce k , both variables were merged into an interaction variable (jk). Such interaction was assumed to be the subject of variation of the intercept random effects u_{jk} placed in equation 3. The random effects u_{jk} are assumed to be normally distributed with mean zero and variance s_{jk}^2 . The errors or residuals ε_{ijk} are also assumed to follow a normal distribution with mean zero and variance s^2 . Using this model design, the estimated value of R_{ijk} represents therefore the overall mean microbial log reduction for the pathogen i attained by a particular sanitizing treatment (C , T , and t), applicable to the entire population of fresh produce and primary studies. Nonetheless, if we wished to estimate the mean microbial log reduction for a particular fresh produce, it can still be done extracting its corresponding random effect u_{jk} and replacing in equation 3.

Since primary studies are expected to differ from each other in the reliability of estimating the true effect of disinfection on the pathogens' numbers on fresh produce, for instance, due to differences in study sizes, analytical methods or experimental designs, a weighted linear mixed

TABLE 1 Microbial mean log reduction observations found in the literature according to type of fresh produce, pathogen, and sanitizer, extracted from published studies

Sanitizer ^a	Type of produce (no. of observations of mean log reduction of indicated organism[s])		
	<i>E. coli</i> O157:H7	<i>L. monocytogenes</i>	<i>Salmonella</i> spp.
AA	Baby spinach (6)		Blueberry (4), rocket leaves (8), spring onion (8)
ASC	Carrot (6), cilantro (4), tatsoi (9)	Carrot (7), cherry tomato (1), cucumber (1)	Carrot (3), cherry tomato (1), cucumber (1), tomato (6)
Bzc	Lettuce (2), tomato (2)		
CA	Baby spinach (9), cilantro (1), lettuce (20), spinach (1)	Lettuce (20), spinach (1)	Blueberry (2), lettuce (20), rocket leaves (8), spring onion (8)
Ca-Oy	Tomato (4)		
Ca-SS	Tomato (4)		
CH	Broccoli (11), lettuce (12), spinach (1)		Spinach (1)
Citrox	Lettuce (9)		
CD	Cabbage (3), cantaloupe (10), carrot (3), lettuce (6), spinach (2), strawberry (6)	Cabbage (3), cantaloupe (10), carrot (3), lettuce (6), strawberry (5)	Apple (3), cabbage (3), carrot (3), cantaloupe (9), lettuce (6), onion (3), peach (3), spinach (2), strawberry (6), tomato (29)
DA	Romaine lettuce (2)		
DioW	Baby spinach (6), broccoli (3), lettuce (6), mung bean (4), mung bean sprouts (4), romaine lettuce (2), spinach (1)	Lettuce (2), spinach (1)	Blueberry (2), lettuce (2), mung bean (4), mung bean sprouts (4), pepper (2)
DisW	Buckwheat (1), Chinese cabbage (3), lettuce (5), sesame (1), spinach (4), tomato (6), cabbage (2), apple (3), mung bean sprouts (2)	Broccoli (1), cabbage (1), mung bean sprouts (3), Chinese cabbage (1), lettuce (4), sesame leaf (1), spinach (1), cucumber (1)	Buckwheat (1), apple (3), blueberry (3), spinach (1), mung bean sprouts (2), lettuce (1), cucumber (3)
EHS	Romaine lettuce (2)		
HP	Baby spinach (9)	Cucumber (1)	Blueberry (5), cantaloupe (2), honeydew (2)
LA	Apple (2), baby spinach (9), lettuce (33), spinach (1), tomato (4)	Lettuce (20)	Apple (2), blueberry (2), lettuce (20), spinach (1)
MA	Baby spinach (9), lettuce (20)	Lettuce (20)	Lettuce (20)
Nisin		Broccoli (1), cabbage (1), mung bean sprouts (1)	
OW	Cabbage (2), Chinese cabbage (2), lettuce (2), spinach (5)	Spinach (1)	Spinach (2)
Oz	Cabbage (2), Chinese cabbage (2), lettuce (2), spinach (2)		
PAA	Carrot (4), lettuce (4), mung bean sprouts (7), rocket leaves (1), spinach (1), tomato (3)	Carrot (4), mung bean sprouts (7)	Carrot (4), green onion (4), lettuce (4), mung bean sprouts (7), spinach (1), tomato (8)
Pediocin		Broccoli (1), cabbage (1), mung bean sprouts (1)	
Phy	Tomato (4)		
SAEW	Chinese cabbage (2), lettuce (2), daikon lettuce (2), mung bean (16), mung bean sprouts (16), sesame leaf (2), spinach (4)	Chinese cabbage (2), lettuce (2), sesame leaf (2), spinach (4)	Chinese celery (1), daikon lettuce (1), lettuce (1), mung bean (16), mung bean sprouts (16)
SC	Cilantro (1)		
SDS			Blueberry (11)
SH	Baby spinach (3), cabbage (2), carrot (4), Chinese cabbage (2), Chinese celery (1), cilantro (1), daikon lettuce (1), lettuce (12), mung bean sprouts (7), romaine lettuce (4), rocket leaves (1), spinach (3), tatsoi (1), tomato (9)	Carrot (6), cherry tomato (2), cucumber (3), lettuce (2), mung bean sprouts (7), spinach (1)	Blueberry (7), carrot (6), cherry tomato (2), Chinese celery (1), cucumber (2), daikon lettuce (1), green onion (4), lettuce (7), mung bean sprouts (7), pepper (2), tomato (20)
TA	Baby spinach (6)		
TSP		Lettuce (4)	Lettuce (4), pepper (4)
Tsunami 100	Lettuce (6)		
Water	Carrot (4), daikon lettuce (1), lettuce (3), tatsoi (1), tomato (1)	Carrot (4), cucumber (1)	Cantaloupe (1), carrot (4), Chinese celery (1), daikon honeydew (1), lettuce (2), tomato (8)

^a Abbreviations: AA, acetic acid; ASC, acidified sodium chlorite; Bzc, benzalkonium chloride; CA, citric acid; Ca-Oy, calcined oyster shell; Ca-SS, calcined Sakhalin surf clam; CH, calcium hypochlorite; CD, chlorine dioxide gas; DA, dodecyl-benzenesulfonic acid; DioW, deionized water; DisW, distilled water; EHS, sodium 2-ethylhexyl-sulfate; HP, hydrogen peroxide; LA, lactic acid; MA, malic acid; OW, ozonized water; Oz, ozone gas; PAA, peroxyacetic acid; Phy, phytic acid; SAEW, slightly acidic electrolyzed water; SC, sodium chlorite; SDS, sodium dodecyl sulfate; SH, sodium hypochlorite; TA, tartaric acid; TSP, trisodium phosphate.

model was preferred, with weights representing the precision in estimating the true microbial log reduction. In meta-analysis, it is common practice to use the standard error of the effect size as a measure of precision to assign weights to each of the primary studies. However, in the present meta-analysis, it was not possible to obtain the standard errors of the log reductions *R* for all primary studies; and consequently, the precision was instead redefined as some measure proportional to the sample size *N* used in every primary study. Hence, the weight—the level of confidence on each of the measured log reductions *R*—was given by the sample size. A weighted mixed-effects linear model (equation 3) was adjusted to each of the 10 selected sanitizers.

Assessing differences among fresh produce. By partitioning data by sanitizer, it is also possible to appraise whether the same disinfection treatment would achieve variable effects depending upon the type of fresh produce. However, to carry out this assessment, we need to choose sanitizers that have been tested in a wide range of fresh produce, and we needed to ensure that these types of fresh produce are roughly the same at least across two pathogens. For instance, observing the data dispersion shown in Table 1, using the data from the gaseous chlorine dioxide (CD) is a good option because it was tested on cabbage, cantaloupe, lettuce, and strawberry for the three pathogens and tested on spinach for *E. coli* O157:H7 and *Salmonella*. Following this reasoning, the sanitizers ASC,

CD, SAEW, and SH were considered suitable for this analysis, and the following meta-analysis model was adjusted to each of the four data sets:

$$R_{ijk} = \beta_{0k} + \beta_1 \log C + \beta_2 T + \beta_3 t + u_{i(j)} + \varepsilon_{ijk} \quad (4)$$

Now, β_0 is the fixed effect of the type of fresh produce k , and $u_{i(j)}$ are the intercept random effects with subject of variation pathogen i nested in the primary study j . The nested random effects $u_{i(j)}$ are assumed to be normally distributed with means zero and variances s^2 and s^2 . With such a model design, the variability due to pathogens is extracted, and the response variable R_{ijk} can be thought of the overall mean log reduction in the entire population of pathogens, from treating a fresh produce k by a particular sanitizing treatment (C , T , and t). In a similar fashion, a weighted regression was opted for, in order to account for the differences in precision among primary studies. The sample size N was used as the weight of each log reduction observation.

Meta-analysis models by pathogen. The microbial data were also partitioned by pathogen, producing three data sets for *E. coli* O157:H7, *L. monocytogenes*, and *Salmonella* spp. Separate meta-analyses were then performed by pathogen, so as to compare the bactericidal efficacies of sanitizers for a common treatment (C , T , and t). For each of the pathogen's data sets, a mixed-effects linear model of the type

$$R_{jkl} = \beta_{0l} + \beta_1 \log C + \beta_2 T + \beta_3 t + u_{jk} + \varepsilon_{jkl} \quad (5)$$

was fitted, where β_0 now represents the fixed effect of the type of sanitizer l and u_{jk} are intercept random effects, whose subject of variation is the interaction study \times fresh produce, which account for the variability due to both the different primary studies j and the different fresh produce k . The regression models were fitted using the sample size N as the weight of each of the observations.

For each of the models explained under “Description of meta-analytical data set” and “Assessing differences among pathogens” above, the regression's basic assumption of data normality was verified: the normality of residuals was first assessed, and then the studentized residuals were examined for identifying spurious data points lower than -3.0 and higher than 3.0 (i.e., outliers). In the case that outliers were present, they were removed from the data and regression models refitted. In addition, for the regression models, a measure of between-study heterogeneity, I^2 , was calculated. The I^2 statistics, or intraclass correlation, estimates the proportion of between-study variance from the total variance. If the intraclass variance is higher than 25% of the total variance, the variance between studies can be deemed large enough to attempt to model it using available study characteristics. Measures of goodness of fit such as Akaike information criterion (AIC) and Bayesian information criterion (BIC) were also computed for each of the meta-analysis models.

The mixed-effects linear models were also used to construct meta-analytical forest plots (63), in order to allow a better visualization of the difference in the effect of a given sanitizer and disinfection treatment among fresh produce (from equation 4) and the difference among sanitizers for a given disinfection treatment (from equation 5). The weighted mixed-effects linear models were fitted in R version 2.14.2 (R Development Core Team) using the “lme” function from the “nlme” package (64). Forest plots were built using the “metafor” package (65).

Cluster analysis of sanitizers. In order to examine similarities and dissimilarities among sanitizers in their bactericidal effects, so that clusters of sanitizers could be identified and separated from others, a hierarchical cluster analysis was performed on the microbial data from selected sanitizers. Since cluster analysis is a multivariate data analysis technique—hence, it requires continuous variables as inputs—it is necessary to first obtain some measurements of the characteristics of each sanitizer in the form of a continuous variable. In a regression analysis of the type

$$R_{ijk} = \beta_0 + \beta_1 \log C + \beta_2 T + \beta_3 t + u_{jk} + \varepsilon_{ijk} \quad (6)$$

adjusted to the whole data from a given sanitizer, the parameter estimates β_0 , β_1 , β_2 , and β_3 can be thought of as the continuous variables characterizing the disinfectant capacity of the sanitizer. This is because for a sanitizer, the higher the intercept β_0 (representing the mean log reduction

R at the mean log concentration C , the mean temperature T , and the mean time t), the higher the microbial mean log reduction R . Similarly, a sanitizer with higher slopes β_1 , β_2 , and β_3 will produce a greater mean log reduction R for a given log concentration C , temperature T , and time t , respectively. Thus, for the cluster analysis, the sanitizers selected needed to be those presenting microbial log reduction observations measured over a wide range of temperatures, concentrations, and times so that precise slope estimates could be computed. Suitable sanitizers for analysis were AA, ASC, CA, CD, CH, HP, LA, MA, PAA, SAEW, and SH. The water types, W, DisW and DioW, were also included in the list of sanitizers as a mechanism for testing the performance of the clustering algorithm to build meaningful groups (said otherwise, because it is known *a priori* that the water types are not sanitizers and their bactericidal effect is the lowest of all, they should be grouped together by the clustering method chosen).

Equation 6 was then fitted to each of the sanitizers l , and the parameter estimates β_{0l} , β_{1l} , β_{2l} , and β_{3l} for $l = \{1, 2, \dots, 14\}$ were organized in an 14-by-4 matrix, where the rows corresponded to the sanitizers and the columns to the four parameter estimates. As the next step, the Euclidean distance between each pair of sanitizers was computed and arranged in a distance matrix. The clustering was performed using a hierarchical algorithm whereby the partition with the cluster in which $k = 1$ (all sanitizers are together in the same cluster) is part of the output, and also the situation with $k = j$ (each sanitizer forms a separate cluster with only a single element). In between, all values of $k = 2, 3, \dots, j - 1$ are covered in a kind of gradual transition: the only difference between $k = r$ and $k = r + 1$ is that one of the r clusters splits up in order to obtain $r + 1$ clusters (i.e., two of the $r + 1$ clusters combine to yield r clusters). The clustering method chosen was Ward's method, which is a minimum-variance method aiming at finding compact and spherical clusters (for further information on hierarchical clustering and clustering methods, see reference 66). The distance matrix was computed using the “dist” function, and the agglomerative hierarchical cluster analysis producing the dendrogram was using the “hclus” function, both from the R “stats” package.

RESULTS

Meta-analysis models by sanitizer. Table 2 shows the results from fitting equation 3 to nine sanitizers plus water ($n = 10$) studied. These are the overall mean log reductions for the specific pathogens caused by each sanitizer treatment applied to fruits and vegetables. From Table 2, it can be seen that for most sanitizers, concentration, temperature, and time have direct effects on the microbial log reduction, even though for water and ASC a quadratic effect of temperature on mean log reduction was also identified. The covariate was not included for temperature when the treatment was SAEW, as data were available only for ambient temperature. The concentration covariate was not included when treatment was water, as it had no meaning for this treatment.

Through the meta-analytical model, it was possible to find that the pathogens studied may differ in terms of their resistance depending on the sanitizers. *L. monocytogenes* presented the lower intercept, meaning that it may be more resistant to CA, PA, and ASC treatments. On the other hand, pathogenic *E. coli* seems to be more resistant to MA, CD, AA, and HP treatments (Table 2). *Salmonella* presented the lowest intercepts (higher resistance) only when the treatment was done with water, while *L. monocytogenes* and pathogenic *E. coli* presented similar resistances to this treatment (Table 2). SH had a similar impact on *L. monocytogenes* and pathogenic *E. coli* inactivation, whereas *L. monocytogenes* and *Salmonella* were equally resistant to CD. SAEW was the only sanitizer for which no differences in inactivation resistance were found for the three pathogens studied (Table 2). It should be highlighted that for some sanitizers, such as AA and HP, no data on *L. monocytogenes* inactivation was available. Therefore, for these

TABLE 2 Parameter estimates of the individual meta-analysis mixed-effects linear models by sanitizer, predicting the microbial mean log reduction in fresh produce as a function of microorganism, sanitizer concentration, and washing time and temperature^a

Sanitizer	Parameters	Mean	SE	Pr > t	AIC/BIC
AA	Predictors of R				
	<i>E. coli</i>	1.234 X	0.909	0.307	27/31
	<i>Listeria</i> ^b	—	—	—	
	<i>Salmonella</i>	1.325 Y	0.565	0.143	
	Concn	0.103	0.072	0.181	
	Temp	0.022	0.004	<0.0001	
	Time	0.029	0.007	<0.0001	
	Variances				
	s^2_u (intercept)	0.705			$I^2 = 99\%$
	s^2 (residual)	0.004			
CA	Predictors of R				
	<i>E. coli</i>	2.576 Y	0.429	<0.0001	102/121
	<i>Listeria</i>	2.337 X	0.430	<0.0001	
	<i>Salmonella</i>	2.868 Z	0.425	<0.0001	
	Concn	0.500	0.057	<0.0001	
	Temp	0.032	0.008	<0.0001	
	Time	0.043	0.003	<0.00001	
	Variances				
	s^2_u (intercept)	0.436			$I^2 = 82\%$
	s^2 (residual)	0.095			
MA	Predictors of R				
	<i>E. coli</i>	4.223 X	0.207	<0.0001	5/18
	<i>Listeria</i>	4.538 Z	0.212	<0.0001	
	<i>Salmonella</i>	4.444 Y	0.207	<0.0001	
	Concn	0.692	0.041	<0.0001	
	Temp ^c	—	—	—	
	Time	0.051	0.003	<0.0001	
	Variances				
	s^2_u (intercept)	0.0007			$I^2 = 12\%$
	s^2 (residual)	0.0053			
SAEW	Predictors of R				
	<i>E. coli</i>	11.34 X	0.340	<.00001	32/49
	<i>Listeria</i>	11.45 X	0.359	<.00001	
	<i>Salmonella</i>	11.31 X	0.341	<.00001	
	Concn	1.573	0.039	<.00001	
	Temp ^c	—	—	—	
	Time	0.019	0.005	<.00001	
	Variances				
	s^2_u (intercept)	0.591			$I^2 = 94\%$
	s^2 (residual)	0.037			
PAA	Predictors of R				
	<i>E. coli</i>	3.843 Z	0.655	<.00001	62/74
	<i>Listeria</i>	3.295 X	0.658	<.00001	
	<i>Salmonella</i>	3.535 Y	0.653	<.00001	
	Concn	0.491	0.116	<.00001	
	Temp	NS	—	—	
	Time	0.057	0.030	0.067	
	Variances				
	s^2_u (intercept)	0.568			$I^2 = 89\%$
	s^2 (residual)	0.067			
SH	Predictors of R				
	<i>E. coli</i>	3.143 X	0.712	<.00001	289/307
	<i>Listeria</i>	3.233 X	0.719	<.00001	
	<i>Salmonella</i>	3.861 Y	0.719	<.00001	
	Concn	0.383	0.148	0.012	
	Temp	NS	—	—	
	Time	0.097	0.047	0.043	
	Variances				
	s^2_u (intercept)	1.040			$I^2 = 94\%$
	s^2 (residual)	0.061			

TABLE 2 (Continued)

Sanitizer	Parameters	Mean	SE	Pr > t	AIC/BIC
CD	Predictors of R				
	<i>E. coli</i>	8.450 X	1.085	<0.0001	359/378
	<i>Listeria</i>	8.914 Y	1.084	<0.0001	
	<i>Salmonella</i>	8.855 Y	1.066	<0.0001	
	Concn	0.854	0.115	<0.0001	
	Temp	NS	—	—	
	Time	0.054	0.010	<0.0001	
	Variances				
	s^2_u (intercept)	1.413			$I^2 = 92\%$
	s^2 (residual)	0.119			
Water	Predictors of R				
	<i>E. coli</i>	1.765 Y	0.318	<0.0001	27/40
	<i>Listeria</i>	1.754 Y	0.331	<0.0001	
	<i>Salmonella</i>	1.653 X	0.322	<0.0001	
	Concn ^d	—	—	—	
	Temp	-0.080	0.022	0.001	
	Temp ^c	0.001	0.000	<0.0001	
	Time	-0.003	0.005	0.600	
	Variances				
	s^2_u (intercept)	0.181			$I^2 = 98\%$
s^2 (residual)	0.003				
ASC	Predictors of R				
	<i>E. coli</i>	2.053 Y	0.525	<0.0001	88/100
	<i>Listeria</i>	1.625 X	0.628	0.015	
	<i>Salmonella</i>	2.686 Z	0.682	<0.0001	
	Concn	0.493	0.119	<0.0001	
	Temp	0.472	0.059	0.001	
	Temp ^c	-0.011	0.002	0.001	
	Time	-0.254	0.046	<0.0001	
	Variances				
	s^2_u (intercept)	0.052			$I^2 = 48\%$
s^2 (residual)	0.056				
HP	Predictors of R				
	<i>E. coli</i>	1.739 X	0.751	0.146	8/14
	<i>Listeria</i> ^b	—	—	—	
	<i>Salmonella</i>	2.906 Y	0.601	0.040	
	Concn	0.348	0.137	0.027	
	Temp	0.031	0.007	0.001	
	Time	0.059	0.043	0.197	
	Variances				
	s^2_u (intercept)	0.052			$I^2 = 66\%$
	s^2 (residual)	0.026			

^a Different letters (X, Y, and Z) indicate significant differences in log reduction among microorganisms. NS, not significant. —, ●●.

^b No data for *L. monocytogenes* were available.

^c As data were available only for ambient temperature, a temperature covariate could not be included.

^d The concentration covariate was not included as it has no meaning for water.

sanitizers, only *Salmonella* and pathogenic *E. coli* were considered, with the latter being more resistant than the former (Table 2).

The I^2 intraclass correlation values obtained were generally >48%, except for the treatment with MA (12%) (Table 2), which suggest that for most sanitizers, there may be other moderating variables explaining the remaining between-study variability that were not codified in the present meta-analysis study. Despite the above, for each of the sanitizers, a reasonable agreement was shown between the observed mean log reduction values extracted from the primary studies, and those fitted by the models from Table 2, with coefficients of correlation ranging from 0.842 to 0.986 (Fig. 1).

Taking into account that the inactivation of *L. monocytogenes*, *Salmonella*, and pathogenic *E. coli* by four sanitizers (ASC, CD,

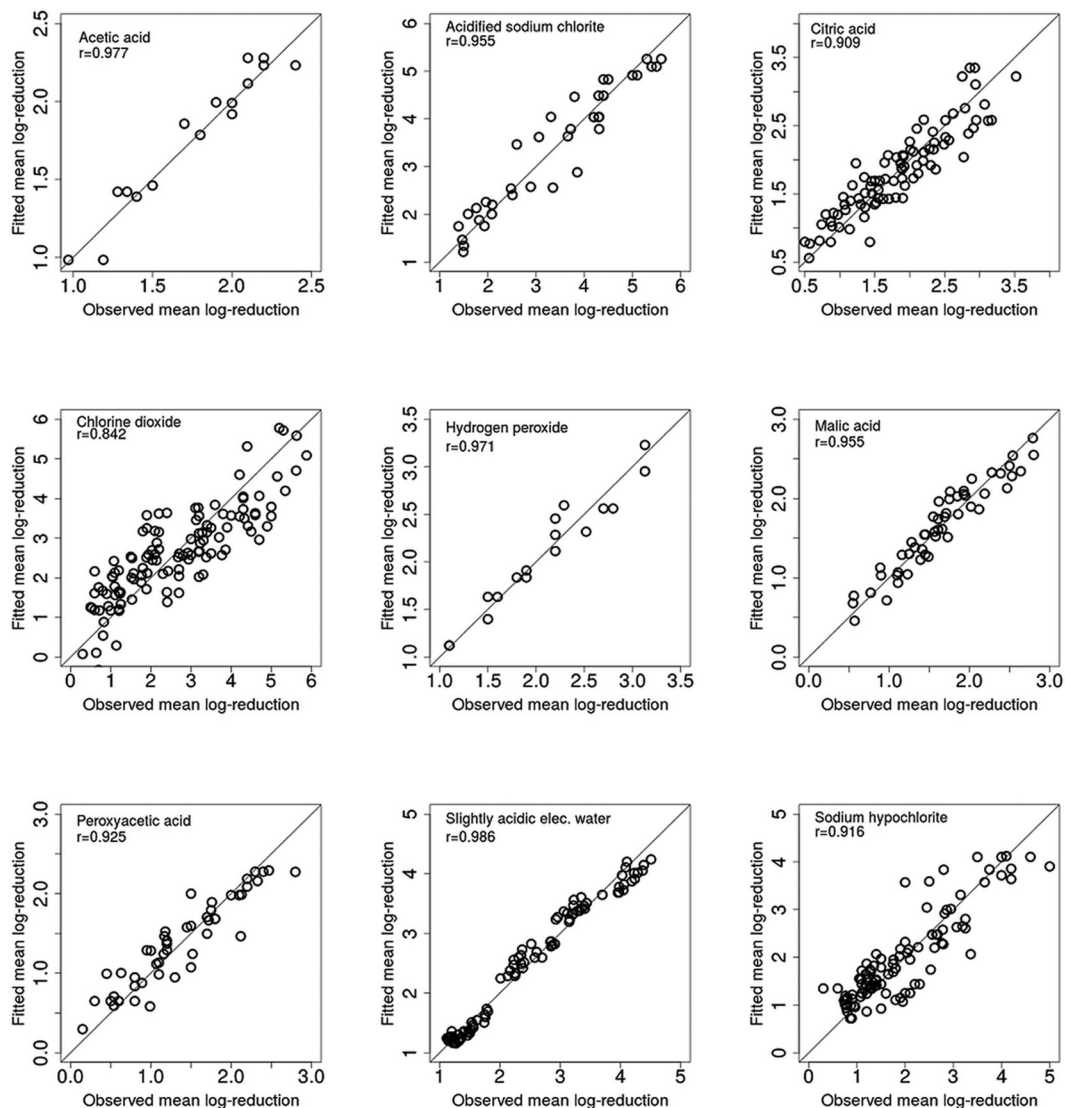


FIG 1 Scatter plots of mean microbial log reduction values fitted by the independent meta-analysis linear mixed models by sanitizer (from Table 2) in comparison with the observed data.

SAEW, and SH) was assessed in at least four different types of produce (Table 1), equation 4 has been used to assess whether the inactivation effectiveness of the same washing treatment would be affected by the type of fresh produce. Forest plots were constructed for each of the four sanitizing treatments using realistic sanitizer concentration and washing time as shown in Fig. 2, 3, 4, and 5. Also, it should be highlighted that these forest plots were constructed based on the meta-analytical model (equation 4) fitted to ASC, CD, SAEW, and SH. For example, data on *L. monocytogenes*, *Salmonella*, and pathogenic *E. coli* inactivation by ASC were available only for six types of fresh produce, namely, cucumber, cherry tomato, tatsoi, cilantro, tomato, and carrots (Fig. 2). Based on the model represented by equation 4, it can be seen that different mean log reductions were achieved according to the type of fresh produce studied. When ASC was the sanitizer used, mean log reductions varied from 1.68 for cucumber to 5.38 for carrots (Fig. 2), while log reductions varied from 0.68 to 3.61, 2.04 to 3.68, and 0.91 to 3.38 for CD, SAEW, and SH treatments, respectively

(Fig. 3 to 5). The data obtained suggest, in general, that sanitizing treatments were less effective (achieved lower log reductions of pathogens) when applied in leafy vegetables than for other fresh produce (Fig. 2, 3, 4, and 5).

Meta-analysis models by pathogen. A further approach taken was to build separate meta-analytical inactivation models for each of the pathogens studied, i.e., *L. monocytogenes*, *Salmonella*, and pathogenic *E. coli*. This was done with the aim to comparing the bactericidal effects of sanitizers for a common treatment (C , T , and t). Tables 3, 4 and 5 show the parameter estimates obtained using equation 5 fitted to each of the pathogens predicting their log reductions. The I^2 values were $>60\%$ for the three models predicting the inactivation of *E. coli* O157:H7, *L. monocytogenes*, and *Salmonella*, suggesting significant remaining heterogeneity in the outcomes from the primary studies (Tables 3, 4, and 5). Considering that the meta-analysis models were fitted by pathogen, it is now possible to see the differences among sanitizers and rank them from the lowest to the highest effects on pathogens inactiva-

F2-4
F5

T3-5

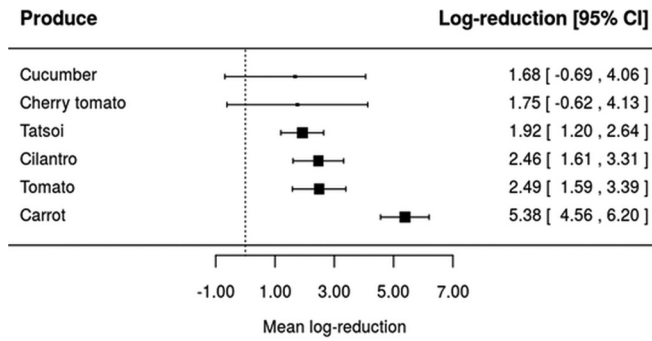


FIG 2 Forest plot of the overall mean log reduction of pathogens (*E. coli* O157:H7, *L. monocytogenes*, and *Salmonella* spp.) on different fresh produce achieved by sanitizing washing with 0.04 g/100 ml of acidified sodium chloride (ASC) at a time and temperature of 3 min and 25°C. CI, confidence interval.

tion (log reduction) (Tables 3, 4, and 5). It should be underlined that the sanitizers listed in these tables are not the same for the three pathogens because of the data sparseness. Totals of 15, 12, and 8 different sanitizers were used for *E. coli* O157:H7, *L. monocytogenes*, and *Salmonella*, respectively. Despite this, it was possible to find some similarity regarding the log reductions caused by the sanitizers over the three pathogens studied. Comparing the sanitizers' intercept values, SAEW, ASC, and CD appeared as the most effective sanitizers against *E. coli* O157:H7, *L. monocytogenes*, and *Salmonella* (Tables 3, 4, and 5). On the other hand, Oz, HP, and AA, CA, SH, and LA caused lower log reductions in *E. coli* O157:H7 and *Salmonella* (Tables 3 and 5), while, AA, LA, and SDS caused lower log reductions in *L. monocytogenes* (Table 4). The fitted intercepts presented in Tables 3, 4, and 5 also suggest that *E. coli* O157:H7 seems to be more resistant to the most effective sanitizers. While SAEW, ASC, and CD caused 3.4, 5.1, and 3.6 mean log reductions in *E. coli* O157:H7 (Table 3), the numbers of log reductions caused in *L. monocytogenes* and *Salmonella* were 6.9, 8.0, and 7.0 and 5.1, 5.4, and 7.0, respectively (Tables 4 and 5). Among these three sanitizers, ASC and CD were more effective against *L. monocytogenes* and *Salmonella*, while SAEW was the less effective against *E. coli* O157:H7 (Tables 3, 4, and 5).

A way to visualize the effect of type of sanitizer on microbial log reduction is through the construction of forest plots. To this end,

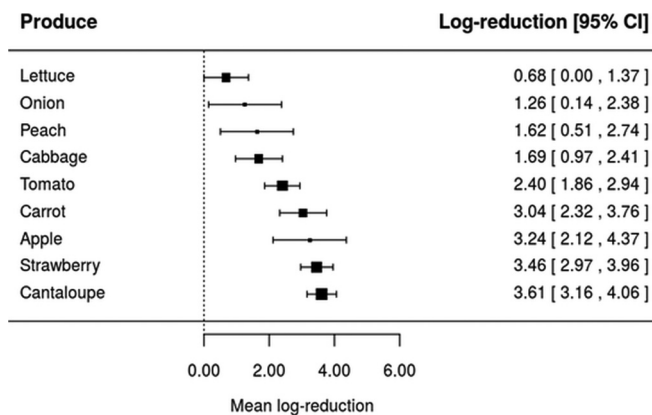


FIG 3 Forest plot of the overall mean log reduction of pathogens (*E. coli* O157:H7, *L. monocytogenes*, and *Salmonella* spp.) on different fresh produce achieved by sanitizing treatment with 0.00033 g/100 ml of gaseous chlorine dioxide (CD) at a time and temperature of 10 min and 25°C.

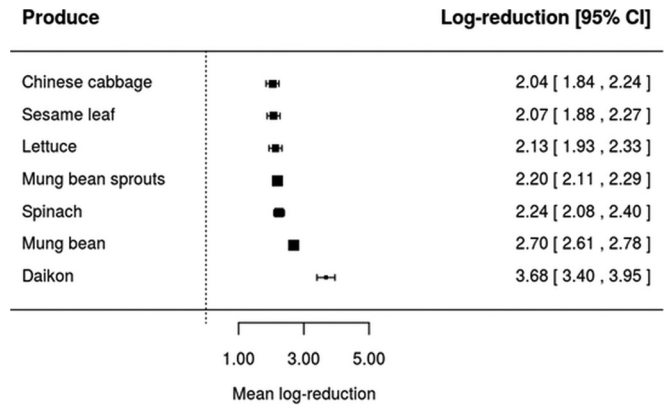


FIG 4 Forest plot of the overall mean log reduction of pathogens (*E. coli* O157:H7, *L. monocytogenes*, and *Salmonella* spp.) on different fresh produce achieved by sanitizing washing with 0.005 g/100 ml of slightly acidic electrolyzed water (SAEW) at a time and temperature of 3 min and 25°C.

the fitted meta-analysis models from Tables 3, 4, and 5 were solved for a hypothetical treatment with sanitizer concentration of 0.001 g/100 ml and exposure time of 3 min at ambient temperature in order to predict the mean log reductions of *E. coli* O157:H7, *L. monocytogenes*, and *Salmonella*. These predicted values are illustrated as forest plots in Fig. 6, 7, and 8. Under the hypothetical treatment conditions, it was found that Oz and ASC resulted in the lowest (0.14 [-0.50, 0.78]) and highest (3.86 [3.24, 4.49]) log reductions of *E. coli* O157:H7, respectively (Fig. 6). For *L. monocytogenes*, the lowest and highest log reductions would be obtained with the use of CA (0.37 [-0.77, 1.50]) and ASC (2.47 [1.41, 3.53]) as sanitizers (Fig. 7), while for *Salmonella*, AA and ASC led to the lowest (0.49 [-0.58, 1.57]) and highest (4.40 [3.40, 5.40]) log reductions, respectively (Fig. 8).

Cluster analysis of sanitizers. Through a hierarchical clustering analysis, the sanitizers were grouped in four clusters according to their bactericidal efficacies. The meta-analytical dendrogram of sanitizers is shown in Fig. 9. Through this approach it was possible to find four groups: waters (blanks) and groups with low bacteri-

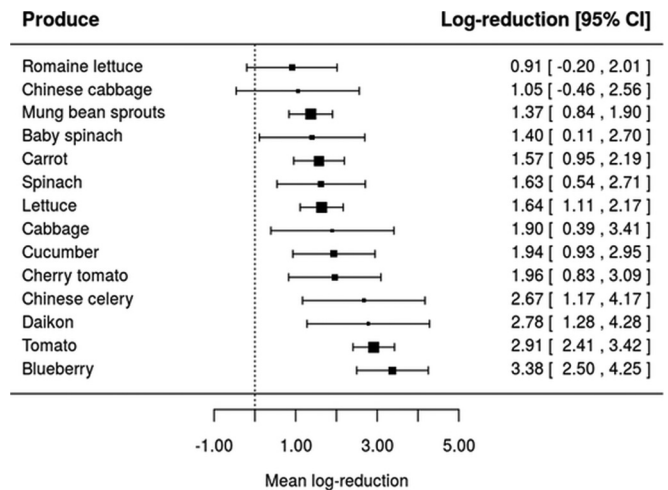


FIG 5 Forest plot of the overall mean log reduction of pathogens (*E. coli* O157:H7, *L. monocytogenes*, and *Salmonella* spp.) on different fresh produce achieved by sanitizing washing with 0.012 g/100 ml of sodium hypochlorite (SH) at a time and temperature of 3 min and 25°C.

TABLE 3 Parameter estimates of the meta-analysis model predicting the mean log reduction of *E. coli* O157:H7 in fresh produce as a function of sanitizer, sanitizer concentration, and washing time and temperature

Parameter	Mean ^a	SE	Pr > t	AIC/BIC
Predictors of R				
Sanitizer				
AA	2.094 B	0.535	<0.0001	742/816
ASC	5.103 G	0.419	<0.0001	
CA	2.362 C	0.440	<0.0001	
CH	3.192 E	0.526	<0.0001	
CD	3.555 F	0.615	<0.0001	
HP	1.987 B	0.507	<0.0001	
LA	2.537 D	0.425	<0.0001	
MA	2.429 CD	0.445	<0.0001	
OW	2.060 B	0.436	<0.0001	
Oz	1.628 A	0.490	0.001	
PAA	3.167 E	0.441	<0.0001	
SAEW	3.455 F	0.485	<0.0001	
SH	2.677 D	0.393	<0.0001	
TA	2.369 C	0.535	<0.0001	
Tsunami 100	2.976 E	0.472	<0.0001	
Log concn	0.367	0.053	<0.0001	
Temp	0.019	0.009	0.049	
Time	0.071	0.007	<0.0001	
Variations				
s^2_u (intercept)	0.689			$I^2 = 64\%$
s^2 (residual)	0.392			

^a Different letters indicate that sanitizers have significantly different effects.

cidal efficacy, medium bactericidal efficacy, and high bactericidal efficacy.

DISCUSSION

Microbial safety is a major concern for fresh produce industry because of the recurrent implication of fruits and vegetables in foodborne disease outbreaks (3, 5, 6). The use of sanitizers during

TABLE 4 Parameter estimates of the meta-analysis model predicting the mean log reduction of *L. monocytogenes* in fresh produce as a function of sanitizer, sanitizer concentration, and washing time and temperature

Parameter	Mean ^a	SE	Pr > t	AIC/BIC	
Predictors of R					
Sanitizer					
ASC	5.442 F	0.493	<0.0001	327/361	
CA	3.282 A	0.696	<0.0001		
CD	7.058 G	0.857	<0.0001		
LA	3.620 B	0.711	<0.0001		
MA	4.218 D	0.711	<0.0001		
PAA	3.980 C	0.622	<0.0001		
SAEW	5.147 E	0.678	<0.0001		
SH	3.594 B	0.502	<0.0001		
Log concn	0.617	0.091	<0.0001		
Temp	ND				
Time	0.081	0.008	<0.0001		
Variations					
s^2_u (intercept)	0.613				$I^2 = 60\%$
s^2 (residual)	0.397				

^a Different letters indicate that sanitizers have significantly different effects. ND, not determined. Temperature effect could not be estimated, as log reduction values for *L. monocytogenes* were mostly obtained at ambient temperature (mean = 21.0°C).

TABLE 5 Parameter estimates of the meta-analysis model predicting the mean log reduction of *Salmonella* spp. in fresh produce as a function of sanitizer, sanitizer concentration, and washing time and temperature

Parameter	Mean ^a	SE	Pr > t	AIC/BIC	
Predictors of R					
Sanitizer					
AA	3.622 A	0.504	<0.0001	850/908	
ASC	8.095 G	0.517	<0.0001		
CA	4.010 B	0.500	<0.0001		
CD	8.225 G	0.672	<0.0001		
HP	4.451 C	0.553	<0.0001		
LA	3.797 A	0.515	<0.0001		
MA	4.000 B	0.532	<0.0001		
PAA	6.675 EF	0.504	<0.0001		
SAEW	6.901 F	0.755	<0.0001		
SDS	3.779 A	0.582	<0.0001		
SH	5.980 D	0.436	<0.0001		
TSP	6.315 E	0.535	<0.0001		
Log concn	0.771	0.070	<0.0001		
Temp	ND				
Time	0.049	0.007	<0.0001		
Variations					
s^2_u (intercept)	1.269			$I^2 = 80\%$	
s^2 (residual)	0.313				

^a Different letters indicate that sanitizers have significantly-different effects. ND, not determined. Temperature effect could not be estimated, as log reduction values for *Salmonella* spp. were mostly obtained at ambient temperature (mean = 22.5°C).

the washing step comprises the main measure aiming to safeguard the safety of fruits and vegetables at postharvest steps. It is recognized that the effectiveness of washing procedures applied during processing of ready-to-eat fruits and vegetables is affected by several factors, such as washing conditions (temperature, time, water circulation, etc.), type of produce (whole, pieces, leafy, etc.) and sanitizers (chemical principle, concentration, etc.) (15). In view of this, models predicting the global effectiveness of sanitizers used in washing treatments of minimally processed vegetables are not available in the literature. In order to contribute to the field, in this

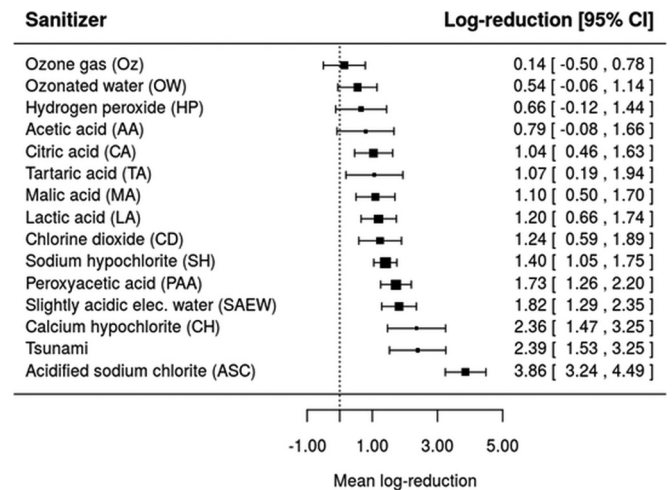


FIG 6 Forest plot of the overall mean log reduction of *E. coli* O157:H7 on a population of fresh fruits and vegetables that would be achieved by different sanitizers using a common hypothetical treatment at a concentration of 0.001 g/100 ml and a washing or exposure time and temperature of 3 min and 25°C.

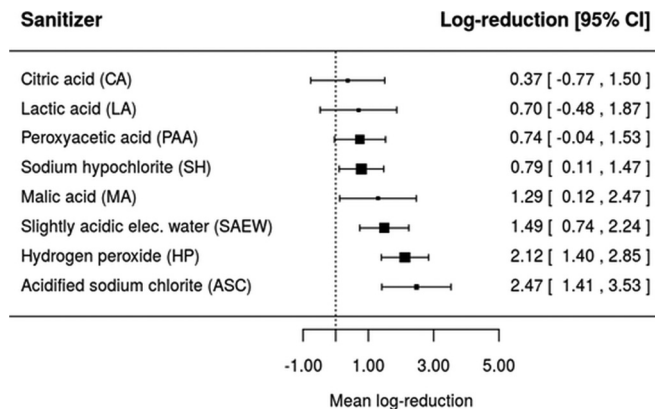


FIG 7 Forest plot of the overall mean log reduction of *L. monocytogenes* on a population of fresh fruits and vegetables that would be achieved by different sanitizers using a common hypothetical treatment at a concentration of 0.001 g/100 ml and a washing or exposure time and temperature of 3 min and 21.0°C.

study we applied a meta-analysis approach to assess *E. coli* O157:H7, *L. monocytogenes*, and *Salmonella* inactivation by sanitizers during washing of fresh produce. We were able to collect data on the microbial log reduction achieved by the sanitizers used during washing treatment of minimally processed fruits and vegetables from 55 primary studies. This first resulted in data on inactivation of *E. coli* O157:H7, *L. monocytogenes*, and *Salmonella* by 27 sanitizers. These records were further refined, leading to data for 18 sanitizers, 30 types of fruits and vegetables, and 1,025 microbial log reduction values (Table 1).

Meta-analysis models by sanitizer. Our first approach in this study was to construct a meta-analysis model by sanitizer, which allowed us to compare the effectiveness of treatments among pathogens and fruits and vegetables (Table 2). From Table 2 it can be seen that log reduction for pathogens was affected by the sanitizer concentration, washing water temperature, and increase in time. This is particularly true for treatments with AA, CA, water, ASC, and HP, but the opposite was found for treatments with SH and CD (Table 2). The higher sensitivity of chlorine-based solu-

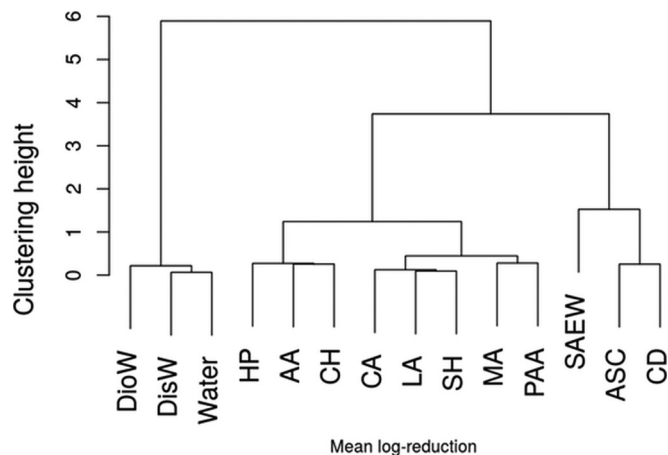


FIG 9 Dendrogram of sanitizers clustered hierarchically showing four main groups according to bactericidal efficacy.

tions to increase of washing water temperature is well known and is deemed one of the major limitations for a wider application of these compounds during sanitation of fresh produce (14, 15, 67).

A major finding of the meta-analytical model by sanitizer (equation 3) is that we found the susceptibility of the three pathogens studied to the treatments (Table 2). The pathogens tend to be more resistant when lower intercepts are obtained in a specific treatment, while when higher intercepts are attained the pathogens tend to be less resistant. The fact that *E. coli* O157:H7 and *L. monocytogenes* presented the lower intercepts for treatments such AA, CA, MA, SAEW, PAA, SH, CD, ASC, and HP indicates that these pathogens are the most resistant to sanitizing treatments applied during washing of fruits and vegetables (Table 2). The higher global resistance of *E. coli* O157:H7 to sanitizers widely used by the fresh-produce industry, such as SH and CD, can provide further insights on the reasons why this pathogen is commonly involved in fresh-produce disease outbreaks (68). Despite the fact that *L. monocytogenes* presented similar resistances to almost all the sanitizers deemed also ineffective against *E. coli* O157:H7 (Table 2), it is known that the former pathogen is more susceptible to inhibition in vegetables (69, 70). *Salmonella* was found to be more resistant than *E. coli* O157:H7 and *L. monocytogenes* only when water was assessed as a washing treatment (Table 2). Nonetheless, as we used water (tap water [W], distilled water [DisW], and deionized water [DioW]) as blanks (concentration of 0 g/100 ml), the inactivation effect, even low, is due to factors such as temperature and time. If one considers the application of water at increasing time and temperature (Table 1), the inactivation of *Salmonella* and other pathogens will be higher. SAEW seemed to be the most effective sanitizer for the inactivation of the three pathogens studied (Table 2). The fact that temperature and concentration covariates were not included in the meta-analysis model for some sanitizing treatments indicates the need for these data to be generated. This will further allow the improvement of meta-analysis predictions (equation 3) (Table 2).

Despite the fact that data were gathered from different primary studies (Table 1), scatter plots show that there is still a reasonable agreement between observed and predicted values for each of the sanitizers (Fig. 1). The fitted versus observed plots highlight how good the meta-analysis models represented the data. This is a great

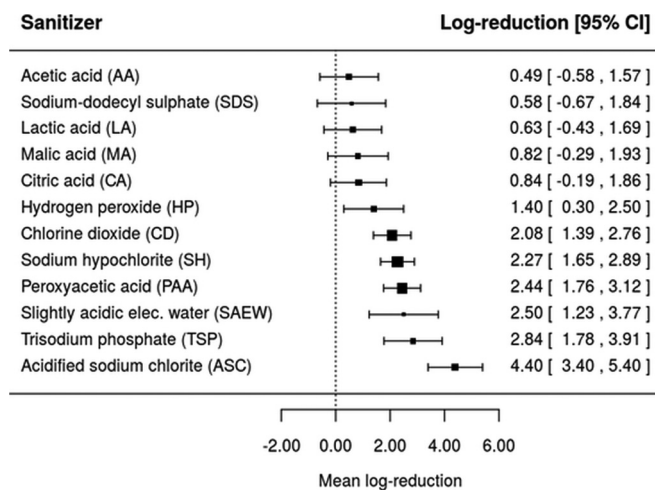


FIG 8 Forest plot of the overall mean log reduction of *Salmonella* spp. on a population of fresh fruits and vegetables that would be achieved by different sanitizers using a common hypothetical treatment at a concentration of 0.001 g/100 ml and a washing or exposure time and temperature of 3 min and 22.5°C.

accomplishment of this study, as we were able to combine data from 40 primary studies (Table 1).

Further, when the impact of a realistic sanitizing treatment (equal sanitizer concentration and washing time) was assessed, we found that the same sanitizing treatment would achieve a different log reduction, which is dependent upon the type of produce (Fig. 2 to 5). The fact that sanitizing treatments applied to leafy greens seemed to result in lower log reductions than for other types of produce (for instance, carrot, tomato, and cantaloupe,) might be related to the physicochemical nature of leafy green surfaces (71). Besides, when leafy vegetables are diced, chopped, or shredded, plant tissue damage takes place, which may result in their increased attachment (72, 73) and even growth in vegetable tissues (74). Another possible reason for the differences obtained between the inactivation of pathogens during washing of leafy greens and other produce, such as carrot, tomato, and cantaloupe, could be related to the different mechanics during washing (brushing can be applied) or even the physicochemical nature of plant surfaces (71, 75, 76).

Meta-analysis models by pathogen. Data partitioned by pathogen were used to make comparisons among the bactericidal efficacy of sanitizers (Tables 3 to 5). It should be highlighted that sanitizers listed in these tables are not the same for the three pathogens considering these data were not available. Thus, because of data sparseness, we fitted the model to data available. The fact that more sanitizing treatment data ($n = 15$) on the inactivation of *E. coli* O157:H7 during fresh-produce washing were available, followed by their effects on *Salmonella* ($n = 12$) and *L. monocytogenes* ($n = 8$), can reflect the relative concern of commodity-pathogen combinations. It is noteworthy that *E. coli* O157:H7 and *Salmonella* are the pathogens most frequently associated with foodborne disease outbreaks linked to fresh produce (3, 5, 6, 68). Therefore, one would expect to find more sanitizer options to be applied during washing aiming to ensure effective disinfection of fresh produce and safeguard public health.

From Tables 3 to 5, it can be seen that weak organic acids, such as CA, AA, and LA, presented the lowest effect on microbial log reduction. The antimicrobial efficiency of weak organic acids is highly dependent on the pH of the final solution applied for fresh produce disinfection, as pH affects the concentration of undissociated acid formed (77, 78). Moreover, it is known that the antimicrobial activity of organic acids is highly dependent on the type of acid (77, 79). This limitation, plus the fact that depending on the organic acid, there might be impact on food taste and flavor and that high BOD and COD values may be found in wastewater, will certainly limit their application in washing water of fresh produce industry (78).

SH was another class of sanitizers that appeared among those compounds with the lowest effects on microbial log reduction (Tables 4 and 5). SH is a highly used chemical principle for sanitization of fresh produce, given its high cost benefit (14, 15), despite the fact that SH solutions are highly affected by the organic matter concentration and pH of the washing water (14, 15, 78). Another weakness of SH application as a sanitizer for fresh produce is the concern with the formation of compounds with potentially carcinogenic or mutagenic effects, such as chloramines and trihalomethanes (14, 80, 81). Because of these risks, the use of SH for fresh-produce sanitation has been prohibited in some parts of the world, such as Europe (15, 77, 78). From the data presented in Tables 3 to 5, it becomes clear that SH presented a higher effect on

microbial log reductions only for *E. coli* O157:H7 (Table 3). Nonetheless, this should be carefully interpreted, as the mean effects of all sanitizers tested against *E. coli* O157:H7 were lower than those found for *Salmonella* and *L. monocytogenes* (Tables 4 and 5). This may reinforce the hypothesis that *E. coli* O157:H7 presents an intrinsic higher resistance to sanitizing agents commonly used for fresh-produce sanitation.

In contrast to SH and organic acid sanitizers, SAEW, ASC, and PAA were found to be the mostly highly efficient chemicals in reducing microbial contamination during washing of fresh produce (Tables 3 to 5). PAA is a chemical successfully used in sanitation of equipment used in food industry (82, 83). PAA can be applied in a wide range of temperature and water physicochemical parameters (including pH and calcium and magnesium contents), presence of organic matter (15, 77, 78). On the other hand, SAEW is deemed highly effective in sanitizing and less inexpensive, with ease of application and of handling (84). Nonetheless, SAEW has some limitations concerning equipment corrosion and low stability of the antimicrobial solution (15, 77, 78, 85, 86). ASC was approved for application in fresh produce sanitation 15 years ago (87). It has been proved to be a highly efficient antimicrobial treatment when applied in the range of 0.5 to 1.2 g liter⁻¹ (50, 88). Nonetheless, ASC has been found to cause physiological damages in fresh produce even when used at concentrations to 1.2 g liter⁻¹, allowed by the FDA (31, 89).

Although data presented in Tables 3 to 5 already suggest the range of efficiencies of the assessed sanitizing treatments over the three pathogens studied, we further established a hypothetical treatment (0.001 g/100 ml; washing time and temperature of 3 min and 25°C, respectively) to be able to visualize, through forest plots, the log reductions caused by each sanitizer for each pathogen. As seen in Fig. 6 to 8, Oz, CA, and AA would cause the lowest log reductions on *E. coli* O157:H7, *L. monocytogenes*, and *Salmonella*, respectively. Oz effects on the microbial log reductions for *E. coli* O157:H7 were found to be lower than those of the other sanitizers (Table 3). The antimicrobial efficacy of Oz is known to be highly influenced by the level of O₃ soluble in the washing water, contact time, water agitation, water pH, and organic matter content (78, 90–92). Although Oz has been reported as a highly antimicrobial agent for fresh-produce washing applications (15), its application in high concentrations (>1 ppm) is not feasible because of likely damages prone to be caused in fresh produce as well as the corrosion potential of equipment (77, 78). This can reinforce that the antimicrobial effectiveness of these compounds seems to be highly dependent upon factors such as time and temperature (78).

On the other hand, ASC was consistently the most effective sanitizer for the three pathogens studied (Fig. 6 to 8). Nonetheless, these findings should be interpreted with care because the rankings given in Fig. 6 to 8 were created for a constant sanitizer concentration (0.001 g/100 ml; washing time and temperature of 3 min and 25°C), when in fact each sanitizer operates at a recommended and proper concentration. These rankings are useful to illustrate the power of sanitizers, but in practice, the use of some of these sanitizers may require specific time and temperature conditions and specific concentrations. For example, it is known that chlorine-based sanitizers have a range of increased antimicrobial activities and that above a certain pH, the increase in chlorine concentration will not result in any further gain from the antimicrobial point of view (14).

Cluster analysis of sanitizers. A further assessment of the meta-analysis models was the use of cluster analysis to group the sanitizers according to bactericidal efficacy by means of hierarchical clustering analysis (Fig. 9). This is a better approach than the previous forest plots (Fig. 6, 7, and 8), because the clustering method instead takes into account the slope of the sanitizer concentration (equation 6) and implicitly considers the specific range of concentrations at which each sanitizer operates (*viz.*, the sanitizer concentration is not assumed to be constant for all the chemicals). Moreover, the clustering method combines the log reduction data for all three pathogens. Because the clustering distance (Fig. 9) is calculated from the variables β_0 , β_1 , β_2 , and β_3 , which characterize the disinfectant capacity of the sanitizers, the elements will be grouped together based on similar antimicrobial activity. From Fig. 9, four clusters can be identified. All the waters (DioW, DisW, and W) have been grouped together, indicating that their bactericidal power is comparable and the lowest of all. A second group with slightly higher bactericidal efficacy is that formed by HP, AA, and CH (for the concentrations recommended for fresh produce washing), and we can label the group as having low bactericidal efficacy. A second category, medium bactericidal efficacy, is given by the organic acids CA, LA, MA, and PAA and the inorganic SH. Although SH apparently should have a stronger bactericidal effect, it is grouped with the organic acids because for the low concentrations allowed for produce washing, its effect is comparable to that of the organic acids. The fourth group can be labeled as having high bactericidal efficacy and includes SAEW, ASC and the gaseous CD. While ASC and CD have comparable bactericidal effects, SAEW has the highest effect of all (Fig. 9).

Conclusions. Through a meta-analysis approach, we were able to assess more than 1,000 data on log reduction of the three main bacterial pathogens impairing the safety of fresh produce. We were able to build predictive models by sanitizer and by pathogen. The study is the first to gather data from a great number of papers ($n = 40$) and packed in such a way that the outputs could be compared. The absence of such a comparison has been cited as one of the major limitations of the works in this field (15). Furthermore, through the hierarchical clustering analysis performed, we were able to classify sanitizers by their bactericidal efficacies.

The findings of this study can be seen as an achievement of very practical relevance, as they can serve regulators to rank sanitizers based on their antimicrobial efficiencies. For example, depending on pathogen of greatest concern in a specific produce item, a sanitizer with the highest bactericidal power could be suggested as preferential for use. Altogether, the outcomes of the present study can serve as scientific information for decision-making (risk-benefit analysis). Regulations can be further harmonized and developed taking into account the findings reported herein.

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