

***STAPHYLOCOCCUS AUREUS* COLONIZATION AND FAMILIAL TRANSMISSION OVER A ONE YEAR PERIOD**

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Abstract

Methods: 177 adults and 86 minors comprising 95 family units were enrolled from two counties in Iowa and followed up for 52 weeks. Random effects logistic regression was used to test the effect of different risk factors on the probability of an individual falling into a different *S. aureus* colonization categories. Additionally, the frequency of *S. aureus* colonization events and familial transmission events were calculated.

Results: The number of positive environmental sites within a participant's house was associated with being a persistent carrier compared to being a non-carrier or intermittent carrier. Age, sharing bath towels, and the number of positive environmental sites within a participant's house were associated with being a persistent or intermittent carrier. Colonization events per year were 3.95 for adults and 3.04 for minors. Duration of colonization was longest for persistent carriers (92.3 days for adults and 97.8 days for minors), and intermittent carriers had the most colonization events.

Conclusions: The average duration of colonization was significantly different when comparing intermittent carriers and non-carriers. We have also established estimates of the duration of colonization and the frequency of transmission events among family units in a non-healthcare population.

Introduction

Staphylococcus aureus (*S. aureus*) is a commensal bacterium, with the nares historically considered the most common colonization site (1, 2). About 30% of the population is colonized with *S. aureus* (3), and most colonized persons are asymptomatic carriers. Asymptomatic carriage may not harm the host, but is a known risk for subsequent symptomatic infections in both the healthcare and community settings (4).

Numerous prior studies have assessed *S. aureus* colonization of the nares and oropharynx in the healthcare setting (5-9), while few have assessed colonization of these sites in a population of healthy community members (10-12). Moreover, we did not identify any published studies that estimated the frequency of colonization events or familial transmission of *S. aureus* in the absence of an infected family member.

This study assessed data on *S. aureus* colonization collected from a longitudinal cohort of 95 family units that was recruited from rural and urban Iowa and had minimal healthcare contact. Utilizing data from this cohort, we determined the incidence of colonization with *S. aureus*, the duration of colonization, the frequency of familial transmission, and the incidence of infections. We also, characterized the colonizing *S. aureus* strains, and we assessed risk factors for colonization.

Methods

Study design and sample collection

Participants were enrolled in a prospective cohort study. Ninety-five families from two counties in Iowa, Johnson County and Keokuk County, were enrolled between October 6th, 2011 and January 4th 2012. These two counties were chosen to reflect an urban (Johnson County) and a rural population (Keokuk County) in Iowa, as defined by the US Census Bureau (13).

County-specific methodologies were employed to recruit potential participants. Residents in Johnson County were recruited via advertisements in a local newspaper and

by mailing lists. Interested persons contacted our study team via email. Participants from Keokuk County were recruited from the Keokuk Rural Health Study, a previously existing cohort (14), and were contacted via letter; the enrollment study visit was scheduled via subsequent phone call.

Trained study team members traveled to each family's home to enroll the family members. During the study visit, potential participants were screened to assess enrollment criteria, instructed on sample collection techniques, and completed enrollment interviews. Inclusion criteria for enrollment in the cohort were: participant must be able to provide consent, assent, or have parents willing to provide consent; and must be willing to participate by completing enrollment and weekly follow-up questionnaires. All participants signed an informed consent document or assent document prior to completing any study activities. The University of Iowa institutional review board approved all study protocols prior to recruitment.

Participants completed questionnaires collecting data on demographics, healthcare exposure, farming exposure, medical conditions, and other risk factors for *S. aureus* colonization and infection. When collecting weekly samples, participants also completed a questionnaire about potential skin and soft tissue infections.

Isolation of *S. aureus*

While at each family's home, study team members instructed each enrolled family member on the proper technique for collecting nasal samples. Adult participants were instructed on providing oropharyngeal samples as well. Participants were also given BBL CultureSwabs so they could sample possible *S. aureus* infections. Samples were collected with BBL CultureSwabs that included Liquid Stuart Medium (Becton, Dickinson and Company, Sparks MD, USA). Samples were transported through the US Postal system with icepacks to the Center for Emerging Infectious Diseases (CEID) laboratory facilities.

Study team members collected environmental swabs during the enrollment visit using 3-inch by 4-inch sterile duster cloths. Six commonly touched surfaces (living room television remote, main bathroom toilet flush lever, main bathroom light switch, kitchen sink handle, refrigerator door handle, and oven knobs) were sampled by wiping the surfaces in all directions for one minute. Each site was sampled with a new duster cloth, and the cloth was placed into a dry sterile bag. The samples were transported to the CEID on ice.

Nasal and oropharyngeal swabs were inoculated separately into 5mL of Baird-Parker medium and incubated at 35°C for 24 hours. Environmental samples were processed by adding 25mL of 1.0% peptone to the bag and homogenizing by hand for one minute. 5mL of the 1.0% peptone was added to 5mL of 2x Baird-Parker medium and incubated at 35°C for 24 hours. Following incubation, isolates were plated onto Baird-Parker agar and BBL CHROMagar MRSA II (Becton, Dickinson and Company, La Jolla CA, USA) and incubated for 48 hours at 35°C. Presumptive *S. aureus* colonies (mauve colonies on CHROMagar MRSA II plates, and black colonies with shiny halos on Baird-Parker agar) were streaked onto Columbia CNA agar with 5% sheep's blood (Becton, Dickinson and Company, Sparks MD, USA) and incubated for 24 hours at 35°C.

Characterization of *S. aureus*

Presumptive *S. aureus* colonies grown on Columbia CNA agar with 5% sheep's blood were assessed with the catalase test, coagulase test, and Pastorex Staph Plus rapid latex agglutination assay (Bio-Rad, Redmond WA, USA). All isolates confirmed to be *S. aureus* had genomic DNA extracted for molecular testing.

Antimicrobial susceptibility testing

Phenotypic resistance to antimicrobials was assessed through minimum inhibitory concentration (MIC) testing following the Clinical and Laboratory Standards Institute's standards (15). Isolates were tested for susceptibility to oxacillin, tetracycline,

erythromycin, clindamycin, trimethoprim-sulfamethoxazole, gentamycin, levofloxacin, vancomycin, daptomycin, quinupristin/dalfopristin, linezolid, and rifampin.

Molecular testing

Bacterial genomic DNA was extracted with the Promega Wizard Genomic DNA purification kit (Promega Corporation, Madison WI, USA). Polymerase chain reaction (PCR) was used to assess the presence of the *mecA* gene (16), and the genes encoding Panton-Valentine leukocidin (PVL) (17), and to amplify the protein A (*spa*) gene (18). Strain type was identified through typing the *spa* gene using the Ridom StaphType software (Ridom GmbH, Germany). Isolates that could not be typed three times were labeled as non-typable (NT) (19). The Based Upon Repeat Pattern (BURP) algorithm was used to identify and characterize genetic clusters (20). All molecular procedures included known positive and negative controls.

Statistical analysis

Data were analyzed using SAS software version 9.3, and R version 2.15.3. Demographic data were assessed for differences between the two counties using the Student's T-Test to assess age, Chi-Squared Test without Yate's Continuity Correction to assess gender, and Fisher's Exact Test to assess dichotomous variables with small sample sizes.

Participant carrier status was determined by taking the number of nares and oropharyngeal swabs that grew *S. aureus* divided by the total number of swabs over the study period, excluding the baseline cultures and culture results. The cutoffs utilized in this manuscript were previously described by van Belkum et al., but were modified to include the oropharynx in the definition (21). Participants were classified as: 'non-carriers' if $\leq 10\%$ of samples from both the nares and the oropharynx grew *S. aureus*, 'intermittent carriers' if between 10% and 80% of samples from either the nares or the oropharynx grew this organism, and 'persistent carriers' if $\geq 80\%$ of samples from either the nares or the oropharynx grew *S. aureus*.

Logistic regression with receiver operating curves (ROC) was used to determine the lowest number of nares and oropharynx swabs necessary to accurately predict each participant's carrier status. For adults, the number of culture positive nares swabs and the number of culture positive oropharynx swabs were modeled as separate variables. The ROC analysis was completed separately for adults and minors, as minors submitted nares swabs only, to generate true positive rates (TPR) and false positive rates (FPR).

Random effects logistic regression using SAS proc glimmix with Satterthwaite's approximation for degrees of freedom was used to test the effect of risk factors on: the probability of being an intermittent carrier given one is not a persistent carrier, the probability of being a persistent carrier compared with non-carriers and intermittent carriers, and the probability of being an intermittent carrier or persistent carrier compared with non-carriers. Number of children, age, and house size were modeled as continuous variables. All other risk factors were modeled as dichotomous or continuous variables. Risk models for adults were adjusted for age; models for minors were adjusted for age, gender, and an age and gender interaction term. Random effects logistic regression was used to assess the random effects of both family units and county in addition to the effects of specific risk factors. Some models had a non-positive variance for county after adjustment for the random effect of family and the risk factor, indicating that the remaining variance was not attributable to county, and thus the variance was set to zero in these instances (22).

Colonization events and familial transmission events were identified based upon the following definitions. A colonization event was defined as a positive nares or oropharyngeal culture (i.e., acquisition of carriage), when the participant's samples from the week prior were negative in both the nares and the oropharynx. For each colonization event, the duration of colonization was calculated as the time from the beginning of the colonization event (i.e., the first positive culture after a prior negative set of nares and oropharyngeal cultures) until the time cultures of both the nares and oropharynx swabs

were negative concurrently. If a participant was colonized during the first week of observation, or during the final week of observation, left truncation and right truncation of the interval were used, respectively. All other colonization durations were calculated with interval censoring. For example, if a participant's nares and the oropharyngeal cultures were negative during one sampling, and were positive at the next two samplings seven days apart, and were negative seven days after that, half of the first interval and half of the last interval (3.5 days each) would be added to the 14 days between the two sets of positive cultures for a total colonization duration of 21 days. If a participant began the study with a positive culture, had a positive culture seven days later, and had two negative cultures seven days after that, the colonization duration would be 17.5 days. The patient would accrue seven days for the positive culture at enrollment, seven days for the second positive culture, and 3.5 days for the right truncation. The duration would be left truncated because we do not know how many weeks prior to our study the participant had positive cultures. We used these censoring and truncation methodologies estimate the duration of colonization most accurately without overestimating this time period. Familial transmission was defined as the acquisition of colonization with a specific strain of *S. aureus* found concurrently within a family member. Concurrent colonization was defined as a current culture from a family member that grew a specific strain of *S. aureus* that was also currently colonizing a different family member, indicating two family members were colonized by the same strain at the same time. The familial transmission event occurs when the second family member acquires the strain.

Results

The logistic regression models with ROC curves indicated the minimum number of swabs necessary to maximize the TPR while minimizing the FPR for adults was 14 consecutive sets of swabs for adults, and a single swab for minors. Of the 177 adults, 154 (87.0%) submitted 14 or more swabs after the baseline swab and were included in this

analysis. Of the 86 minors, 77 (89.5%) continued past the baseline and were included in analysis. Adults submitted an average of 34.71 swabs (range: 14-52) and minors submitted an average of 30.87 swabs (range: 2-46). The average age of the adult participants was 44.5 (23.10-67.72 years), while the average age of minors was 9.65 years (0.42-18.18 years). Additional demographics information is included in the supplemental Tables C2 and C3.

A total of 10,987 swabs were processed during follow-up, 3,119 of which were culture positive for *S. aureus*. Among adults, 29.8% of cultures from at least one anatomical site grew methicillin-susceptible *S. aureus* (MSSA), and 1.3% of cultures grew MRSA. Among minors, 30.4% of nares cultures grew MSSA, and 0.6% of nares cultures grew MRSA. For additional details on prevalence, see Table 1. Of the adults, 47.4% met the definition of non-carrier, 32.2% met the definition of intermittent carrier, and 20.4% met the definition of persistent carrier (Table 3, Figure 1). Additional details on site-specific prevalence can be found in Table S1.

One hundred twenty-five *spa* types were identified amongst the 3,119 *S. aureus* isolates, of which, the most common were t002 (n=345), t008 (n=248), and t216 (n=180). Adjustment for family clonal isolates was done by taking a single representative sample of each clonal strain from each family unit. Following adjustment, the number of unique isolates was reduced to 278. The most common *spa* types following adjustment were t002 (n=23), t084 (n=17), t008 (n=16), and t216 (n=16). The BURP algorithm identified 10 clusters in Keokuk County and seven clusters in Johnson County, but the number of *spa* types per cluster was higher in Johnson County than in Keokuk County (Appendix Figures S1 and S2). After adjustment, *mecA* prevalence was 5.4%, and PVL prevalence was 1.8% (Table 2).

Antimicrobial susceptibility testing was performed on a subset of 725 isolates obtained from samples received during the first week of each study month. Observed rates of resistance were zero for most tested antibiotics. However, 25.9% of isolates were

resistant to erythromycin after adjustment and 17.7% were intermediately resistant to quinupristin/dalfopristin. Resistance rates to other agents were all under 10% (Figure 3).

Random effects logistic regression was used to assess the probability of being an intermittent carrier given that one was not a persistent carrier, and found significant associations with age (p-value: 0.0145) and with the number of positive environmental sites (p-value: 0.0137). Farming exposure did not quite meet the 0.05 significance level (p-value of 0.0733). The number of positive environmental sites was the only factor associated with being a persistent carrier compared with being a non-carrier or an intermittent carrier (p-value: 0.0137). Age (p-value: 0.0353), sharing bath towels (p-value: 0.0338), and the number of positive environmental sites (p-value: 0.0026) were associated with being either a persistent carrier or an intermittent carrier, compared with being a non-carrier. Data on the directionality of associations are presented in Tables 5 and 6.

The average number of observed colonization events was similar for adults (3.95) and minors (3.04). Among both adults and minors, intermittent carriers had the greatest number of observed colonization events. Duration of colonization was longest among persistent carriers: 92.3 and 97.8 days for adults and minors, respectively. Both adults and minors acquired strains carried by a family member (familial transmission), with an average of 0.77 events per person year of follow-up for adults, and 1.12 events for minors (Table 4). Participants reported skin and soft tissue infections during study follow-up, with six participants (four adults and two minors) reporting eight potential infections. Five wound swabs were submitted for culture, one (20%) of which grew *S. aureus*. The strain isolated from the wound and the participant's current colonizing strain had the same spa type. The observed rate of putative infections per person year of follow-up was 0.049 for adults and 0.060 for minors.

Discussion

Most investigators assessing *S. aureus* colonization have utilized data on nasal colonization to assign individual persons to one of three categories with regard to carriage: persistent carriers, intermittent carriers, and non-carriers (4, 23, 24). Investigators who utilize only nasal colonization to assess carrier state could misclassify oropharynx-only carriers as non-carriers. In addition, investigators could misclassify individual persons if the time period over which cultures are obtained is too short. Using the data from this manuscript, we estimate that 9 persistent carriers and 23 intermittent carriers (32/154, 20.8%) would have been misclassified if nasal cultures alone were used to identify *S. aureus* carriage (Figure 1). Unrecognized *S. aureus* carriers may transmit *S. aureus* to other persons and, thus, may ‘render infection-control programs futile’ (25). Additionally, Robinson et al. found that knowing the colonization status of a patient from a prior visit was associated with a reduced 30-day mortality from MRSA bacteremia (26), which the investigators felt might reflect the earlier use of glycopeptides in the group with previous MRSA colonization. Thus, their data support the idea that screening some patients to determine their *S. aureus* colonization status may improve outcomes in some patient Populations.

All cultures from 19 of 152 adults (12.5%) assessed in this study were negative for *S. aureus*. However, most (87.5%) participants submitted at least one sample that grew *S. aureus*, indicating most (if not all) people could be colonized. A person’s colonization status may depend upon the extent of contamination in their environment, or on exposure to persons carrying *S. aureus*, rather than the person’s biological susceptibility to the pathogen. The number of positive environmental sites, and farming exposure, (which nearly met the 0.05 significance level) both assess environmental pressures and level of exposure to *S. aureus*. The number of contaminated environmental sites was significantly associated with *S. aureus* carriage in all analyses for adults, and with being an intermittent carrier when compared with non-carriers for minors. Both of

these variables support the hypothesis that that given adequate exposure, any person could be colonized.

Van Belkum et al. inoculated *S. aureus* into participant's nares and observed that those classified as non-carriers and intermittent carriers lost *S. aureus* colonization on average 4 and 14 days post colonization event, respectively (21). Their results suggest that persons we classified as intermittent or non-carriers may have acquired and cleared *S. aureus* between study samplings. We also found that non-carriers had the longest time without colonization, but also had the fewest colonization events, suggesting that environmental pressure for these participants was low.

Given our definitions, non-carriers must have infrequent colonization events and a short duration of colonization for each event, and persistent carriers must have extended durations of colonization, and fewer colonization events. Of interest, however, is the number of colonization events and average duration of colonization for intermittent carriers. Van Belkum et al. previously showed that non-carriers and intermittent carriers have similar levels of antistaphylococcal antibodies, and that the duration of colonization is not statistically different for the two groups following nasal challenge with *S. aureus* (21). However, in our study, the average duration of colonization for intermittent carriers and non-carriers were significantly different for both adults and minors (Table 4). A distribution demonstrated that the duration of colonization for a majority of participants was clustered between 0 and 25 days, which suggests that a majority of intermittent carriers may be similar to non-carriers (Figure S3). In contrast, some intermittent carriers had an average colonization duration longer than 25 days and may be similar to persistent carriers. For example, one participant had 3 colonization events with an average duration of 85.8 days. This participant did not meet the definition of persistent carriage because he/she had at least one positive culture at 79% of samplings, thus not meet the cutoff of $\geq 80\%$ required to be categorized as a persistent carrier. Thus, our results suggest that the participants we categorized as intermittent carriers may be a mixed population, some

whose carriage dynamics are more like non-carriers and some whose colonization dynamics are more like persistent carries.

Only one swab obtained from putative infections grew *S. aureus* when cultured. This sample was the only one collected by a participant's physician, suggesting that participants may have obtained inadequate samples because we did not teach participants to obtain samples from possible infections. Alternative explanations for these results are that these participants had infections caused by pathogens other than *S. aureus*, or that they had non-infectious skin lesions.

The observed oxacillin resistance levels using genotypic and phenotypic assays are similar for both unadjusted (2.7 vs. 2.07) and adjusted methodologies (5.4 vs. 4.71), indicating the subset of isolates submitted for MIC testing was likely representative of the entire set of 3,185 isolates. Overall, the levels of antimicrobial resistance for the adjusted isolate set were low, with resistance levels of less than 10% for all but two antimicrobials.

The minimum spanning tree revealed that patterns of *spa* types were similar for the two counties (Figure 2). We did not observe differences amongst the lineages, suggesting the same strains circulate within both populations. This observation is similar to the observation by Melles et al. that the *S. aureus* population structures in the United States and in Denmark are similar (27).

We found a significant inverse relationship of age with intermittent carriage given one is not a persistent carrier. Thus, older participants had a lower probability of being an intermittent carrier. For example, the probability of a 30-year-old participant being colonized (probability: 0.5439) was higher than that a participant who was 60 (probability: 0.2657). We also found a similar inverse association of age and colonization (defined as being an intermittent carrier or persistent carrier).

This study had several strengths. We enrolled family units and we enrolled persons from urban and rural settings, which allowed us to adjust for random variability,

and to observe familial transmission events. Longitudinal follow-up, with an average of 34.7 swab sets for adults and 30.9 nasal swabs for minors, allowed us to estimate the frequency of transmission events and of symptomatic infection. Delacour et al. previously showed that *S. aureus* can be viable 18 days after collection in non-charcoal liquid transport medium (28). In addition, we trained participants to swab their nares and oropharynges. Thus, we maximized our ability to detect *S. aureus* colonization and minimized participant inconvenience.

This study had potential limitations. Patients reported symptomatic SSTIs and they obtained wound cultures. Additionally, only one of five patients who submitted wound cultures had a confirmed *S. aureus* infection. Thus, we may have misclassified noninfectious skin lesions as infections or we may have misclassified infections caused by other organisms as *S. aureus* infections. However, *S. aureus* is the most common cause of SSTIs seen in emergency departments (29), suggesting our self-reported SSTIs may be classified correctly as *S. aureus* infections.

An additional limitation is that our study duration of up to 52 weeks may not have been adequate to observe colonization and transmission events for persistent carriers. Our follow-up period far exceeds the timeframe observed by van Belkum et al. that colonization duration was 4 days for non-carriers and 14 days for intermittent carrier (21), allowing us to observe multiple colonization and decolonization events for both non-carriers and intermittent carriers. We censored and truncated data when we calculated the duration of colonization so our estimates of duration of colonization would be as accurate as possible while accounting for time between samples. Because we utilized left and right truncation, we most likely underestimated the true duration of colonization but this method minimized the assumptions required for the calculations and allowed us to establish the most accurate estimates of duration of colonization with the data available.

In conclusion, 20.8% of participants would have been misclassified if oropharyngeal samples were excluded, indicating that the oropharynx is an important screening site for persons from the healthy community who present to a healthcare setting and may be at risk of a *S. aureus* infection. We also found that the average duration of colonization were significantly different for intermittent carriers and non-carriers, which is in contrast to the findings by van Belkum et al. (21). However, intermittent carriers may be a mixed population with some people having carriage dynamics more like non-carriers and some having carriage dynamics more like persistent carriers. Further studies are needed to assess the frequency of colonization at other anatomical sites among healthy persons in the community. In addition, a study that uses biologic markers and does not involve decolonization and subsequent challenge with *S. aureus* could help elucidate the differences between non-carriers, intermittent carriers, and persistent carriers.

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Declaration on interests

The authors have no conflicts of interest.

Table 1: Total number of sets of oropharyngeal and nasal cultures that grew MSSA or MRSA, age grouped by county

	MSSA	MRSA*	Total
Adults			
Johnson Co.	820/2228 (36.8%)	12/2228 (0.5%)	832/2228 (37.3%)
Keokuk Co.	849/2904 (29.2%)	55/2904 (1.9%)	904/2904 (31.1%)
Total	1669/5132 (32.5%)	67/5132 (1.3%)	1736/5132 (33.8%)
Minors			
Johnson Co.	405/1292 (31.3%)	12/1292 (0.9%)	417/1292 (32.3%)
Keokuk Co.	294/1052 (27.9%)	2/1052 (0.2%)	296/1052 (28.1%)
Total	699/2344 (29.8%)	14/2344 (0.6%)	713/2344 (30.4%)

Note: The numerator for adults was the number of participants with *S. aureus* isolated from the nares, the oropharynx, or both. The numerator for minors was the number of participants with *S. aureus* isolated from the nares. The denominator is the total number of participants by county for adults and minors.

*MRSA isolates are designated by the presence of the *mecA* gene.

Table 2: Molecular characteristics of *S. aureus* isolates

	<i>mecA</i>		PVL	
	Adjusted	Unadjusted	Adjusted	Unadjusted
Johnson Co.	5/156 (3.2%)	24/1569 (1.5%)	0/156 (0%)	0/1569 (0%)
Keokuk Co.	10/122 (8.2%)	57/1550 (3.7%)	5/122 (4.1%)	26/1550 (1.7%)
Total	15/278 (5.4%)	81/3119 (2.6%)	5/278(1.8%)	26/3119 (0.8%)

Note: Unadjusted values include all *S. aureus* isolates from participants broken down by county. Adjusted values include only a single copy of isolates with the same molecular profile (same *spa* type, and presence/absence of *mecA* and PVL) to account for familial clustering and the presence of clonal isolates in the nares and oropharynx of adult participants.

Table 3: Participants by carrier group

	Johnson Co. (n=73)	Keokuk Co. (n=79)	Total (n=152)
Non-carrier	28 (38.36%)	44 (55.70%)	72 (47.37%)
Intermittent carrier	29 (39.73%)	20 (25.32%)	49 (32.24%)
Persistent carrier	16 (21.92%)	15 (18.99%)	31 (20.39%)

Note: The cutoffs used are as follows: non-carrier - $\leq 10\%$ of cultures from both the nares and the oropharynx were positive for *S. aureus*. Intermittent carrier - between 10% and 80% of cultures were positive from both the nares and the oropharynx. Persistent carrier - $\geq 80\%$ of cultures were positive from either the nares or the oropharynx.

Table 4: Colonization events and familial transmission events for adults and minors

	Number of Colonization Events per Person Year Follow-up	Average Colonization Duration (days)	Familial Transmission Events per Person Year Follow-up
Adults			
Non-carrier	1.56	10.77	0.33
Intermittent Carrier	7.72	20.69	1.31
Persistent Carrier	3.64	92.30	0.92
All Adults	3.95	31.71	0.77
Minors			
Non-carrier	1.24	7.07	0.62
Intermittent Carrier	5.39	25.71	1.89
Persistent Carrier	3.37	97.79	1.57
All Minors	3.04	35.31	1.22

Note: The cutoffs used are as follows: non-carrier - $\leq 10\%$ of cultures from both the nares and the oropharynx were positive for *S. aureus*. Intermittent carrier - between 10% and 80% of cultures were positive from both the nares and the oropharynx. Persistent carrier - $\geq 80\%$ of cultures were positive from either the nares or the oropharynx.

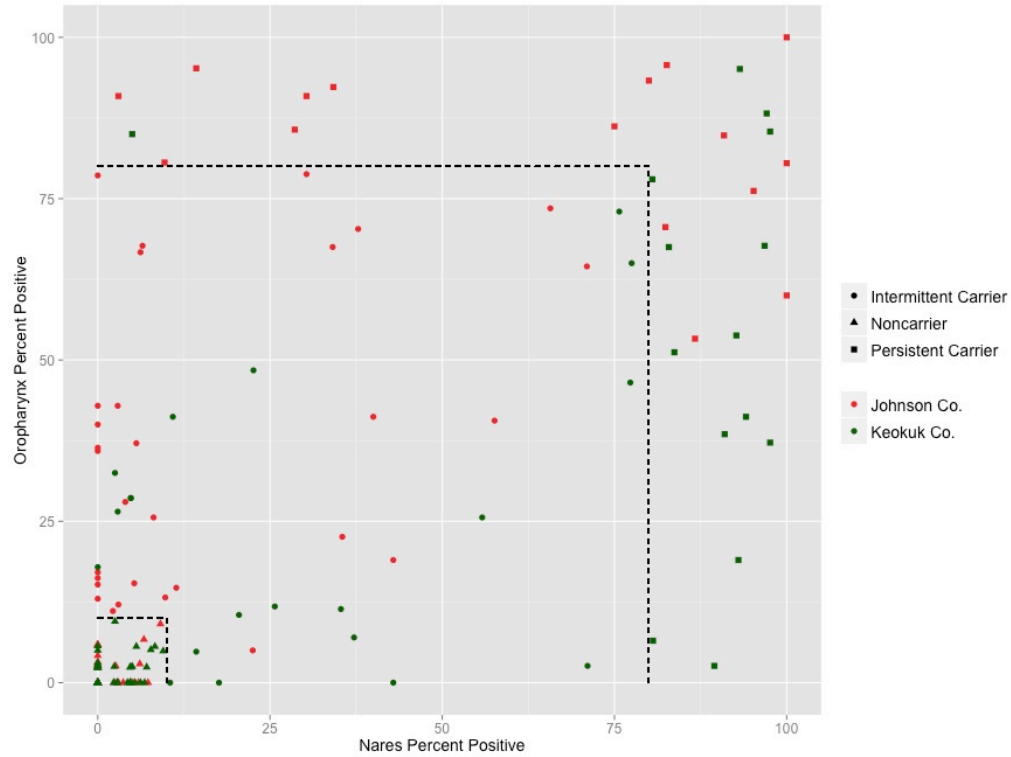


Figure 1: Scatterplot of participant nares and oropharynx colonization rates. Each circle denotes a single participant. Black-hashed lines indicate cutoffs between non-carriers, intermittent carriers, and persistent carriers (21).

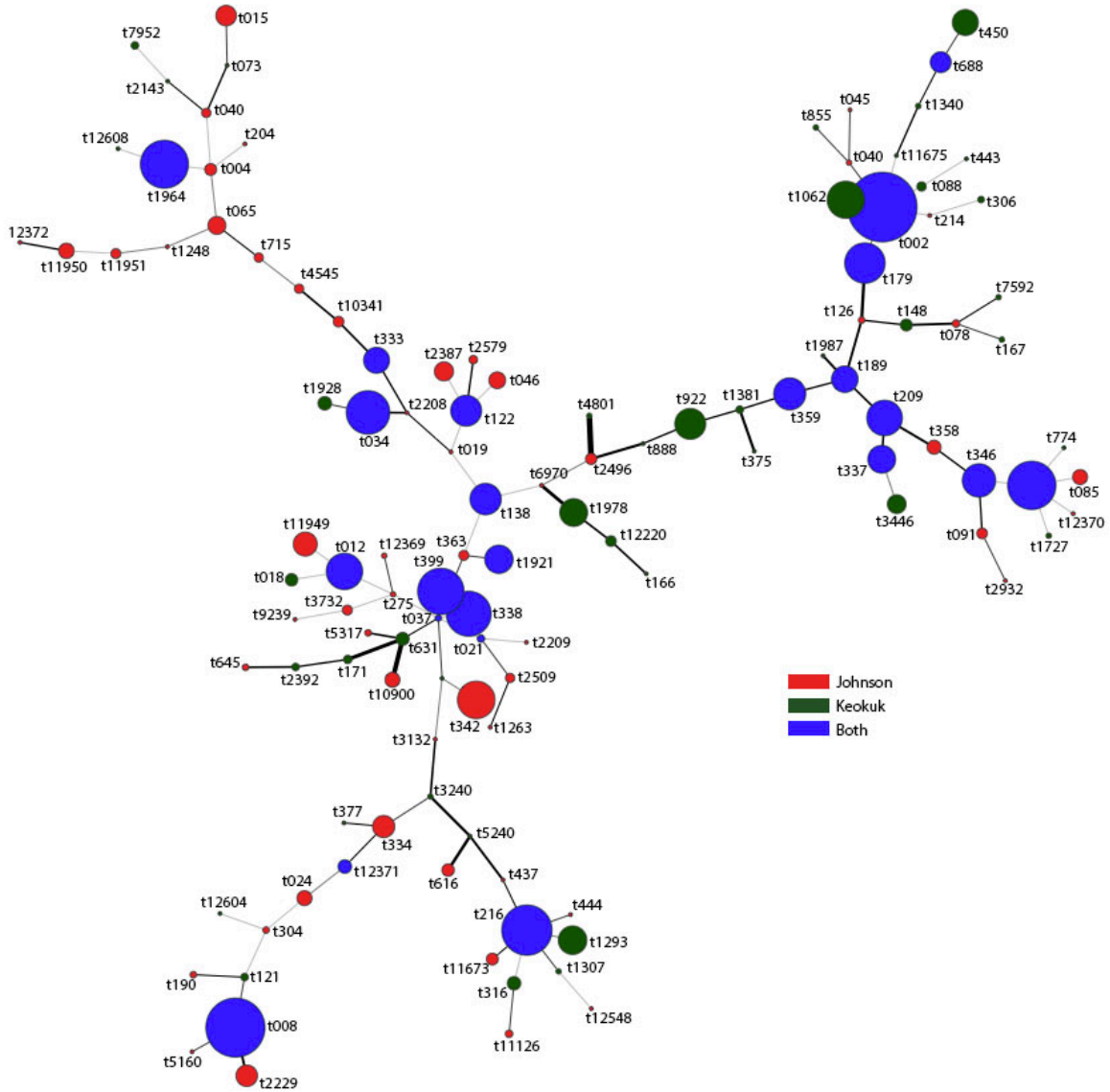


Figure 2: Minimum spanning tree of *spa* type diversity among 147 isolates comprising 57 *spa* types by county. The size of the circle is proportional to the number of isolates within the set; the thickness of connecting lines is proportional to the genetic distance.

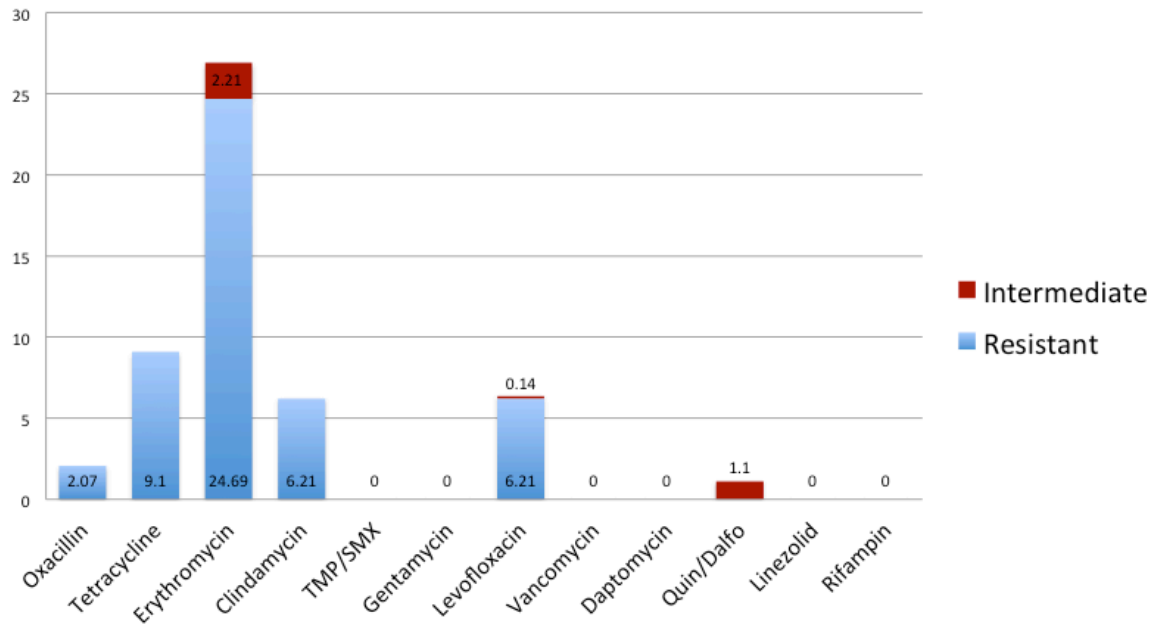


Figure 3a

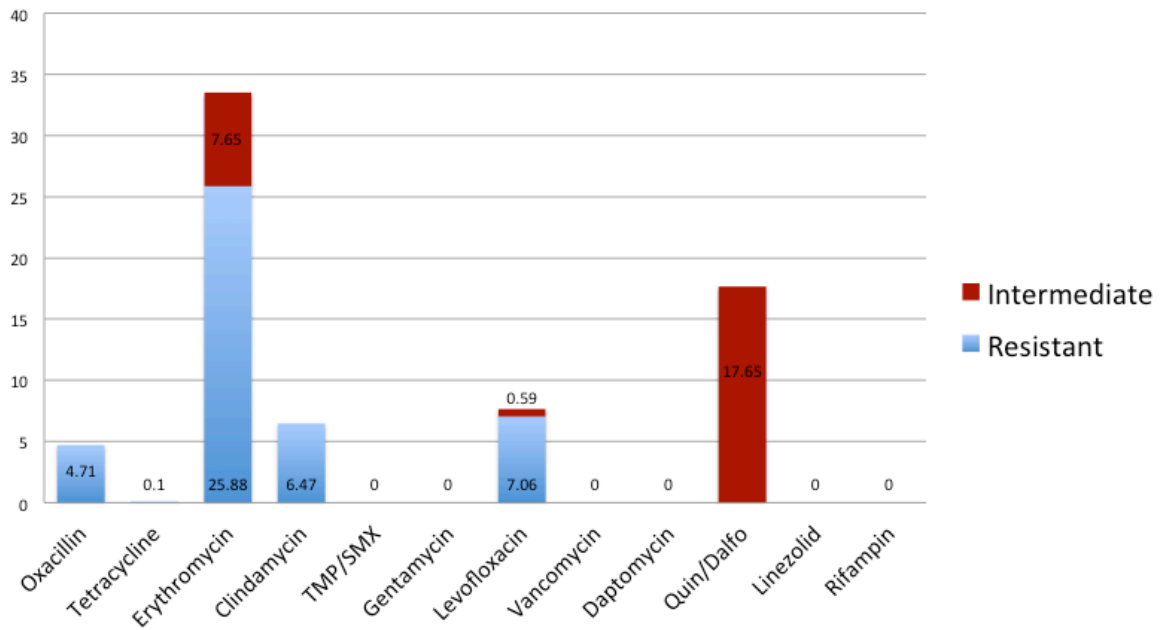


Figure 3b

Figure 3: Antimicrobial susceptibility to a panel of antimicrobials tested via minimum inhibitory concentration. A. Unadjusted results (n=725). B. Adjusted results (n=170).

Table 5: Trending and significant participant demographics and risk factors

Risk Factor [†]	Number (Percent %)		Total (N)	Probability	95% CI	p-value
	Non-carrier	Intermittent Carrier				
Intermittent Carrier vs. Non-carrier				Intermittent Carrier (at age=44.8)		
Age*						
Continuous	---	---	119	---	---	0.015
Any Family Member Play Team Sports						
No	47 (66.20)	24 (33.80)	71	0.33	0.22-0.47	
Yes	19 (47.50)	21 (52.50)	40	0.53	0.36-0.69	0.081
Number of Positive Environmental Sites						
Zero	57 (64.77)	31 (35.23)	88	0.16	0.10-0.22	
One	10 (55.56)	8 (44.44)	18	0.23	0.16-0.31	
Two	4 (40.00)	6 (60.00)	10	0.32	0.20-0.47	
Three	0 (0.00)	1 (100.00)	1	0.43	0.23-0.66	
Four	0 (0.00)	1 (100.00)	1	0.55	0.25-0.82	
Five	0 (0.00)	1 (100.00)	1	0.66	0.28-0.91	0.014

Table 5 Continued

Farming Exposure ^s							
	No Exposure	42 (58.33)	30 (41.67)	72	0.39	0.004-0.99	
	Non-occupational Exposure	14 (77.78)	4 (22.22)	18	0.20	0.05-0.55	
	Non-large Facility Occupational Exposure	2 (66.67)	1 (33.33)	3	0.42	0.05-0.92	
	Large Facility Occupational Exposure	2 (25.00)	6 (75.00)	8	0.82	0.39-0.97	0.073
Non- or Intermittent Carrier vs. Persistent Carrier		Non- or Intermittent Carrier	Persistent Carrier		Persistent Carrier (at age=44.8)		
Number of Positive Environmental Sites							
	Zero	88 (83.81)	17 (16.19)	105	0.16	0.10-0.23	
	One	18 (85.71)	3 (14.29)	21	0.23	0.16-0.31	
	Two	10 (62.50)	6 (37.50)	16	0.32	0.20-0.47	
	Three	1 (33.33)	2 (66.67)	3	0.43	0.23-0.66	
	Four	1 (50.00)	1 (50.00)	2	0.55	0.25-0.82	
	Five	1 (50.00)	1 (50.00)	2	0.66	0.28-0.91	0.014

Table 5 Continued

Non-carrier vs. Intermittent and Persistent Carrier	Non-carrier	Any Carrier	Any Carrier (at age=44.8)			
Age*						
Continuous	---	---	149	---	---	0.035
Gender						
Females	43 (53.09)	38 (46.91)	81	0.46	0.34-0.58	
Males	26 (39.39)	40 (60.61)	66	0.62	0.48-0.73	0.072
Share Bath Towels						
No	41 (41.84)	57 (58.16)	98	0.60	0.48-0.70	
Yes	30 (58.82)	21 (41.18)	51	0.38	0.24-0.54	0.034
Number of Positive Environmental Sites						
Zero	57 (54.29)	48 (45.71)	105	0.43	0.33-0.53	
One	10 (47.62)	11 (52.38)	21	0.63	0.51-0.74	
Two	4 (25.00)	12 (75.00)	16	0.80	0.61-0.91	
Three	0 (0.00)	3 (100.00)	3	0.90	0.68-0.98	
Four	0 (0.00)	2 (100.00)	2	0.96	0.75-0.99	
Five	0 (0.00)	2 (100.00)	2	0.98	0.81-0.99	0.003

Note: Bivariable analysis was conducted using random-effects logistic regression, with random effects assessed at both the family and county level. Only significant variables and those trending towards significance (<0.09) we included in this table.

¶All risk factors adjusted for age at 44.8 years

*Mean age for non-carriers is 47.34. Mean age for intermittent carriers is 41.09. Mean age for persistent carriers is years 44.3.

§Farming exposure refers to contact with livestock (swine, cattle, poultry, etc.)

Table 6: Trending and significant participant demographics and risk factors for minors

Risk Factor [¶]	Number (Percent %)		Total (N)	Probability	95% CI	P-value
	Non-carrier	Intermittent Carrier				
Non-carriers vs. Intermittent Carriers				Intermittent Carrier (at age=9.65)		
Race						
	Caucasian	29 (50.90)	28 (49.10)	57	---	---
	Other	6 (100.00)	0 (0.00)	6	---	0.033 [#]
Non- or Intermittent Carrier vs. Persistent Carrier	Non- or Intermittent Carrier		Persistent Carrier		Persistent Carrier (at age=9.65)	
Age [*]						
	Continuous	---	---	74	---	0.033

Table 6 Continued

Number of Positive Environmental Sites						
Zero	45 (91.84)	4 (8.16)	49	0.024	0.006-0.53	
One	7 (77.78)	2 (22.22)	9	0.046	0.001-0.78	
Two	5 (71.43)	2 (28.57)	7	0.085	0.001-0.926	
Three	2 (100.00)	0 (0.00)	2	0.15	0.001-0.96	
Four	3 (75.00)	1 (25.00)	4	0.26	0.005-0.961	
Five	1 (33.33)	2 (66.67)	3	0.40	0.014-0.97	0.041

Note: Bivariable analysis was conducted using random-effects logistic regression, with random effects assessed at both the family and county level. Only significant variables and those trending towards significance (<0.09) we included in this table.

[¶]All risk factors adjusted for age (at 9.65 years), gender, and an age-gender interaction term.

*Mean age for non-carriers is 9.35. Mean age for intermittent carriers is 8.80. Mean age for persistent carriers is 12.79.

#Fisher’s Exact Test used to determine p-value. Probabilities and 95% CIs could not be determined for these risk factors due to small sample sizes.

Works Cited

1. Kluytmans J, van Belkum A, Verbrugh H. Nasal carriage of *Staphylococcus aureus*: epidemiology, underlying mechanisms, and associated risks. *Clin Microbiol Rev.* 1997 Jul;10(3):505-20. PubMed PMID: 9227864. Pubmed Central PMCID: PMC172932. Epub 1997/07/01. eng.
2. Nouwen JL, Ott A, Kluytmans-Vandenbergh MF, Boelens HA, Hofman A, van Belkum A, et al. Predicting the *Staphylococcus aureus* nasal carrier state: derivation and validation of a "culture rule". *Clin Infect Dis.* 2004 Sep 15;39(6):806-11. PubMed PMID: 15472812. Epub 2004/10/09. eng.
3. Gorwitz RJ, Kruszon-Moran D, McAllister SK, McQuillan G, McDougal LK, Fosheim GE, et al. Changes in the prevalence of nasal colonization with *Staphylococcus aureus* in the United States, 2001-2004. *J Infect Dis.* 2008 May 1;197(9):1226-34. PubMed PMID: 18422434. Epub 2008/04/22. eng.
4. Wertheim HF, Melles DC, Vos MC, van Leeuwen W, van Belkum A, Verbrugh HA, et al. The role of nasal carriage in *Staphylococcus aureus* infections. *Lancet Infect Dis.* 2005 Dec;5(12):751-62. PubMed PMID: 16310147. Epub 2005/11/29. eng.
5. Bignardi GE, Lowes S. MRSA screening: throat swabs are better than nose swabs. *J Hosp Infect.* 2009 Apr;71(4):373-4. PubMed PMID: 19215999. Epub 2009/02/14. eng.
6. Bitterman Y, Laor A, Itzhaki S, Weber G. Characterization of the best anatomical sites in screening for methicillin-resistant *Staphylococcus aureus* colonization. *Eur J Clin Microbiol Infect Dis.* 2010 Apr;29(4):391-7. PubMed PMID: 20111880. Epub 2010/01/30. eng.
7. Collins J, Raza M, Ford M, Hall L, Brydon S, Gould FK. Review of a three-year methicillin-resistant *Staphylococcus aureus* screening programme. *J Hosp Infect.* 2011 Jun;78(2):81-5. PubMed PMID: 21507518. Epub 2011/04/22. eng.
8. Harbarth S, Schrenzel J, Renzi G, Akakpo C, Ricou B. Is throat screening necessary to detect methicillin-resistant *Staphylococcus aureus* colonization in patients upon admission to an intensive care unit? *J Clin Microbiol.* 2007 Mar;45(3):1072-3. PubMed PMID: 17229852. Pubmed Central PMCID: PMC1829122. Epub 2007/01/19. eng.
9. Marshall C, Spelman D. Re: is throat screening necessary to detect methicillin-resistant *Staphylococcus aureus* colonization in patients upon admission to an intensive care unit? *J Clin Microbiol.* 2007 Nov;45(11):3855. PubMed PMID: 17728478. Pubmed Central PMCID: PMC2168532. Epub 2007/08/31. eng.
10. Hamdan-Partida A, Sainz-Espunes T, Bustos-Martinez J. Characterization and persistence of *Staphylococcus aureus* strains isolated from the anterior nares and throats of healthy carriers in a Mexican community. *J Clin Microbiol.* 2010 May;48(5):1701-5. PubMed PMID: 20335416. Pubmed Central PMCID: PMC2863913. Epub 2010/03/26. eng.
11. Wardyn SE, Forshey BM, Smith TC. High prevalence of Panton-Valentine leukocidin among methicillin-sensitive *Staphylococcus aureus* colonization isolates in

- rural Iowa. Microbial drug resistance (Larchmont, NY). 2012 Aug;18(4):427-33. PubMed PMID: 22533373. Pubmed Central PMCID: PMC3462407. Epub 2012/04/27. eng.
12. Smith TC, Forshey BM, Hanson BM, Wardyn SE, Moritz ED. Molecular and epidemiologic predictors of Staphylococcus aureus colonization site in a population with limited nosocomial exposure. *Am J Infect Control*. 2012 Dec;40(10):992-6. PubMed PMID: 22418604. Epub 2012/03/16. eng.
 13. Urban and Rural Classification - Geography - U.S. Census Bureau 2013. Available from: <http://www.census.gov/geo/reference/urban-rural.html>.
 14. Stromquist AM, Merchant JA, Burmeister LF, Zwerling C, Reynolds SJ. The Keokuk County Rural Health Study. *Journal of Agromedicine*. 1997 1997/08/08;4(3-4):243-8.
 15. CLSI. Performance Standards for Antimicrobial Susceptibility Testing: Twenty-Second Informational Supplement: Clinical & Laboratory Standards Institute; 2012.
 16. Bosgelmez-Tinaz G, Ulusoy S, Aridogan B, Coskun-Ari F. Evaluation of different methods to detect oxacillin resistance in Staphylococcus aureus and their clinical laboratory utility. *Eur J Clin Microbiol Infect Dis*. 2006 Jun;25(6):410-2. PubMed PMID: 16767493. Epub 2006/06/13. eng.
 17. Lina G, Piemont Y, Godail-Gamot F, Bes M, Peter MO, Gauduchon V, et al. Involvement of Panton-Valentine leukocidin-producing Staphylococcus aureus in primary skin infections and pneumonia. *Clin Infect Dis*. 1999 Nov;29(5):1128-32. PubMed PMID: 10524952. Epub 1999/10/19. eng.
 18. Shopsis B, Gomez M, Montgomery SO, Smith DH, Waddington M, Dodge DE, et al. Evaluation of protein A gene polymorphic region DNA sequencing for typing of Staphylococcus aureus strains. *J Clin Microbiol*. 1999 Nov;37(11):3556-63. PubMed PMID: 10523551. Pubmed Central PMCID: PMC85690. Epub 1999/10/19. eng.
 19. Baum C, Haslinger-Loffler B, Westh H, Boye K, Peters G, Neumann C, et al. Non-spa-typeable clinical Staphylococcus aureus strains are naturally occurring protein A mutants. *J Clin Microbiol*. 2009 Nov;47(11):3624-9. PubMed PMID: 19759222. Pubmed Central PMCID: PMC2772612. Epub 2009/09/18. eng.
 20. Mellmann A, Weniger T, Berssenbrugge C, Rothganger J, Sammeth M, Stoye J, et al. Based Upon Repeat Pattern (BURP): an algorithm to characterize the long-term evolution of Staphylococcus aureus populations based on spa polymorphisms. *BMC Microbiol*. 2007;7:98. PubMed PMID: 17967176. Pubmed Central PMCID: PMC2148047. Epub 2007/10/31. eng.
 21. van Belkum A, Verkaik NJ, de Vogel CP, Boelens HA, Verveer J, Nouwen JL, et al. Reclassification of Staphylococcus aureus nasal carriage types. *J Infect Dis*. 2009 Jun 15;199(12):1820-6. PubMed PMID: 19419332. Epub 2009/05/08. eng.
 22. Kiernan K, Tao J, Gibbs P. Tips and Strategies for Mixed Modeling with SAS/STAT® Procedures 2012 [cited 2013]. Available from: <http://support.sas.com/resources/papers/proceedings12/332-2012.pdf>.
 23. Eriksen NH, Espersen F, Rosdahl VT, Jensen K. Carriage of Staphylococcus aureus among 104 healthy persons during a 19-month period. *Epidemiol Infect*. 1995 Aug;115(1):51-60. PubMed PMID: 7641838. Pubmed Central PMCID: PMC2271555. Epub 1995/08/01. eng.

24. Hu L, Umeda A, Kondo S, Amako K. Typing of *Staphylococcus aureus* colonising human nasal carriers by pulsed-field gel electrophoresis. *J Med Microbiol*. 1995 Feb;42(2):127-32. PubMed PMID: 7869348. Epub 1995/02/01. eng.
25. Mertz D, Frei R, Jaussi B, Tietz A, Stebler C, Fluckiger U, et al. Throat swabs are necessary to reliably detect carriers of *Staphylococcus aureus*. *Clin Infect Dis*. 2007 Aug 15;45(4):475-7. PubMed PMID: 17638197. Epub 2007/07/20. eng.
26. Robinson JO, Phillips M, Christiansen KJ, Pearson JC, Coombs GW, Murray RJ. Knowing prior methicillin-resistant *Staphylococcus aureus* (MRSA) infection or colonization status increases the empirical use of glycopeptides in MRSA bacteraemia and may decrease mortality. *Clin Microbiol Infect*. 2013 Sep 5. PubMed PMID: 24224545. Epub 2013/11/15. Eng.
27. Melles DC, Tenover FC, Kuehnert MJ, Witsenboer H, Peeters JK, Verbrugh HA, et al. Overlapping population structures of nasal isolates of *Staphylococcus aureus* from healthy Dutch and American individuals. *J Clin Microbiol*. 2008 Jan;46(1):235-41. PubMed PMID: 17977984. Pubmed Central PMCID: PMC2224299. Epub 2007/11/06. eng.
28. Delacour H, Van Cuyck H, Dubrous P, Soullie B, Leroy P, Koeck JL. Efficacy of a swab transport system in maintaining long-term viability of *Staphylococcus aureus*. *Diagn Microbiol Infect Dis*. 2009 Nov;65(3):345-6. PubMed PMID: 19729263. Epub 2009/09/05. eng.
29. Moran GJ, Krishnadasan A, Gorwitz RJ, Fosheim GE, McDougal LK, Carey RB, et al. Methicillin-resistant *S. aureus* infections among patients in the emergency department. *N Engl J Med*. 2006 Aug 17;355(7):666-74. PubMed PMID: 16914702. Epub 2006/08/18. eng.

SUPPLEMENTAL

Table S1: Site Specific Prevalence of Nares and Oropharynx Anatomical Sites by County for Adults

	Johnson Co.	Keokuk Co.	Total
Nares	458/2225 (20.6%)	756/2085 (36.3%)	1214/4310 (28.2%)
Oropharynx	694/2226 (31.2%)	498/2087 (23.9%)	1192/4313 (27.6%)

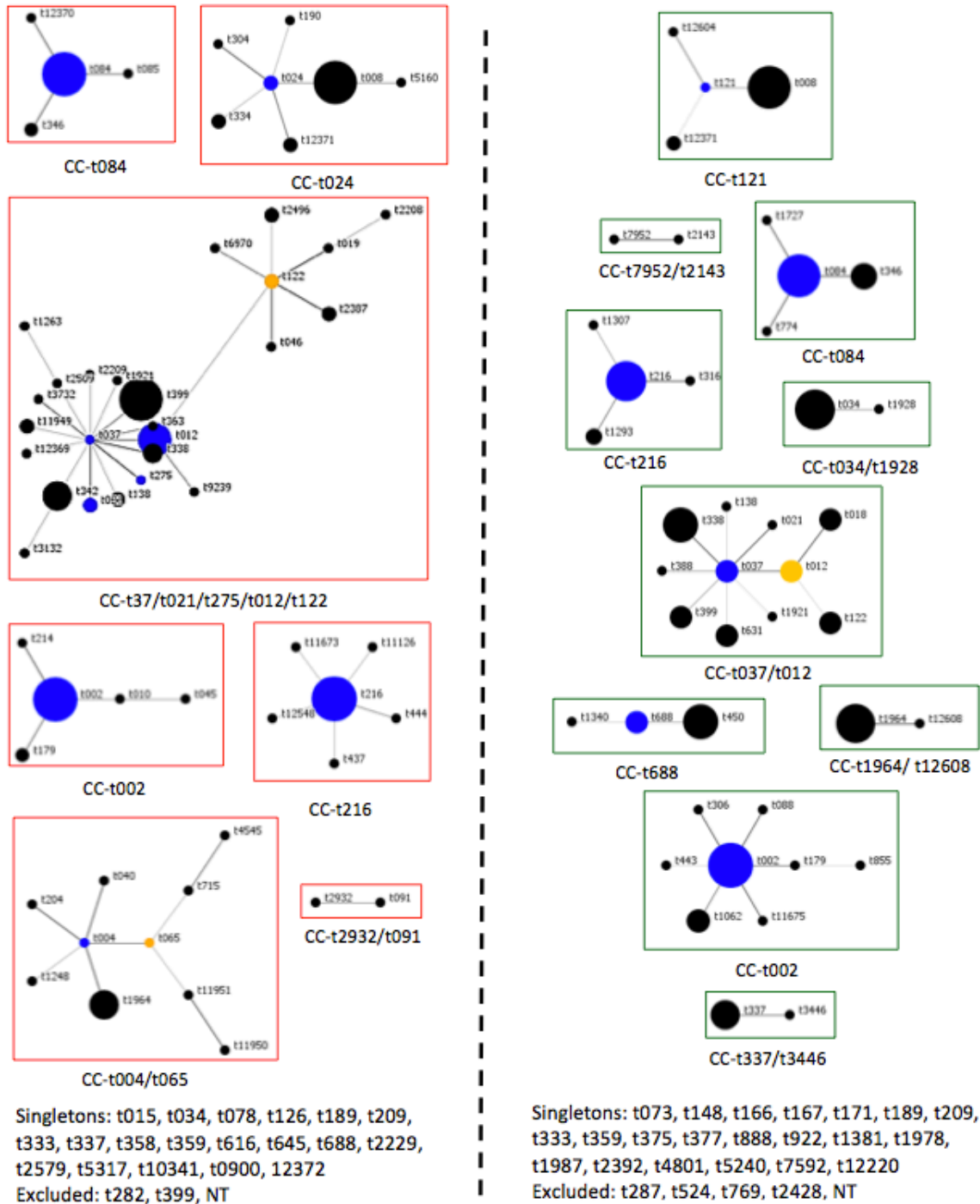


Figure S1: BURP analysis of spa typing data by county, adjusted to control for familial clustering of clonal isolates. Each circle represents a single *spa* type and size of the circle is proportionate to the relative number of isolates of that *spa* type. Blue circles indicate the putative founder, yellow circles are secondary putative founders.

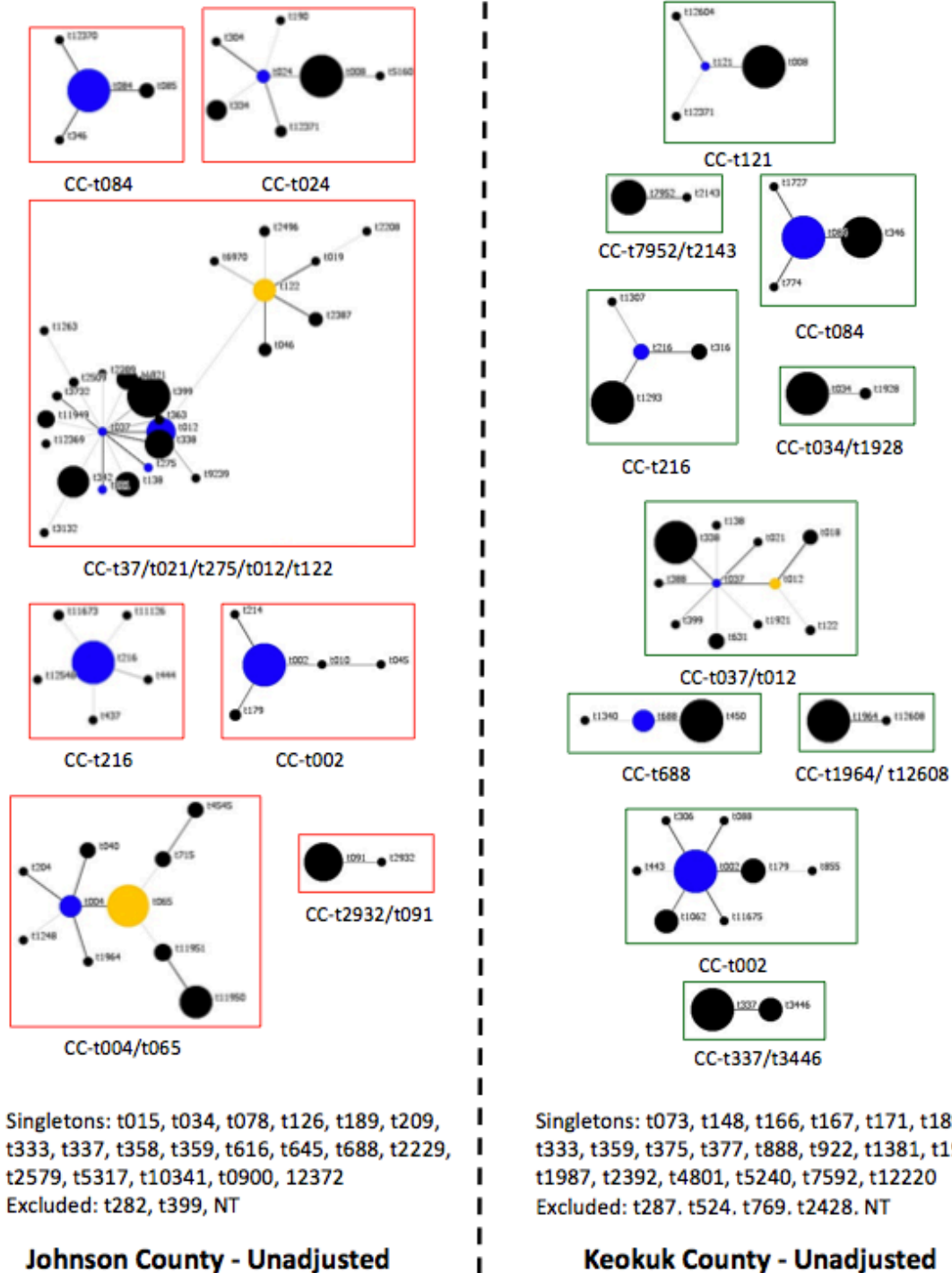


Figure C2: BURP analysis of spa typing data by county. Unadjusted proportions represent all 3184 isolates. Each circle represents a single *spa* type and size of the circle is proportionate to the relative number of isolates of that *spa* type. Blue circles indicate the putative founder, yellow circles are secondary putative founders.

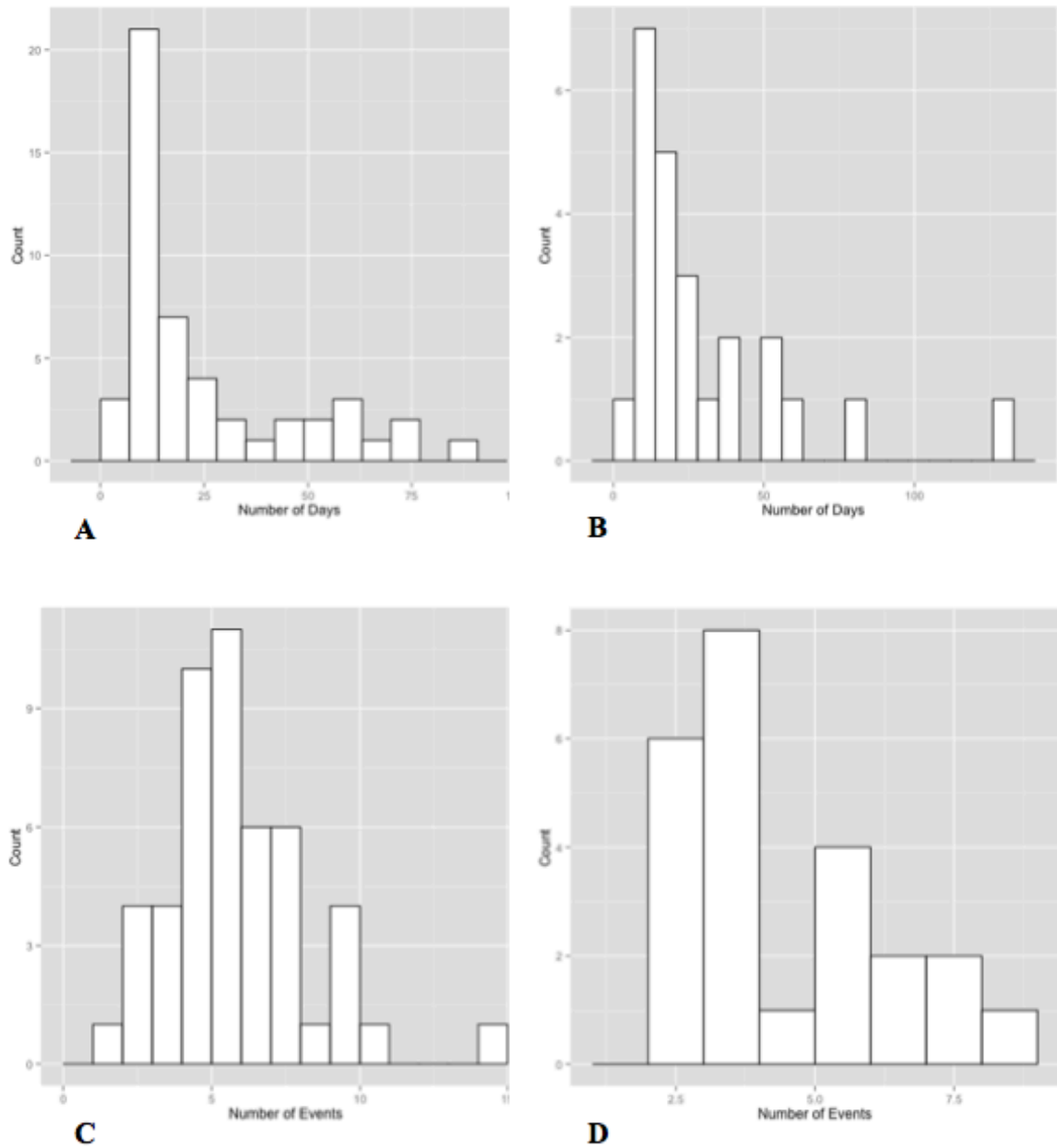


Figure C3: Frequency distributions of the average number of days of colonization and the number of colonization events for intermediate colonizers. A. The average number of days, grouped into 7-day intervals, of adult intermediate colonizers. B. The average number of days, grouped into 7-day intervals, of minor intermediate colonizers. C. The number of colonization events for adult intermediate colonizers. D. The number of colonization events for minor intermediate colonizers.

Table C2: All assessed risk factors for adults

Risk Factor[¶]	Number (Percent %)		Total (N)	Probability	95% CI	p-value
Intermittent carrier vs. Non-carrier	Non-carrier	Intermittent Carrier		Intermittent Carrier (at age=44.8)		
Age*						
Continuous	---	---	119	---	---	0.015
Gender						
Females	43 (64.18)	24 (35.82)	67	0.33	0.23-0.47	
Males	26 (52.00)	24 (48.00)	50	0.50	0.35-0.64	0.12
Race						
Caucasian	68 (61.26)	43 (38.74)	111	0.61	0.25-0.88	
Other	3 (37.50)	5 (62.50)	8	0.38	0.29-0.49	0.25
House Size						
One	1 (100.00)	0 (0.00)	1	0.23	0.11-0.44	
Two	35 (62.50)	21 (37.50)	56	0.28	0.16-0.43	
Three	21 (70.00)	9 (30.00)	30	0.32	0.21-0.46	
Four	9 (47.37)	10 (52.63)	19	0.37	0.23-0.55	
Five	5 (41.67)	7 (58.33)	12	0.43	0.22-0.67	
Six	0 (---)	0 (---)	0	0.48	0.20-0.78	
Seven	0 (0.00)	1 (100.00)	1	0.54	0.18-0.86	0.25

Table C2 Continued

Number of Children							
	Zero	42 (64.62)	23 (35.38)	65	0.27	0.16-0.43	
	One	13 (65.00)	7 (35.00)	20	0.32	0.21-0.47	
	Two	12 (52.17)	11 (47.83)	23	0.38	0.23-0.55	
	Three	4 (40.00)	6 (60.00)	10	0.43	0.22-0.67	
	Four	0 (---)	0 (---)	0	0.49	0.21-0.79	
	Five	0 (0.00)	1 (100.00)	1	0.55	0.19-0.87	0.218
Child in Daycare							
	No	58 (59.79)	39 (40.21)	97	0.42	0.32-0.53	
	Yes	13 (59.09)	9 (40.91)	22	0.31	0.14-0.54	0.36
Gym Exposure							
	No	52 (63.41)	30 (36.59)	82	0.40	0.29-0.52	
	Yes	18 (52.94)	16 (47.06)	34	0.37	0.21-0.56	0.77
Play Team Sports							
	No	68 (60.18)	45 (39.82)	113	0.39	0.30- 0.49	
	Yes	3 (60.00)	2 (40.00)	5	0.38	0.08-0.80	0.94
Any Family Member Play Team Sports							
	No	47 (66.20)	24 (33.80)	71	0.33	0.22-0.47	
	Yes	19 (47.50)	21 (52.50)	40	0.53	0.36-0.69	0.081

Table C2 Continued

Share Bath Towels							
No	41 (53.95)	35 (46.05)	76	0.47	0.35-0.59		
Yes	30 (69.77)	13 (30.23)	43	0.28	0.16-0.44	0.77	
Share Hand Towels							
No	12 (66.67)	6 (33.33)	18	0.37	0.17-0.64		
Yes	59 (58.42)	42 (41.58)	101	0.43	0.30-0.51	0.82	
Use Soap with Antimicrobials							
No	17 (60.71)	11 (39.29)	28	0.33	0.17-0.54		
Yes	54 (59.34)	37 (40.66)	91	0.42	0.31-0.53	0.45	
Number of Positive Environmental Sites							
Zero	57 (64.77)	31 (35.23)	88	0.16	0.10-0.22		
One	10 (55.56)	8 (44.44)	18	0.23	0.16-0.31		
Two	4 (40.00)	6 (60.00)	10	0.32	0.21-0.47		
Three	0 (0.00)	1 (100.00)	1	0.43	0.23-0.66		
Four	0 (0.00)	1 (100.00)	1	0.55	0.25-0.82		
Five	0 (0.00)	1 (100.00)	1	0.66	0.28-0.91	0.014	

Table C2 Continued

Farming Exposure ^s							
No Exposure	42 (58.33)	30 (41.67)	72	0.40	0.004-0.99		
Non-occupational Exposure	14 (77.78)	4 (22.22)	18	0.20	0.05-0.55		
Non-large Facility Occupational Exposure	2 (66.67)	1 (33.33)	3	0.42	0.05-0.92		
Large Facility Occupational Exposure	2 (25.00)	6 (75.00)	8	0.82	0.39-0.97	0.073	
Healthcare Exposure							
No Exposure	32 (59.26)	22 (40.74)	54	0.38	0.25-0.53		
Non-occupational Exposure	24 (61.54)	15 (38.46)	39	0.42	0.26-0.60		
Non-large Facility Occupational Exposure	5 (71.43)	2 (28.57)	7	0.29	0.06-0.70		
Large Facility Occupational Exposure	4 (50.00)	4 (50.00)	8	0.39	0.12-0.75	0.94	

Table C2 Continued

Any Healthcare Occupational Exposure							
No	55 (59.78)	37 (40.22)	92	0.40	0.30-0.52		
Yes	15 (60.00)	10 (40.00)	25	0.36	0.19-0.58	0.74	
Visited a Hospital							
No	40 (57.97)	29 (42.03)	69	0.40	0.28-0.53		
Yes	31 (63.27)	18 (36.73)	49	0.39	0.25-0.54	0.92	
Admitted to an Outpatient Center							
No	65 (59.63)	44 (40.37)	109	0.39	0.30-0.50		
Yes	5 (62.50)	3 (37.50)	8	0.40	0.12-0.76	0.99	
Family Member was Hospitalized							
No	70 (60.87)	45 (39.13)	115	0.39	0.29-0.49		
Yes	1 (25.00)	3 (75.00)	4	0.74	0.21-0.97	0.25	
Hospitalized ^o							
No	71 (60.68)	46 (39.32)	117	---	---		
Yes	0 (---)	0 (---)	0	---	---	---	
Have Asthma							
No	64 (58.18)	46 (41.82)	110	0.42	0.32-0.52		
Yes	6 (75.00)	2 (25.00)	8	0.24	0.06-0.62	0.35	

Table C2 Continued

Have Eczema							
No	65 (60.19)	43 (39.81)	108	0.40	0.30-0.50		
Yes	6 (54.55)	5 (45.45)	11	0.42	0.17-0.72	0.90	
Have Skin Condition Other Than Eczema							
No	65 (60.75)	42 (39.25)	107	0.38	0.29-0.49		
Yes	6 (50.00)	6 (50.00)	12	0.52	0.25-0.79	0.39	
Have Diabetes							
No	67 (58.77)	47 (41.23)	114	0.40	0.30-0.51		
Yes	4 (80.00)	1 (20.00)	5	0.30	0.04-0.81	0.70	
Have an Immune Disorder Other Than Diabetes							
No	66 (61.68)	41 (38.32)	107	0.38	0.29-0.48		
Yes	1 (16.67)	5 (83.33)	6	0.80	0.30-0.97	0.11	
Have Heart Condition							
No	67 (58.77)	47 (41.23)	114	0.41	0.31-0.51		
Yes	4 (80.00)	1 (20.00)	5	0.25	0.03-0.77	0.54	
Ever Had Cancer							
No	65 (58.56)	46 (41.44)	111	0.41	0.31-0.51		
Yes	6 (75.00)	2 (25.00)	8	0.30	0.07-0.70	0.61	

Table C2 Continued

Took Immuno-compromising Medicines							
	No	70(60.87)	45 (39.13)	115	0.38	0.30-0.49	
	Yes	1 (25.00)	3 (75.00)	4	0.72	0.20-0.97	0.24
Used Antibiotics							
	No	60 (56.60)	46 (43.40)	106	0.43	0.33-0.53	
	Yes	11 (84.62)	2 (15.38)	13	0.16	0.04-0.47	0.093
Skin or Soft Tissue Infection							
	No	69 (58.97)	48 (41.03)	117	---	---	
	Yes	2 (100.00)	0 (0.00)	2	---	---	0.51 [#]
Family with Skin or Soft Tissue Infection							
	No	62 (57.41)	46 (42.59)	108	0.42	0.32-0.52	0.45
	Yes	6 (75.00)	2 (25.00)	8	0.27	0.06-0.67	
Smoke any Tobacco Product							
	Never	49 (57.65)	36 (42.35)	85	0.40	0.30-0.54	0.48
	Past Usage	15 (60.00)	10 (40.00)	25	0.43	0.02-0.65	
	Current Usage	6 (85.71)	1 (14.29)	7	0.15	0.02-0.62	

Table C2 Continued

Non- or intermittent carrier vs. Persistent Carrier	Non- or Intermittent Carrier	Persistent Carrier		Persistent Carrier (at age=44.8)			
Age*							
Continuous	---	---	149	---	---		0.86
Gender							
Females	67 (82.72)	14 (17.28)	81	0.17	0.10-0.27		
Males	50 (75.76)	16 (24.24)	66	0.24	0.15-0.36		0.30
Race							
Caucasian	111 (79.29)	29 (20.71)	140	0.21	0.15-0.29		
Other	8 (88.89)	1 (11.11)	9	0.11	0.02-0.51		0.51
House Size							
One	1 (100.00)	0 (0.00)	1	0.18	0.09-0.34		
Two	56 (78.87)	15 (21.13)	71	0.19	0.12-0.29		
Three	30 (85.71)	5 (14.29)	35	0.20	0.14-0.28		
Four	19 (73.08)	7 (26.92)	26	0.21	0.13-0.32		
Five	12 (85.71)	2 (14.29)	14	0.22	0.11-0.41		
Six	0 (---)	0 (---)	0	0.24	0.08-0.51		
Seven	1 (50.00)	1 (50.00)	2	0.25	0.06-0.62		0.73
Number of Children							
Zero	65 (79.27)	17 (20.73)	82	0.19	0.12-0.29		
One	20 (86.96)	3 (13.04)	23	0.20	0.14-0.28		
Two	23 (76.67)	7 (23.33)	30	0.21	0.13-0.33		
Three	10 (83.33)	2 (16.67)	12	0.22	0.10-0.41		
Four	0 (---)	0 (---)	0	0.23	0.08-0.51		
Five	1 (50.00)	1 (50.00)	2	0.24	0.06-0.61		0.77

Table C2 Continued

Child in Daycare							
No	97 (77.60)	28 (22.40)	125	0.23	0.16-0.31		
Yes	22 (91.67)	2 (8.33)	24	0.08	0.02-0.27	0.11	
Gym Exposure							
No	82 (79.61)	21 (20.39)	103	0.20	0.13-0.30		
Yes	34 (79.07)	9 (20.93)	43	0.21	0.10-0.37	0.97	
Play Team Sports							
No	113 (79.58)	29 (20.42)	142	0.20	0.14-0.28		
Yes	5 (83.33)	1 (16.67)	6	0.17	0.02-0.65	0.84	
Any Family Member Play Team Sports							
No	71 (77.17)	21 (22.83)	92	0.23	0.15-0.33		
Yes	40 (81.63)	9 (18.37)	49	0.18	0.10-0.32	0.53	
Share Bath Towels							
No	76 (77.55)	22 (22.45)	98	0.23	0.15-0.32		
Yes	43 (84.31)	8 (15.69)	51	0.16	0.08-0.29	0.33	
Share Hand Towels							
No	18 (75.00)	6 (25.00)	24	0.26	0.11-0.48		
Yes	101 (80.80)	24 (19.20)	125	0.19	0.13-0.27	0.50	

Table C2 Continued

Use Soap with Antimicrobials							
	No	28 (80.00)	7 (20.00)	35	0.20	0.09-0.37	
	Yes	91 (79.82)	23 (20.18)	114	0.20	0.14-0.29	0.96
Number of Positive Environmental Sites							
	Zero	88 (83.81)	17 (16.19)	105	0.16	0.10-0.23	
	One	18 (85.71)	3 (14.29)	21	0.23	0.16-0.31	
	Two	10 (62.50)	6 (37.50)	16	0.32	0.20-0.47	
	Three	1 (33.33)	2 (66.67)	3	0.43	0.23-0.66	
	Four	1 (50.00)	1 (50.00)	2	0.55	0.25-0.81	
	Five	1 (50.00)	1 (50.00)	2	0.66	0.28-0.92	0.014
Farming Exposure [§]							
	No Exposure	72 (80.90)	17 (19.10)	89	---	---	
	Non-occupational Exposure	18 (78.26)	5 (21.74)	23	---	---	
	Non-large Facility Occupational Exposure	3 (100.00)	0 (0.00)	3	---	---	
	Large Facility Occupational Exposure	8 (72.73)	3 (27.27)	11	---	---	0.88 [#]

Table C2 Continued

Healthcare Exposure							
No Exposure	54 (80.60)	13 (19.40)	67	0.20	0.11-0.31		
Non-occupational	39 (82.98)	8 (17.02)	47	0.16	0.07-0.31		
Non-large Facility Occupational Exposure	7 (87.50)	1 (12.50)	8	0.13	0.02-0.57		
Large Facility Occupational Exposure	8 (72.73)	3 (27.27)	11	0.30	0.092-0.65	0.79	
Any Healthcare Occupational Exposure							
No	92 (80.70)	22 (19.30)	114	0.19	0.13-0.28		
Yes	25 (75.76)	8 (24.24)	33	0.25	0.12-0.43	0.51	
Visited a Hospital							
No	69 (81.18)	16 (18.82)	85	0.18	0.11-0.29		
Yes	49 (77.78)	14 (22.22)	63	0.23	0.14-0.36	0.52	
Admitted to an Outpatient Center							
No	109 (78.42)	30 (21.58)	139	---	---		
Yes	8 (100.00)	0 (0.00)	8	---	---	0.21 [#]	

Table C2 Continued

Family Member was Hospitalized							
No	115 (79.31)	30 (20.69)	145	---	---		
Yes	4 (100.00)	0 (0.00)	4	---	---	0.58 [#]	
Hospitalized ^o							
No	117 (80.14)	29 (19.86)	146	---	---		
Yes	0 (---)	0 (---)	0	---	---	---	
Have Asthma							
No	110 (79.71)	28 (20.29)	138	0.20	0.14-0.28		
Yes	8 (80.00)	2 (20.00)	10	0.20	0.05-0.55	0.96	
Have Eczema							
No	108 (79.41)	28 (20.59)	136	0.21	0.14-0.29		
Yes	11 (84.62)	2 (15.38)	13	0.15	0.04-0.46	0.66	
Have Skin Condition Other Than Eczema							
No	107 (80.45)	26 (19.55)	133	0.20	0.14-0.28		
Yes	12 (75.00)	4 (25.00)	16	0.25	0.09-0.51	0.63	
Have Diabetes							
No	114 (80.85)	27 (19.15)	141	0.19	0.13-0.27		
Yes	5 (71.43)	2 (28.57)	7	0.29	0.07-0.69	0.56	

Table C2 Continued

Have an Immune Disorder Other Than Diabetes							
No	107 (80.45)	26 (19.55)	133	0.20	0.14-0.27		
Yes	6 (85.71)	1 (14.29)	7	0.14	0.02-0.59		0.74
Have Heart Condition							
No	114 (81.43)	26 (18.57)	140	0.18	0.13-0.26		
Yes	5 (62.50)	3 (37.50)	8	0.40	0.12-0.76		
Ever Had Cancer							
No	111 (80.43)	27 (19.57)	138	0.19	0.13-0.27		
Yes	8 (72.73)	3 (27.27)	11	0.28	0.09-0.62		0.51
Took Immuno-compromising Medicines							
No	115 (80.42)	28 (19.58)	143	0.20	0.14-0.27		
Yes	4 (66.67)	2 (33.33)	6	0.33	0.08-0.74		0.44
Used Antibiotics							
No	106 (79.70)	27 (20.30)	133	0.20	0.14-0.28		
Yes	13 (81.25)	3 (18.75)	16	0.20	0.06-0.46		0.90
Skin or Soft Tissue Infection							
No	117 (79.59)	30 (20.41)	147	---	---		
Yes	2 (100.00)	0 (0.00)	2	---	---		1.0

Table C2 Continued

Family Member with Skin or Soft Tissue Infection							
	No	108 (79.41)	28 (20.59)	136	0.20	0.14-0.29	
	Yes	8 (88.89)	1 (11.11)	9	0.11	0.015-0.52	0.53
Smoke any Tobacco Product							
	Never	85 (80.19)	21 (19.81)	106	0.20	0.13-0.29	
	Past Usage	25 (80.65)	6 (19.35)	31	0.20	0.09-0.38	
	Current Usage	7 (87.50)	1 (12.50)	8	0.13	0.02-0.55	0.90
Non-carrier vs. Intermittent or Persistent Carrier		Non-carrier	Any Carrier		Any Carrier (at age =44.8)		
Age*							
	Continuous	---	---	149	---	---	0.035
Gender							
	Females	43 (53.09)	38 (46.91)	81	0.46	0.34-0.58	
	Males	26 (39.39)	40 (60.61)	66	0.62	0.48-0.73	0.072
Race							
	Caucasian	68 (48.57)	72 (51.43)	140	0.51	0.42-0.61	
	Other	3 (33.33)	6 (66.67)	9	0.67	0.30-0.90	0.43

Table C2 Continued

House Size						
One	1 (100.00)	0 (0.00)	1	0.43	0.26-0.61	
Two	35 (49.30)	36 (50.00)	71	0.48	0.36-0.60	
Three	21 (60.00)	14 (40.00)	35	0.53	0.43-0.62	
Four	9 (34.62)	17 (65.38)	26	0.58	0.44-0.70	
Five	5 (35.71)	9 (64.29)	14	0.63	0.42-0.79	
Six	0 (---)	0 (---)	0	0.67	0.39-0.87	
Seven	0 (0.00)	2 (100.00)	2	0.72	0.36-0.92	0.25
Number of Children						
Zero	42 (51.22)	40 (48.78)	82	0.49	0.37-0.60	
One	13 (56.52)	10 (43.48)	23	0.53	0.44-0.62	
Two	12 (40.00)	18 (60.00)	30	0.58	0.44-0.70	
Three	4 (33.33)	8 (66.67)	12	0.62	0.41-0.79	
Four	0 (---)	0 (---)	0	0.67	0.38-0.87	
Five	0 (0.00)	2 (100.00)	2	0.71	0.35-0.92	0.29
Child in Daycare						
No	58 (46.40)	67 (53.60)	125	0.56	0.45-0.66	
Yes	13 (54.17)	11 (45.83)	24	0.36	0.18-0.60	0.15
Gym Exposure						
No	52 (50.49)	51 (49.51)	103	0.52	0.41-0.63	
Yes	18 (41.86)	25 (58.14)	43	0.51	0.34-0.68	0.93

Table C2 Continued

Play Team Sports							
No	68 (47.89)	74 (52.11)	142	0.52	0.43-0.61		
Yes	3 (50.00)	3 (50.00)	6	0.47	0.14-0.84	0.83	
Any Family Member Play Team Sports							
No	47 (51.09)	45 (48.91)	92	0.49	0.38-0.62		
Yes	19 (38.78)	30 (61.22)	49	0.61	0.44-0.76	0.27	
Share Bath Towels							
No	41 (41.84)	57 (58.16)	98	0.60	0.48-0.70		
Yes	30 (58.82)	21 (41.18)	51	0.38	0.24-0.54	0.034	
Share Hand Towels							
No	12 (50.00)	12 (50.00)	24	0.55	0.32-0.76		
Yes	59 (47.20)	66 (52.80)	125	0.52	0.42-0.62	0.81	
Use Soap with Antimicrobials							
No	17 (48.57)	18 (51.43)	35	0.48	0.30-0.67		
Yes	54 (47.37)	60 (52.63)	114	0.54	0.43-0.64	0.61	

Table C2 Continued

Number of Positive Environmental Sites							
	Zero	10 (47.62)	11 (52.38)	21	0.43	0.33-0.54	
	One	4 (25.00)	12 (75.00)	16	0.63	0.51-0.74	
	Two	0 (0.00)	3 (100.00)	3	0.80	0.61-0.91	
	Three	0 (0.00)	2 (100.00)	2	0.90	0.68-0.98	
	Four	0 (0.00)	2 (100.00)	2	0.96	0.75-0.99	
	Five	10 (47.62)	11 (52.38)	21	0.98	0.80-1.00	0.003
Farming Exposure [§]							
	No Exposure	42 (47.19)	47 (52.81)	89	0.51	0.01-0.99	
	Non-occupational Exposure	14 (60.87)	9 (39.13)	23	0.40	0.11-0.79	
	Non-large Facility Occupational Exposure	2 (66.67)	1 (33.33)	3	0.41	0.04-0.92	
	Large Facility Occupational Exposure	2 (18.18)	9 (81.82)	11	0.88	0.51-0.98	0.11

Table C2 Continued

Healthcare Exposure							
No Exposure	32 (47.76)	35 (52.24)	67	0.50	0.36-0.64		
Non-occupational	24 (51.06)	23 (48.94)	47	0.53	0.37-0.69		
Non-large Facility Occupational Exposure	5 (62.50)	3 (37.50)	8	0.38	0.11-0.75		
Large Facility Occupational Exposure	4 (36.36)	7 (63.64)	11	0.61	0.28-0.86	0.84	
Any Healthcare Occupational Exposure							
No	55 (48.25)	59 (51.75)	114	0.52	0.42-0.63		
Yes	15 (45.45)	18 (54.55)	33	0.53	0.34-0.70	0.97	
Visited a Hospital							
No	40 (47.06)	45 (52.94)	85	0.50	0.38-0.62		
Yes	31 (49.21)	32 (50.79)	63	0.55	0.40-0.68	0.64	
Admitted to an Outpatient Center							
No	65 (46.76)	74 (53.24)	139	0.53	0.44-0.63		
Yes	5 (62.50)	3 (37.50)	8	0.38	0.11-0.74	0.43	

Table C2 Continued

Family Member was Hospitalized							
	No	70 (48.28)	75 (51.72)	145	0.52	0.42-0.61	
	Yes	1 (25.00)	3 (75.00)	4	0.73	0.19-0.97	0.45
Hospitalized ^o							
	No	71 (48.63)	75 (51.37)	146	---	---	
	Yes	0 (---)	0 (---)	0	---	---	---
Have Asthma							
	No	64 (46.38)	74 (53.62)	138	0.54	0.44-0.63	
	Yes	6 (60.00)	4 (40.00)	10	0.37	0.13-0.70	0.33
Have Eczema							
	No	65 (47.79)	71 (52.21)	136	0.52	0.43-0.62	
	Yes	6 (46.15)	7 (53.85)	13	0.53	0.25-0.79	0.97
Have Skin Condition Other Than Eczema							
	No	65 (48.87)	68 (51.13)	133	0.51	0.41-0.61	
	Yes	6 (37.50)	10 (62.50)	16	0.63	0.36-0.84	0.39
Have Diabetes							
	No	67 (47.52)	74 (52.48)	141	0.52	0.43-0.62	
	Yes	4 (57.14)	3 (42.86)	7	0.48	0.15-0.83	0.84

Table C2 Continued

Have an Immune Disorder Other Than Diabetes							
No	66 (49.62)	67 (50.38)	133	0.51	0.41-0.60		
Yes	1 (14.29)	6 (85.71)	7	0.84	0.36-0.98	0.15	
Have Heart Condition							
No	67 (47.86)	73 (52.14)	140	0.52	0.42-0.61		
Yes	4 (50.00)	4 (50.00)	8	0.58	0.23-0.87	0.74	
Ever Had Cancer							
No	65 (47.10)	73 (52.90)	138	0.52	0.43-0.62		
Yes	6 (54.55)	5 (45.45)	11	0.52	0.22-0.80	0.97	
Took Immuno-compromising Medicines							
No	70 (48.95)	73 (51.05)	143	0.51	0.42-0.60		
Yes	1 (16.67)	5 (83.33)	6	0.81	0.32-0.98	0.22	
Used Antibiotics							
No	60 (45.11)	73 (54.89)	133	0.55	0.45-0.64		
Yes	11 (68.75)	5 (31.25)	16	0.33	0.14-0.61	0.15	

Table C2 Continued

Skin or Soft Tissue Infection							
No	69 (46.94)	78 (53.06)	147	---	---		
Yes	2 (100.00)	0 (0.00)	2	---	---		0.22 [#]
Family Member with Skin or Soft Tissue Infection							
No	62 (45.59)	74 (54.41)	136	0.54	0.44-0.64		
Yes	6 (66.67)	3 (33.33)	9	0.34	0.10-0.72		0.31
Smoke any Tobacco Product							
Never	49 (46.23)	57 (53.77)	106	0.52	0.41-0.63		
Past Usage	15 (48.39)	16 (51.61)	31	0.54	0.35-0.73		
Current Usage	6 (75.00)	2 (25.00)	8	0.27	0.06-0.67		0.44

Note: [¶]All risk factors adjusted for age at 44.8 years ^{*} Mean age for non-carriers is 47.34. Mean age for intermittent carriers is 41.09. Mean age for persistent carriers is years 44.3. [§]Farming exposure refers to contact with livestock (swine, cattle, poultry, etc.) [°]Probability, 95% Confidence Interval, and p-value not calculable due to multiple cells with cell counts of zero [#]Fisher's Exact test done because of small sample sizes.

Table C3: All assessed risk factors for minors

Risk Factor [¶]	Number (Percent %)		Total (N)	Probability	95% CI	p-value
	Non-carrier	Intermittent Carrier				
Non-carriers vs. Intermittent Carriers						
Age*						
Continuous	---	---	63	---	---	0.28
Gender						
Females	20 (60.60)	13 (39.40)	33	0.39	0.001-0.10	
Males	15 (50.00)	15 (50.00)	30	0.467	0.002-0.10	0.62
Race						
Caucasian	29 (50.90)	28 (49.10)	57	---	---	
Other	6 (100.0)	0 (0.0)	6	---	---	0.033 [#]
House Size						
Two	2 (100.00)	0 (0.00)	2	0.48	0.04-0.96	
Three	11 (68.75)	5 (31.25)	16	0.46	0.03-0.97	
Four	8 (30.77)	18 (69.23)	26	0.44	0.01-0.99	
Five	12 (80.00)	3 (20.00)	15	0.42	0.001-0.10	
Six	0 (---)	0 (---)	0	0.40	0.003-0.99	
Seven	2 (50.00)	2 (50.00)	4	0.38	0.02-0.96	0.77
Number of Children						
One	12 (80.00)	3 (20.00)	15	0.40	0.02-0.95	
Two	11 (34.38)	21 (65.63)	32	0.41	0.01-0.99	
Three	10 (83.33)	2 (16.67)	12	0.42	0.001-0.10	
Four	0 (---)	0 (---)	0	0.42	0.003-0.99	
Five	2 (50.00)	2 (50.00)	4	0.43	0.02-0.97	0.91

Table C3 Continued

Child in Daycare							
	No	23 (57.50)	17 (42.50)	40	0.38	0.002-0.10	
	Yes	12 (52.20)	11 (47.80)	23	0.48	0.01-0.98	0.61
Gym Exposure							
	No	20 (58.80)	14 (41.20)	34	0.31	0.001-0.10	
	Yes	15 (51.70)	14 (48.30)	29	0.54	0.002-0.10	0.15
Play Team Sports							
	No	20 (58.80)	14 (41.20)	34	0.38	0.01-0.99	
	Yes	15 (53.60)	13 (46.40)	28	0.46	0.01-0.99	0.62
Any Family Member Play Team Sports							
	No	11 (52.40)	10 (47.60)	21	0.44	0.02-0.98	
	Yes	23 (56.10)	18 (43.90)	41	0.40	0.003-0.99	0.79
Share Bath Towels							
	No	17 (50.00)	17 (50.00)	34	0.41	0.001-0.10	
	Yes	17 (60.70)	11 (39.30)	28	0.44	0.004-0.99	0.84
Share Hand Towels							
	No	5 (71.40)	2 (28.60)	7	0.38	0.03-0.93	
	Yes	30 (53.60)	26 (46.40)	56	0.42	0.001-0.10	0.89
Use Soap with Antimicrobials							
	No	6 (50.00)	6 (50.00)	12	0.54	0.03-0.98	
	Yes	29 (56.90)	22 (43.10)	51	0.39	0.001-0.10	0.427

Table C3 Continued

Number of Positive Environmental Sites							
	Zero	25 (55.56)	20 (44.44)	45	0.38	0.001-0.10	
	One	5 (71.43)	2 (28.57)	7	0.42	0.0004-0.10	
	Two	3 (60.00)	2 (40.00)	5	0.47	0.003-0.10	
	Three	0 (0.00)	2 (100.00)	2	0.51	0.02-0.99	
	Four	1 (33.33)	2 (66.67)	3	0.55	0.05-0.97	
	Five	1 (100.00)	0 (0.00)	1	0.59	0.06-0.97	0.48
Farming Exposure ^s							
	No Exposure	24 (58.50)	17 (41.50)	41	0.43	0.001-0.10	
	Non-occupational Exposure	9 (56.30)	7 (43.80)	16	0.41	0.01-0.99	
	Large Facility Occupational Exposure	2 (50.00)	2 (50.00)	4	0.20	0.01-0.90	0.72
Healthcare Exposure							
	No Exposure	28 (58.33)	20 (41.67)	48	0.37	0.001-0.10	
	Non-occupational	7 (46.67)	8 (53.33)	15	0.55	0.03-0.98	0.31
Visited a Hospital							
	No	30 (58.82)	21 (41.18)	51	0.38	0.001-0.10	
	Yes	5 (41.67)	7 (58.33)	12	0.58	0.04-0.98	0.28

Table C3 Continued

Admitted to an Outpatient Center							
No	33 (55.00)	27 (45.00)	60	0.42	0.001-0.10		
Yes	2 (66.67)	1 (33.33)	3	0.42	0.02-0.96		0.10
Family Member was Hospitalized							
No	32 (56.14)	25 (43.86)	57	0.40	0.00003-0.10		
Yes	3 (50.00)	3 (50.00)	6	0.73	0.06-0.99		0.24
Hospitalized							
No	34 (55.74)	27 (44.26)	61	0.41	0.001-0.10		
Yes	1 (50.00)	1 (50.00)	2	0.58	0.03-0.99		0.71
Have Asthma							
No	32 (56.14)	25 (43.86)	57	0.43	0.0004-0.10		
Yes	3 (50.00)	3 (50.00)	6	0.29	0.02-0.91		0.53
Have Eczema							
No	33 (57.89)	24 (42.11)	57	0.39	0.001-0.10		
Yes	2 (33.33)	4 (66.67)	6	0.66	0.08-0.98		0.28
Have Skin Condition Other Than Eczema							
No	32 (59.26)	22 (40.74)	54	0.40	0.001-0.10		
Yes	3 (33.33)	6 (66.67)	9	0.63	0.10-0.97		0.30

Table C3 Continued

Immune Disorder Other Than Diabetes							
No	35 (57.38)	26 (42.62)	61	---	---		
Yes	0 (0.00)	2 (100.00)	2	---	---	0.18 [#]	
Took Immuno-compromising Medicines							
No	32 (53.33)	28 (46.67)	60	---	---		
Yes	3 (100.00)	0 (0.00)	3	---	---	0.25 [#]	
Used Antibiotics							
No	26 (50.98)	25 (49.02)	51	0.45	0.001-0.10		
Yes	9 (75.00)	3 (25.00)	12	0.20	0.01-0.87	0.17	
Had Skin or Soft Tissue Infection							
No	34 (54.84)	28 (45.16)	62	---	---		
Yes	1 (100.00)	0 (0.00)	1	---	---	1.000 [#]	
Family Member with Skin or Soft Tissue Infection							
No	31 (54.39)	26 (45.61)	57	0.44	0.001-0.10		
Yes	3 (60.00)	2 (40.00)	5	0.33	0.02-0.92	0.69	

Table C3 Continued

Non- or Intermittent Carrier vs. Persistent Carrier	Non- or Intermittent Carrier	Persistent Carrier		Persistent Carrier (at age=9.65)			
Age*							
Continuous	---	---	74	---	---	---	0.033
Gender							
Females	33 (82.50)	7 (17.50)	40	0.09	0.0002-0.98		
Males	30 (88.24)	4 (11.76)	34	0.11	0.001-0.96		0.75
Race							
Caucasian	57 (87.69)	8 (12.31)	65	0.07	0.001-0.85		
Other	6 (66.67)	3 (33.33)	9	0.18	0.01-0.85		0.34
House Size							
Two	2 (66.67)	1 (33.33)	3	0.18	0.01-0.85		
Three	16 (94.12)	1 (5.88)	17	0.07	0.001-0.85		
Four	26 (96.30)	1 (3.70)	27	0.18	0.01-0.85		
Five	15 (65.22)	8 (34.78)	23	0.07	0.001-0.85		
Six	0 (---)	0 (---)	0	0.18	0.01-0.85		
Seven	4 (100.00)	0 (0.00)	4	0.07	0.001-0.85		0.35
Number of Children							
Zero**	0 (0.00)	1 (100.00)	1	0.07	0.003-0.68		
One	15 (88.24)	2 (11.76)	17	0.08	0.002-0.74		
Two	32 (96.97)	1 (3.03)	33	0.08	0.001-0.93		
Three	12 (63.16)	7 (36.84)	19	0.09	0.001-0.88		
Four	0 (---)	0 (---)	0	0.01	0.004-0.77		
Five	4 (100.00)	0 (0.00)	4	0.11	0.003-0.82		0.8

Table C3 Continued

Child in Daycare							
No	40 (85.11)	7 (14.89)	47	0.05	0.003-0.52		
Yes	23 (85.19)	4 (14.81)	27	0.16	0.01-0.84	0.293	
Gym Exposure							
No	34 (94.44)	2 (5.56)	36	0.05	0.003-0.48		
Yes	29 (78.38)	8 (21.62)	37	0.16	0.01-0.82	0.22	
Play Team Sports							
No	34 (91.89)	3 (8.11)	37	0.08	0.002-0.81		
Yes	28 (77.78)	8 (22.22)	36	0.10	0.003-0.82	0.84	
Any Family Member Play Team Sports							
No	21 (100.00)	0 (0.00)	21	---	---		
Yes	41 (78.85)	11 (21.15)	52	---	---	0.093 [#]	
Share Bath Towels							
No	34 (85.00)	6 (15.00)	40	0.11	0.00003-0.10		
Yes	28 (84.85)	5 (15.15)	33	0.04	0.0001-0.95	0.29	
Share Hand Towels							
No	7 (70.00)	3 (30.00)	10	0.12	0.01-0.80		
Yes	56 (87.50)	8 (12.50)	64	0.08	0.001-0.91	0.69	
Use Soap with Antimicrobials							
No	12 (85.71)	2 (14.29)	14	0.11	0.004-0.78		
Yes	51 (85.00)	9 (15.00)	60	0.08	0.001-0.90	0.80	

Table C3 Continued

Number of Positive Environmental Sites							
	Zero	45 (91.84)	4 (8.16)	49	0.02	0.001-0.53	
	One	7 (77.78)	2 (22.22)	9	0.05	0.001-0.78	
	Two	5 (71.43)	2 (28.57)	7	0.09	0.001-0.93	
	Three	2 (100.00)	0 (0.00)	2	0.15	0.001-0.96	
	Four	3 (75.00)	1 (25.00)	4	0.26	0.01-0.96	
	Five	1 (33.33)	2 (66.67)	3	0.40	0.01-0.97	0.041
Farming Exposure ^s							
	No Exposure	41 (87.23)	6 (12.77)	47	0.04	0.00002-0.99	
	Non-occupational Exposure	16 (84.21)	3 (15.79)	19	0.10	0.0001-0.99	
	Large Facility Occupational Exposure	4 (80.00)	1 (20.00)	5	0.15	0.001-0.95	0.64
Healthcare Exposure							
	No Exposure	49 (87.5)	7 (12.5)	56	0.07	0.002-0.79	
	Non-occupational	15 (75.0)	5 (25.0)	20	0.13	0.01-0.81	0.44
Visited a Hospital							
	No	48 (88.89)	6 (11.11)	54	0.08	0.001-0.88	
	Yes	15 (75.00)	5 (25.00)	20	0.14	0.01-0.82	0.45

Table C3 Continued

Admitted to an Outpatient Center							
	No	60 (85.71)	10 (14.29)	70	0.08	0.001-0.94	
	Yes	3 (75.00)	1 (25.00)	4	0.09	0.002-0.81	0.94
Family Member was Hospitalized							
	No	57 (85.07)	10 (14.93)	67	0.09	0.0001-0.99	
	Yes	6 (85.71)	1 (14.29)	7	0.02	0.0003-0.65	0.40
Hospitalized							
	No	61 (84.72)	11 (15.28)	72	---	---	
	Yes	2 (100.00)	0 (0.00)	2	---	---	1.00 [#]
Have Asthma							
	No	57 (87.69)	8 (12.31)	65	0.076	0.001-0.93	
	Yes	6 (66.67)	3 (33.33)	9	0.27	0.01-0.91	0.17
Have Eczema							
	No	57 (86.36)	9 (13.64)	66	0.079	0.001-0.90	
	Yes	6 (75.00)	2 (25.00)	8	0.20	0.01-0.87	0.38
Have Skin Condition Other Than Eczema							
	No	54 (84.38)	10 (15.63)	64	0.09	0.001-0.92	
	Yes	9 (90.00)	1 (10.00)	10	0.05	0.002-0.65	0.68

Table C3 Continued

Have an Immune Disorder Other Than Diabetes							
No	61 (84.72)	11 (15.28)	72	---	---		
Yes	2 (100.00)	0 (0.00)	2	---	---		1.00 [#]
Took Immuno-compromising Medicines							
No	60 (85.71)	10 (14.29)	70	---	---		
Yes	3 (100.00)	0 (0.00)	3	---	---		1.00 [#]
Used Antibiotics							
No	51 (86.44)	8 (13.56)	59	0.09	0.001-0.94		
Yes	12 (80.00)	3 (20.00)	15	0.08	0.003-0.72		0.87
Had Skin or Soft Tissue Infection							
No	62 (86.11)	10 (13.89)	72	---	---		
Yes	1 (50.00)	1 (50.00)	2	---	---		0.29 [#]
Family Member with Skin or Soft Tissue Infection							
No	57 (85.07)	10 (14.93)	67	0.07	0.0001-0.98		
Yes	5 (83.33)	1 (16.67)	6	0.20	0.01-0.93		0.45

Table C3 Continued

Non-carrier vs. Intermittent or Persistent Carrier	Non-carrier	Any Carrier	Any Carrier (at age =9.65)				
Age*							
Continuous	---	---	74	---	---	0.74	
Gender							
Females	20 (50.00)	20 (50.00)	40	0.50	0.14-0.86		
Males	15 (44.12)	19 (55.88)	34	0.58	0.22-0.86	0.54	
Race							
Caucasian	29 (44.62)	36 (55.38)	65	0.55	0.25-0.81		
Other	6 (66.67)	3 (33.33)	9	0.34	0.09-0.72	0.27	
House Size							
Two	2 (66.67)	1 (33.33)	3	0.50	0.19-0.81		
Three	11 (64.71)	6 (35.29)	17	0.51	0.21-0.80		
Four	8 (29.63)	19 (70.37)	27	0.52	0.17-0.85		
Five	12 (52.17)	11 (47.83)	23	0.53	0.21-0.83		
Six	0 (---)	0 (---)	0	0.54	0.22-0.83		
Seven	2 (50.00)	2 (50.00)	4	0.55	0.18-0.87	0.85	
Number of Children							
Zero**	0 (0.00)	1 (100.00)	1	0.50	0.19-0.82		
One	12 (70.59)	5 (29.41)	17	0.51	0.22-0.80		
Two	11 (33.33)	22 (66.67)	33	0.52	0.17-0.85		
Three	10 (52.63)	9 (47.37)	19	0.53	0.21-0.83		
Four	2 (50.00)	2 (50.00)	4	0.54	0.21-0.83		
Five	0 (0.00)	1 (100.00)	1	0.54	0.17-0.88	0.90	

Table C3 Continued

Child in Daycare							
	No	23 (48.94)	24 (51.06)	47	0.47	0.13-0.84	
	Yes	12 (44.44)	15 (55.56)	27	0.61	0.24-0.89	0.36
Gym Exposure							
	No	20 (55.56)	16 (44.44)	36	0.42	0.08-0.87	
	Yes	15 (40.54)	22 (59.46)	37	0.64	0.18-0.93	0.12
Play Team Sports							
	No	20 (54.05)	17 (45.95)	37	0.47	0.19-0.77	
	Yes	15 (41.67)	21 (58.33)	36	0.58	0.27-0.84	0.45
Any Family Member Play Team Sports							
	No	11 (52.38)	10 (47.62)	21	0.48	0.19-0.78	
	Yes	23 (44.23)	29 (55.77)	52	0.54	0.20-0.85	0.62
Share Bath Towels							
	No	17 (42.50)	23 (57.50)	40	0.56	0.27-0.81	
	Yes	17 (51.52)	16 (48.48)	33	0.46	0.18-0.77	0.45
Share Hand Towels							
	No	5 (50.00)	5 (50.00)	10	0.53	0.18-0.85	
	Yes	30 (46.88)	34 (53.13)	64	0.52	0.18-0.84	0.98
Use Soap with Antimicrobials							
	No	6 (42.86)	8 (57.14)	14	0.61	0.26-0.88	
	Yes	29 (48.33)	31 (51.67)	60	0.51	0.14-0.86	0.50

Table C3 Continued

Number of Positive Environmental Sites							
	Zero	25 (51.02)	24 (48.98)	49	0.44	0.12-0.83	
	One	5 (55.56)	4 (44.44)	9	0.52	0.10-0.91	
	Two	3 (42.86)	4 (57.14)	7	0.59	0.18-0.90	
	Three	0 (0.00)	2 (100.00)	2	0.66	0.29-0.90	
	Four	1 (25.00)	3 (75.00)	4	0.72	0.34-0.93	
	Five	1 (33.33)	2 (66.67)	3	0.78	0.35-0.96	0.13
Farming Exposure ^s							
	No Exposure	24 (51.06)	23 (48.94)	47	0.51	0.13-0.87	
	Non-occupational Exposure	9 (47.37)	10 (52.63)	19	0.51	0.18-0.84	
	Large Facility Occupational Exposure	2 (40.00)	3 (60.00)	5	0.43	0.07-0.89	0.96
Healthcare Exposure							
	No Exposure	28 (51.85)	26 (48.15)	54	0.48	0.11-0.87	
	Non-occupational	7 (35.00)	13 (65.00)	20	0.66	0.29-0.90	0.19
Visited a Hospital							
	No	30 (51.72)	28 (48.28)	58	0.49	0.13-0.86	
	Yes	5 (31.25)	11 (68.75)	16	0.68	0.32-0.91	0.20

Table C3 Continued

Admitted to an Outpatient Center							
No	33 (47.14)	37 (52.86)	70	0.52	0.17-0.85		
Yes	2 (50.00)	2 (50.00)	4	0.58	0.14-0.92		0.83
Family Member was Hospitalized							
No	32 (47.76)	35 (52.24)	67	0.52	0.13-0.89		
Yes	3 (42.86)	4 (57.14)	7	0.58	0.16-0.91		0.77
Hospitalized							
No	34 (47.22)	38 (52.78)	72	0.52	0.18-0.85		
Yes	1 (50.00)	1 (50.00)	2	0.58	0.06-0.97		0.88
Have Asthma							
No	32 (49.23)	33 (50.77)	65	0.51	0.17-0.84		
Yes	3 (33.33)	6 (66.67)	9	0.61	0.22-0.89		0.62
Have Eczema							
No	33 (50.00)	33 (50.00)	66	0.50	0.11-0.89		
Yes	2 (25.00)	6 (75.00)	8	0.78	0.34-0.96		0.15
Have Skin Condition Other Than Eczema							
No	32 (50.00)	32 (50.00)	64	0.51	0.21-0.80		
Yes	3 (30.00)	7 (70.00)	10	0.75	0.36-0.95		0.18

Table C3 Continued

Have an Immune Disorder Other Than Diabetes							
No	35 (48.61)	37 (51.39)	72	---	---		
Yes	0 (0.00)	2 (100.00)	2	---	---		0.49 [#]
Took Immuno-compromising Medicines							
No	32 (45.71)	38 (54.29)	70	---	---		
Yes	3 (100.00)	0 (0.00)	3	---	---		0.11 [#]
Used Antibiotics							
No	26 (44.07)	33 (55.93)	59	0.55	0.19-0.86		
Yes	9 (60.00)	6 (40.00)	15	0.37	0.12-0.72		0.25
Had Skin or Soft Tissue Infection							
No	34 (47.22)	38 (52.78)	72	0.52	0.18-0.85		
Yes	1 (50.00)	1 (50.00)	2	0.55	0.06-0.96		0.93
Family Member with Skin or Soft Tissue Infection							
No	31 (46.27)	36 (53.73)	67	0.54	0.14-0.89		
Yes	3 (50.00)	3 (50.00)	6	0.49	0.12-0.87		0.82

Note: [¶]All risk factors adjusted for age at 9.65 years ^{*}Mean age for non-carriers is 9.35. Mean age for intermittent carriers is 8.80. Mean age for persistent carriers is years 12.79. [#]Fisher's Exact Test used to determine p-value. Probabilities and 95% CIs could not be determined for these risk factors due to small sample sizes. [§]Farming exposure refers to contact with livestock (swine, cattle, poultry, etc.) ^{**}One participant was enrolled as a minor, but was over 18 years of age. This participant is included in this table because the minor enrollment procedure was used.

