1	Running head: pathogen growth on cheese
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3	Growth of Listeria monocytogenes, Salmonella spp., Escherichia coli O157:H7, and
4	Staphylococcus aureus on Cheese during Extended Storage at 25°C
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20 ABSTRACT

A potentially hazardous food (PHF) requires time/temperature control to maintain safety. 21 The US Food and Drug Administration would classify most cheeses as PHF based on pH and a_w 22 and a product assessment would be required to evaluate safety for >6 h storage at 21°C. We 23 tested the ability of 67 market cheeses to support growth of *Listeria monocytogenes* (LM), 24 Salmonella spp. (SALM), Escherichia coli O157:H7 (EC), and Staphylococcus aureus (SA) over 25 15-day storage at 25°C. Hard (Asiago and Cheddar), semi-hard (Colby and Havarti), and soft 26 27 cheeses (Mozzarella and Mexican-style) were among types tested, and included some reducedsodium and reduced-fat types. Single-pathogen cocktails were prepared and individually 28 inoculated onto cheese slices ($\sim 10^5$ CFU/g). Cocktails were comprised of 10 strains of LM, six of 29 SALM, or five of EC or SA. Inoculated slices were vacuum packaged and stored at 25° C for < 30 15 days, with surviving inocula enumerated every three days. Salt-in-the-moisture phase 31 (% SMP), calculated from measured moisture (%) and salt (%), titratable acidity (%), pH, and a_w 32 were measured. Pathogens did not grow on 53 cheeses, while 14 cheeses supported growth of 33 SA, six of SALM, four of LM, and three of EC. Of the cheeses supporting pathogen growth, all 34 supported growth of SA, ranging from 0.57 to 3.08 log CFU/g (avg. 1.70 log CFU/g). Growth of 35 SALM, LM, and EC ranged from 1.01 to 2.05 log CFU/g (avg. 2.05 log CFU/g), 0.60 to 2.68 log 36 CFU/g (avg. 1.60 CFU/g), and 0.41 to 2.90 log CFU/g (avg. 1.69 CFU/g), respectively. Cheese 37 38 pH and %SMP most affected pathogen growth, with pH having a dominant effect. Pathogen growth/no-growth varied within some cheese types or lots. Except for Swiss-type cheeses, mold-39 or bacterial-ripened cheeses, and cheeses made with non-bovine milk where insufficient data 40 exists, the pathogen growth/no-growth interface could be modeled and boundary conditions 41 established for safe, extended storage (<25°C) of cheeses based on pH and %SMP. 42

43 **INTRODUCTION**

Temperature-dependent storage of most cheeses has three major roles – to allow for 44 curing/ripening of cheeses that contain added or indigenous bacteria and enzymes, to prevent 45 quality defects, and to control pathogen growth (3). The 2009 US Food and Drug Administration 46 (FDA) Food Code (40) defines a potentially hazardous food as a food that requires 47 time/temperature control to limit the growth of pathogenic microorganisms or toxin formation. In 48 49 this publication, potentially hazardous foods are also designated as Time/Temperature Control for Safety (TCS) foods. This latter designation has been adopted in the 2013 Food Code (45). In 50 both versions of the Food Code, foods with a pH of <4.2 and any a_w or a_w of <0.88 and any pH 51 52 are not considered potentially hazardous. Foods considered potentially hazardous, unless shown to be safe by a product assessment, fall into one of the following categories: $a_w \ge 0.88$ and pH 53 >5.0, $a_w > 0.90-0.92$ and pH >4.6, or $a_w > 0.92$ and pH >4.2. The Food Code indicates that TCS 54 55 foods must be maintained at $<5^{\circ}$ C, or, if placed outside refrigeration, can be stored for up to 6 h at a temperature no greater than 21°C, after which the product must be discarded. 56

57 The composition of many cheeses, when evaluated using the Food Code criteria, places them into the category of TCS foods, thus limiting the ability of retailers to market the cheeses 58 59 under room-temperature conditions which could enhance cheese flavor and aroma (12). The Food Code-mandated time and temperature control may also limit industry flexibility in the 60 transportation, handling, and storage of cheeses. It has, however, been suggested that the 61 biochemical changes that occur during cheese ripening create an environment hostile for 62 pathogen growth, and that time/temperature control of some cheese is primarily needed to 63 maintain the organoleptic quality of cheese, not to maintain safety (3). Bishop and Smukowski 64 conducted a thorough review of the literature available up until 2006 and recommended that 65

66 cheeses meeting certain criteria, e.g. cheeses manufactured in the US with pasteurized or heattreated milk (>63°C for >16 sec), cheeses manufactured following Good Manufacturing 67 Practices and under the principles of HACCP (Hazard Analysis and Critical Control Points), and 68 69 cheeses manufactured meeting standards of identity outlined in 21 CFR (Code of Federal Regulations) part 133 (43), should be exempted from refrigeration requirements during ripening, 70 storage, shipping, and display (3). Bishop and Smukowski recommended that the following 71 cheeses could meet these criteria: Asiago (medium and old), Cheddar, Colby, Feta, Monterey 72 Jack, Muenster, Parmesan, Pasteurized process, Provolone, Romano, and Swiss/Emmentaler. 73

In order to establish whether a particular food, e.g. cheese, can be exempted from TCSrequirements, the Food Code allows processors or retailers to conduct a microbial challenge study in order to assess the ability of a food product to inhibit pathogenic bacterial growth or inactivate these microorganisms. The Food and Drug Administration (FDA) has outlined parameters for conducting such challenge studies (*44*).

79 When experts consider the major microbiological hazards across the food supply, the risk of bacterial illness from dairy products such as milk and cheese can be attributed primarily to 80 Listeria monocytogenes, Yersinia entercolitica, Campylobacter spp., and non-typhoidal 81 82 Salmonella spp. (2). Between 1990 and 2011, there were 105 reported foodborne illness outbreaks in the US, with over 2000 illnesses, linked to cheese/cheese products in the US (11). 83 Pathogens linked to these cheese-related outbreaks included Salmonella spp. (37 outbreaks), 84 Listeria monocytogenes (16 outbreaks), pathogenic Escherichia coli (6 outbreaks), 85 Staphylococcus aureus (4 outbreaks), Norovirus (21 outbreaks), Campylobacter spp. (9 86 outbreaks), and Brucella spp. (5 outbreaks) (11). Among the 105 outbreaks, 17 were linked to 87 cheeses made with pasteurized milk, 30 were linked to cheese made with raw milk, and the 88

89 pasteurization status of cheeses involved in the remaining 58 outbreaks was unspecified. The pathogenic bacteria primarily responsible for foodborne illness outbreaks linked to cheese 90 manufactured with pasteurized milk were L. monocytogenes, Salmonella spp., and E. coli 91 92 O157:H7. Cheeses implicated in these outbreaks included process cheese, Mozzarella, and Mexican-style cheeses (7, 11). The low incidence of S. aureus-linked outbreaks related to cheese 93 is presumed to be due to the low incidence of this pathogen in pasteurized milk, and the growth 94 characteristics of this bacterium (21). However, S. aureus is commonly carried by humans and 95 thus could contaminate cheese during post-pasteurization handling (16). S. aureus is also the 96 bacterial pathogen considered to have the highest tolerance to reduced-moisture conditions or 97 increased salt concentration (22), and therefore could be considered a target pathogen in 98 determining the safety of cheese contaminated post-processing and stored for extended periods 99 100 of time at room temperature.

101 The goal of this project was to evaluate survival of strains of *L. monocytogenes*, 102 *Salmonella* spp., *E. coli* O157:H7, and *S. aureus* on natural market cheeses during extended 103 storage at 25°C, and to determine the effect of cheese compositional factors such as pH, a_w, and 104 %salt on pathogen survival. Pathogen-survival data from laboratory research and data from 105 published literature were then combined in order to model the boundary conditions for pathogen 106 growth / no-growth during storage of cheese at room temperature.

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108 MATERIALS AND METHODS

109 Cheeses. Sixty-seven cheeses were purchased from local retail establishments or obtained directly from the manufacturer and stored at 4°C. Cheeses studied were Asiago (aged, 110 young), Brick (2 brands), Cheddar (mild, regular, sharp), Cheddar-Mozzarella, Colby, Colby-111 112 Jack, Farmer's, Feta, Gouda, Gruyere, Havarti (2 brands), Jack (goats' milk), Monterey Jack, Muenster (2 brands), Parmesan, Pepper Jack (2 brands), Provolone (mild, regular; 2 brands 113 sharp), Provolone-Mozzarella, Queso Blanco, Queso Fresco, Queso Quesadilla, String cheese (2 114 brands), Swiss (Baby, 2 brands; Lacey, regular), reduced-fat cheeses (Cheddar, Colby-Jack, 115 116 Provolone) and reduced-sodium cheeses (Colby Jack, Provolone). Where a type of cheese was tested more than once, tested cheeses were from different brands and/or from different 117 production dates of the same brand. All cheeses were manufactured in the United States from 118 pasteurized milk (Table 1, 2). 119

Proximate analysis. The cheeses tested in this study were characterized by % moisture, % salt, and a_w at the beginning of each trial. Changes in both % titratable acidity (%TA) and pH were anticipated over time; thus pH was measured on pathogen-inoculated cheeses at every sampling time (days 0, 3, 6, 9, 12 and 15), and %TA was measured on un-inoculated cheeses on days 0, 6 and 15. Duplicate trials were performed for each compositional analysis, and average values were reported.

Moisture (%) was determined using a standard method (4) by drying a representative 3-g sample at 100°C for 5 h in a vacuum oven maintained at -98kPa throughout the drying process (M.D.O. Vacuum Oven, Model 3623, Lab-Line Instrument Inc., Melrose Park, IL). Salt (%) was determined by titration of chloride using the silver titration standard method (4). For each trial, a

representative 5-g sample was diluted with distilled water 1:20 (w/v) and % chloride was determined according to the standard method using a Model M926 Chloride Analyzer (Nelson Jameson, Marshfield, WI). The % chloride content was automatically calculated by the analyzer and expressed as mg%/liter of sodium chloride, which was converted to % salt by multiplying the appropriate dilution and conversion factors. Salt (%) and moisture (%) of an individual cheese sample were used to calculate % salt-in-moisture-phase (%SMP) using Equation 1:

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$$\%$$
SMP = (% salt x 100) / (% salt + % moisture) (1)

Water activity (a_w) was determined for each cheese at the beginning of each trial using an 137 AquaLab LITE water activity meter (Decagon Devices Inc., Pullman, WA) according to a 138 standard method (1). Titratable acidity (%) was monitored during storage (days 0, 6, and 15) 139 according to a standard method (4). Briefly, for each cheese/trial, one sample (10.0 + 0.5 g) that 140 had been manually crumbled was automatically blended with 50-ml distilled water and titrated 141 using a Model DL22 Automatic Titrator (Mettler Toledo, Schwerzenbach, Switzerland), which 142 was set to calculate % titratable acidity (%TA) using the molecular weight of lactic acid. To 143 determine the impact, if any, of the presence of inoculum bacteria or growth of indigenous 144 bacteria on cheese pH, the surface pH was measured for individual inoculated cheese slices at 145 each sampling time (days 0, 3, 6, 9, 12, and 15) using an Accumet AB15 pH meter equipped 146 with a flat surface combination electrode (Fisher Scientific, Itasca, IL). 147

Inoculum preparation. Ten strains of *L. monocytogenes*, six strains of *Salmonella* spp., five strains of *E. coli* O157:H7, and five strains of *S. aureus*, representing a wide variety of sources and serotypes, were used in this study (Table 3). Stock cultures were maintained at -20°C in brain heart infusion broth (BHIB; Difco, Becton Dickinson, Sparks, MD) with 10% 152 (wt/vol) added glycerol (Fisher). Fresh working cultures were prepared monthly by thawing stock cultures and streaking for isolation as follows: L. monocytogenes on Listeria Selective agar 153 (LSA; Oxoid, Ogdensburg, NY) with added Listeria Selective Supplement (Oxford formulation, 154 Oxoid), Salmonella and E. coli O157:H7 on modified Levine's Eosin Methylene Blue agar (m-155 LEMB), prepared from lactose-free LEMB agar (Difco) with the addition of 10 g/liter D-sorbitol 156 (Fisher) and 5 g/liter NaCl (Fisher); and S. aureus on Baird-Parker agar (BP; Difco) with added 157 egg yolk Tellurite enrichment (Difco). Working culture plates were incubated for 24 h at 35°C 158 for Salmonella spp. and E. coli O157:H7, and 48 h at 35°C for L. monocytogenes and S. aureus, 159 whereupon all cultures were observed for consistent colony morphology and stored at 4°C for 160 <40 days. Inoculation cultures were prepared for individual strains by transferring a single 161 colony of each strain into a separate tube containing 9 ml of Nutrient broth (NB; Difco) for L. 162 monocytogenes, or BHIB for Salmonella spp., E. coli O157:H7 and S. aureus. Preliminary 163 cheese challenge studies showed better survival of L. monocytogenes over 15 days at 25°C on 164 Cheddar and Swiss cheeses when inocula had been grown in NB, while the other three pathogens 165 166 survived better on cheeses when inocula had been grown in BHIB (n=2, data not shown). Following incubation for 20 to 24 h at 35°C, 1 ml of stationary-phase culture of each strain for a 167 designated pathogen (10^8 CFU/ml for L. monocytogenes, and 10^9 CFU/ml for Salmonella spp., E. 168 coli O157:H7, and S. aureus) was transferred to a sterile 9-ml tube to produce a single-pathogen, 169 multi-strain cocktail. Each pathogen cocktail was mixed by vortexing and diluted, as necessary, 170 to produce a starting inoculum cocktail of 10^7 CFU/ml. Pathogen levels in the cocktails were 171 estimated by plating the inocula on brain heart infusion agar (BHIA; Difco) and incubating at 172 35°C for 24 h. 173

Sample inoculation. The working surface of a biosafety cabinet was sterilized with 70% 174 (v/v) ethanol and covered with aluminum foil prior to cheese inoculation. Cheese slices (approx. 175 25-30 g, approx 70-80 cm²) were placed on the aluminum foil aseptically, six cheese slices per 176 trial. An aliquot (0.1ml) of a single-pathogen cocktail (10^7 CFU/ml) was pipetted onto each of 177 the six cheese slices. An L-shaped spreader was used to evenly distribute the inoculum over the 178 surface of the six slices, then samples were left to air-dry under the hood for 15 min to allow 179 180 bacterial attachment and evaporation of excess liquid. The aw values of control and air-dried inoculated samples were not significantly different (n=3; p>0.05; data not shown). Inoculated 181 cheese slices were folded into half, with the inoculated cheese surfaces facing inward. Folded 182 cheese samples were weighed, then individually vacuum-packaged in standard retail barrier bags 183 (B-2175; Cryovac Food Packaging and Food Solutions, Duncan, SC) and stored at 25°C for up 184 to 15 days. Oxygen transmission rate for the bags was 3-6 cm³/m² at 40°F in 24 h. The initial 185 inoculum level on each cheese slice was $\sim 10^5$ CFU/g. 186

Sampling and enumeration. Packaged cheese samples were analyzed following 187 inoculation (time 0) and throughout storage for up to 15 days. Every three days, one cheese slice 188 189 per pathogen was removed from incubation, the storage/barrier bag was aseptically opened, and Butterfield's phosphate diluent (BPD; Nelson Jameson, Marshfield, WI) was added to create a 190 1:10 (w/w) dilution. The cheese/diluent mixture was stomached in the bag (AES Smasher, AES 191 Chemunex, Bruz, France) for 2 min at high speed. Stomached samples were serially diluted in 192 BPD, and 0.1-ml portions were spread-plated onto LSA, m-LEMB, m-LEMB, and BP for 193 cheeses inoculated with L. monocytogenes, Salmonella spp., E. coli O157:H7, and S. aureus, 194 respectively. A preliminary trial confirmed better recovery of Salmonella spp. by plating on m-195 LEMB rather than on Xylose Lysine Desoxycholate agar (XLD; Difco), and better recovery of E. 196

197 coli O157:H7 by plating on m-LEMB rather than on Sorbitol MacConkey agar (SMAC; Difco). Inoculated samples were also spread-plated on deMan-Rogosa-Sharpe agar (MRS; Difco) at 0, 6, 198 and 15 days to monitor changes in lactic acid bacteria (LAB) populations during storage, and to 199 200 thereby investigate the impact, if any, of indigenous, starter, or adjunct bacterial growth on inoculum survival. The m-LEMB spread-plates were incubated 24 h at 35°C, LSA and BP plates 201 48 h at 35°C, and MRS plates 72 h at 35°C, after which time counts were recorded for each plate, 202 203 with countable plate counts converted to log CFU/g. On m-LEMB, typical colonies of E. coli O157:H7 appear colorless to pink, while colonies of Salmonella spp. are dark red-black with 204 metallic green sheen. Colonies of S. aureus are typically shiny black and surrounded with clear 205 zone on BP agar. L. monocytogenes colonies are normally grey in color surrounded by black halo 206 on LSA. Data were used to calculate $\Delta \log$ CFU/g, relative to time 0, over the 15-day storage 207 208 period for each pathogen/cheese combination.

209 Literature data search and selection. To provide additional data to augment our product assessment, data from published literature were combined with data from this study. In 210 searching for relevant published studies, keywords including, but not limited to, "pathogen, 211 212 survival, cheeses, temperature, pH, salt" were entered into online scientific databases. Reference lists of publications were also screened for relevant studies with appropriate data. Published 213 challenge studies that met the following criteria were selected: (i) the inoculated cheeses were 214 made with pasteurized cow's milk, (ii) the cheeses were inoculated with at least one of the 215 pathogens: L. monocytogenes, Salmonella, E. coli O157:H7, or S. aureus, (iii) the pathogen(s) 216 was inoculated on the finished cheese, and (iv) inoculated cheeses were stored at 20-30°C. 217 Studies with surface-ripened, mold-ripened, Swiss, or processed cheeses, or cheese made with 218 non-bovine milk were excluded. Of 155 studies published between 1959 and 2012 and which 219

investigated pathogen behavior in or on cheeses, six published studies met the criteria (*14, 24, 25, 33, 34, 39*). From each publication, the following information was extracted: type of cheese,
temperature and length of storage, type and number of pathogen strains, composition (all available information for pH, a_w, % moisture, %SMP, % TA) of cheeses and behavior (growth vs. no-growth) of pathogen(s) (Table 4).

Evaluating compositional characteristics affecting pathogen growth. The relationship 225 between compositional factors and behavior of pathogens on cheeses was explored. 226 Compositional factors of cheese: moisture (%), initial pH, %SMP, a_w, and initial TA (%), were 227 paired, i.e. one compositional factor as "x" and one as "y", and a growth vs. no-growth outcome 228 was plotted for each cheese as a function of the x and y values to analyze the influences of the 229 paired compositional factors on pathogen growth. Values of compositional factors were 230 normalized to a 100-point scale before plotting as follows: for each compositional factor, the 231 232 minimum value of the data set was subtracted from the observed value and the total was divided by the range of the values and multiplied by 100 to obtain the normalized value, as shown in 233 Equation 2. 234

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Normalized value =
$$[(value - minimum value)/range] \times 100$$
 (2)

In this analysis, a "growth" result was indicated for a cheese when the Δ -log CFU/g for any cheese/pathogen combination over the 15-day storage period was a positive value that exceeded the pathogen-specific plating variability: 0.39, 0.41, 0.27, and 0.25 log CFU/g for *L. monocytogenes*, *Salmonella* spp. *E. coli* O157:H7 and *S. aureus*, respectively. The growth / nogrowth outcome plot from each pair of compositional factors was inspected and compared with predictions from a logistic regression equation (SAS 9.2, SAS Institute, Cary, NC). A model at 242 *P*=0.05 based on the variables pH and %SMP was generated according to the method of
243 McMeekin et al. (29) (Figure 1).

244 **RESULTS AND DISCUSSION**

In this study, 67 cheese samples, representing a variety of national brands, were tested for 245 their ability to support pathogen growth during extended storage at 25°C (Table 1, 2). Cheeses 246 were manufactured using pasteurized milk in facilities meeting applicable federal and state food 247 safety regulatory requirements. Cheeses met a standard of identity, where applicable. Among the 248 67 cheese samples tested, 52 were duplicate samples of cheeses from different lots/production 249 dates of the same brand. The majority of cheeses that were tested in this study would be labeled 250 as 'hard' or 'semi-hard' cheeses according to FDA classification (43), and were expected to be 251 252 safe for extended room-temperature storage due to reduced moisture level and low pH. Trials on 'soft' cheeses with higher moisture were also included in this study in order to clarify 253 compositional differences affecting pathogen growth/no-growth outcomes. Inoculated cheeses 254 255 were vacuum packaged to prevent mold growth and moisture loss which could inhibit pathogen 256 growth.

The FDA, in its guide to microbial challenge testing, notes that it can be important to evaluate a range of intrinsic factors which can influence the safety of a food during its intended shelf life (*44*). Compositional factors in cheese that could influence pathogen behavior were analyzed: surface pH (Day 0, 3, 6, 9, 12, 15), % moisture, % salt, and a_w (Day 0); and % TA (Day 0, 6, 15). Change in lactic acid bacteria (LAB) count was determined on Day 0, 6, and 15. Across all cheese samples, moisture content ranged from a low of 32.07% to a high of 57.64%, for one lot of Gruyere and Feta cheese, respectively. Salt content ranged from 0.33% for one lot

of Lacey Swiss to 3.30% for Queso Blanco. Salt-in-moisture phase (%SMP) was calculated from % moisture and % salt (Equation 1) with values ranging from 0.73% for one lot of Lacey Swiss to 7.21% for one lot of Parmesan. Water activity (a_w) varied little across the cheese samples tested, ranging from 0.96 to 0.99, except for Parmesan (average a_w =0.93) (Table 1).

Cheese pH measured at the surface, ranged from 4.33 to 6.49 for Feta (average of two 268 lots) and Queso Fresco, respectively (Table 1, 2) on Day 0. Over the 15-day storage period, 269 270 change in pH ranged from -1.44 to +0.53 pH units, for Queso Fresco and Baby Swiss (average of 4 lot), respectively, with most cheeses exhibiting only slight change in pH. To quantify the 271 272 amount of organic acid present in each cheese at the beginning of storage and to determine the effect, if any, of storage on changes in organic acid level, % TA was measured (Table 1, 2). The 273 %TA across the cheeses tested ranged from 0.26% to 2.83% for Queso Blanco and Feta (average 274 of 2 lots), respectively, at the beginning of storage. Change in %TA over storage was not clearly 275 276 linked with change in pH and bacterial survival (data not shown). Change in LAB count in cheese samples was estimated during extended storage at 25°C storage (Table 1, 2). LAB count 277 on Day 0 across the cheeses ranged from 2.00 to 8.08 log CFU/g for one lot of Pepper Jack and 278 279 Monterey Jack, respectively. Initial LAB counts on similar cheese samples from different brands, or different lots of the same brand, could vary widely. The Day 0 count for LAB on different 280 lots of Provolone (reduced-fat) (Brand 3) varied by 3.25 log CFU/g between purchase dates. 281 Similarly, one sample of Provolone (Brand 3) had one of the lowest Day 0 LAB counts, 2.70 log 282 CFU/g, while another sample of a different brand of Provolone (Brand 4) had one of the highest 283 initial LAB counts, 7.70 log CFU/g. The Day 0 LAB counts for the two samples of Brand 3 284 Provolone were 2.70 log CFU/g and 3.78 log CFU/g, and these rose to 5.40 log CFU/g and 7.19 285 log CFU/g, respectively; equivalent to a Δ -log of 2.70 and 3.41 log CFU/g, respectively. The 286

LAB count for the one lot of Brand 4 Provolone increased by one order of magnitude, from 7.70 log CFU/g (Day 0) to 8.70 log CFU/g (Day 15). Throughout the storage period and across all cheese samples tested, changes in LAB count ranged from -2.92 CFU/g for one lot of Parmesan to +5.66 log CFU/g for one lot of Pepper Jack (Brand 4). Of the 67 cheese samples tested, LAB population increased on storage in 47 cheese samples tested. LAB count was relatively constant $(0 < \Delta \log \le 0.3 \log CFU/g)$ in 7 cheese samples tested, and declined ($\Delta \log \ge -0.3 \log CFU/g$) in 13 other cheese samples during storage.

Cheeses were tested for their ability to support growth of L. monocytogenes, Salmonella 294 spp., E. coli O157:H7, and S. aureus (Table 1, 2). Pathogens did not grow on 53 cheese samples 295 over the 15 days (Table 1), while 14 cheese samples supported growth of S. aureus, six of 296 Salmonella, four of L. monocytogenes, and three of E. coli O157:H7 (Table 2). The pattern of 297 pathogen survival for each cheese lot was consistent over storage except for Queso Quesadilla 298 299 (Table 2). We observed growth of S. aureus (+0.57 log CFU/g) at Day 6 on Queso Quesadilla however by Day 15 we noted a decrease in pathogen population (overall $\Delta \log = -0.40 \log$ 300 CFU/g). Of the cheese samples which did support pathogen growth, all supported growth of S. 301 302 aureus, ranging from 0.57 to 3.08 log CFU/g (avg. 1.62 log CFU/g across all 14 cheeses). Growth of L. monocytogenes, Salmonella spp., and E. coli O157:H7, ranged from 0.60 to 2.68 303 log CFU/g (avg. 1.60 log CFU/g), 1.01 to 3.02 log CFU/g (avg. 2.05 log CFU/g), and 0.41 to 304 2.90 log CFU/g (avg. 1.69 CFU/g), respectively. Cheese samples which supported growth of S. 305 aureus included Farmer's, Gruyere (2 lots), Jack (goat's milk), Muenster (Brand 6), Provolone 306 (Brand 3; 2 lots), reduced-sodium Provolone (2 lots), Queso Blanco, Queso Fresco, and 2 brands 307 of String cheese. The six cheeses that supported growth of Salmonella spp. included: Gruyere (2 308 lots), Jack (goats' milk), Muenster (Brand 6), Queso Fresco, and one brand of String cheese 309

(Brand 14). The four cheeses that supported growth of *L. monocytogenes* included: Gruyere (one
lot), Queso Blanco, Queso Fresco, one brand of String cheese (Brand 14), and the three cheeses
that supported growth of *E. coli* O157:H7 included: Muenster (Brand 6), Queso Fresco, and
String (Brand 14).

Among the cheeses which supported pathogen growth at some point during the 15-day 314 storage period, seven supported only the growth of S. aureus: Farmer's, Provolone (Brand 3; 2) 315 trials), reduced-sodium Provolone (Brand 6; 2 trials), String cheese (Brand 6) and Queso 316 Ouesadilla (at Day 6 sampling point only) (Table 2). Three cheeses supported the growth of S. 317 aureus and one other pathogen: one lot of Gruyere and Jack (goats' milk) cheese each supported 318 the growth of S. aureus and Salmonella, while Queso Blanco supported the growth of S. aureus 319 and L. monocytogenes. Two cheeses supported the growth of three pathogens: one lot of Gruyere 320 supported the growth of L. monocytogenes, Salmonella spp, and S. aureus, and one lot of 321 322 Muenster (Brand 6) supported the growth of Salmonella spp., E. coli O157:H7, and S. aureus (Table 2). There were two cheeses which supported growth of all four pathogens, Queso Fresco 323 and one brand of String cheese (Brand 14). 324

Salmonella spp., L. monocytogenes and E. coli O157:H7 have, in recent years, been 325 326 implicated in foodborne illness outbreaks linked to cheeses made with pasteurized milk (7, 11). S. aureus has not often been associated with foodborne illness outbreaks linked to cheese, even 327 though this pathogen is generally linked to foods, like cheese, which are often hand-manipulated 328 during processing and packaging (8, 42). We included S. aureus in the study design not only 329 because of its link to poor sanitation and post-processing contamination but also because it is the 330 pathogen most likely to grow in or on foods with reduced moisture and/or low a_w (21). For 331 ready-to-eat food products, the FDA has established a zero-tolerance policy for L. 332

333 monocytogenes, Salmonella spp., and E. coli O157:H7, due to the potentially low infectious dose of E. coli O157:H7 and Salmonella spp, and the high mortality rate (15-30%) associated with L. 334 monocytogenes infections (41). Although none of these pathogens should be present in finished 335 336 cheeses made from pasteurized or heat-treated milk, the composition of a cheese supporting growth of any of these bacteria during extended room-temperature storage presents an 337 unacceptable risk. A zero-tolerance policy is not in place for S. aureus in ready-to-eat foods 338 339 because staphylococcal food poisoning occurs as a result of ingestion of a preformed enterotoxin which is only produced in amounts sufficient to cause illness as a result of extended temperature 340 abuse and growth of the pathogen to a high concentration ($\sim 10^5$ CFU) (30). Thus a cheese with 341 compositional characteristics allowing growth of S. aureus during storage is also an unacceptable 342 risk. For these reasons, growth of four target pathogens: L. monocytogenes, Salmonella spp., E. 343 coli O157:H7, and S. aureus, as post-processing contaminants on cheeses was investigated. 344

345 Pathogen strains used in this study represented a variety of sources and serotypes (Table 3). The strains of *L. monocytogenes* and *Salmonella* spp. had been screened in previous research 346 in our laboratory to confirm tolerance to salt and pH conditions typical of cheese (13). Strains of 347 348 E. coli O157:H7, Salmonella spp., and S. aureus were exposed to acid during inoculum preparation in BHIB, as a pH drop of ~ 1 unit was observed during overnight incubation. L. 349 monocytogenes was grown in NB, with no acid production or pH drop during inoculum 350 preparation. Where it occurred, the slight exposure to acid during inoculum preparation was 351 unlikely to have led to acid adaptation of strains. Therefore, the key characteristic of strains 352 selected for use in this study was their human or animal/animal-product origin, making these 353 strains perhaps representative of organisms to be found in a food processing or handling 354 environment. 355

356 Growth of L. monocytogenes was observed on four cheese samples: Gruvere, Queso Blanco, Queso Fresco, and one brand of String cheese (Brand 14) (Table 2), ranging from 0.60 to 357 2.68 log CFU/g. Growth of L. monocytogenes on Muenster (Brand 6, 0.17 log CFU/g) and 358 359 String (Brand 6, 0.22 log CFU/g) did not exceed the plating variability for the pathogen (0.39 log CFU/g) and 'growth' was not declared. Genigeorgis et al. studied the survival of L. 360 monocytogenes on 11 different types of market cheeses stored at 30°C (14). Pathogen growth 361 was observed only on Hispanic-style cheeses: Queso Fresco, Queso Ranchero, and Queso 362 Panela, and ranged from 0.38 to 3.18 log CFU/g (14). Uhlich et al. observed an increase of more 363 than 5 log CFU/g of L. monocytogenes on Queso Blanco stored at 25°C for up to 6.25 days (39). 364 In the present study, we observed growth of L. monocytogenes on one brand of String cheese 365 (Brand 14) that slightly exceeded the plating variability, i.e. the observed growth of 0.60 log 366 CFU/g exceeded the plating variability of 0.39 log CFU/g. Genigeorgis et al. (14) did not 367 observe growth of L. monocytogenes on String cheese, instead noting a drop in L. 368 monocytogenes population of 2.36 log CFU/g over 9 days at 30°C. The String cheese that 369 370 Genigeorgis et al. tested had similar pH and %SMP values to the cheese sample that we evaluated, but an unknown level of LAB. The String cheese sample in our study allowing some 371 growth of pathogen simultaneously supported a dramatic increase in LAB population, from 4.87 372 log CFU/g at Day 0 to 8.86 log CFU/g by Day 15 (Table 2). 373

Growth of *L*. monocytogenes was not observed on 63 samples of cheese tested (Table 1, 2), many of the cheeses which did not support pathogen growth would be classified as 'hard' or 'semi-hard' cheeses based on FDA classification (*43*) and may be suitable for extended room temperature storage. Shrestha et al. (*33*) did not observe growth of *L. monocytogenes* on a range of Cheddar-type cheeses stored at 21°C for 30 days, with counts of *L. monocytogenes* dropping

379 by $< 1.1 \log CFU/g$ during storage. We also observed a slight decrease in the population of L. monocytogenes on mild, reduced-fat, and sharp Cheddar cheeses during storage at 25° (Table 1). 380 Pathogen populations decreased from 0.00 to 0.76 log CFU/g across samples and Cheddar 381 382 cheese-type tested. Genigeorgis et al. also reported a decrease of L. monocytogenes population on mild Cheddar cheeses during storage (14). Similarly, Genigeorgis et al. evaluated the growth 383 of L. monocytogenes on Monterey Jack, Colby, Provolone, Muenster, and Feta cheeses during 384 storage, and observed a decrease in pathogen population of >1-2 log CFU/g in all cases. In our 385 study, we noted an average decrease in pathogen population of 0.2 log CFU/g for Colby, 4.74 log 386 CFU/g for Feta, 1.83 log CFU/g for Monterey Jack, 0.25 log CFU/g for Muenster (Brand 3), and 387 0.99 log CFU/g for several different types of Provolone (regular, mild, sharp) (Table 1, 2). Two 388 lots of Provolone (Brand 3) which supported growth of S. aureus did not support the growth of 389 390 L. monocytogenes (Table 2). One brand of Muenster (Brand 6) appeared to support a slight growth of L. monocytogenes during storage (0.17 log CFU/g), but this was found not to exceed 391 the plating variability associated with this pathogen (0.39 log CFU/g), and thus 'no growth' was 392 declared. 393

394 Growth of Salmonella was observed on six cheeses: Gruyere, Jack (goat's milk), Muenster (Brand 6), Queso Fresco, and String (Brand 14), ranging from 1.01 to 3.02 log CFU/g 395 over 15 days. Slight growth of Salmonella was also observed for Brand 6 of String cheese (0.39 396 log CFu/g) but this was below the plating variability for this pathogen (0.41 log CFU/g), and 397 therefore counted as 'no growth.' Kasrazadeh and Genigeorgis (25) studied the growth of 398 Salmonella inoculated onto sliced Queso Fresco stored at 20°C. They noted rapid growth, a lag 399 time of 2.5-3.5 h and a generation time of 1.65-2.17 h, for Salmonella on Queso Fresco. We 400 observed an increase in Salmonella concentration of 3.02 log CFU/g on Queso Fresco stored at 401

402 25°C over 15 days. This was the highest level of *Salmonella* growth observed over all 67 cheese
403 samples tested.

There were 61 cheeses which did not support the growth of *Salmonella* in this study. 404 Shrestha et al. (34) examined the survival of Salmonella on a range of Cheddar-type cheeses 405 406 stored for up to 30 days at 21°C. Cheddar cheese manufactured to standards of pH and salt was comminuted, inoculated with Salmonella spp., and stored at 21°C for up to 30 days. Salmonella 407 spp. counts decreased significantly at 21°C for all cheese-types. We evaluated the survival of 408 Salmonella spp. on mild, reduced-fat, and sharp Cheddar cheeses and observed average 409 decreases of 0.3, 1.12, and 1.26 log CFU/g, respectively, for the brands tested. Growth of E. coli 410 O157:H7 was observed on three cheeses: Muenster (Brand 6), Queso Fresco, and String (Brand 411 14), ranging from 0.41 log CFU/g (Muenster) to 2.90 log CFU/g (Queso Fresco) over 15 days. 412 Kasrazadeh and Genigeorgis (24) also observed rapid growth of E. coli O157:H7 on Queso 413 414 Fresco stored at 20° C. There were 64 cheese samples in this study which did not support the 415 growth of this pathogen.

The survival pattern for pathogens on cheeses was consistent during storage, with the 416 exception of the survival of S. aureus on Queso Quesadilla which increased by 0.57 log CFU/g 417 on Day 6 of storage, but decreased by 0.40 log CFU/g relative to the time-zero level by Day 15. 418 In all other cases, pathogen growth/no-growth was consistent, displaying an increase or decrease 419 over the 15-day storage period. LAB count increased in 47 of 67 cheeses tested in this study. 420 With one exception, cheeses which supported pathogen growth also supported LAB growth. 421 LAB count decreased in Jack (goat's milk) cheese which supported growth of Salmonella (+2.50 422 log CFU/g) and S. aureus (+1.62 log CFU/g); otherwise LAB count increased from 1.54 to 4.47 423 log CFU/g in cheeses which supported pathogen growth. The level of inoculum on each cheese 424

slice at time 0 averaged 4.7 log CFU/g (n=268). This level allowed for accurate enumeration of 425 growth or death without reaching the limits of research methodology. This inoculum level could 426 have placed pathogens at a level to effectively compete with active indigeneous organisms. LAB 427 428 count on Day 0 averaged 5.03 log CFU/g for cheeses which supported pathogen growth (n=14, Table 2). While previous studies have shown that initial inoculum level does not affect the 429 survivability or growth kinetics of pathogens (6, 26, 46), a higher proportion of S. aureus 430 compared to LAB may aid in the survival of this particular pathogen (17). Although growth of S. 431 *aureus* is reported to be weak when a high load of competitive bacteria, e. g. lactic acid bacteria 432 (LAB) is present, increasing the proportion of S. aureus to LAB has been shown to aid in 433 survival of this pathogen (17, 23). 434

The change in pH on storage among cheeses that supported pathogen growth showed no 435 clear trend, remaining the same (Δ pH < 0.3 units) in 7 samples, and increasing in 6 samples 436 437 (Table 2). Cheese samples that supported pathogen growth had %TA which ranged from 0.26 % to 1.67 % at the beginning of storage (Table 2); while cheese samples which did not support 438 pathogen growth had %TA ranging from 0.66% to 2.86% at the beginning of storage (Table 1). 439 440 Change in %TA over storage (data not shown) had no apparent relationship with the change of pH and LAB count. Among cheeses that supported pathogen growth, LAB count increased in all 441 but one sample (Jack (goats' milk cheese)), with an increase ranging from 1.54 log CFU/g for 442 one lot of reduced sodium Provolone to 4.43 log CFU/g for Farmer's cheese. Correlation 443 between changes in pH and LAB count in cheeses was weak ($r^2=0.25$). 444

A total of 53 cheeses did not support the growth of any pathogen tested. These cheeses were most notably characterized by lower pH; there was little difference in % moisture and %SMP between these cheeses and those that supported pathogen growth. When cheese samples 448 were separated into roughly equal groups by initial pH value: 4.29 - 5.20 (29 cheeses), 5.21 -5.40 (18 cheeses), and 5.41 - 6.50 (20 cheeses), it was readily apparent that pathogen growth 449 was better supported on higher pH cheeses. With the exception of Provolone (Brand 3; pH 5.15) 450 451 and reduced-sodium Provolone (pH 5.15), cheeses with Day 0 pH ranging from 4.8-5.2 did not support growth of any pathogens (Table 1). Feta was the most acidic cheese tested (average pH 452 4.33, n=2 lots), and pathogen viability on this cheese type decreased over time more than for any 453 other cheese with average reductions of 4.74 log CFU/g for L. monocytogenes, 4.82 log CFU/g 454 for Salmonella spp., 4.34 log CFU/g for E. coli O157:H7, and 3.84 log CFU/g for S. aureus. As 455 pH increased to 5.21 - 5.40, four of 18 cheeses supported growth: Provolone (Brand 3; 1 lot), 456 reduced-sodium Provolone (1 lot), String cheese (Brand 6) and Queso Quesadilla; all supporting 457 the growth of S. aureus (average 1.14 log CFU/g across all 3 cheeses), but no other pathogen 458 459 (Table 2). In the pH range 5.41 - 6.50, eight cheeses supported pathogen growth: Jack (goats' milk) (pH 5.41), String (Brand 14, pH 5.44), Farmer's (pH 5.46), Muenster (pH 5.48), Gruyere 460 (2 lots; pH 5.68; 6.28), Queso Blanco (pH 6.37), and Queso Fresco (pH 6.49). Pathogen growth 461 462 on Queso Fresco was the greatest across all cheeses - tested; this was also the cheese with the highest initial pH. Cheeses with an initial pH >5.46 supported growth of at least one pathogen, 463 with the exception of Swiss-style cheeses (Baby Swiss, Swiss, Lacey Swiss – pH range 5.50 – 464 6.02), and one lot of Havarti (pH 5.49) which did not support growth. Optimal pH for growth of 465 S. aureus is between pH 6.0 and 7.0, with pH 4.0 as the reported minimum for growth (20). 466 Minimum pH values for growth that have been reported for L. monocytogenes, Salmonella spp., 467 E. coli O157:H7 are 4.39, 4.20, and 4.40, respectively (20). Only Feta cheese (pH 4.29, 4.38) 468 was below the reported minimum pH for growth of any of the pathogens tested. 469

470 The average moisture content for cheese samples which supported growth (43.11%)varied little from moisture content for cheese samples which did not support growth (40.38%) 471 (Table 1,2). An even narrower difference in -average SMP was observed between cheeses which 472 473 supported growth (3.76%) and cheeses which did not support growth (3.52%) (Table 1, 2), however the range of values in each category (growth/no-growth) was much wider, ranging from 474 0.73 to 7.21 %SMP for cheese samples which did not support growth, and from 2.26 to 6.56 475 %SMP for cheese samples that did. The greater growth potential that we observed for *S. aureus* 476 on cheeses could be attributed, in part, to the high salt-tolerance of this pathogen. Nunheimer and 477 Fabian reported that some strains of S. aureus are able to tolerate up to 20% NaCl (31). 478 Sutherland et al. (36) reported growth of S. aureus in BHIB with pH 4.48 and 8.5% NaCl at 479 25°C. Ingham et al. reported greater tolerance of S. aureus than of L. monocytogenes to high salt-480 481 and low a_w in meat products stored at 21°C (19).

Where applicable, we tested cheeses from different brands, or from different lots within 482 the same brand, to allow us to determine lot-to-lot or brand-to-brand variation for a similar type 483 484 of cheese. For example, Muenster cheese from two manufacturers was tested; cheese from one brand (Brand 6, Table 2) supported growth of three pathogens, S. aureus, Salmonella spp., and E. 485 coli O157:H7 (+0.41 to +1.77 log CFU/g;), while Muenster cheese from a different brand (Brand 486 3, Table 1) did not support growth of any pathogen (-0.00 to -0.75 log CFU/g). Among ten 487 Provolone cheeses tested (mild, sharp, regular (3 lots from 2 brands), reduced-fat (2 lots), 488 reduced-sodium (2 lots), and a Provolone-Mozzarella blend), six cheeses (2 lots of reduced-fat, 489 regular, sharp, mild, and Provolone-Mozzarella blend) did not support growth of any pathogen 490 (Table 1). The contribution of pH, %SMP, and other inhibitory compounds present in cheese, 491 492 such as metabolites of LAB and the presence of free fatty acids may have varied from lot-to-lot,

brand-to-brand, and between cheese types, resulting in differences in pathogen growth during non-refrigerated storage. The effect of these factors on microbial survival has been shown to be highly dependent on the concentration of inhibitory compound and the species and strain of both LAB and pathogen (*10*, *15*, *17*, *35*). The apparent inconsistencies in pathogen growth patterns observed for cheeses of a similar type supports the assertion that it may be compositional characteristics, more than cheese type, that determine the likelihood of pathogen growth on a sample of cheese.

The compositional factors of pH, %SMP, a_w, and %TA were paired in all combinations 500 501 and a pathogen growth / no-growth outcome for each cheese was plotted as a function of each pair of factors. Plotting growth / no-growth outcome as a function of pH and %SMP, combined 502 with logistic regression, created a growth / no-growth interface that could be used to clearly 503 differentiate cheeses which inhibited pathogen growth from those that allowed pathogen growth 504 505 (Figure 1). A similar approach using other pairs of compositional factors was not successful in generating a clear growth / no-growth interface (data not shown). These results are consistent 506 with those of Oh et al. who evaluated the effect of compositional factors of low-sodium Cheddar 507 508 cheeses on the growth of strains of Salmonella spp., L. monocytogenes, S. aureus, and Shiga toxin-producing E. coli (STEC). In a model low-sodium Cheddar-cheese extract, STEC survived 509 significantly better than the other three pathogens. Principal component analysis indicated that 510 STEC survival was primarily determined by pH, and not by % salt or % lactate (32). 511

The eight Swiss-style cheese samples tested did not fit the pattern established by data from the other cheeses tested. These Swiss-style cheeses had the lowest %SMP (0.73-1.87%) of all cheeses tested, a relatively high pH (5.36-6.02), and a high a_w (0.98-0.99). Despite compositional factors which seem to be permissive for growth, none of the Swiss-style cheeses 516 supported pathogen growth. Lever and Johnson reported poorer survival of Salmonella spp. on 517 Swiss cheeses than on Cheddar and Mozzarella (27). Swiss-style cheeses are unique among the types of cheeses that we tested due to the addition of propionic acid bacteria added as an adjunct 518 519 culture in cheese manufacture. The added propionic acid bacteria can produce metabolites with antimicrobial properties, such as propionic acid, acetic acid, and diacetyl (9). Studies have shown 520 greater antimicrobial properties linked to propionic acid (pK_a=4.87) as compared to lactic acid 521 $(pK_a=3.86)$ (37). The results of our study would suggest that target pathogens will not grow on 522 Swiss-style cheeses during extended storage at 25° C, but the safety of such cheeses should be 523 evaluated independently from cheeses which are fermented using only lactic acid-producing 524 bacteria. Similarly, research suggests that the ability of pathogens to grow on bacterial surface-525 ripened or mold-ripened should be evaluated independently from cheeses manufactured without 526 527 these ripening adjuncts. Bacterial surface-ripened and mold-ripened cheeses have added cultures that are capable of growing and altering the environment for pathogen growth. Growth of added 528 bacterial and/or mold cultures can result in the production of antimicrobial compounds (e.g. 529 530 bacteriocins) which could hinder pathogen growth, but can also lead to lactate metabolism which can subsequently increase cheese pH and enhance pathogen growth (5). Genigeorgis et al. found 531 a significant reduction of L. monocytogenes (> $-2.36 \log \text{ CFU/g}$) when inoculated onto 532 Limburger, a bacterial surface-ripened cheese (14). While the high pH of Limburger (pH 7.2) 533 would suggest that this cheese could support pathogen growth, the growth of smear bacteria 534 results in extensive lipolysis which produces a high concentration of free fatty acids, which are 535 compounds known to have antimicrobial activity (35). Goats' milk cheese may also contain high 536 levels of free fatty acids. Woo et al. evaluated the free fatty acid content in a variety of cheeses 537 and concluded that Blue, Swiss, Limburger, and goats' milk cheeses contained high 538

concentrations of free fatty acids (47). Thus, we conclude that surface-ripened cheeses, moldripened cheeses, and non-cow's milk cheeses, along with Swiss-style cheeses should be evaluated separately, perhaps by group, and more data gathered in order to assess their suitability for extended non-refrigerated storage.

Water activity (a_w) and pH are the two criteria used in the FDA Food Code to determine 543 the shelf stability of food products (45). However, %SMP can be seen as a more appropriate 544 factor than a_w in assessing the likelihood of pathogen survival on cheese. In addition to salt, other 545 solutes in cheese such as non-protein nitrogen-containing compounds and products released 546 during proteolysis, could contribute to the reduction of a_w, yet these compounds may not play a 547 role in inhibiting pathogen growth (28). Tapia et al. (38) suggested that the usefulness of 548 measured a_w as an indicator of microbial safety or stability is diminished by the 'specific solute 549 550 effect'; that is that the solute in the food matrix dramatically alters the minimum a_w for microbial growth. Hilderbrand (18) supported %SMP as a more reliable factor than a_w in determining 551 bacterial growth in smoked fish. In addition, %SMP is routinely determined and has historically 552 553 been used in the cheese industry as a measure of product quality. Our search of published literature indicated that other researchers investigating survival of pathogens as post-processing 554 contaminants on cheese routinely monitored %SMP (14, 24, 25, 33, 34, 39), while only a few 555 studies investigating pathogen survival on cheese considered the impact of product a_w (33, 34, 556 39). Furthermore we identified that pH and %SMP were the two compositional factors which 557 could be used to clearly differentiate cheeses which supported pathogen growth from those that 558 559 inhibited growth (Figure 1), while the compositional factors of pH and a_w were not similarly effective. 560

561 Of the 67 market cheeses studied, 53 did not support the growth of L. monocytogenes, Salmonella, E. coli O157:H7, or S. aureus and could safely be kept at < 25°C for an extended 562 period of time. The risk of pathogen growth for those cheeses which supported growth can be 563 characterized as follows: S. aureus (growth on 14 of 14 cheeses supporting pathogen growth) >> 564 Salmonella (growth on 6 of 14) > L. monocytogenes (growth on 4 of 14) > E. coli O157:H7 565 (growth on 3 of 14). Of several intrinsic compositional factors associated with cheese, i.e. pH, a_w. 566 %SMP, and %TA, cheese pH has the clearest effect on pathogen growth. Laboratory data was 567 combined with relevant published research in order to expand our product assessment. Pathogen 568 growth/no-growth outcomes for 82 cheeses, 56 cheeses tested in our laboratory and 26 cheeses 569 for which published results were available in the literature, were plotted on a graph with axes of 570 pH and %SMP. Logistic regression analysis generated a P=0.05 boundary line, which indicated a 571 572 clear differentiation between cheese compositions (in terms of pH and %SMP) which supported pathogen growth and those which did not. Data from Swiss-type cheeses, mold-or bacterial-573 ripened cheeses, or cheeses made with non-bovine milk were excluded from this analysis due to 574 575 insufficient data or lack-of-fit. The growth/no-growth interface established by the logistic regression line clearly shows that many common cheese types, if made from pasteurized cows' 576 milk in compliance with US regulatory standards, can safely be considered non-TCS foods. Non-577 TCS cheeses should be described in terms of pH and %SMP rather than cheese-type or brand, 578 and would include cheeses with pH/%SMP values more restrictive than any of the following 579 combinations drawn from Figure 1 (in order of increasing pH): <4.60/>0.24; 4.61-4.70/0.25-580 0.91; 4.71-4.80/0.92-1.58; 4.81-4.90/1.59-2.24; 4.91-5.00/2.25-2.91; 5.01-5.10/2.92-3.58; 5.11-581 5.20/3.59-4.25; 5.21-5.30/4.26-4.92; 5.31-5.40/4.93-5.59; 5.41-5.50/5.60-6.26; and 5.51-582 583 5.60/6.27-6.93. More research would be necessary to develop boundary conditions for safe,

584	extended room-temperature storage of cheeses not covered in this model, including Swiss-type
585	cheeses, bacterial surface-ripened or mold-ripened cheeses, cheeses made from non-bovine milk,
586	or cheeses made from unpasteurized milk.
587	
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Table 1. Composition of natural cheeses that did not support growth of *L. monocytogenes, Salmonella* spp., *E. coli* O157:H7, and *S.*

746 *aureus* and pathogen survival ($\Delta \log CFU/g$) during storage for 15-days at 25°C.

	Duand	%	%	%SMP ^d	,	$\mathbf{p}\mathbf{H}^{f}$		%TA ^g	LAB	count ^h	Patho	Pathogen survival ($\Delta \log CFU/g$) ⁱ			
Cheese ^{<i>a</i>}	Brand	Moisture ^b	Salt ^c	%SIMP	$\mathbf{a_w}^e$	0 d	15 d	0 d	0 d	15 d	LM	SALM	EC	SĂ	
Asiago (Young)	1	36.00	1.71	4.53	0.96	5.36	5.21	2.16	7.36	6.87	-2.05	-3.74	-2.12	-1.13	
Asiago (Young)	1	38.63	1.83	4.52	0.96	5.12	5.01	2.82	7.79	7.40	-2.26	-2.12	-0.68	-1.07	
Asiago (Aged)	10	38.84	0.96	2.41	0.97	5.15	4.98	1.78	6.22	7.16	-2.92	-2.79	-3.70	-3.53	
Asiago (Aged)	10	43.30	1.94	4.29	0.97	5.09	5.06	2.02	6.02	6.94	-3.84	-3.63	-1.59	-2.67	
Baby Swiss	5	38.36	0.61	1.57	0.98	5.77	6.28	_j	7.07	7.50	-0.71	-1.38	-1.98	-0.62	
Baby Swiss	5	36.25	0.69	1.87	0.98	5.79	6.32	-	6.94	8.05	-0.67	-0.76	-0.75	-1.15	
Baby Swiss	9	37.21	0.62	1.64	0.99	5.55	6.04	-	7.19	7.72	-1.00	-2.43	-1.27	-0.79	
Baby Swiss	9	35.58	0.65	1.79	0.99	5.71	6.27	-	7.35	7.72	-0.39	-1.45	-0.61	-1.02	
Brick	11	40.39	1.52	3.63	0.96	5.43	4.90	1.29	7.23	7.66	-0.40	-0.71	-0.38	-0.74	
Brick	11	41.21	1.95	4.52	0.97	5.30	4.98	0.90	7.19	7.82	-0.32	-0.70	-0.40	-0.98	
Brick	2	38.28	1.52	3.82	-	5.25	5.37	1.07	6.33	8.08	-0.09	-0.22	-0.42	-0.79	
Cheddar (Mild)	3	37.34	1.57	4.04	0.96	5.09	5.00	1.89	7.41	6.78	-0.70	-0.88	-0.30	-0.43	
Cheddar (Mild)	3	36.59	1.77	4.61	0.97	5.09	5.06	1.44	7.39	6.81	-0.76	-1.00	-0.80	-0.17	
Cheddar (Reduced-Fat)	6	40.26	1.60	3.82	0.97	5.19	5.11	1.15	5.35	6.21	-0.13	-0.65	-0.43	-1.28	
Cheddar (Reduced-Fat)	6	44.00	1.66	3.64	0.98	4.99	5.27	0.90	5.52	5.79	-0.69	-0.57	-0.55	-0.97	
Cheddar (Sharp)	3	36.34	1.78	4.67	0.96	5.27	5.27	1.69	4.30	6.39	-0.35	-0.75	-0.96	-1.19	
Cheddar (Sharp)	3	36.57	1.32	3.48	0.97	5.19	5.28	1.71	4.63	5.84	0.00	-1.03	-0.59	-1.34	
Cheddar-Mozzarella	6	40.09	1.62	3.88	-	5.19	5.33	1.42	6.99	6.24	-0.09	-0.27	-0.31	-0.48	
Colby	4	35.96	1.61	4.28	0.96	5.45	5.61	1.09	5.76	7.39	-0.39	-0.50	-0.21	-0.57	
Colby	4	40.14	1.60	3.83	0.97	5.30	5.47	1.78	5.91	6.38	-0.11	-0.63	-0.24	-0.39	
Colby Jack	5	36.13	1.42	3.78	0.96	5.17	5.10	1.26	7.19	7.19	-0.20	-0.97	-0.80	-0.46	
Colby Jack	5	36.85	1.35	3.53	0.98	5.01	5.40	1.37	7.70	7.38	-0.44	-0.59	-0.08	-0.46	
Colby Jack (Reduced-Fat)	6	43.96	1.64	3.60	0.97	5.29	5.00	1.09	5.79	7.68	0.02	-0.90	-0.76	-1.12	
Colby Jack (Reduced-Fat)	6	46.00	1.76	3.69	0.97	5.08	5.11	1.39	4.52	6.91	-0.56	-0.74	-0.73	-1.05	
Colby Jack (Reduced-Na)	6	36.30	1.26	3.35	0.97	5.11	5.03	1.48	4.52	6.91	-0.17	-0.46	-1.03	-1.09	
Colby Jack (Reduced-Na)	6	36.45	1.13	3.01	0.98	5.09	5.17	0.89	4.12	5.40	-0.69	-0.96	-0.39	-0.64	
Feta	3	57.10	2.35	3.95	0.99	4.29	4.60	2.80	4.80	6.57	-4.58	-4.71	-4.60	-2.93	
Feta	3	57.64	1.72	2.90	0.98	4.38	4.53	2.86	3.30	3.40	-4.89	-4.94	-4.07	-4.74	
Gouda	6	41.15	1.62	3.79	0.97	5.28	5.25	0.88	7.29	7.38	-0.51	-0.32	-0.23	-0.83	

Gouda	6	41.08	1.39	3.27	0.97	5.30	5.28	1.24	7.40	7.48	-0.44	-0.46	-0.34	-0.79
Havarti	3	37.79	1.33	3.40	0.97	5.49	5.52	1.08	6.88	7.26	-0.25	-0.61	-0.21	-0.73
Havarti	3	38.17	1.20	3.05	0.98	5.34	5.59	0.66	6.88	7.20	-0.51	$+0.21^{k}$	-0.29	$+0.01^{k}$
Havarti	6	41.32	1.27	2.98	-	5.11	5.26	1.40	8.28	7.75	-0.16	-0.61	-0.37	-0.70
Monterey Jack	5	45.10	1.87	3.98	0.98	5.15	5.20	2.41	8.08	8.16	-1.03	-0.91	-0.33	-0.37
Monterey Jack	5	35.45	1.64	4.42	0.97	5.08	5.11	2.28	8.06	7.98	-2.63	-1.17	-0.91	-0.66
Muenster	3	42.20	1.63	3.72	0.97	5.20	5.28	1.27	6.90	7.80	-0.49	-0.25	-0.24	0.00
Muenster	3	41.94	1.75	4.01	0.98	5.29	5.12	0.74	7.11	6.26	-0.10	-0.75	-0.45	-0.46
Parmesan	8	32.44	2.52	7.21	0.93	5.41	5.36	1.40	6.92	4.00	-0.88	-1.45	-1.25	-0.59
Parmesan	8	32.70	2.35	6.70	0.92	5.45	5.40	1.48	5.31	7.23	-1.51	-1.66	-1.86	-1.80
Pepper Jack	4	36.13	1.58	4.19	0.98	5.11	4.76	2.12	2.00	7.66	-0.85	-3.87	-0.81	-1.09
Pepper Jack	4	38.69	1.60	3.97	0.97	4.93	5.12	1.94	4.69	7.14	-2.86	-3.40	-3.25	-3.58
Pepper Jack	3	40.42	1.64	3.90	0.97	5.14	5.12	1.53	4.65	7.30	-2.39	-2.32	-2.10	-0.72
Pepper Jack	3	38.27	1.54	3.87	0.97	5.21	5.15	1.45	8.25	7.39	-0.62	-0.73	-0.35	-0.30
Provolone	4	42.15	1.38	3.17	-	5.24	4.97	1.81	7.70	8.70	-1.34	-0.97	-0.16	-0.72
Provolone (Mild)	8	43.05	2.08	4.61	-	5.18	5.22	1.80	5.53	6.70	-0.50	-1.84	-0.57	-0.71
Provolone (Sharp)	10	40.02	1.72	4.12	-	5.09	5.17	2.20	6.43	7.45	-1.59	-2.83	-1.27	-1.73
Provolone (Reduced-Fat)	3	48.98	1.43	2.84	0.97	4.97	4.67	1.83	6.95	7.98	-2.80	-2.23	-0.62	-1.55
Provolone (Reduced-Fat)	3	52.71	1.35	2.50	0.98	4.98	4.94	-	3.70	7.94	-0.56	-0.95	-0.24	-0.97
Provolone-Mozzarella	6	42.26	1.68	3.82	-	5.38	5.33	1.61	7.67	7.28	-0.25	-0.19	-0.17	-0.68
Swiss	6	38.57	0.52	1.33	0.98	5.36	5.50	-	5.95	6.59	-1.20	-1.11	-0.73	-2.32
Swiss	6	36.91	0.64	1.70	0.99	5.50	5.80	-	5.28	6.19	-0.93	-1.30	-0.36	-1.20
Swiss (Lacey)	5	45.17	0.33	0.73	0.99	6.02	5.87	-	7.00	8.18	-0.43	-1.19	-0.46	-1.02
Swiss (Lacey)	5	45.92	0.37	0.80	0.99	5.65	5.94	-	7.92	5.70	-1.83	-1.21	-0.31	-1.06

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^a Cheeses were national brands obtained from local retail outlets or directly from manufacturers. Qualifying descriptive information, e.g. 'mild,'

⁷⁴⁹ 'sharp' is reproduced where provided on the package.

750 ^{*b*}Moisture content (%) of cheese sample on Day 0, n=2.

751 ^{*c*} % salt of cheese sample on Day 0, n=2.

^{*d*}% salt-in-moisture phase (% SMP) of cheese sample on Day 0. Calculated from % moisture and % salt of the same cheese.

- 753 ^e Water activity (a_w) of cheese sample on Day 0.
- ^{*f*} pH of cheese slice surface on Day 0 and Day 15, n=2.
- 755 g % titratable acidity (%TA) of cheese sample on Day 0 and Day 15, n=2.
- ^{*h*} DeMan-Rogosa-Sharpe (MRS) agar count for lactic acid bacteria (LAB) on Day 0 and Day 15 (log CFU/g), n=2.
- *i* Survival of pathogen LM=*L. monocytogenes*, SALM=*Salmonella* spp., EC=*E. coli* O157:H7, and SA=*S. aureus*. (+) indicates
- 758 growth, (-) indicates no-growth.
- 759 j not determined.
- ^{*k*} Growth of pathogen did not exceed plating variability: 0.39, 0.41, 0.27, 0.25 log CFU/g for *L. monocytogenes, Salmonella* spp., *E.*
- 761 *coli* O157:H7and S. *aureus*, respectively.

762

Table 2. Composition of natural cheeses that supported growth of L. monocytogenes, Salmonella spp., E. coli O157:H7, and/or S. aureus and

Cheese ^a	Brand	%	%		a _w ^e	р	\mathbf{H}^{f}	% TA ^g	LA cou	AB int ^h	Pathog	en surviva	l (Δlog C	FU/g) ⁱ
		Moisture ^b	Salt ^c	\mathbf{SMP}^d		0 d	15 d	0 d	0 d	15 d	LM	SALM	EC	SA
Farmer's	12	39.85	1.71	4.11	_j	5.46	4.99	1.14	4.63	9.10	-0.41	-0.10	-0.39	+1.48 ^l
Gruyere	7	34.25	1.01	2.86	0.97	5.68	5.74	1.04	5.70	7.40	+1.01	+1.01	-0.40	+3.08
Gruyere	7	32.07	1.41	4.21	0.98	6.28	5.78	1.55	5.04	6.70	-0.54	+2.13	-0.67	+2.19
Jack (goats' milk)	13	45.20	2.33	4.90	-	5.41	5.24	1.44	7.74	6.88	-0.40	+2.50	-0.62	+1.62
Muenster	6	41.58	1.21	2.83	-	5.48	5.53	0.66	4.85	7.67	$+0.17^{k}$	+1.65	+0.41	+1.77
Provolone	3	43.17	1.03	2.33	0.97	5.29	4.78	1.36	2.70	5.40	-1.10	-0.40	-0.88	+0.80
Provolone	3	44.08	1.58	3.46	0.98	5.15	5.19	1.55	3.78	7.19	-0.40	-0.80	-0.52	+0.81
Provolone (Reduced-Na)	6	42.93	1.05	2.39	0.98	5.15	4.95	1.24	6.25	7.79	-1.20	-0.31	-0.30	+0.62
Provolone (Reduced-Na)	6	44.09	1.02	2.26	0.98	5.28	5.12	1.62	5.73	7.39	-0.29	-0.27	-0.63	+1.59
Queso Blanco	7	47.02	3.30	6.56	0.96	6.37	6.11	0.26	4.38	6.78	+2.68	-1.07	-2.11	+2.57
Queso Fresco	7	51.19	1.85	3.49	0.98	6.49	5.05	0.31	4.86	8.68	+2.09	+3.02	+2.90	+1.55
String	14	47.91	1.98	3.97	-	5.44	4.96	1.59	4.87	8.86	+0.60	+2.00	+1.75	+2.39
String	6	47.07	2.18	4.43	-	5.33	5.02	1.67	4.85	8.65	$+0.22^{k}$	$+0.39^{k}$	-0.38	+1.58
Queso Quesadilla	7	43.10	2.18	4.81	0.97	5.35	5.39	1.21	4.57	6.29	-0.01	-0.57	-0.48	-0.40 ^m

pathogen survival ($\Delta \log CFU/g$) during storage for 15-days at 25°C.

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⁷⁶⁴ ^{*a*} Cheeses were national brands obtained from local retail outlets or directly from manufacturers. Qualifying descriptive information,

- res. 'mild,' 'sharp' is reproduced where provided on the package.
- 766 ^{*b*} Moisture content (%) of cheese sample on Day 0, n=2.
- 767 c % salt of cheese sample on Day 0, n=2.
- d % salt-in-moisture phase (%SMP) of cheese sample on Day 0. Calculated from % moisture and % salt of the same cheese.
- ^{*e*} Water activity (a_w) of cheese sample on Day 0.

- ^f pH of cheese slice surface on Day 0 and Day 15, n=2.
- ^{*g*} % titratable acidity (%TA) of cheese sample on Day 0 and Day 15, n=2.
- ^hDeMan-Rogosa-Sharpe (MRS) agar count for lactic acid bacteria (LAB) on Day 0 and Day 15 (log CFU/g), n=2.
- *i* Survival of pathogen LM=*L. monocytogenes*, SALM=*Salmonella* spp., EC=*E. coli* O157:H7, and SA=*S. aureus*. (+) indicates
 growth, (-) indicates no-growth.
- 775 j not determined.
- ^k Growth of pathogen did not exceed plating variability: 0.39, 0.41, 0.27, 0.25 log CFU/g for *L. monocytogenes, Salmonella* spp., *E. coli* O157:H7 and S. *aureus*, respectively
- ⁷⁷⁸ ^{*l*}Bolded numbers indicate growth beyond the pathogen-plating variability.
- ^{*m*} Growth (+ 0.57 log CFU/g) at Day 6 sampling; no net growth over 15-day storage period

780	Table 3.	Pathogen	strains	used in	laboratory	cheese cl	hallenge studie	s.

Inoculum	Serotype	Strain ^a	Collection ^b	Source ^c
Listeria	4b	LM 101	FRI	Hard salami
monocytogenes	4b	LM 310	FRI	Goat cheese
	4b	ATCC 43256	ATCC	Mexican-style cheese, Calif. (1985 outbreak strain)
	4b	ATCC 43257	ATCC	Mexican-style cheese, Calif. (1985 outbreak strain)
	4b	ATCC 51414	ATCC	Raw milk, Massachusetts
	4b	ATCC 51776	ATCC	Cheese, Belgium
	4b	ATCC 51777	ATCC	Cheese, Belgium
	4b	ATCC 51778	ATCC	Cheese, Belgium
	4b	Scott A	FRI	Clinical
	1/2a	V7	FRI	Raw milk
Salmonella spp.	Cerro	FSL R8-370	FSL	Bovine
	Typhimurium	FSL S5-433	FSL	Bovine
	Newport	FSL S5-436	FSL	Bovine
	Agona	FSL S5-517	FSL	Human
	Typhimurium	FSL S5-536	FSL	Human
	Newport	FSL S5-639	FSL	Human
Escherichia coli	O157:H7	FRIK 22	FRI	Unknown
O157:H7	O157:H7	FRIK 2000	FRI	Bovine
	O157:H7	F5854	FRI	Cheese curds (1998 outbreak strain)
	O157:H7	039732	NMDH	Gouda cheese (2010 outbreak strain)
	O157:H7	CWD EC1	VT	Farmstead goat cheese
Staphylococcus		Ι	FPL	Raw milk
aureus		J	FPL	Raw milk
		FRI 100	FRI	Cake
		FRI 1007	FRI	Genoa sausage
		ATCC 25923	ATCC	Clinical

781

^aStrain designation provided by Collection.

^bCollection: FRI = Food Research Institute, University of Wisconsin-Madison, Madison, Wisc.;

ATCC = American Type Culture Collection, Manassas, Va.; FSL = Food Safety Laboratory, Dr.

Katherine Boor, Cornell University, Ithaca, N.Y.; NMDH = New Mexico Department of Health,

Santa Fe, N.M.; VT = Vermont Institute for Artisan Cheese, Dr. D.J. D'Amico, University of

787 Vermont, Burlington, Vt.; FPL= Food Pathogen Laboratory, Dr. Barbara Ingham, University of

788 Wisconsin-Madison, Madison, Wisc.

789	^c Source provided by Collection.
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Reference	Pathogen	No. of strains	Cheese	Storage (days)	Temp (°C)	$\mathbf{p}\mathbf{H}^{a}$	%SMP ^b	a _w	Growth/Death
31	Salmonella	9	Queso Fresco	_d	20	6.60	1.64	-	LT ^e :2.5 - 3.5h GT: 1.65 - 2.17
30	<i>E. coli</i> O157:H7	2	Queso Fresco	-	20	6.60	1.61	-	LT: 3 - 3.45 h GT: 2.33 - 2.56
						5.06	1.70	0.98	-1.11
42	L.	F		20	21	5.30	1.80	0.97	-0.48
42	monocytogenes	5	Cheddar	30	21	5.66	5.00	0.95	-0.14
						5.28	4.80	0.95	-0.96
						5.06	1.70	0.98	-3.2
12	Salmonella	~		20	2.1	5.30	1.80	0.97	-3.9
43	spp.	5	Cheddar	30	21	5.66	5.00	0.95	-3.8
						5.28	4.80	0.95	-3.5
48	L. monocytogenes	5	Queso Blanco	6.25	25	6.80	4.53	0.97	> 5.00
			Queso Fresco	3	30	6.60	6.60	-	+0.39
			Queso Fresco	6	30	6.60	4.50	-	+0.95
			Queso Fresco	3	30	6.50	6.15	-	+0.74
			Queso Ranchero	1	30	6.20	4.10	-	+2.60
			Queso Panela	3	30	6.20	2.50	-	+1.81
			Queso Panela	1	30	6.70	3.95	-	+3.18
			Queso Panela	3	30	6.60	3.48	-	+0.79
			Cotija	8	30	5.60	9.60	-	> -2.00
	L.		Cotija	6	30	5.50	12.50	-	> -2.00
17	monocytogenes	5	Monterey Jack	4	30	5.00	3.00	-	> -1.40
			Monterey Jack	13	30	5.20	2.72	-	> - 2.09
			Mild Cheddar	4	30	4.90	2.60	-	> -1.26
			Mild Cheddar	7	30	5.20	4.49	-	> -2.09
			Colby	9	30	5.50	4.93	-	> -2.36
			String Cheese	9	30	5.50	4.24	-	> -2.36
			Provolone	9	30	5.60	4.62	-	> -2.36
			Muenster	9	30	5.50	3.80	-	> -2.36
			Domestic Feta	4	30	4.30	7.50	-	> -2.04
			Domestic Feta	4	30	4.30	2.20	-	> -2.04

807 Table 4. Data from published research selected to augment laboratory product assessment.

^{*a*} pH values of cheeses at initial sampling point of experiment

809	^b Certain publications stated %SMP as % brine, which was calculated using the same equation as
810	in this study (Equation 1). For publications that included both % moisture and % salt, % SMP
811	was calculated using Equation 1.
812	^c Behavior of pathogen over storage, expressed as $\Delta \log CFU/g$ or LT/GT.
813	^d Not specified.
814	^{<i>e</i>} LT: Lag time (h); GT: generation time (h).
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831 Figure legend.

- 832 Figure 1. Growth (Δ) or No-Growth (x) of Listeria monocytogenes, Salmonella spp.,
- 833 Escherichia coli O157:H7, and Staphylococcus aureus on cheeses stored at 20-30°C based on
- cheese pH (Day 0) and %SMP (salt-in-moisture-phase). Data from published research (n=26;
- Table 4) and this study (n=55). Solid line represents the growth/no-growth interface (P=0.05).

