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# THE NHSN STANDARDIZED INFECTION RATIO (SIR)

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*A Guide to the SIR*  
(Based on 2015 National Baseline)

*Updated March 2024*



The Standardized Infection Ratio (SIR) is the primary summary measure used by the National Healthcare Safety Network (NHSN) to track healthcare-associated infections (HAIs). As NHSN grows, both in its user-base and surveillance capability, the SIR continues to evolve. Highlighting the SIR and changes resulting from an updated baseline, this document is intended to serve both as guidance for those who are new to this metric as well as a useful reference for more experienced infection prevention professionals.

## CORRECTIONS AND UPDATES AS OF March 2024

*Recent changes to this document are listed here:*

- Page 3: Corrections were made to the Table of Contents
- Page 11: The picture and description of the NHSN analysis reports treeview were updated to reflect the current version of the NHSN application
- Page 33: Updated language referring to the CMS Care Compare website
- Page 34: Updated reference to the CMS Hospital Inpatient Prospective Payment System (IPPS)

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## Overview of the Standardized Infection Ratio (SIR)

### What is the SIR?

The standardized infection ratio (SIR) is a summary measure used to track HAIs at a national, state, or local level over time. The SIR adjusts for various facility and/or patient-level factors that contribute to HAI risk within each facility. The method of calculating an SIR is similar to the method used to calculate the Standardized Mortality Ratio (SMR), a summary statistic widely used in public health to analyze mortality data. In HAI data analysis, the SIR compares the actual number of HAIs reported to the number that would be predicted, given the standard population (i.e., NHSN baseline), adjusting for several risk factors that have been found to be significantly associated with differences in infection incidence. In other words, an SIR greater than 1.0 indicates that more HAIs were observed than predicted; conversely, an SIR less than 1.0 indicates that fewer HAIs were observed than predicted. SIRs are currently calculated in NHSN for the following HAI types: central line-associated bloodstream infections (CLABSI), mucosal barrier injury laboratory-confirmed bloodstream infections (MBI-LCBI), catheter-associated urinary tract infections (CAUTI), surgical site infections (SSI), *Clostridioides difficile* infections (CDI), methicillin-resistant *Staphylococcus aureus* bloodstream infections (MRSA), and ventilator-associated events (VAE).

### Why not rates?

In the past, NHSN has published annual HAI rates for device-associated infections. These rates, or pooled means, were calculated using aggregate data reported to NHSN. The total number of infections was divided by the applicable number of device days for that time period. However, a problem with strictly using pooled mean rates is that they cannot reflect differences in risk between populations, and therefore lose comparability over time and across entities. For example, calculating rates from two facilities serving entirely different patient populations can lead to an unfair comparison. One solution to this problem is the stratification of pooled means, as was done with location-stratified CLABSI and CAUTI pooled means. However, this method only allows for comparison of rates within strata and does not lend itself to calculating an overall performance metric for a facility.

Instead, the SIR allows users to summarize data by more than a single stratum (e.g., location or procedure category), adjusting for differences in the incidence of infection among the strata. For example, NHSN allows users to obtain one CLABSI SIR for their facility, adjusting for all locations reported. Similarly, users can also obtain one CLABSI SIR for all intensive care units in their facility.

Additionally, the SIR allows for a comparison to the national benchmark from a baseline time period, and can be used to measure progress from a single point in time. In other words, the SIR permits comparisons between the number of infections experienced by a facility, group, or state to the number of infections that were predicted to have occurred based on national data (i.e., baseline data).

## How is the SIR calculated?

The SIR is calculated by dividing the number of observed infections by the number of predicted infections. The number of predicted infections is calculated using multivariable regression models generated from nationally aggregated data during a baseline time period. These models are applied to a facility's denominator and risk factor data to generate a predicted number of infections. Please refer to [the SIR Guide Supplement on page 14](#) for more details regarding the models.

$$SIR = \frac{\text{Observed (O) HAIs}}{\text{Predicted (P) HAIs}}$$

In order to enforce a minimum precision criterion, **SIRs are currently not calculated when the number of predicted infections is less than 1.0**. This rule was instituted to avoid the calculation and interpretation of statistically imprecise SIRs, which typically have extreme values.

## Calculating the Number of Predicted Infections

The number of predicted infections in NHSN is calculated based on the 2015 national HAI aggregate data and is adjusted for each facility using variables found to be significant predictors of HAI incidence. NHSN uses either a logistic regression model or a negative binomial regression model to perform this calculation. Logistic regression models are used when there is an opportunity for a single outcome for each exposure (e.g., SSI following a procedure). Negative binomial regression models are used when estimating incidence from a summarized population (e.g., CLABSIs in a Medical ICU). Examples in applying each model type are provided below.

### ❖ Example: Logistic Regression Model (SSI)

The logistic regression model is the specific type of model used for surgical site infection risk adjustment. At a high level, the model uses a set of fixed parameters (adjustment variables) to predict the log-odds of a surgical site infection following an inpatient procedure. To obtain the total number of predicted SSIs, the following steps are completed in NHSN:

1. Determine the log-odds for each procedure
2. Convert the log-odds into a probability, or risk of infection ( $\hat{p}$ ), for each procedure
3. Sum the risk of infections across all procedures in a given timeframe

The sum of the risks from a set of procedures will amount to the total number of predicted infections for that same set of procedures. *Table 1* below shows the risk factors found to be significant for abdominal hysterectomy (HYST) procedures (Complex 30-Day model) in NHSN. Note that each risk factor's contribution to the SIR varies, as represented by the parameter estimate for each factor. Parameter estimates describe the relationship between the variable and the risk of SSI; positive parameter estimates indicate that the risk of SSI increases with

increasing values of the variable. Negative parameter estimates indicate that the risk of SSI decreases with increasing values of the variable.

**Table 1. Risk Factors for SSI HYST: Complex 30-Day Model (2015 Baseline)**

<u>Factor</u>	<u>Parameter Estimate</u>	<u>P-value</u>	<u>Variable Coding</u>
<b>Intercept</b>	-5.1801	-	-
<b>Diabetes</b>	0.3247	<0.0001	<b>Yes= 1</b> <b>No= 0</b>
<b>ASA Score</b>	0.4414	<0.0001	<b>1= 1</b> <b>2= 2</b> <b>3= 3</b> <b>4/5= 4</b>
<b>Body Mass Index (BMI)</b>	0.1106	0.0090	<b>≥ 30= 1</b> <b>&lt; 30= 0</b>
<b>Patient Age</b>	-0.1501	<0.0001	<b>Patient’s age/10</b>
<b>Oncology Hospital</b>	0.5474	0.0005	<b>Oncology hospital= 1</b> <b>Non-oncology hospital= 0</b>

The parameter estimates from *Table 1* can be plugged into the following general logistic regression formula:

$$\text{logit}(\hat{p}) = \alpha + \beta_1 X_1 + \beta_2 X_2 + \dots + \beta_i X_i, \text{ where:}$$

- $\alpha$  = Intercept
- $\beta_i$  = Parameter Estimate
- $X_i$  = Value of Risk Factor (Categorical variables= 1 if present, 0 if not present. Refer to “Variable Coding” column in Table 1 above.)
- $i$  = Number of Predictors

The probability of SSI is calculated using the logistic regression model above, by utilizing the relationship between the log-odds and the probability (risk). Let’s say we have a patient (Patient 1) who is 32 years old, has diabetes, and a BMI score of 29. She had an ASA score of 2 and her procedure took place in an oncology hospital. We can use the model above to plug in these values as shown below:

$$\text{logit}(\hat{p}) = -5.1801 + 0.3247(\text{DIABETES}) + 0.4414(\text{ASA}) + 0.1106(\text{BMI}) - 0.1501(\text{AGE}) + 0.5474(\text{ONCOLOGY HOSPITAL})$$

$$\text{logit}(\hat{p}) = -5.1801 + 0.3247(1) + 0.4414(2) + 0.1106(0) - 0.1501(3.2) + 0.5474(1) = -3.9055$$

The value -3.9055 is the log-odds of SSI for Patient 1. To convert this value into the risk of SSI ( $\hat{p}$ ), we must use the logit function below:

$$\hat{p} = \frac{e^{\text{logit}(\hat{p})}}{1 + e^{\text{logit}(\hat{p})}}$$

$$\hat{p} = \frac{e^{-3.9055}}{1 + e^{-3.9055}}$$

$$\hat{p} = 0.020$$

Note that this can also be interpreted as a 2.0% risk of SSI for Patient 1. The probability of SSI is calculated for each procedure and then summed across all procedures to give the total number of predicted SSIs for this population. *Table 2* provides a partial list of 100 hypothetical patients who have undergone this particular procedure type and demonstrates how the total number of predicted SSIs is calculated.

**Table 2. Risk Factors for 100 Patients Undergoing a HYST Procedure (Complex 30-Day model)**

Patient	Diabetes	ASA score	BMI	Age	Oncology Hospital	SSI Identified?	Probability of SSI ( $\hat{p}$ )
1	Y	2	29	32	Y	1	0.020
2	N	3	35	49	Y	0	0.019
3	N	5	20	51	Y	1	0.026
.	.	.	.	.	.	.	.
.	.	.	.	.	.	.	.
100	N	4	27	27	Y	0	0.037
<b>TOTAL</b>	.	.	.	.	.	<b>8 (observed SSIs)</b>	<b>6.750 (predicted SSIs)</b>

Notice in the above table that the probability of SSI is different for each patient, given the risk factors present during the reported procedure.

The SIR can now be calculated for those 100 procedures as follows:

$$SIR = \frac{\text{Observed (O) HAIs}}{\text{Predicted (P) HAIs}} = \frac{8}{6.750} = 1.190$$

❖ Example: Negative Binomial Regression Model

Negative binomial regression models are used to calculate the number of predicted events for CLABSI, MBI-LCBI, CAUTI, VAE, MRSA bacteremia LabID, and *C. difficile* (CDI) LabID under the 2015 baseline. Below is a general formula for a negative binomial regression model.

$$\log(\lambda) = \alpha + \beta_1 X_1 + \beta_2 X_2 + \dots + \beta_i X_i, \text{ where:}$$

- $\alpha$  = Intercept
- $\beta_i$  = Parameter Estimate
- $X_i$  = Value of Risk Factor (Categorical variables: 1 if present, 0 if not present)
- $i$  = Number of Predictors

As an example, *Table 3* below represents the negative binomial regression model used to calculate the number of predicted healthcare facility-onset (HO) CDI LabID events in acute care hospitals under the 2015 baseline.

**Table 3. Risk Factors Used in the Acute Care Hospital CDI LabID Event Model**

<u>Factor</u>	<u>Parameter Estimate</u>	<u>P-value</u>
<b><i>Intercept</i></b>	-8.9463	<0.0001
<b>Inpatient community-onset (CO) admission prevalence rate</b>	0.7339	<0.0001
<b>CDI test type= EIA</b>	-0.1579	<0.0001
<b>CDI test type= NAAT</b>	0.1307	<0.0001
<b># ICU beds: ≥ 43</b>	0.7465	<0.0001
<b># ICU beds: 20-42</b>	0.7145	<0.0001
<b># ICU beds: 10-19</b>	0.6261	<0.0001
<b># ICU beds: 5-9</b>	0.4394	<0.0001
<b>Oncology hospital (facility type = HOSP-ONC)</b>	1.2420	<0.0001
<b>General acute care hospital (facility type = HOSP-GEN)</b>	0.3740	<0.0001
<b>Total facility bed size</b>	0.0003	<0.0001
<b>CDI LabID surveillance in ED or 24-hour observation location(s)</b>	0.1119	<0.0001
<b>Teaching facility (major, graduate, or undergraduate)</b>	0.0331	0.0028

The SIR for *C. difficile* LabID events in an acute care hospital is calculated on the facility-wide inpatient (FacWideIN) level for each quarter. More information on the details of the LabID Event SIR calculations can be found in the [SIR Guide Supplement on page 14](#).

We can input the model details from *Table 3* into the general negative binomial regression formula for CDI in acute care hospitals:



# predicted HO CDI =

$$\begin{aligned} & \text{Exp [ -8.9463} \\ & \quad + 0.7339 \text{ (CO prevalence rate)} \\ & \quad - 0.1579 \text{ (CDI test type = EIA)} \\ & \quad + 0.1307 \text{ (CDI test type = NAAT)} \\ & \quad + 0.7465 \text{ (ICU beds } \geq 43) \\ & \quad + 0.7145 \text{ (ICU beds: 20 – 42)} \\ & \quad + 0.6261 \text{ (ICU beds: 10-19)} \\ & \quad + 0.4394 \text{ (ICU beds: 5-9)} \\ & \quad + 1.2420 \text{ (Oncology hospital)} \\ & \quad + 0.3740 \text{ (General hospital)} \\ & \quad + 0.0003 \text{ (Total facility bed size)} \\ & \quad + 0.1119 \text{ (Reporting from ED or 24 hr. Obs)} \\ & \quad + 0.0331 \text{ (Teaching hospital) ] } \times \text{ CDI patient days} \end{aligned}$$

For most variables shown in parentheses in the equation above, you would replace the variable name (and therefore, multiply each parameter estimate) with a “1” or “0” depending on whether that factor is present in your facility (Yes= “1”, No= “0”). The inpatient CO prevalence rate and total number of beds are continuous variables and should be replaced with the actual values of the inpatient CO prevalence rate and total number of beds. The last step in the equation is to multiply the resulting value by the appropriate HAI denominator (i.e., patient days for MRSA/CDI events, or device days for CLABSI/MBI/CAUTI/VAE). In this example, we multiply by CDI patient days.

*Note: in NHSN, “CDI patient days” refers to the patient days entered on Row 3 of the FacWideIN monthly denominator forms, for an entire quarter. This value represents that total number of patient days from all inpatient units within the facility, with the exception of NICUs, well-baby units, and CMS-certified rehab and psych units.*

Let’s walk through an example of calculating the number of predicted CDI events for an acute care hospital for 2015 Q1. The facility in our example has reported 5,000 CDI patient days and 5 healthcare facility-onset CDI LabID events in 2015 Q1. After running the CDI rate tables in NHSN, the facility records that their 2015 Q1 CO admission prevalence rate was 1.25 per 100 admissions. The facility was using a NAAT CDI test type, has 5 ICU beds, is enrolled in NHSN as a children’s non-teaching hospital, and has 100 total beds. The facility has an Emergency Department and is thus reporting CDI data from this location per NHSN protocol.

In our example hospital, the completed formula looks like this:

$$\begin{aligned} & \text{Exp } [-8.9463 \\ & \quad + 0.7339 \text{ (1.25)} \\ & \quad - 0.1579 \text{ (0)} \\ & \quad + 0.1307 \text{ (1)} \\ & \quad + 0.7465 \text{ (0)} \\ & \quad + 0.7145 \text{ (0)} \\ & \quad + 0.6261 \text{ (0)} \\ & \quad + 0.4394 \text{ (1)} \\ & \quad + 1.2420 \text{ (0)} \\ & \quad + 0.3740 \text{ (0)} \\ & \quad + 0.0003 \text{ (100)} \\ & \quad + 0.1119 \text{ (1)} \\ & \quad + 0.0331 \text{ (0)} ] \times 5,000 = \boxed{3.321 \text{ predicted CDI LabID events}} \end{aligned}$$

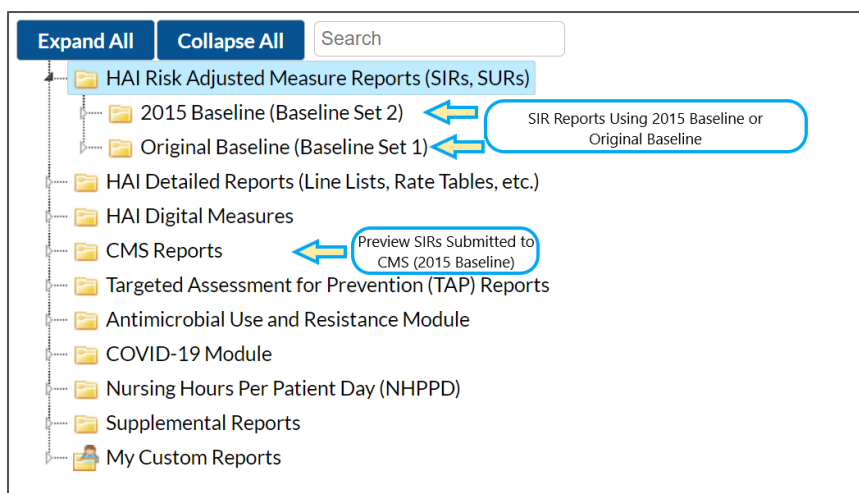
Because the facility was not using EIA test type, was not a general or oncology hospital, and was not a teaching hospital, the associated parameters in the model were not met. Therefore, the parameter estimates for each of those variables were multiplied by 0 and fell out of the equation.

To calculate the CDI LabID SIR, divide the number of observed HO CDI LabID events by the number of predicted HO CID LabID events. In our example:

$$SIR = \frac{5 \text{ observed HO CDI LabID events}}{3.321 \text{ predicted HO CDI LabID events}} = 1.506$$

### What SIR reports are available?

To run analysis reports in NHSN, users must first generate analysis data sets (Analysis > Generate Data Sets). NHSN recommends users regenerate data sets after entering new data into the application or before creating new reports. After data sets have been regenerated, users can select Analysis > Reports from the NHSN homepage to view all of the available reports. SIR Reports can be found under the "HAI Risk Adjusted Measure Reports" tab, and either the 2015 baseline or the original baseline can be selected. Each baseline folder contains HAI-specific subfolders with all of the available SIR reports, rate tables, and run charts for the chosen baseline. In addition, SIR reports are available that mirror the data submitted to the Centers for Medicare & Medicaid Services (CMS) Quality Reporting Programs. These reports can be found in the analysis folder titled "CMS Reports".



SIRs can be generated for data through 2016 using the original NHSN baselines by running reports in the "Baseline Set 1" reports folder. Data representing a later time period (i.e., starting in January 2017) can only be analyzed in NHSN using the new 2015 rebaseline models. Year 2016 is the final year of data that can use the original models to calculate SIRs. See [Additional Resources](#) for information about the original SIR baselines.



Note: SIRs calculated under the original baseline cannot be directly compared to SIRs calculated under the updated baseline. Additional information about NHSN Re-baseline can be found here:

[https://www.cdc.gov/nhsn/pdfs/training/2017/Dudeck\\_March21.pdf](https://www.cdc.gov/nhsn/pdfs/training/2017/Dudeck_March21.pdf);

[https://www.cdc.gov/nhsn/pdfs/training/2017/Dudeck\\_March22.pdf](https://www.cdc.gov/nhsn/pdfs/training/2017/Dudeck_March22.pdf).

## ***How do I Interpret the SIRs?***

### **SIR**

- If the SIR > 1.0, then more HAIs were observed than predicted, based on the 2015 national aggregate data.
- If the SIR < 1.0, then fewer HAIs were observed than predicted, based on the 2015 national aggregate data.
- If the SIR= 1.0, then the same number of HAIs were observed as predicted, based on the 2015 national aggregate data.
- Remember, the SIR is only calculated when the number of predicted infections is at least 1.0. When the predicted number of infections is less than 1.0, facilities have a few options for reviewing and interpreting HAI data in NHSN:
  - A longer time period can be included in the SIR calculation in order to reach the threshold of 1.0 predicted infection.
  - Infection rates can be used to track internal HAI incidence over time.
  - Run the TAP Reports to review the CAD (cumulative attributable difference, which is the difference between the number of observed infections and the number of predicted infections, multiplied by the SIR goal). Information and guidance about running TAP reports can be found in [Additional Resources](#).

### **P-value**

- In the context of the SIR, the p-value is a statistical measure that tells us whether the number of observed infections is statistically significantly different than the number of predicted infections (i.e., whether the SIR is significantly different from 1.0). NHSN calculates p-values using a mid-P exact test.
- Given the typical cutoff value of 0.05, if the p-value  $\leq$  0.05, we can conclude that the number of observed infections is statistically significantly different than the number of predicted infections.
- If the p-value > 0.05, then we can conclude that the number of observed infections is not statistically significantly different than the number of predicted infections.

### **95% Confidence Interval**

- The 95% confidence interval is a statistical range of values in which we have a high degree of confidence that the true SIR lies.

- If the confidence interval does not include the value of 1, then the SIR is significantly different than 1 (i.e., the number of observed infections is significantly different than the number of predicted infections).
  - Example: 95% confidence interval= (0.85, 0.92)
- If the confidence interval includes the value of 1, then the SIR is not significantly different than 1 (i.e., the number of observed infections is not significantly different than the number of predicted infections).
  - Example: 95% confidence interval= (0.85, 1.24)
- If the SIR is 0.000 (i.e., the observed infection count is 0 and the number of predicted infections is  $\geq 1.0$ ), then the lower bound of the 95% confidence interval will not be calculated.

As an example, let's take a look at the CLABSI SIR output. Below is a table showing the overall CLABSI SIR for a hospital during the first quarter of 2015.

orgID	summaryYQ	infCount	numPred	numcldays	SIR	SIR_pval	sir95ci
10018	2015Q1	5	2.365	1850	2.114	0.1251	0.775, 4.686

- During the first quarter (January– March) of 2015 (“summaryYQ”), there were 5 CLABSIs identified in our facility (“infCount”), and we observed a total of 1,850 central line days (“numcldays”) from the locations under surveillance.
- Based on the NHSN 2015 baseline data, 2.365 CLABSIs were predicted (“numPred”) in our facility.
- This results in an SIR of 2.114 (5/2.365), signifying that during this time period, our facility identified more CLABSIs than were predicted.
- Because the p-value (“SIR\_pval”) is above the significance level of 0.05 and the 95% confidence interval (“sir95ci”) includes the value of 1, we can conclude that our facility’s SIR is not statistically significant; in other words, our facility did not observe a statistically significantly different number of CLABSIs than predicted.

When analyzing these data as a Group user, an additional overall SIR will be calculated for all facilities in the Group. More information about using the Group function in NHSN can be found here:

<https://www.cdc.gov/nhsn/group-users/index.html>.

## SIR Guide Supplement: Risk Adjustment Factors Included in the SIR Calculations, 2015 Baseline

### Introduction to the SIR Guide Supplement

The following pages contain information on the risk factors used in the calculation of the number of predicted events for each HAI and facility type under the 2015 SIR baseline. This information is provided in order to aid in the interpretation of the SIR calculations produced by NHSN. The tables displayed in this document list the variables included in each risk adjustment model, as well as parameter estimates and standard errors. Some risk adjustment variables are broken into different levels, or categories (i.e., categorical variables), while other variables are treated as continuous variables without any categorization. Standard errors reflect the precision of the parameter estimate.

- **Categorical variables:**

*Example: “medical school affiliation” in the CAUTI Acute Care Hospital model, [page 23](#)*

Variables are categorized based on significant differences in HAI risk between the categories. Parameter estimates reflect the nature of the relationship between the variable and the risk of HAI. In the case of categorical variables, the risk of HAI in an individual category is compared to the risk of HAI in the “referent” category. A positive parameter estimate indicates that the risk of HAI in that category (and therefore, the number of predicted HAIs) is higher compared to the risk of HAI in the referent category. A negative parameter estimate indicates that the HAI risk in that category is lower compared to the HAI risk in the “referent” category.

- **Continuous variables:**

*Example: “facility bed size” in the CDI Acute Care Hospital model, [page 39](#)*

Parameter estimates reflect the nature of the relationship between the variable and the risk of HAI (and therefore, the number of predicted HAIs). For continuous variables, a positive parameter estimate indicates that the risk of HAI increases as the variable increases, while negative parameter estimates indicate that the risk of HAI decreases as the variable increases.

- **Derived variables:**

*Example: The proportion of admissions with traumatic and non-traumatic spinal cord dysfunction in the CAUTI IRF model, [page 26](#)*

Derived variables are variables created from two or more variables, and may involve summation, division, or multiplication. They may be categorical or continuous. Parameter estimates are interpreted as above if the derived variables are categorical or continuous.

## Risk Adjustment Factors Included in the SIR Calculation: 2015 Baseline

### CLABSI – Central Line-Associated Bloodstream Infection

The number of predicted CLABSIs is calculated using a negative binomial regression model (see [page 8](#) above for more information). Inpatient locations that were previously excluded from the original baseline are now included in the SIR under the 2015 baseline (e.g., Telemetry Ward, Mixed Acuity Ward). Refer to Table 1 below for a list of location types included in the Acute Care Hospital CLABSI SIR; if an inpatient location is not included in the model table below, then data from that location type will be excluded from the SIR due to insufficient 2015 baseline data. In addition, data from Governmental and Non-governmental Public Health Emergency (PHE) Facilities (facType as HOSP-PHE/G or HOSP-PHE/NG), will also be excluded from the SIR. In cases when the number of predicted events is less than 1.0, the SIR will not be calculated in NHSN. CLABSI events reported to NHSN as mucosal barrier injury (MBI-LCBI), or with extracorporeal life support (ECMO), or a ventricular assist device (VAD) (2019 events and later) are excluded from the numerator of the CLABSI SIR.

The number of predicted CLABSIs calculated under the 2015 baseline is risk adjusted based on the following variables found to be statistically significant predictors (*risk adjustment updated August 2018*):

**Table 1. CLABSI in Acute Care Hospitals (non-NICU locations)**

Parameter	Parameter Estimate	Standard Error	P-value
<i>Intercept</i>	-7.6325	0.0606	<0.0001
<u>CDC Location Code: Adult Critical Care Units, Oncology Critical Care Units</u> <i>Medical Cardiac Critical Care</i> <i>Surgical Cardiothoracic Critical Care</i> <i>Medical Critical Care</i> <i>Medical/Surgical Critical Care</i> <i>Neurologic Critical Care</i> <i>Neurosurgical Critical Care</i> <i>Medical Oncology Critical Care</i> <i>Medical/Surgical Oncology Critical Care</i> <i>Pediatric Oncology Critical Care</i> <i>Surgical Oncology Critical Care</i> <i>Prenatal Critical Care</i> <i>Respiratory Critical Care</i> <i>Surgical Critical Care</i>	0.3257	0.0435	<0.0001
<u>CDC Location Code: Pediatric Critical Care</u> <i>Pediatric Burn Critical Care</i> <i>Pediatric Cardiothoracic Critical Care</i> <i>Pediatric Medical/Surgical Critical Care</i> <i>Pediatric Medical Critical Care</i> <i>Pediatric Neurosurgical Critical Care</i> <i>Pediatric Surgical Critical Care</i> <i>Pediatric Trauma Critical Care</i>	0.5695	0.0699	<0.0001
CDC Location Code: Burn Critical Care (Adult)	1.4269	0.1125	<0.0001
CDC Location Code: Trauma Critical Care (Adult)	0.6287	0.0835	<0.0001
<u>CDC Location Code: Specialty Care Areas</u> <i>Inpatient Dialysis</i> <i>Solid Organ Transplant (adult)</i>	0.3766	0.1304	0.0039

<b>Parameter</b>	<b>Parameter Estimate</b>	<b>Standard Error</b>	<b>P-value</b>
<i>Solid Organ Transplant (pediatric)</i>			
<b>CDC Location Code: Step-down Units</b> <i>Adult Step-down Unit</i> <i>Oncology Step-down Unit</i> <i>Pediatric Step-down Unit</i> <i>Step-down Neonatal Nursery (Level II)</i>	0.2155	0.0521	<0.0001
<b>CDC Location Code: Select Adult Wards</b> <i>Medical Ward</i> <i>Medical/Surgical Ward</i> <i>Neurology Ward</i> <i>Neurosurgical Ward</i> <i>Surgical Ward</i> <i>Telemetry Ward</i>	0.1797	0.0427	<0.0001
<b>CDC Location Code: Oncology Wards</b> <i>ONC General Hematology/Oncology Ward</i> <i>ONC Pediatric General Hematology/Oncology Ward</i> <i>ONC Leukemia Ward</i> <i>ONC Leukemia/Lymphoma Ward</i> <i>ONC Lymphoma Ward</i> <i>ONC Solid Tumor Ward</i>	0.3698	0.0550	<0.0001
<b>CDC Location Code: Oncology Stem Cell Transplant Wards</b> <i>ONC Hematopoietic Stem Cell Transplant Ward (adult)</i> <i>ONC Pediatric Hematopoietic Stem Cell Transplant Ward</i>	0.6876	0.0816	<0.0001
<b>CDC Location Code: Pediatric Wards &amp; Nurseries</b> <i>Pediatric Behavioral Health Ward</i> <i>Pediatric Burn Ward</i> <i>Pediatric Medical Ward</i> <i>Pediatric Medical/Surgical Ward</i> <i>Pediatric Neurosurgical Ward</i> <i>Well Baby Nursery (Level I)</i> <i>Pediatric Neurology Ward</i> <i>Pediatric Orthopedic Ward</i> <i>Pediatric Rehabilitation Ward (non-CMS)</i> <i>Pediatric Surgical Ward</i>	0.1912	0.0704	0.0066
<b>CDC Location Code: All Other Wards</b> <i>Adult Mixed Acuity</i> <i>Mixed Age Mixed Acuity</i> <i>Pediatric Mixed Acuity</i> <i>Oncology Mixed Acuity</i> <i>Antenatal Care Ward</i> <i>Burn Ward</i> <i>Behavioral Health/Psych Ward</i> <i>Adolescent Behavioral Health Ward</i> <i>Ear/Nose/Throat Ward</i> <i>Gastrointestinal Ward</i> <i>Gerontology Ward</i> <i>Genitourinary Ward</i>	REFERENT	-	-



Parameter	Parameter Estimate	Standard Error	P-value
<i>Gynecology Ward</i>	REFERENT (continued from previous page)	-	-
<i>Jail Unit</i>			
<i>Labor and Delivery Ward</i>			
<i>Labor, Delivery, Recovery, Postpartum Suite (LDRP)</i>			
<i>Orthopedic Ward</i>			
<i>Plastic Surgery Ward</i>			
<i>Postpartum Ward</i>			
<i>Pulmonary Ward</i>			
<i>Rehabilitation Ward (non-CMS)</i>			
<i>Stroke (Acute) Ward</i>			
<i>Orthopedic Trauma Ward</i>			
<i>Vascular Surgery Ward</i>			
<i>Chronic Care Unit</i>			
<i>Chronic Behavioral Health/Psychiatric Unit</i>			
<i>Inpatient Hospice</i>			
<i>Chronic Ventilator Dependent</i>			
<i>Chronic Rehabilitation Unit</i>			
Facility bed size*: $\geq 224$ beds	0.2571	0.0471	<0.0001
Facility bed size*: 94 - 223 beds	0.1160	0.0493	0.0187
Facility bed size*: $\leq 93$ beds	REFERENT	-	-
Medical school affiliation*: Major	0.2627	0.0211	<0.0001
Medical school affiliation*: Graduate	0.1494	0.0244	<0.0001
Medical school affiliation*: Undergraduate/Non-teaching	REFERENT	-	-
<u>Facility type: (based on NHSN enrollment)</u>	0.1429	0.0526	0.0066
<i>Children's</i>			
<i>Military</i>			
<i>Veterans' Affairs</i>			
<i>Women's</i>			
<i>Women's and Children's</i>			
<u>Facility type: (based on NHSN enrollment)</u>	REFERENT	-	-
<i>General Acute Care</i>			
<i>Oncology</i>			
<i>Orthopedic</i>			
<i>Psychiatric</i>			
<i>Surgical</i>			

\* Facility bed size and medical school affiliation are taken from the [Annual Hospital Survey](#).

Note: For data from Specialty Care Areas and Oncology locations, CLABSI events are eligible for inclusion in the SIR regardless of the type of central line (temporary or permanent). Similarly, total central line days (numcldays) used in the SIR calculation are summed from temporary and permanent central line days reported. If a CLABSI event is reported from a month/location with missing denominator data for either temporary or permanent central lines, then the data for that month/location will be excluded from SIR calculations.

**Table 2. CLABSI in Acute Care Hospital NICUs (Level II/III, Level III, and Level IV NICU locations)**

<b>Parameter</b>	<b>Parameter Estimate</b>	<b>Standard Error</b>	<b>P-value</b>
<i>Intercept</i>	-7.2573	0.0553	<0.0001
Birthweight A: ≤ 750 grams	1.2780	0.0745	<0.0001
Birthweight B: 751-1000 grams	0.9780	0.0791	<0.0001
Birthweight C: 1001-1500 grams	0.4579	0.0843	<0.0001
Birthweight D & E: 1501-2500 grams and > 2500 grams	REFERENT	-	-

Note: For NICUs, CLABSI events as well as central line days are reported by birthweight category. If a CLABSI event is reported from a month/location with missing denominator data for one of the birthweight categories, then data for that month/location will be excluded from SIR calculations.

**Table 3. CLABSI in Critical Access Hospitals (CAHs)**

<b>Parameter</b>	<b>Parameter Estimate</b>	<b>Standard Error</b>	<b>P-value</b>
<i>Intercept*</i>	-8.2066	0.1967	<0.0001

\* None of the variables investigated were statistically significantly associated with CLABSIs in CAHs. The predicted number of CLABSI events for CAHs is calculated using the 2015 national CAH CLABSI pooled mean (i.e., intercept-only model).

**Table 4. CLABSI in Long-Term Acute Care Hospitals (LTACHs)**

<b>Parameter</b>	<b>Parameter Estimate</b>	<b>Standard Error</b>	<b>P-value</b>
<i>Intercept</i>	-7.8328	0.1307	<0.0001
Location Type: ICU	0.6716	0.1031	<0.0001
Location Type: Ward	REFERENT	-	-
Facility bed size*: ≥ 45 beds	0.2819	0.0686	<0.0001
Facility bed size*: < 45 beds	REFERENT	-	-
Average length of stay*: ≥ 28 days	0.1481	0.0708	0.0365
Average length of stay*: < 28 days	REFERENT	-	-
Proportion of admissions on a ventilator*: ≥ 0.328	0.3907	0.0971	<0.0001
Proportion of admissions on a ventilator*: ≥ 0.125 and < 0.328	0.2127	0.0859	0.0133
Proportion of admissions on a ventilator*: < 0.125	REFERENT	-	-
Proportion of admissions on hemodialysis*: ≥ 0.138	0.5785	0.1341	<0.0001
Proportion of admissions on hemodialysis*: ≥ 0.008 and < 0.138	0.5090	0.1296	<0.0001
Proportion of admissions on hemodialysis*: < 0.008	REFERENT	-	-

\* Facility bed size, average length of stay, and admission proportions are taken from the [Annual LTACH Survey](#). Average length of stay is calculated as: total # of annual patient days / total # of annual admissions.

**Table 5. CLABSI in Inpatient Rehabilitation Facilities (IRFs): Free-standing Rehabilitation Hospitals and CMS-Certified IRF Units Within a Hospital**

<b>Parameter</b>	<b>Parameter Estimate</b>	<b>Standard Error</b>	<b>P-value</b>
<i>Intercept</i>	-8.6717	0.3579	<0.001
Proportion of admissions with stroke*: ≥ 0.135	0.7707	0.3222	0.0168
Proportion of admissions with stroke*: < 0.135	REFERENT	-	-

<b>Parameter</b>	<b>Parameter Estimate</b>	<b>Standard Error</b>	<b>P-value</b>
Proportion of admissions in other non-specific diagnostic categories*: $\geq 0.197$	0.4452	0.2051	0.0300
Proportion of admissions in other non-specific diagnostic categories*: $< 0.197$	REFERENT	-	-

\* Admission proportions are taken from the [Annual IRF Survey](#). “Other non-specific diagnostic categories” include all other primary diagnoses not listed specifically on the Annual IRF Survey.



## Risk Adjustment Factors Included in the SIR Calculation: 2015 Baseline

### MBI-LCBI – Mucosal Barrier Injury Laboratory-Confirmed Bloodstream Infection

The number of predicted MBI-LCBIs is calculated using a negative binomial regression model (see [page 8](#) above for more information) and is only available for acute care hospitals. Only CLABSI events reported to NHSN as mucosal barrier injury (MBI-LCBI) are included in the numerator of the MBI-LCBI SIR. Refer to Table 1 below for a list of location types included in the Acute Care Hospital MBI-LCBI SIR; if an inpatient location is not included in the model table below, then data from that location type will be excluded from the SIR due to insufficient 2015 baseline data. In addition, data from Governmental and Non-governmental Public Health Emergency (PHE) Facilities (facType as HOSP-PHE/G or HOSP-PHE/NG) will also be excluded from the SIR. In cases when the number of predicted events is less than 1.0, the SIR will not be calculated in NHSN.

*\*Note: The variables included in the MBI risk adjustment model for acute care hospitals are shown below. The MBI-LCBI SIR is not submitted to CMS.*

The number of predicted MBI-LCBI events calculated under the 2015 baseline is risk adjusted based on the following variables found to be statistically significant predictors:

**Table 1. MBI-LCBI in Acute Care Hospitals**

<b>Parameter</b>	<b>Parameter Estimate</b>	<b>Standard Error</b>	<b>P-value</b>
<i>Intercept</i>	-10.9284	0.1397	<0.0001
<b>CDC Location Code: Adult Critical Care Units</b>			
<i>Burn Critical Care</i>			
<i>Medical Cardiac Critical Care</i>			
<i>Surgical Cardiothoracic Critical Care</i>			
<i>Medical Critical Care</i>			
<i>Medical-Surgical Critical Care</i>	-0.5102	0.0938	<0.0001
<i>Neurologic Critical Care</i>			
<i>Neurosurgical Critical Care</i>			
<i>Prenatal Critical Care</i>			
<i>Respiratory Critical Care</i>			
<i>Surgical Critical Care</i>			
<i>Trauma Critical Care</i>			
<b>CDC Location Code: Oncology Critical Care Units</b>			
<i>Oncology Medical Critical Care</i>			
<i>Oncology Medical-Surgical Critical Care</i>	2.6269	0.4176	<0.0001
<i>ONC Pediatric Critical Care</i>			
<i>Oncology Surgical Critical Care</i>			
<b>CDC Location Code: Pediatric Critical Care Units</b>			
<i>Pediatric Burn Critical Care</i>			
<i>Pediatric Surgical Cardiothoracic Critical Care</i>			
<i>Pediatric Medical-Surgical Critical Care</i>			
<i>Pediatric Medical Critical Care</i>			
<i>Pediatric Neurosurgical Critical Care</i>	0.7732	0.1803	<0.0001
<i>Pediatric Surgical Critical Care</i>			
<i>Pediatric Trauma Critical Care</i>			
<b>CDC Location Code: Step-down Units</b>			
<i>Adult Step-down Unit</i>	-0.7252	0.2004	0.0003

<b>Parameter</b>	<b>Parameter Estimate</b>	<b>Standard Error</b>	<b>P-value</b>
<i>Step-down Neonatal Nursery (Level II)</i> <i>Pediatric Step-down Unit</i>			
<b>CDC Location Code: Oncology Wards and Step-down Units</b> <i>ONC Step-down Unit</i> <i>ONC General Hematology-Oncology Ward</i> <i>ONC Hematopoietic Stem Cell Transplant Ward</i> <i>ONC Leukemia Ward</i> <i>ONC Leukemia/Lymphoma Ward</i> <i>ONC Lymphoma Ward</i> <i>ONC Solid Tumor Ward</i>	3.1224	0.0901	<0.0001
<b>CDC Location Code: Pediatric Oncology Wards</b> <i>ONC Pediatric General Hematology/Oncology Ward</i> <i>ONC Pediatric Hematopoietic Stem Cell Transplant Ward</i>	3.1967	0.1710	<0.0001
<b>CDC Location Code: Pediatric Wards</b> <i>Pediatric Behavioral Health Ward</i> <i>Pediatric Burn Ward</i> <i>Pediatric Medical-Surgical Ward</i> <i>Pediatric Medical Ward</i> <i>Pediatric Neurosurgical Ward</i> <i>Pediatric Neurology Ward</i> <i>Pediatric Orthopedic Ward</i> <i>Pediatric Rehabilitation Ward (within Hospital)</i> <i>Pediatric Surgical Ward</i>	1.3335	0.1464	<0.0001
<b>CDC Location Code: All Other Wards</b> <i>Mixed Age Mixed Acuity Unit</i> <i>Adult Mixed Acuity Unit</i> <i>Pediatric Mixed Acuity Unit</i> <i>Oncology Mixed Acuity Unit</i> <i>Dialysis Specialty Care Area</i> <i>Solid Organ Transplant Specialty Care Area</i> <i>Pediatric Solid Organ Transplant Specialty Care Area</i> <i>Antenatal Care Ward</i> <i>Burn Ward</i> <i>Behavioral Health/Psych Ward</i> <i>Adolescent Behavioral Health Ward</i> <i>Ear, Nose, Throat Ward</i> <i>Gastrointestinal Ward</i> <i>Gerontology Ward</i> <i>Genitourinary Ward</i> <i>Gynecology Ward</i> <i>Jail Unit</i> <i>Labor and Delivery Ward</i> <i>Labor, Delivery, Recovery, Postpartum Suite</i> <i>Medical Ward</i> <i>Medical-Surgical Ward</i> <i>Neurology Ward</i> <i>Neurosurgical Ward</i> <i>Well Baby Nursery (Level I)</i> <i>Orthopedic Ward</i>	REFERENT	-	-

Parameter	Parameter Estimate	Standard Error	P-value
<i>Plastic Surgery Ward</i>			
<i>Postpartum Ward</i>			
<i>Pulmonary Ward</i>			
<i>Rehabilitation Ward (within Hospital)</i>			
<i>Surgical Ward</i>			
<i>Stroke (Acute) Ward</i>			
<i>Telemetry Ward</i>			
<i>Orthopedic Trauma Ward</i>			
<i>Vascular Surgery Ward</i>			
<i>Chronic Care Unit</i>			
<i>Chronic Behavioral Health/Psych Unit</i>			
<i>Inpatient Hospice</i>			
<i>Ventilator Dependent Unit</i>			
<i>Chronic Rehabilitation Unit</i>			
Facility bed size*: $\geq 149$ beds	0.5422	0.1389	<0.0001
Facility bed size*: <149 beds	REFERENT	-	-
Medical school affiliation*: Major	0.4113	0.0699	<0.0001
Medical school affiliation*: Graduate/Undergraduate/Non-teaching	REFERENT	-	-

\* Facility bed size and medical school affiliation are taken from the [Annual Hospital Survey](#).

## Risk Adjustment Factors Included in the SIR Calculation: 2015 Baseline

### CAUTI – Catheter-Associated Urinary Tract Infection

The number of predicted CAUTIs is calculated using a negative binomial regression model (see [page 8](#) above for more information). Previously excluded inpatient locations from the original baseline are included in the SIR under the 2015 baseline (e.g., Telemetry Ward, Mixed Acuity Ward). Refer to Table 1 below for a list of location types included in the Acute Care Hospital CAUTI SIR; if an inpatient location is not included in the model table below, then data from that location type will be excluded from the SIR due to insufficient 2015 baseline data. In addition, data from Governmental and Non-governmental Public Health Emergency (PHE) Facilities (facType as HOSP-PHE/G or HOSP-PHE/NG) will also be excluded from the SIR. In cases when the number of predicted events is less than 1.0, the SIR will not be calculated in NHSN.

The number of predicted CAUTIs calculated under the 2015 baseline is risk adjusted based on the following variables found to be statistically significant predictors (*risk adjustment updated July 2017*):

**Table 1. CAUTI in Acute Care Hospitals**

<b>Parameter</b>	<b>Parameter Estimate</b>	<b>Standard Error</b>	<b>P-value</b>
<i>Intercept</i>	-10.2667	0.1618	<0.0001
CDC Location Code: Burn Critical Care	3.3318	0.1580	<0.0001
CDC Location Code: Cardiac Critical Care	2.5703	0.1301	<0.0001
CDC Location Code: Medical Critical Care	2.3834	0.1250	<0.0001
CDC Location Code: Neurologic Critical Care and Neurosurgical Critical Care	3.3675	0.1285	<0.0001
CDC Location Code: Surgical Critical Care	2.7034	0.1270	<0.0001
CDC Location Code: Trauma Critical Care	3.1104	0.1344	<0.0001
<u>CDC Location Code: Other Critical Care</u> <i>Surgical Cardiothoracic Critical Care</i> <i>Medical/Surgical Critical Care</i> <i>Prenatal Critical Care</i> <i>Respiratory Critical Care</i>	2.3661	0.1214	<0.0001
<u>CDC Location Code: Oncology Critical Care/Step-down</u> <i>Oncology Medical Critical Care</i> <i>Oncology Medical/Surgical Critical Care</i> <i>Surgical Oncology Critical Care</i> <i>Pediatric Oncology Critical Care</i> <i>Oncology Mixed Acuity Unit</i> <i>Oncology Step-Down Unit</i>	2.2171	0.2239	<0.0001
CDC Location Code: Pediatric Cardiothoracic Critical Care	2.0965	0.2322	<0.0001
<u>CDC Location Code: Other Pediatric Critical Care</u> <i>Pediatric Burn Critical Care</i> <i>Pediatric Medical/Surgical Critical Care</i> <i>Pediatric Medical Critical Care</i> <i>Pediatric Neurosurgical Critical Care</i> <i>Pediatric Surgical Critical Care</i> <i>Pediatric Trauma Critical Care</i>	2.6419	0.1461	<0.0001
<u>CDC Location Code: Mixed Acuity</u> <i>Adult Mixed Acuity Unit</i>	2.3378	0.1416	<0.0001

<b>Parameter</b>	<b>Parameter Estimate</b>	<b>Standard Error</b>	<b>P-value</b>
<i>Pediatric Mixed Acuity Unit</i> <i>Mixed Age Mixed Acuity Unit</i>			
<b>CDC Location Code: Adult Step-down Unit</b>	2.4800	0.1235	<0.0001
<b>CDC Location Code: Pediatric Step-down Unit</b> <i>Neonatal Step-down Nursery (Level II)</i> <i>Pediatric Step-down Unit</i>	2.3616	0.5351	<0.0001
<b>CDC Location Code: Solid Organ Transplant</b> <i>Solid Organ Transplant SCA</i> <i>Pediatric Solid Organ Transplant SCA</i>	2.3900	0.1979	<0.0001
<b>CDC Location Code: Adult Burn Ward</b>	2.4564	0.3396	<0.0001
<b>CDC Location Code: Behavioral/Psychiatric Ward</b>	3.2503	0.2207	<0.0001
<b>CDC Location Code: Pulmonary Ward</b>	2.5024	0.1664	<0.0001
<b>CDC Location Code: Rehabilitation Ward (non-CMS)</b>	3.3578	0.2700	<0.0001
<b>CDC Location Code: Neurology and Stroke</b> <i>Neurologic Ward</i> <i>Neurosurgical Ward</i> <i>Stroke Ward</i>	2.8223	0.1314	<0.0001
<b>CDC Location Code: Orthopedic Ward</b> <i>Orthopedic Ward</i> <i>Orthopedic Trauma Ward</i>	1.9992	0.1300	<0.0001
<b>CDC Location Code: Other Wards</b> <i>Inpatient Dialysis SCA</i> <i>Gerontology Ward</i> <i>Jail Unit</i> <i>Medical Ward</i> <i>Telemetry Ward</i>	2.3576	0.1216	<0.0001
<b>CDC Location Code: Other Wards</b> <i>Ear, Nose, Throat Ward</i> <i>Gastroenterology Ward</i> <i>Genitourinary Ward</i> <i>Medical/Surgical Ward</i> <i>Plastic Surgery Ward</i> <i>Surgical Ward</i> <i>Vascular Surgery Ward</i>	2.2532	0.1210	<0.0001
<b>CDC Location Code: Hematology</b> <i>General Hematology/Oncology Ward</i> <i>Hematopoietic Stem Cell Transplant Ward</i>	2.6125	0.1315	<0.0001
<b>CDC Location Code: Pediatric Oncology</b> <i>Pediatric Hematology/Oncology Ward</i> <i>Pediatric Hematopoietic Stem Cell Transplant Ward</i>	2.7077	0.2915	<0.0001
<b>CDC Location Code: Adult Oncology Wards</b> <i>Leukemia Ward</i> <i>Lymphoma Ward</i> <i>Leukemia/Lymphoma Ward</i> <i>Solid Tumor Ward</i>	2.2253	0.2001	<0.0001
<b>CDC Location Code: Pediatric Wards</b> <i>Adolescent Behavioral Ward</i> <i>Pediatric Behavioral Ward</i>	1.8899	0.1712	<0.0001



<b>Parameter</b>	<b>Parameter Estimate</b>	<b>Standard Error</b>	<b>P-value</b>
<i>Pediatric Burn Ward</i> <i>Pediatric Medical/Surgical Ward</i>			
<i>Pediatric Medical Ward</i> <i>Pediatric Neurosurgical Ward</i> <i>Pediatric Neurologic Ward</i> <i>Pediatric Orthopedic Ward</i> <i>Pediatric Rehabilitation Ward (non-IRF)</i> <i>Pediatric Surgical Ward</i> <i>Well-baby Nursery</i>			
<b>CDC Location Code: Chronic Care</b> <i>Chronic Care Unit</i> <i>Chronic Behavioral Health/Psychiatric Unit</i> <i>Chronic Rehabilitation Unit</i> <i>Inpatient Hospice</i> <i>Ventilator Dependent Unit</i>	2.7695	0.1855	<0.0001
<b>CDC Location Code: Labor and Delivery, Gynecology</b> <i>Antenatal Ward</i> <i>Gynecology Ward</i> <i>Labor and Delivery Ward</i> <i>Labor, Delivery, Postpartum Ward</i> <i>Postpartum Ward</i>	REFERENT	-	-
Medical school affiliation*: Major	0.3744	0.0195	<0.0001
Medical school affiliation*: Graduate	0.1313	0.0220	<0.0001
Medical school affiliation*: Undergraduate/Non-teaching	REFERENT	-	-
Facility bed size*: ≥ 215 beds	0.4901	0.0429	<0.0001
Facility bed size*: 87-214 beds	0.2871	0.0445	<0.0001
Facility bed size*: ≤ 86 beds	REFERENT	-	-
<b>Facility type: (based on NHSN enrollment)</b> <i>General Acute Care Hospital</i> <i>Military Hospital</i> <i>Psychiatric Hospital</i> <i>Oncology Hospital</i> <i>Veterans' Affairs Hospital</i>	0.3927	0.1069	0.0002
Facility type: Children's Hospital	0.4888	0.1556	0.0017
<b>Facility type: (based on NHSN enrollment)</b> <i>Orthopedic Hospital</i> <i>Surgical Hospital</i> <i>Women's Hospital</i> <i>Women's and Children's Hospital</i>	REFERENT	-	-

\* Medical school affiliation and facility bed size are taken from the [Annual Hospital Survey](#).

**Table 2. CAUTI in Critical Access Hospitals (CAHs)**

<b>Parameter</b>	<b>Parameter Estimate</b>	<b>Standard Error</b>	<b>P-value</b>
<i>Intercept</i>	-7.3337	0.0970	<0.0001
Medical school affiliation*: Undergraduate	1.3191	0.4744	0.0054
Medical school affiliation*: Major/Graduate/Non-teaching	REFERENT	-	-

\* Medical school affiliation is taken from the [Annual Hospital Survey](#).

**Table 3. CAUTI in Long-Term Acute Care Hospitals (LTACHs)**

<b>Parameter</b>	<b>Parameter Estimate</b>	<b>Standard Error</b>	<b>P-value</b>
<i>Intercept</i>	-6.8683	0.0773	<0.0001
Average length of stay*: $\geq 29.33$ days	0.5379	0.0837	<0.0001
Average length of stay*: 26.42 – 29.32 days	0.2779	0.0876	0.0015
Average length of stay*: $\leq 26.41$ days	REFERENT	-	-
Setting**: Freestanding	0.1700	0.0716	0.0176
Setting**: Within a Hospital	REFERENT	-	-
Location Type: ICU	0.3153	0.1072	0.0033
Location Type: Ward	REFERENT	-	-

\* Average length of stay is taken from the [Annual LTACH Survey](#). It is calculated as: total # of annual patient days / total # of annual admissions.

\*\* LTACH Setting (free-standing vs. within a hospital) is taken from the [Annual LTACH Survey](#).

**Table 4. CAUTI in Inpatient Rehabilitation Facilities (IRFs): Free-standing Rehabilitation Hospitals and CMS-Certified IRF Units Within a Hospital**

<b>Parameter</b>	<b>Parameter Estimate</b>	<b>Standard Error</b>	<b>P-value</b>
<i>Intercept</i>	-6.8305	0.0848	<0.0001
Setting*: Within a Hospital	0.2897	0.0841	0.0006
Setting*: Freestanding	REFERENT	-	-
Proportion of admissions with traumatic and non-traumatic spinal cord dysfunction**: $\geq 0.05$	0.3603	0.0832	<0.0001
Proportion of admissions with traumatic and non-traumatic spinal cord dysfunction**: $< 0.05$	REFERENT	-	-
Proportion of admissions with stroke**: $\geq 0.24$	0.2750	0.0798	0.0006
Proportion of admissions with stroke**: $< 0.24$	REFERENT	-	-

\* IRF Setting is taken from the [Annual IRF Survey](#) and NHSN enrollment/location mapping data. “Within a hospital” includes CMS-certified IRF units mapped as locations within a hospital, as well as Rehabilitation hospitals enrolled as unique facilities in NHSN in which the facility indicated “healthcare facility-based” on their annual IRF survey.

\*\* Proportion of annual admissions with primary diagnoses are taken from the Annual IRF Survey and are calculated as: # of admissions with the primary diagnosis (stroke, or traumatic/non-traumatic spinal cord dysfunction) / total # of annual admissions.

## Risk Adjustment Factors Included in the SIR Calculation: 2015 Baseline

### VAE – Ventilator-Associated Events

#### A. Total VAE in Long-Term Acute Care Hospitals (LTACHs)

The number of predicted VAE events is calculated using a negative binomial regression model (see [page 8](#) above for more information). Separate VAE SIRs are available for “Total VAE” and “IVAC Plus”. The “Total VAE” SIR includes events identified as ventilator-associated condition (VAC), infection-related ventilator-associated complication (IVAC), and possible ventilator-associated pneumonia (pVAP). In cases when the number of predicted events is less than 1.0, the SIR will not be calculated in NHSN.

The number of predicted “Total VAE” events calculated under the 2015 baseline is risk adjusted based on the following variables found to be statistically significant predictors of Total VAE incidence:

**Table 1. Total VAE in Long-Term Acute Care Hospitals (LTACHs)**

Parameter	Parameter Estimate	Standard Error	P-value
Intercept	-8.3689	0.3361	<0.0001
Facility bed size <sup>†</sup> : ≥ 32 beds	0.4645	0.1562	0.0030
Facility bed size <sup>†</sup> : < 32 beds	REFERENT	-	-
Proportion of admissions on hemodialysis*: > 0.11	-0.4098	0.1190	0.0006
Proportion of admissions on hemodialysis*: ≤ 0.11	REFERENT	-	-
Proportion of admissions on ventilator*: > 0.18	0.9313	0.1813	<0.0001
Proportion of admissions on ventilator*: ≤ 0.18	REFERENT	-	-
Location type: ICU	0.4118	0.1598	0.0099
Location type: Ward	REFERENT	-	-
Average length of stay <sup>**</sup> : ≥ 25 days	1.0940	0.2602	<0.0001
Average length of stay <sup>**</sup> : < 25 days	REFERENT	-	-

† Facility bed size is taken from the [Annual LTACH Survey](#).

\* Proportion of annual admissions on a ventilator (or hemodialysis) is taken from the [Annual LTACH Survey](#). It is calculated as: number of admissions on a ventilator (or hemodialysis) / total # of annual admissions.

\*\*Average length of stay is taken from the [Annual LTACH Survey](#). It is calculated as: # annual patient days / # annual admissions.

#### B. Infection-related Ventilator-Associated Complication (IVAC) Plus in Long-Term Acute Care Hospitals (LTACHs)

The number of predicted VAE events is calculated using a negative binomial regression model (see [page 8](#) above for more information). Separate VAE SIRs are available for “Total VAE” and “IVAC Plus”. The “IVAC Plus” SIR includes events identified as IVAC and possible ventilator-associated pneumonia (pVAP). In cases when the number of predicted events is less than 1.0, the SIR will not be calculated in NHSN.

The number of predicted “IVAC Plus” events calculated under the 2015 baseline is risk adjusted based on the following variables found to be statistically significant predictors of “IVAC Plus” incidence:

**Table 1. IVAC Plus in Long-Term Acute Care Hospitals**

<b>Parameter</b>	<b>Parameter Estimate</b>	<b>Standard Error</b>	<b>P-value</b>
<i>Intercept</i>	-9.9593	0.5891	< 0.0001
Facility bed size <sup>†</sup> : ≥ 32 beds	1.1201	0.3633	0.0020
Facility bed size <sup>†</sup> : < 32 beds	REFERENT	-	-
Proportion of admissions on a ventilator*: > 0.18	0.7130	0.3151	0.0236
Proportion of admissions on a ventilator*: ≤ 0.18	REFERENT	-	-
Average length of stay <sup>**</sup> : ≥ 25 days	0.8166	0.4157	0.0495
Average length of stay <sup>**</sup> : < 25 days	REFERENT	-	-

† Facility bed size is taken from the [Annual LTACH Survey](#).

\* Proportion of annual admissions on a ventilator is taken from the [Annual LTACH Survey](#). It is calculated as: number of admissions on a ventilator / total # of annual admissions.

\*\* Average length of stay is taken from the [Annual LTACH Survey](#). It is calculated as: total # of annual patient days / total # of annual admissions.

### C. Total VAE in Acute Care Hospitals (ACHs)

The number of predicted VAE events is calculated using a negative binomial regression model (see [page 8](#) above for more information). Separate VAE SIRs are available for “Total VAE” and “IVAC Plus”. The “Total VAE” SIR includes events identified as Ventilator-Associated Condition (VAC), Infection-related Ventilator-Associated Complication (IVAC), and Possible Ventilator-Associated Pneumonia (PVAP). Refer to Table 1 below for a list of location types included in the Acute Care Hospital Total VAE SIR; if an inpatient location is not included in the model table below, then data from that location type will be excluded from the SIR due to insufficient 2015 baseline data. In addition, data from Governmental and Non-governmental Public Health Emergency (PHE) Facilities (facType as HOSP-PHE/G or HOSP-PHE/NG) will also be excluded from the SIR. In cases when the number of predicted events is less than 1.0, the SIR will not be calculated in NHSN.

The number of predicted “Total VAE” events calculated under the 2015 baseline is risk adjusted based on the following variables found to be statistically significant predictors of Total VAE incidence:

**Table 1. Total VAE in Acute Care Hospitals (ACHs)**

<b>Parameter</b>	<b>Parameter Estimate</b>	<b>Standard Error</b>	<b>P-value</b>
<i>Intercept</i>	-6.8748	0.1407	<0.0001
<u>CDC Location Code: Adult Critical Care Units, Oncology Critical Care Units</u> <i>Oncology Medical Critical Care</i> <i>Oncology Surgical Critical Care</i> <i>Oncology Medical-Surgical Critical Care</i> <i>Prenatal Critical Care</i> <i>Respiratory Critical Care</i>	0.5009	0.1810	0.0057
CDC Location Code: <u>Surgical Cardiothoracic Critical Care</u>	0.9418	0.0862	<.0001



<b>Parameter</b>	<b>Parameter Estimate</b>	<b>Standard Error</b>	<b>P-value</b>
CDC Location Code: <i>Medical-Surgical Critical Care</i>	1.0161	0.0822	<.0001
CDC Location Code: <u>Adult Critical Care Units</u> <i>Burn Critical Care</i> <i>Medical Cardiac Critical Care</i> <i>Medical Critical Care</i> <i>Neurologic Critical Care</i> <i>Neurosurgical Critical Care</i> <i>Surgical Critical Care</i>	1.1140	0.0820	<.0001
CDC Location Code: <i>Adult Mixed Acuity Unit</i>	1.3225	0.1296	<.0001
CDC Location Code: <i>Trauma Critical Care</i>	1.4320	0.0882	<.0001
CDC Location Code: <u>Step-down Units</u> <i>Adult Step-down Unit</i> <i>Oncology Step-down Unit</i>	0.4096	0.1060	0.0001
CDC Location Code: <u>Wards, Solid Organ Transplant Specialty Care Area</u> <i>Antenatal Care Ward</i> <i>Behavioral Health/Psychology Ward</i> <i>Burn Ward</i> <i>Ear, Nose, Throat Ward</i> <i>Gastrointestinal Ward</i> <i>Genitourinary Ward</i> <i>Gerontology Ward</i> <i>Gynecology Ward</i> <i>Jail Unit</i> <i>Labor and Delivery Ward</i> <i>Labor, Delivery, Recovery, Postpartum Suite</i> <i>Medical Ward</i> <i>Medical-Surgical Ward</i> <i>Neurology Ward</i> <i>Neurosurgical Ward</i> <i>Oncology Leukemia Ward</i> <i>Oncology Lymphoma Ward</i> <i>Oncology Leukemia/Lymphoma Ward</i> <i>Oncology Solid Tumor Ward</i> <i>Oncology Hematopoietic Stem Cell Transplant Ward</i> <i>Oncology General Hematology-Oncology Ward</i> <i>Ophthalmology Ward</i> <i>Orthopedic Ward</i> <i>Orthopedic Trauma Ward</i>	REFERENT	-	-



<b>Parameter</b>	<b>Parameter Estimate</b>	<b>Standard Error</b>	<b>P-value</b>
<i>Plastic Surgery Ward</i>			
<i>Postpartum Ward</i>			
<i>Pulmonary Ward</i>			
<i>Rehabilitation Ward (within Hospital)</i>			
<i>School Infirmary</i>			
<i>Stroke (Acute) Ward</i>			
<i>Surgical Ward</i>			
<i>Telemetry Ward</i>			
<i>Vascular Surgery Ward</i>			
<i>Solid Organ Transplant Specialty Care Area</i>			
Facility bed size*: 85-129 beds	0.1591	0.0787	0.0433
Facility bed size*: 130-425 beds	0.2513	0.0679	0.0002
Facility bed size*: 426-526 beds	0.5123	0.0716	<.0001
Facility bed size*: ≥ 527 beds	0.6471	0.0706	<.0001
Facility bed size*: ≤ 84 beds	REFERENT	-	-
Medical School Affiliation*: Major	0.2905	0.0239	<.0001
Medical School Affiliation*: Graduate/Undergraduate	0.1395	0.0240	<0.0001
Medical School Affiliation*: Non-teaching	REFERENT	-	-
<b>Facility Type (based on NHSN enrollment): GEN-VA</b>			
<i>General Acute Care Hospital</i>	0.2154	0.0987	0.0290
<i>Veterans' Affairs Hospital</i>			
<b>Facility Type (based on NHSN enrollment): Other</b>			
<i>Military Hospital</i>	REFERENT	-	-
<i>Psychiatry Hospital</i>			
<i>Oncology Hospital</i>			
<i>Orthopedic Hospital</i>			
<i>Surgical Hospital</i>			
<i>Women's Hospital</i>			
<i>Women's and Children's Hospital</i>			

\* Facility bed size and medical school affiliation are taken from the [Annual ACH Survey](#)

**Table 2. Summary of Risk Factors in the Total VAE Model for Other Facility Types**

<b>Facility Type</b>	<b>Risk Factors</b>
Critical Access Hospitals (CAH)	Intercept-only model*
Inpatient Rehabilitation Facilities (IRF)	No SIR available^

\* None of the variables investigated were statistically significantly associated with Total VAE in CAHs. These facilities will have the predicted number of events calculated using the 2015 national pooled mean (i.e., intercept-only model).

^ Insufficient data were reported to NHSN. Therefore, SIRs are not available for Total VAE in IRFs.

## D. Infection-related Ventilator-Associated Complication (IVAC) Plus in Acute Care Hospitals (ACHs)

The number of predicted VAE events is calculated using a negative binomial regression model (see [page 8](#) above for more information). Separate VAE SIRs are available for “Total VAE” and “IVAC Plus”. The “IVAC Plus” SIR includes events identified as IVAC and Possible Ventilator-Associated Pneumonia (PVAP). Refer to Table 1 below for a list of location types included in the Acute Care Hospital IVAC Plus SIR; if an inpatient location is not included in the model table below, then data from that location type will be excluded from the SIR due to insufficient 2015 baseline data. In addition, data from Governmental and Non-governmental Public Health Emergency (PHE) Facilities (facType as HOSP-PHE/G or HOSP-PHE/NG) will also be excluded from the SIR. In cases when the number of predicted events is less than 1.0, the SIR will not be calculated in NHSN.

The number of predicted “IVAC Plus” events calculated under the 2015 baseline is risk adjusted based on the following variables found to be statistically significant predictors of “IVAC Plus” incidence:

**Table 1. IVAC Plus in Acute Care Hospitals (ACHs)**

Parameter	Parameter Estimate	Standard Error	P-value
<i>Intercept</i>	-7.4627	0.0925	< 0.0001
<u>CDC Location Code: Adult Critical Care Units</u> <i>Burn Critical Care</i> <i>Neurologic Critical Care</i> <i>Neurosurgical Critical Care</i> <i>Surgical Critical Care</i>	1.1747	0.0922	<.0001
<u>CDC Location Code: Adult Critical Care Units, Oncology Critical Care Units</u> <i>Medical Cardiac Critical Care</i> <i>Medical Critical Care</i> <i>Medical-Surgical Critical Care</i> <i>Oncology Medical Critical Care</i> <i>Oncology Surgical Critical Care</i> <i>Oncology Medical-Surgical Critical Care</i> <i>Prenatal Critical Care</i> <i>Respiratory Critical Care</i> <i>Surgical Cardiothoracic Critical Care</i>	0.9092	0.0889	<.0001
<u>CDC Location Code: Trauma Critical Care</u>	1.5429	0.0984	<.0001
<u>CDC Location Code: Adult Mixed Acuity Unit</u>	1.2291	0.1779	<.0001
<u>CDC Location Code: Wards, Specialty Care Areas, Step-down Units</u> <i>Antenatal Care Ward</i> <i>Behavioral Health/Psychology Ward</i> <i>Burn Ward</i> <i>Ear, Nose, Throat Ward</i> <i>Gastrointestinal Ward</i> <i>Genitourinary Ward</i>	REFERENT	-	-

Parameter	Parameter Estimate	Standard Error	P-value
<i>Gerontology Ward</i>			
<i>Gynecology Ward</i>			
<i>Jail Unit</i>			
<i>Labor and Delivery Ward</i>			
<i>Labor, Delivery, Recovery, Postpartum Suite</i>			
<i>Medical Ward</i>			
<i>Medical-Surgical Ward</i>			
<i>Neurology Ward</i>			
<i>Neurosurgical Ward</i>			
<i>Oncology Leukemia Ward</i>			
<i>Oncology Lymphoma Ward</i>			
<i>Oncology Leukemia/Lymphoma Ward</i>			
<i>Oncology Solid Tumor Ward</i>			
<i>Oncology Hematopoietic Stem Cell Transplant Ward</i>			
<i>Oncology General Hematology-Oncology Ward</i>			
<i>Ophthalmology Ward</i>			
<i>Orthopedic Ward</i>			
<i>Orthopedic Trauma Ward</i>			
<i>Plastic Surgery Ward</i>			
<i>Postpartum Ward</i>			
<i>Pulmonary Ward</i>			
<i>Rehabilitation Ward (within Hospital)</i>			
<i>School Infirmary</i>			
<i>Stroke (Acute) Ward</i>			
<i>Surgical Ward</i>			
<i>Telemetry Ward</i>			
<i>Vascular Surgery Ward</i>			
<i>Solid Organ Transplant Specialty Care Area</i>			
<i>Adult Step-down Unit</i>			
<i>Oncology Step-down Unit</i>			
Facility bed size*: 290-425 beds	0.1540	0.0370	<.0001
Facility bed size*: 426-526 beds	0.4058	0.0433	<.0001
Facility bed size*: >=527 beds	0.5079	0.0385	<.0001
Facility bed size*: ≤ 289 beds	REFERENT	-	-
Medical School Affiliation*: Major	0.3157	0.0354	<.0001
Medical School Affiliation*: Graduate/Undergraduate	0.1630	0.0362	<.0001
Medical School Affiliation*: Non-teaching	REFERENT	-	-

\* Facility bed size and medical school affiliation are taken from the [Annual ACH Survey](#)

**Table 2. Summary of Risk Factors in the IVAC Plus Model for Other Facility Types**

Facility Type	Risk Factors
Critical Access Hospitals (CAH)	No SIR Available <sup>^</sup>
Inpatient Rehabilitation Facilities (IRF)	No SIR Available <sup>^</sup>

<sup>^</sup> Insufficient data were reported to NHSN. Therefore, SIRs are not available for 'IVAC Plus' in CAHs or IRFs.



## Risk Adjustment Factors Included in the SIR Calculation: 2015 Baseline

### SSI – Surgical Site Infections

The number of predicted SSI events is calculated using a logistic regression model (see [page 5](#) above for more information). The SSI SIR is calculated for facilities who enroll in NHSN as acute care hospitals or critical access hospitals. Data from Governmental and Non-governmental Public Health Emergency (PHE) Facilities (facType as HOSP-PHE/G or HOSP-PHE/NG) are excluded from the SIR. Under the 2015 SIR baseline, procedures and associated SSI events occurring in adult and pediatric patients are modeled separately. There are three SSI SIR models available for inpatient adult procedures (and associated SSIs) and two models available for inpatient pediatric procedures (and associated SSIs). Please see *Table 1* below for a summary of the SSI SIR models. Under the 2015 SIR baseline, procedures, regardless of closure methods, are included in the SIR calculation, as long as the inclusion criteria listed below are met and none of the exclusion criteria apply.

**Table 1. Summary of SSI Models**

SSI SIR Model	Inclusion Criteria	Patient Population
All SSI SIR Model	<ul style="list-style-type: none"> <li>Includes <u>only</u> inpatient procedures</li> <li>Includes Superficial, Deep &amp; Organ/Space SSIs</li> <li>Superficial &amp; Deep Incisional SSIs limited to primary incisional SSIs only</li> <li>Includes SSIs identified on admission, readmission &amp; via post-discharge surveillance</li> </ul>	<ul style="list-style-type: none"> <li>Procedures in adult patients</li> <li>Procedures in pediatric patients</li> </ul>
Complex Admission/Readmission (A/R) SSI Model	<ul style="list-style-type: none"> <li>Includes <u>only</u> inpatient procedures</li> <li>Includes <u>only</u> Deep Incisional Primary SSIs &amp; Organ/Space SSIs</li> <li>Includes <u>only</u> SSIs identified on Admission/Readmission to facility where procedure was performed</li> <li>Used for the annual CDC publication of national benchmarks</li> </ul>	<ul style="list-style-type: none"> <li>Procedures in adult patients</li> <li>Procedures in pediatric patients</li> </ul>
Complex 30-Day SSI model (used for CMS IPPS)	<ul style="list-style-type: none"> <li>Includes <u>only</u> in-plan, inpatient COLO and HYST procedures in adult patients (i.e., ≥ 18 years of age)</li> <li>Includes only Deep Incisional Primary SSIs and Organ/Space SSIs with an event date within 30 days of the procedure</li> <li>Includes SSIs regardless of detection method</li> <li>Used only for CMS IPPS reporting and for public reporting on the CMS Care Compare website</li> </ul>	<ul style="list-style-type: none"> <li>Procedures in adult patients</li> </ul>

### Exclusion Criteria

In addition to the above inclusion criteria, there is also a list of exclusion criteria that applies to all the SSI SIR models. This list is often referred to as the universal exclusion criteria. The list of exclusion criteria applies to both procedures and the associated SSI events. Often the reason for excluding procedures and SSI events from the SIR calculation is due to potential data quality issues. It is important that facilities review their data for quality assurance and to determine the reason for exclusion from the SIR calculation.

*Note: When a procedure is excluded from the denominator, the associated SSI event is excluded from the numerator.*

**Table 2. Universal Procedure/SSI Event Exclusions**

<u>General Exclusions</u>
Gender= 'Other'
Outpatient procedures and resulting SSIs
Present at time of surgery (PATOS) is 'Yes'
SSIs that are reported as superficial incisional secondary (SIS) or deep incisional secondary (DIS)
<u>Exclusions due to potential data quality issues or outliers</u>
Age at the time of procedure is greater than 109 years
Closure technique is missing
ASA score is missing
Gender is missing
Adult patients ≥ 18 years: if BMI is less than 12 or greater than 60*
Pediatric patients < 18 years: if BMI less than 10.49 or greater than 65.79**
Procedure duration less than 5 minutes
Procedure duration is greater than IQR5 ( <i>please see <a href="#">Table 4</a> in the SSI Section for more information</i> )
<u>Facility-level Exclusions</u>
Data from ambulatory surgery centers (ASCs) and long-term acute care hospitals (LTACHs)
Medical affiliation is missing or medical affiliation is 'Y' and medical type is missing ( <i>from <a href="#">Annual Facility Survey</a></i> )
Number of beds is missing ( <i>from <a href="#">Annual Facility Survey</a></i> )

\*This BMI exclusion applies to all procedures on adult patients in all 3 SSI models (All SSI, Complex A/R, Complex 30-Day).

\*\*This BMI exclusion applies to all procedures on pediatric patients, in both applicable SSI models (All SSI and Complex A/R). CDC Growth Charts are used to assess BMI in pediatric patients, calculated using height, weight, age, and gender.

**Additional clarification on the BMI exclusion rule for pediatric procedures:** Although there are BMI thresholds for procedures performed on pediatric patients (10.49-65.79), there is an additional level of consideration made for the biological plausibility of a given BMI using the patient's age and gender. After applying the BMI outlier exclusion rule, we review the BMIs for the remaining pediatric procedures to determine if they are biologically plausible based on the patient's age and gender. So essentially, we take age and gender into consideration along with the calculated BMI. Only procedures in which the patient's BMI meets the inclusion rule (10.49-65.79), **and** in which the patient's BMI is biologically plausible based on age and gender, are included in the SIR. The determination of biologically plausible BMIs is made using the macro available at this site: <https://www.cdc.gov/nccdphp/dnpao/growthcharts/resources/sas.htm>.

## Predictive Risk Factors by SSI Models

The number of predicted events calculated under the 2015 baseline for SSI is risk adjusted based on the following variables found to be statistically significant predictors of SSIs. The following tables (*3a-3f*) list the factors included in each procedure-specific model, grouped by the three SSI models outlined above. In some procedure-specific models, the interaction of age and gender is considered as a single factor. It is listed as age-gender interaction. In cases when the number of predicted events is less than 1.0, the SIR will not be calculated in NHSN.

*Note: Parameter estimates are shown for colon (COLO) and abdominal hysterectomy (HYST) procedures under the Complex 30-Day Model used for the CMS Hospital Inpatient Prospective Payment System (IPPS). Full model details for all procedures under the All-SSI Model and the Complex A/R Model are available here:*

<https://www.cdc.gov/nhsn/ps-analysis-resources/sirguide-ssimodels-508.xlsx>

**Table 3a. Colon Procedures, Complex 30-Day Model**

<u>Parameter</u>	<u>Parameter Estimate</u>	<u>Standard Error</u>	<u>P-value</u>
<i>Intercept</i>	-3.6601	0.0678	<0.0001
Diabetes: Yes	0.0821	0.0303	0.0066
Diabetes: No	REFERENT	-	-
ASA score: 1, 2, 3/4/5	0.3028	0.0237	<0.0001
Gender: Male	0.1036	0.0225	<0.0001
Gender: Female	REFERENT	-	-
Age (Patient's age/10)	-0.1396	0.0075	<0.0001
BMI: ≥ 30	0.1259	0.0234	<0.0001
BMI: < 30	REFERENT	-	-
Closure technique: Other (non-Primary)	0.2383	0.0494	<0.0001
Closure technique: Primary	REFERENT	-	-
Oncology Hospital: Yes	0.5437	0.0937	<0.0001
Oncology Hospital: No	REFERENT	-	-

**Table 3b. Abdominal Hysterectomy Procedures, Complex 30-Day Model**

<u>Parameter</u>	<u>Parameter Estimate</u>	<u>Standard Error</u>	<u>P-value</u>
<i>Intercept</i>	-5.1801	0.1057	< 0.0001
Diabetes: Yes	0.3247	0.0605	<0.0001
Diabetes: No	REFERENT	-	-
ASA score: 1, 2, 3, 4/5	0.4414	0.0350	<0.0001
BMI: ≥ 30	0.1106	0.0423	0.0090
BMI: < 30	REFERENT	-	-
Age (Patient's age/10)	-0.1501	0.0180	<0.0001
Oncology Hospital: Yes	0.5474	0.1578	0.0005
Oncology Hospital: No	REFERENT	-	-

**Tables 3c to 3f have been moved from this section and are now available to view (with details) in the addendum of the SIR guide, here: <https://www.cdc.gov/nhsn/ps-analysis-resources/sirguide-ssimodels-508.xlsx>**

### Procedure Duration Outliers

The **IQR5, also called the procedure duration cutoff point**, is used as an indicator of an extreme outlier for procedure durations when calculating the SSI SIRs. The IQR5 is calculated as five times the interquartile range (Q1-Q3) above the 75th percentile. For example, if the interquartile range is 30 minutes, and the 75th percentile is 100 minutes, the IQR5 would be calculated as:  $100 + (30 \times 5) = 250$  minutes. Procedures with a duration greater than the IQR5 were excluded from the baseline data and will be excluded from all SSI SIR calculations for your facility.

**Table 4. IQR5 Values, in Minutes, for NHSN Operative Procedures, Adult and Pediatric Patients**

NHSN Operative Procedure	IQR5 (in minutes)	IQR5 (in hours and minutes)	
	Minutes	Hours	Minutes
AAA	1116	18	36
AMP	300	5	0
APPY	210	3	30
AVSD	471.5	7	51.5
BILI	1295	21	35
BRST	777	12	57
CARD	1001	16	41
CBGB	847	14	7
CBGC	847	14	7
CEA	376	6	16
CHOL	346	5	46
COLO	697	11	37
CRAN	904	15	4
CSEC	170	2	50
FUSN	874	14	34
FX	532	8	52
GAST	489	8	9
HER	521	8	41
HPRO	349	5	49
HTP	1355	22	35
HYST	547	9	7
KPRO	316	5	16
KTP	670	11	10
LAM	687	11	27
LTP	1243	20	43
NECK	1796	29	56
NEPH	774	12	54
OVRV	594	9	56
PACE	311	5	11
PRST	737	12	17
PVBY	850	14	10
REC	1136	18	56
RFUSN	1129	18	49
SB	856	14	16
SPLE	1073	17	53
THOR	721	12	1
THYR	506	8	26
VHYS	506	8	26
VSHN	378	6	18
XLAP	724	12	4

## Risk Adjustment Factors Included in the SIR Calculation: 2015 Baseline

### ***Methicillin-resistant Staphylococcus aureus (MRSA) Bacteremia Laboratory-Identified Events***

The number of predicted MRSA bacteremia LabID events is calculated using a negative binomial regression model (see [page 8](#) above for more information). For most settings, the MRSA bacteremia SIR is only calculated on the facility-wide inpatient, or FacWideIN, level, and cannot be calculated for any individual location (*note: CMS-designated inpatient rehabilitation units within a hospital will receive a separate SIR*). Data from Governmental and Non-governmental Public Health Emergency (PHE) Facilities (facType as HOSP-PHE/G or HOSP-PHE/NG) are excluded from the SIR. In cases when the number of predicted events is less than 1.0, the SIR will not be calculated in NHSN. The SIRs for MRSA bacteremia include only healthcare facility-onset (HO), non-duplicate MRSA blood LabID events in the numerator. Information on which events are counted in the numerator of the MRSA bacteremia SIR can be found here: [http://www.cdc.gov/nhsn/pdfs/ps-analysis-resources/mrsacdi\\_tips.pdf](http://www.cdc.gov/nhsn/pdfs/ps-analysis-resources/mrsacdi_tips.pdf).

The number of predicted events calculated under the 2015 baseline for MRSA bacteremia is risk adjusted based on the following variables found to be statistically significant predictors of MRSA bacteremia incidence:

*Notes for Acute Care Hospitals: MRSA LabID SIRs for acute care hospitals can only be calculated at the quarter-level or higher. This is because two of the risk factors involving the community-onset prevalence rate require that all community-onset data have been entered for an entire quarter. The quarter's community-onset prevalence rates, both inpatient and outpatient, are used to calculate the number of predicted events for the SIR.*

**Table 1. MRSA Bacteremia in Acute Care Hospitals**

<b>Parameter</b>	<b>Parameter Estimate</b>	<b>Standard Error</b>	<b>P-value</b>
<i>Intercept</i>	-11.3759	0.1167	<0.0001
Inpatient community-onset prevalence rate*: > 0.037 per 100 admissions	0.3650	0.0286	<0.0001
Inpatient community-onset prevalence rate*: ≤ 0.037 per 100 admissions	REFERENT	-	-
Average length of stay**: ≥ 5.1 days	0.2787	0.0343	<0.0001
Average length of stay**: 4.3-5.0 days	0.0955	0.0341	0.0050
Average length of stay**: 0-4.2 days	REFERENT	-	-
Medical school affiliation‡: Major	0.2585	0.0334	<0.0001
Medical school affiliation‡: Graduate/undergraduate	0.1166	0.0345	0.0007
Medical school affiliation‡: Non-teaching	REFERENT	-	-
Facility type: Oncology Hospital ( <i>HOSP-ONC</i> )	1.1894	0.2085	<0.0001
Facility type: General Acute Care Hospital ( <i>HOSP-GEN</i> )	0.4355	0.0897	<0.0001
Facility type: Other Specialty Hospital	REFERENT	-	-
Number of ICU beds‡: ≥ 45	0.5650	0.0898	<0.0001
Number of ICU beds‡: 21-44	0.4599	0.0899	<0.0001
Number of ICU beds‡: 11-20	0.3394	0.0922	0.0002
Number of ICU beds‡: 7-10	0.4720	0.0993	<0.0001
Number of ICU beds‡: 0-6	REFERENT	-	-

**Table 1, continued. MRSA Bacteremia in Acute Care Hospitals**

<u>Parameter</u>	<u>Parameter Estimate</u>	<u>Standard Error</u>	<u>P-value</u>
Outpatient community-onset prevalence rate ED/24-hour Observation Unit <sup>^</sup> : > 0.032 per 100 encounters	0.3476	0.0336	<0.0001
Outpatient community-onset prevalence rate ED/24-hour Observation Unit <sup>^</sup> : > 0 and ≤ 0.032 per 100 encounters	0.1048	0.0330	0.0015
Outpatient community-onset prevalence rate ED/24-hour Observation Unit <sup>^</sup> : 0 per 100 encounters, or no applicable locations	REFERENT	-	-

\* Inpatient community-onset prevalence is calculated as the # of inpatient community-onset MRSA blood events, divided by total admissions x 100. (i.e., MRSA\_admPrevBldCount / numadms \* 100).

\*\* Average length of stay is taken from the [Annual Hospital Survey](#). It is calculated as: total # of annual patient days / total # of annual admissions.

‡ Medical school affiliation and number of ICU beds are taken from the [Annual Hospital Survey](#).

<sup>^</sup> Emergency department (ED)/24-hour observation unit prevalence rate combines MRSA bacteremia data from all EDs and/or 24-hour observation units into a single, de-duplicated prevalence rate. This rate is calculated as the # of unique community-onset MRSA blood events that occurred in an ED or 24-hour observation unit / total encounters \* 100. (i.e., MRSA\_EDOBSPrevCount / numTotencounters \* 100). *NOTE: If you do not have an ED or 24-hour observation location that meets the [NHSN location definition](#) and thus are not reporting MRSA bacteremia data from these locations, the number of predicted events will be risk adjusted using the referent level of this variable.*

**Table 2. MRSA Bacteremia in Critical Access Hospitals (CAHs)**

<u>Parameter</u>	<u>Parameter Estimate</u>	<u>Standard Error</u>	<u>P-value</u>
<i>Intercept*</i>	-10.7795	0.2025	<0.0001

\* MRSA LabID SIRs for CAHs can be calculated for any aggregate of time (month, quarter, half-year, or year). None of the variables investigated were statistically significantly associated with healthcare facility-onset MRSA bacteremia in CAHs. The predicted number of events for CAHs will be calculated using the 2015 national CAH MRSA bacteremia pooled mean (i.e., intercept-only model).

**Table 3. MRSA Bacteremia in Long-Term Acute Care Hospitals (LTACHs)**

<u>Parameter</u>	<u>Parameter Estimate</u>	<u>Standard Error</u>	<u>P-value</u>
<i>Intercept*</i>	-9.3095	0.0936	<0.0001
Percent of admissions on ventilator**	0.0160	0.0027	<0.0001

\* MRSA LabID SIRs for LTACHs can be calculated for any aggregate of time (month, quarter, half-year, or year).

\*\* Percent of annual admissions on a ventilator is taken from the [Annual LTACH Survey](#). It is calculated as: # admissions on a ventilator / total # of annual admissions x 100 (i.e., numAdmvent / numAdmitsSurv \* 100).

**Table 4. MRSA Bacteremia in Inpatient Rehabilitation Hospitals (IRFs): Free-standing Rehabilitation Hospitals and CMS-Certified IRF Units Within a Hospital**

<u>Parameter</u>	<u>Parameter Estimate</u>	<u>Standard Error</u>	<u>P-value</u>
<i>Intercept*</i>	-10.8703	0.0890	<0.0001

\* MRSA LabID SIRs for IRFs can be calculated for any aggregate of time (month, quarter, half-year, or year). None of the variables investigated were statistically significantly associated with healthcare facility-onset MRSA bacteremia in IRFs. Free-standing IRFs and CMS-certified IRF units within a hospital will have the predicted number of events calculated using the 2015 national IRF MRSA bacteremia pooled mean (i.e., intercept-only model).

## Risk Adjustment Factors Included in the SIR Calculation: 2015 Baseline

### *Clostridioides difficile* (CDI) Laboratory-Identified Events

The number of predicted CDI LabID events is calculated using a negative binomial regression model (see [page 8](#) above for more information). For most settings, the CDI SIR is only calculated on the facility-wide inpatient, or FacWideIN, level, and cannot be calculated for any individual location (*note: CMS-designated inpatient rehabilitation units within a hospital will receive a separate SIR*). Data from Governmental and Non-governmental Public Health Emergency (PHE) Facilities (facType as HOSP-PHE/G or HOSP-PHE/NG) are excluded from the SIR. In cases when the number of predicted events is less than 1.0, the SIR will not be calculated in NHSN. The FacWideIN SIRs for CDI include only incident, healthcare facility-onset (HO), non-duplicate *C. difficile* LabID events in the numerator. Information on which events are counted in the numerator of the FacWideIN and IRF Unit CDI SIR can be found here: [http://www.cdc.gov/nhsn/pdfs/ps-analysis-resources/mrsacdi\\_tips.pdf](http://www.cdc.gov/nhsn/pdfs/ps-analysis-resources/mrsacdi_tips.pdf).

For all facility types, the CDI LabID SIR can only be calculated at the quarter-level or higher. Monthly SIRs cannot be calculated due to certain risk factors used in each of the models that require complete data entry for a quarter (e.g., CDI test type is reported on the FacWideIN and IRF unit's MDRO denominator form on the 3rd month of each quarter).

The number of predicted events calculated under the 2015 baseline for CDI is risk adjusted based on the following variables found to be statistically significant predictors of CDI incidence:

**Table 1. CDI in Acute Care Hospitals**

Parameter	Parameter Estimate	Standard Error	P-value
Intercept	-8.9463	0.0523	<0.0001
Inpatient community-onset prevalence rate*	0.7339	0.0181	<0.0001
CDI test type <sup>†</sup> : EIA	-0.1579	0.0246	<0.0001
CDI test type <sup>†</sup> : NAAT	0.1307	0.0219	<0.0001
CDI test type <sup>†</sup> : OTHER	REFERENT	-	-
Medical school affiliation <sup>‡</sup> : Major, graduate, or undergraduate	0.0331	0.0111	0.0028
Medical school affiliation <sup>‡</sup> : Non-teaching	REFERENT	-	-
Number of ICU beds <sup>‡</sup> : ≥ 43	0.7465	0.0412	<0.0001
Number of ICU beds <sup>‡</sup> : 20- 42	0.7145	0.0395	<0.0001
Number of ICU beds <sup>‡</sup> : 10-19	0.6261	0.0396	<0.0001
Number of ICU beds <sup>‡</sup> : 5-9	0.4394	0.0420	<0.0001
Number of ICU beds <sup>‡</sup> : 0-4	REFERENT	-	-
Facility type: Oncology Hospital ( <i>HOSP-ONC</i> )	1.2420	0.0765	<0.0001
Facility type: General Acute Care Hospital ( <i>HOSP-GEN</i> )	0.3740	0.0342	<0.0001
Facility type: Other Specialty Hospital	REFERENT	-	-
Facility bed size <sup>‡</sup>	0.0003	0.0000	<0.0001
Reporting from ED or 24-hour observation unit <sup>^</sup> : YES	0.1119	0.0179	<0.0001
Reporting from ED or 24-hour observation unit <sup>^</sup> : NO	REFERENT	-	-

\* Inpatient community-onset (CO) prevalence is calculated as the # of inpatient CO CDI events, divided by total admissions x 100 (i.e.,  $\text{cdif\_admPrevCOcount} / \text{numCdifadms} * 100$ ). The prevalence rate for an entire quarter is used in the risk

**Table 1 Footnotes continued:**

adjustment. An SIR cannot be calculated for any quarter that has an outlier inpatient CO prevalence rate, defined as greater than 2.6 CO events per 100 admissions.

+ CDI test type is reported on the FacWideIN MDRO denominator form on the 3<sup>rd</sup> month of each quarter.

-Starting in 2018 Q1, CDI test type is categorized as:

**Nucleic acid amplification test (NAAT):** This includes NAAT, GDH + NAAT, and GDH + EIA + NAAT.

**Enzyme immunoassay (EIA) for toxin:** This includes EIA for toxin, GDH antigen + EIA for toxin, and **NAAT + EIA**

**Other:** This includes all other CDI test types, including the selection of “Other” and associated free-text entry.

-Prior to 2018 Q1, CDI test type was categorized as: (refer to 2018 NHSN protocol changes for details)

**Nucleic acid amplification test (NAAT):** This includes NAAT, GDH + NAAT, GDH + EIA + NAAT, and NAAT + EIA.

**Enzyme immunoassay (EIA) for toxin:** This includes EIA for toxin, and GDH antigen + EIA for toxin.

**Other:** This includes all other CDI test types, including the selection of “Other” and associated free-text entry.

‡ Medical school affiliation, number of ICU beds, and facility bed size are taken from the [Annual Hospital Survey](#).

^ If your facility has a designated Emergency Department (ED) or 24-hour observation location meeting the standard NHSN definitions, these locations should be mapped, included in your facility’s monthly reporting plan for LabID events, and have appropriate outpatient LabID data reported to NHSN. If you do not have an ED or 24-hour observation location and thus are not reporting CDI data from these locations, your hospital’s # predicted events will be risk adjusted using the referent value for this variable.

**Table 2. CDI in Critical Access Hospitals (CAHs)**

Parameter	Parameter Estimate	Standard Error	P-value
Intercept	-8.4180	0.0879	<0.0001
Inpatient community-onset prevalence rate*: > 0	0.7207	0.1108	<0.0001
Inpatient community-onset prevalence rate*: 0	REFERENT	-	-

\* Inpatient community-onset (CO) prevalence rate is calculated as: # of inpatient CO CDI events, divided by total admissions x 100. (i.e.,  $\text{cdif\_admPrevCOCount} / \text{numCdifadms} * 100$ ). The prevalence rate for an entire quarter is used in the risk adjustment.

**Table 3. CDI in Long-Term Acute Care Hospitals (LTACHs)**

Parameter	Parameter Estimate	Standard Error	P-value
Intercept	-7.3345	0.0507	<0.0001
Inpatient community-onset prevalence rate*: > 0	0.3683	0.0493	<0.0001
Inpatient community-onset prevalence rate*: 0	REFERENT	-	-
Percent of admissions on a ventilator‡: ≥ 27.1%	0.3116	0.0478	<0.0001
Percent of admissions on a ventilator‡: ≥ 18% to < 27.1%	0.1463	0.0590	0.0131
Percent of admissions on a ventilator‡: < 18%	REFERENT	-	-
CDI test type^: NAAT or OTHER	0.1607	0.0444	0.0003
CDI test type^: EIA	REFERENT	-	-
Percent of single occupancy rooms+: ≤ 77%	0.0963	0.0425	0.0235
Percent of single occupancy rooms+: >77%	REFERENT	-	-

\* Inpatient community-onset prevalence is calculated as the # of inpatient community-onset CDI events, divided by total admissions \* 100. (i.e.,  $\text{cdif\_admPrevCOCount} / \text{numCdifadms} * 100$ ). The prevalence rate for an entire quarter is used in the risk adjustment.

‡ Percent of annual admissions on a ventilator is taken from the [Annual LTACH Survey](#). It is calculated as: # admissions on a ventilator / total # annual admissions x 100. (i.e.,  $\text{numAdmVent} / \text{numAdmitsSurv} * 100$ ).



**Table 3 Footnotes continued:**

^ CDI test type is reported on the FacWideIN MDRO denominator form on the 3<sup>rd</sup> month of each quarter.

-Starting in 2018 Q1, CDI test type is categorized as:

Nucleic acid amplification test (NAAT) or Other: This includes NAAT, GDH + NAAT, GDH + EIA + NAAT, and all other (non-EIA) CDI test types, including the selection of “Other” and associated free-text entry.

Enzyme immunoassay (EIA) for toxin: This includes EIA for toxin, GDH antigen + EIA for toxin, and **NAAT + EIA**.

-Prior to 2018 Q1, CDI test type was categorized as: (refer to 2018 NHSN protocol changes for details)

Nucleic acid amplification test (NAAT) or Other: This includes NAAT, GDH + NAAT, GDH + EIA + NAAT, NAAT + EIA, and all other (non-EIA) CDI test types, including the selection of “Other” and associated free-text entry.

Enzyme immunoassay (EIA) for toxin: This includes EIA for toxin, and GDH antigen + EIA for toxin.

+ Percent of beds located in single occupancy rooms is taken from the [Annual LTACH Survey](#). It is calculated as: # of single occupancy rooms / total number of beds x 100. (i.e., numSingOccRm / numbeds \* 100).

**Table 4. CDI in Inpatient Rehabilitation Facilities (IRFs): Free-standing Rehabilitation Hospitals and CMS-Certified IRF Units Within a Hospital**

Parameter	Parameter Estimate	Standard Error	P-value
Intercept	-8.4475	0.0689	<0.0001
CDI test type <sup>^</sup> : NAAT	0.2921	0.0534	<0.0001
CDI test type <sup>^</sup> : OTHER	0.2163	0.0747	0.0038
CDI test type <sup>^</sup> : EIA	REFERENT	-	-
CMS-certified IRF Unit within a hospital	0.2188	0.0495	<0.0001
Free-standing HOSP-REHAB with reported community-onset CDI events	0.4168	0.0803	<0.0001
Free-standing HOSP-REHAB with zero reported community-onset CDI events	REFERENT	-	-
Percent of admissions with orthopedic conditions*: ≤ 23.9%	0.2015	0.0427	<0.0001
Percent of admissions with orthopedic conditions*: > 23.9%	REFERENT	-	-
Percent of admissions with traumatic and non-traumatic spinal cord dysfunction*: > 5.2%	0.1657	0.0437	0.0002
Percent of admissions with traumatic and non-traumatic spinal cord dysfunction*: ≤ 5.2%	REFERENT	-	-
Percent of admissions with stroke*: ≤ 23.8%	0.1965	0.0444	<0.0001
Percent of admissions with stroke*: > 23.8%	REFERENT	-	-

^ CDI test type is reported on the FacWideIN or IRF Unit’s MDRO denominator form on the 3<sup>rd</sup> month of each quarter.

-Starting with 2018 Q1, CDI test type is categorized as:

Nucleic acid amplification test (NAAT): This includes NAAT, GDH + NAAT, and GDH + EIA + NAAT.

Enzyme immunoassay (EIA) for toxin: This includes EIA for toxin, GDH antigen + EIA for toxin, and **NAAT + EIA**.

Other: This includes all other CDI test types, including the selection of “Other” and associated free-text entry.

-Prior to 2018 Q1, CDI test type was categorized as: (refer to 2018 NHSN protocol changes for details)

Nucleic acid amplification test (NAAT): This includes NAAT, GDH + NAAT, GDH + EIA + NAAT, and NAAT + EIA.

Enzyme immunoassay (EIA) for toxin: This includes EIA for toxin, and GDH antigen + EIA for toxin.

Other: This includes all other CDI test types, including the selection of “Other” and associated free-text entry.

^ Percent of annual admissions with primary diagnoses are taken from the [Annual IRF Survey](#), and calculated as the # of admissions with the primary diagnosis / total # of annual admissions x 100.

## Using an Intercept-Only Model to Calculate the Number of Predicted Events Example: MRSA Bacteremia LabID Event

Several regression models from the 2015 national baseline are “intercept-only models”. For example, none of the investigated variables were found to have a significant association with the incidence of healthcare facility-onset (HO) MRSA bacteremia in critical access hospitals or inpatient rehabilitation facilities. Therefore, the number of predicted events is calculated by applying the following intercept-only formula:

$$\text{Number of Predicted Events} = \exp(\text{Intercept Value}) \times \text{Patient Days}$$

Let’s say a critical access hospital had 1,400 total patient days during a select time period. The number of predicted events would be calculated as:

$$\text{Number of Predicted Events} = \exp(-10.7795) \times 1400$$

$$\text{Number of Predicted Events} = 0.029$$

Because the number of predicted events is less than 1.0, an SIR will not be calculated for this facility and time period in NHSN.

## Additional Resources

### ➤ **Information about Transitioning to 2015 SIR Baselines:**

NHSN Rebaseline webpage: <https://www.cdc.gov/nhsn/2015rebaseline/>

Introduction to the NHSN Re-baseline – March 2017

[https://www.cdc.gov/nhsn/pdfs/training/2017/Dudeck\\_March21.pdf](https://www.cdc.gov/nhsn/pdfs/training/2017/Dudeck_March21.pdf)

The NHSN Re-Baseline: In Depth – March 2017

[https://www.cdc.gov/nhsn/pdfs/training/2017/Dudeck\\_March22.pdf](https://www.cdc.gov/nhsn/pdfs/training/2017/Dudeck_March22.pdf)

### ➤ **Original SIR Baselines for Acute Care Hospitals:**

CLABSI (original baseline = 2006-2008): <https://www.cdc.gov/nhsn/PDFs/dataStat/2009NHSNReport.pdf>

CAUTI (original baseline = 2009): [https://www.cdc.gov/nhsn/PDFs/NHSNReport\\_DataSummaryfor2009.pdf](https://www.cdc.gov/nhsn/PDFs/NHSNReport_DataSummaryfor2009.pdf)

SSI (original baseline = 2006-2008): [https://www.cdc.gov/nhsn/PDFs/pscManual/SSI\\_ModelPaper.pdf](https://www.cdc.gov/nhsn/PDFs/pscManual/SSI_ModelPaper.pdf)

MRSA bacteremia and CDI LabID event (original baseline= 2010-2011):

<https://www.cdc.gov/nhsn/pdfs/mrsa-cdi/riskadjustment-mrsa-cdi.pdf>

December 2010 Special Edition NHSN Newsletter - Introduction to SIR (original baseline):

[https://www.cdc.gov/nhsn/pdfs/newsletters/nhsn\\_nl\\_oct\\_2010se\\_final.pdf](https://www.cdc.gov/nhsn/pdfs/newsletters/nhsn_nl_oct_2010se_final.pdf)

### ➤ **Original SIR Baselines for Long-term Acute Care Hospitals (LTACHs) and Inpatient Rehabilitation Facilities (IRFs):**

CLABSI/CAUTI in LTACHs, and CAUTI in IRFs (original baseline = 2013):

<https://www.cdc.gov/nhsn/xls/reportdatatables/nhsn-2013-report.xlsx>

### ➤ **NHSN Analysis Trainings & Other Resources:**

*A comprehensive guide to NHSN's SSI SIR, including risk factors used in the SIR calculations under the 2015 baseline:* <https://www.cdc.gov/nhsn/ps-analysis-resources/sirguide-ssimodels-508.xlsx>

Analysis Resources, Trainings, and NHSN Data Dictionary:

<https://www.cdc.gov/nhsn/ps-analysis-resources/index.html>

Targeted Assessment for Prevention (TAP) General Information: <https://www.cdc.gov/hai/prevent/tap.html>

Quick Reference Guides: How to run and interpret NHSN reports (including SIR and TAP reports):  
<https://www.cdc.gov/nhsn/ps-analysis-resources/reference-guides.html>

Troubleshooting CLABSI and CAUTI SIRs:  
[https://www.cdc.gov/nhsn/pdfs/ps-analysis-resources/clabscauti\\_sirtroubleshooting.pdf](https://www.cdc.gov/nhsn/pdfs/ps-analysis-resources/clabscauti_sirtroubleshooting.pdf)

Troubleshooting SSI SIRs: [https://www.cdc.gov/nhsn/pdfs/ps-analysis-resources/ssi-sir\\_tips.pdf](https://www.cdc.gov/nhsn/pdfs/ps-analysis-resources/ssi-sir_tips.pdf)

Troubleshooting MRSA and CDI LabID Event SIRs:  
[https://www.cdc.gov/nhsn/pdfs/ps-analysis-resources/mrsacdi\\_tips.pdf](https://www.cdc.gov/nhsn/pdfs/ps-analysis-resources/mrsacdi_tips.pdf)

Information related to SIRs used for the Centers for Medicare and Medicaid Services (CMS) Quality Reporting Programs: <https://www.cdc.gov/nhsn/cms/index.html>

NHSN Annual Hospital Survey: [https://www.cdc.gov/nhsn/forms/57.103\\_pshospsurv\\_blank.pdf](https://www.cdc.gov/nhsn/forms/57.103_pshospsurv_blank.pdf)  
• Instructions for NHSN Annual Hospital Survey: [https://www.cdc.gov/nhsn/forms/instr/57\\_103-toi.pdf](https://www.cdc.gov/nhsn/forms/instr/57_103-toi.pdf)

NHSN Annual LTACH Survey: [https://www.cdc.gov/nhsn/forms/57.150\\_ltacfacsurv\\_blank.pdf](https://www.cdc.gov/nhsn/forms/57.150_ltacfacsurv_blank.pdf)  
• Instructions for NHSN Annual LTACH Survey: <https://www.cdc.gov/nhsn/forms/instr/toi-57.150-ltac.pdf>

NHSN Annual IRF Survey: [https://www.cdc.gov/nhsn/forms/57.151\\_rehabfacsurv\\_blank.pdf](https://www.cdc.gov/nhsn/forms/57.151_rehabfacsurv_blank.pdf)  
• Instructions for NHSN Annual IRF Survey: <https://www.cdc.gov/nhsn/forms/instr/toi-57.151-irf.pdf>

NHSN Location Mapping: [https://www.cdc.gov/nhsn/pdfs/pscmanual/15locationsdescriptions\\_current.pdf](https://www.cdc.gov/nhsn/pdfs/pscmanual/15locationsdescriptions_current.pdf)

Keys to Success with NHSN Data: <https://www.cdc.gov/nhsn/ps-analysis-resources/keys-to-success.html>

## ADDENDUM TO THE NHSN GUIDE TO THE SIR

*Hospital Outpatient Department (HOPD)  
Outpatient Procedure Component (OPC)*

## Risk Adjustment Factors Included in the SIR Calculation: 2015 Baseline

### Hospital Outpatient Department (HOPD) Procedure/SSI SIR Model

The number of predicted SSI events is calculated using a logistic regression model (see [page 5](#) above for more information). The SSI SIR is calculated for facilities who enroll in NHSN as acute care hospitals or critical access hospitals. Data from Governmental and Non-governmental Public Health Emergency (PHE) Facilities (facType as HOSP-PHE/G or HOSP-PHE/NG) are excluded from the SIR. Under the 2015 SIR baseline, procedures and associated SSI events occurring in adult and pediatric patients are modeled separately. There is only one SSI SIR Model available for the hospital outpatient procedures (and associated SSIs). Please see *Table 1* below for a summary of the SSI SIR model. Under the 2015 SIR baseline, procedures, regardless of closure methods, are included in the SIR calculation, as long as the inclusion criteria listed below are met and none of the exclusion criteria apply.

**Table 1. Summary of SSI Models**

SSI SIR Model	Inclusion Criteria	Patient Population
All SSI SIR Model	<ul style="list-style-type: none"> <li>Includes <u>only</u> hospital outpatient procedures</li> <li>Includes Superficial, Deep &amp; Organ/Space SSIs</li> <li>Superficial &amp; Deep Incisional SSIs limited to primary incisional SSIs only</li> <li>Includes SSIs identified on admission, readmission &amp; via post-discharge surveillance</li> </ul>	<ul style="list-style-type: none"> <li>Procedures in adult patients</li> <li>Procedures in pediatric patients</li> </ul>

### Exclusion Criteria

The list of exclusion criteria is the same as those listed for the inpatient SSI –Surgical Site Infections–Hospital Inpatient Procedures Models on [page 27](#). The only difference is that this model **includes** outpatient procedures and **excludes** inpatient procedures.

### Predictive Risk Factors for HOPD All SSI SIR Model

The number of predicted events calculated under the 2015 baseline for HOPD SSI is risk adjusted based on the following variables found to be statistically significant predictors of SSIs. The following tables (2a and 2b) list the factors included in each procedure category for adults and pediatrics.

**Table 2a. Predictive Risk Factors from the hospital outpatient procedure department (HOPD) data: All SSI Logistic Regression Model, Adults ≥ 18 years of age<sup>^</sup>**

NHSN Operative Procedure	Risk Factor(s) - All SSI SIR Model, Adults
APPY	<i>Intercept-only model<sup>‡</sup></i>
AVSD	Medical school affiliation*, procedure duration
BRST	Wound class, medical school affiliation*, procedure duration, BMI
CHOL	Diabetes, hospital bed size*, procedure duration
COLO	Procedure duration
FUSN	Spinal level
FX	Procedure duration
HER	Gender, wound class, hospital bed size*, procedure duration, BMI

**Table 2a, Continued<sup>^</sup>**

<b>NHSN Operative Procedure</b>	<b>Risk Factor(s) - All SSI SIR Model, Adults</b>
HPRO	Emergency, hospital bed size*, procedure duration
HYST	Emergency, age, procedure duration, oncology hospital,
KPRO	Medical school affiliation*, age
LAM	Diabetes, medical school affiliation*, procedure duration
OVRY	<i>Intercept-only model<sup>‡</sup></i>
PACE	<i>Intercept-only model<sup>‡</sup></i>
THYR	<i>Intercept-only model<sup>‡</sup></i>
VHYS	<i>Intercept-only model<sup>‡</sup></i>
XLAP	age

\* These risk factors are taken from the [Annual Facility Survey](#).

<sup>^</sup> SIRs are not available for procedure categories that had insufficient data (i.e., < 50 procedures and <1 SSI event) that were reported to NHSN during the baseline period.

<sup>‡</sup> None of the variables investigated were statistically significantly associated with SSI risk in these procedure categories. As a result, the overall pooled mean will be used in the SIR calculation (i.e., intercept-only model).

**Table 2b. Predictive Risk Factors from the hospital outpatient procedure department (HOPD) data: All SSI Logistic Regression Model, Pediatrics <18 years of age\***

<b>NHSN Operative Procedure</b>	<b>Risk Factor(s) - All SSI SIR Model, Adults</b>
FX	Procedure duration
HER, (age >=2)	BMI

\* SIRs are not available for procedure categories that had insufficient data (i.e., < 50 procedures and <1 SSI event) that were reported to NHSN during the baseline period.

## HOPD Procedure Duration Outliers

Please see table 4 on [page 49](#).

## Risk Adjustment Factors Included in the SIR Calculation: 2015 Baseline

### Outpatient Procedure Component Surgical Site Infections (OPC SSI)

The number of predicted SSI events is calculated using a logistic regression model (see [page 5](#) above for more information). The OPC SSI SIR is calculated for facilities who enroll in NHSN as Ambulatory Surgery Centers (ASC). Data from Governmental and Non-governmental Public Health Emergency (PHE) Facilities (facType as HOSP-PHE/G or HOSP-PHE/NG) are excluded from the SIR. Under the 2015 SIR baseline, procedures and associated SSI events occurring in adult outpatients are modeled separately in this new component. There is one SIR model available for outpatient adult procedures (and associated SSIs). Please see *Table 1* below for a summary of the OPC SSI SIR model.

**Table 1. Summary of OPC SSI Model**

OPC SSI SIR Model	Inclusion Criteria	Patient Population
All SSI SIR Model	<ul style="list-style-type: none"> <li>Includes <u>only</u> ambulatory surgery centers procedures</li> <li>Includes Superficial, Deep &amp; Organ/Space SSIs</li> <li>Superficial &amp; Deep Incisional SSIs limited to primary incisional SSIs only</li> <li>Includes SSIs identified on active and passive surveillance</li> </ul>	<ul style="list-style-type: none"> <li>Procedures in adult patients</li> </ul>

### Exclusion Criteria

In addition to the above inclusion criteria, there is also a list of exclusion criteria that applies to the OPC All SSI SIR model. Similar to the PSC, the list of exclusion criteria applies to both procedures and the associated SSI events. Often the reason for excluding procedures and SSI events from the SIR calculation is due to potential data quality issues. It is important that facilities review their data for quality assurance and to determine the reason for exclusion from the SIR calculation.

*Note: When a procedure is excluded from the denominator, the associated SSI event is excluded from the numerator.*

**Table 2. Procedure Exclusions specific to OPC**

<u>General Exclusions</u>
Gender= 'Other'
Inpatient and HOPD procedures and resulting SSIs
SSIs that are reported as superficial incisional secondary (SIS) or deep incisional secondary (DIS)
<u>Exclusions due to potential data quality issues or outliers</u>
Age at the time of procedure is greater than 109 years
Gender is missing
Adult patients ≥ 18 years: if BMI is less than 12 or greater than 60
Procedure duration less than 5 minutes
Procedure duration is greater than IQR5 ( <i>please see <a href="#">Table 4</a> in the OPC SSI Section for more information</i> )

### Predictive Risk Factors in OPC All SSI SIR Model

The number of predicted events, calculated under the 2015 baseline for SSI, is risk adjusted based on the following variables found to be statistically significant predictors of SSIs. The following Table 3 lists the factors included in each procedure with the OPC All SSI model SIR outlined above. Procedure categories that do not have an SIR available will be



revaluated in the future with baseline data to calculate an SIR. In cases when the number of predicted events is less than 1.0, the SIR will not be calculated in NHSN.

**Table 3. Predictive Risk Factors from the OPC All SSI Logistic Regression Model, Adults ≥ 18 years of age\***

<b>NHSN Operative Procedure</b>	<b>Risk Factor(s) - All SSI SIR Model, Pediatrics</b>
BRST	age, anesthesia, BMI
HER	age, BMI, procedure duration
KPRO	<i>Intercept-only model</i> ‡
LAM	<i>Intercept-only model</i> ‡

\*SIRs are not available for procedure categories that had insufficient data (i.e., < 1000 procedures and <1 SSI event) that were reported to NHSN during the baseline period.

‡ None of the variables investigated were statistically significantly associated with SSI risk in these procedure categories. As a result, the overall pooled mean will be used in the SIR calculation (i.e., intercept-only model).

### Procedure Duration Outliers

The **IQR5, also called the procedure duration cutoff point**, is used as an indicator of an extreme outlier for procedure durations when calculating the SSI SIRs. The IQR5 is calculated as five times the interquartile range (Q1-Q3) above the 75th percentile. For example, if the interquartile range is 30 minutes, and the 75th percentile is 100 minutes, the IQR5 would be calculated as:  $100 + (30 \times 5) = 250$  minutes. Procedures with a duration greater than the IQR5 were excluded from the baseline data and will be excluded from all SSI SIR calculations for your facility. **This list of procedure IQR5 apply to both the HOPD Procedure/SSI SIR and the OPC SIR Models.**

**Table 4. IQR5 Values, in Minutes, for NHSN Operative Procedures, Adult Outpatient Procedures**

<b>NHSN Operative Procedure</b>	<b>IQR5 (in minutes)</b>		<b>IQR5 (in hours and minutes)</b>	
	<i>Minutes</i>		<i>Hours</i>	<i>Minutes</i>
AMP	197		3	17
APPY	153		2	33
AVSD	308		5	8
BILI	300		5	0
BRST	355		5	5
CEA	477		7	57
CHOL	223		3	43
COLO	524		8	44
CSEC	166		2	46
FUSN	392		6	32
FX	326		5	26
GAST	326		5	26
HER	249		4	9
HPRO	255		4	15
HYST	452		7	32
KPRO	273		4	33
LAM	307		5	7
NECK	384		6	24
NEPH	296		4	46
OVRY	388		6	28
PACE	228		3	48



**Table 4, Continued**

NHSN Operative Procedure	IQR5 (in minutes)	IQR5 (in hours and minutes)	
	Minutes	Hours	Minutes
PRST	340	5	40
PVBY	627	10	27
REC	228	3	48
RFUSN	542	9	2
SB	669	11	9
SPLE	604	10	4
THOR	414	6	54
THYR	334	8	26
VHYS	433	5	34
VSHN	244	4	4
XLAP	345	5	45

***Outpatient Procedure Component Surgical Site Infections (OPC SSI)***

ASC Surveillance for SSI Events: <https://www.cdc.gov/nhsn/opc/ssi/index.html>

