Data verification log file

Hospital name: Hypothetical Hospital

Country name: Hypothetical Country

Data from:

01 Jan 2016 to 31 Dec 2016

This report is for users to review variable names and data values used by microbiology data file and hospital admission data file saved within the same folder as the application file (AMASS.bat). This report can be used to assist users while completing the data dictionaries for both data files.

Generated on: 16 Jul 2025 08:15:42

Software version: 3.1 released on 16 Jul 2025

Data verification log file

Please review the following information carefully before interpreting the AMR surveillance report generated by AMASS application.

The microbiology data file (stored in the same folder as the application file) had:

50414 records with collection dates ranging from **01 Jan 2016** to **31 Dec 2016** Collection date are in multiple formats.

YMD format. Example : 2016-11-10 00:00:00, 2016-11-10 00:00:00, 2016-11-10 00:00:00

others defined format. Example: 03nov2016, 28apr2016, 04mar2016

Microbiology data's hospital number are from 1 to 9 charecter lengths. The following are most common lengths.

6 charecters long : 91.82% (46290/50414) 3 charecters long : 3.25% (1639/50414) 4 charecters long : 2.62% (1321/50414)

Missing data.

0 records are missing or unknown format of collection date.

0 records are wrong date or wrong date format of collection date.

0 records are missing specimen type.

0 records are missing culture result.

The hospital admission data file (stored in the same folder as the application file) had:

60117 records with admission dates ranging from 01 Jan 2016 to 31 Dec 2016

Admission date are in **others** format. Example : **17may2016**, **11sep2016**, **29may2016**

Discharge date are in **others** format. Example : **01nov2016**, **28sep2016**, **01oct2016**

hospital admission data's hospital number are from **1** to **7** charecter lengths. The following are most common lengths.

6 charecters long : 76.99% (46287/60117) 7 charecters long : 17.74% (10665/60117) 3 charecters long : 2.72% (1634/60117)

Missing data.

0 records are missing or unknown format of admission date.

0 records are missing or unknown format of discharge date.

0 records are wrong date or wrong date format of admission date.

0 records are wrong date or wrong date format of discharge date

0 records are missing outcome.

0 records are missing age.

0 records are missing gender.

The merged data set between microbiology data and hospital admission data had:

Microbiology data records unable to merged with hospital admission data records.

944 records are unable to merge due to no hospital number found in hospital admission data.

367 records are unable to merge due to have hospital number found in hospital admission data but collection date not in admission period.

Merged data.

49103 records are merged (All specimen type).

14833 records are merged (Only blood specimen).

13975 records are merged (Only BSI specimen).

Table S1A: List of data values of the variable recorded for "organism" in your microbiology data file, which are mainly used for the main report

| Data values of variable "organism" in AMASS, which are mainly used for the main report | Data values of variable recorded for "organism" in your microbiology data file | Number of observations |
|--|--|------------------------|
| organism_acinetobacter_anitratus | | 0 |
| organism_acinetobacter_baumannii | Acinetobacter baumannii | 422 |
| organism_acinetobacter_calcoaceticus | | 0 |
| organism_acinetobacter_haemolyticus | | 0 |
| organism_acinetobacter_johnsonii | | 0 |
| organism_acinetobacter_junii | | 0 |
| organism_acinetobacter_lwoffii | | 0 |
| organism_acinetobacter_nosocomialis | | 0 |
| organism_acinetobacter_pittii | | 0 |
| organism_acinetobacter_spp | Acinetobacter spp | 18 |
| organism_enterococcus_avium | | 0 |
| organism_enterococcus_casseliflavus | | 0 |
| organism_enterococcus_dispar | | 0 |
| organism_enterococcus_durans | | 0 |
| organism_enterococcus_faecalis | | 0 |
| organism_enterococcus_faecium | | 0 |
| organism_enterococcus_flavescens | | 0 |
| organism_enterococcus_gallinarum | | 0 |
| organism_enterococcus_hirae | | 0 |
| organism_enterococcus_malodoratus | | 0 |
| organism_enterococcus_mundtii | | 0 |
| organism_enterococcus_pseudoavium | | 0 |
| organism_enterococcus_raffinosus | | 0 |
| organism_enterococcus_seriolicida | | 0 |
| organism_enterococcus_solitarius | | 0 |
| organism_enterococcus_spp | Enterococcus spp | 163 |
| organism_escherichia_coli | Escherichia coli | 963 |
| organism_klebsiella_pneumoniae | Klebsiella pneumoniae | 440 |
| organism_no_growth | no growth | 44018 |
| organism_pseudomonas_aeruginosa | Pseudomonas aeruginosa | 175 |

| Data values of variable "organism" in AMASS, which are mainly used for the main report | Data values of variable recorded for "organism" in your microbiology data file | Number of observations |
|--|--|------------------------|
| organism_salmonella_agona | | 0 |
| organism_salmonella_amsterdam | | 0 |
| organism_salmonella_anatum | | 0 |
| organism_salmonella_arechavaleta | | 0 |
| organism_salmonella_bareilly | | 0 |
| organism_salmonella_blockley | | 0 |
| organism_salmonella_bongori | | 0 |
| organism_salmonella_bonn | | 0 |
| organism_salmonella_bovismorbificans | | 0 |
| organism_salmonella_braenderup | | 0 |
| organism_salmonella_brandenburg | | 0 |
| organism_salmonella_bredeney | | 0 |
| organism_salmonella_cerro | | 0 |
| organism_salmonella_chester | | 0 |
| organism_salmonella_choleraesuis | | 0 |
| organism_salmonella_copenhagen | | 0 |
| organism_salmonella_derby | | 0 |
| organism_salmonella_dublin | | 0 |
| organism_salmonella_emek | | 0 |
| organism_salmonella_enteritidis | | 0 |
| organism_salmonella_falkensee | | 0 |
| organism_salmonella_gallinarum | | 0 |
| organism_salmonella_give | | 0 |
| organism_salmonella_goldcoast | | 0 |
| organism_salmonella_hadar | | 0 |
| organism_salmonella_heidelberg | | 0 |
| organism_salmonella_hirschfeldii | | 0 |
| organism_salmonella_infantis | | 0 |
| organism_salmonella_java | | 0 |

| Data values of variable "organism" in AMASS, which are mainly used for the main report | Data values of variable recorded for "organism" in your microbiology data file | Number of observations |
|--|--|------------------------|
| organism_salmonella_javiana | | 0 |
| organism_salmonella_kaapstad | | 0 |
| organism_salmonella_kedougou | | 0 |
| organism_salmonella_kentucky | | 0 |
| organism_salmonella_kottbus | | 0 |
| organism_salmonella_krefeld | | 0 |
| organism_salmonella_lexington | | 0 |
| organism_salmonella_litchfield | | 0 |
| organism_salmonella_livingstone | | 0 |
| organism_salmonella_lomita | | 0 |
| organism_salmonella_london | | 0 |
| organism_salmonella_manhattan | | 0 |
| organism_salmonella_mbandaka | | 0 |
| organism_salmonella_montevideo | | 0 |
| organism_salmonella_muenchen | | 0 |
| organism_salmonella_muenster | | 0 |
| organism_salmonella_narashino | | 0 |
| organism_salmonella_newport | | 0 |
| organism_salmonella_ohio | | 0 |
| organism_salmonella_oranienburg | | 0 |
| organism_salmonella_orion | | 0 |
| organism_salmonella_ouakam | | 0 |
| organism_salmonella_panama | | 0 |
| organism_salmonella_paratyphi | | 0 |
| organism_salmonella_poona | | 0 |
| organism_salmonella_potsdam | | 0 |
| organism_salmonella_pullorum | | 0 |
| organism_salmonella_reading | | 0 |
| organism_salmonella_rissen | | 0 |

| Data values of variable "organism" in AMASS, which are mainly used for the main report | Data values of variable recorded for "organism" in your microbiology data file | Number of observations |
|--|--|------------------------|
| organism_salmonella_rissen | | 0 |
| organism_salmonella_saintpaul | | 0 |
| organism_salmonella_schottmuelleri | | 0 |
| organism_salmonella_schwarzengrund | | 0 |
| organism_salmonella_senftenberg | | 0 |
| organism_salmonella_spp | Salmonella serogroup O:13 | 9 |
| organism_salmonella_spp | Salmonella serogroup C | 36 |
| organism_salmonella_spp | Salmonella serogroup B | 12 |
| organism_salmonella_spp | Salmonella serogroup D | 73 |
| organism_salmonella_spp | Salmonella serogroup E | 6 |
| organism_salmonella_stanley | | 0 |
| organism_salmonella_tennessee | | 0 |
| organism_salmonella_thompson | | 0 |
| organism_salmonella_typhi | | 0 |
| organism_salmonella_typhimurium | | 0 |
| organism_salmonella_typhisuis | | 0 |
| organism_salmonella_virchow | | 0 |
| organism_salmonella_virginia | | 0 |
| organism_salmonella_wandsworth | | 0 |
| organism_salmonella_weltevreden | | 0 |
| organism_salmonella_weybridge | | 0 |
| organism_salmonella_worthington | | 0 |
| organism_staphylococcus_aureus | Staphylococcus aureus | 415 |
| organism_streptococcus_pneumoniae | Streptococcus pneumoniae | 71 |

Table S1B: List of data values of the variable recorded for "organism" in your microbiology data file, which are mainly used for the annex

| Optional: Data values of variable "organism" in AMASS, which are mainly used for the annex | Data values of the variable recorded for "organism" in your microbiology data file | Number of observations |
|--|--|------------------------|
| organism_achromobacter_spp | | 0 |
| organism_achromobacter_xylosoxidans | | 0 |
| organism_aeromonas_caviae | | 0 |
| organism_aeromonas_dhakensis | | 0 |
| organism_aeromonas_hydrophila | | 0 |
| organism_aeromonas_spp | | 0 |
| organism_aeromonas_veronii | | 0 |
| organism_arcanobacterium_haemolyticum | | 0 |
| organism_arcanobacterium_spp | | 0 |
| organism_arthrobacter_spp | | 0 |
| organism_bacillus_alvei | | 0 |
| organism_bacillus_anthracis | | 0 |
| organism_bacillus_brevis | | 0 |
| organism_bacillus_cereus | | 0 |
| organism_bacillus_circulans | | 0 |
| organism_bacillus_coagulans | | 0 |
| organism_bacillus_firmus | | 0 |
| organism_bacillus_laterosporus | | 0 |
| organism_bacillus_lentus | | 0 |
| organism_bacillus_licheniformis | | 0 |
| organism_bacillus_macerans | | 0 |
| organism_bacillus_megaterium | | 0 |
| organism_bacillus_mycoides | | 0 |
| organism_bacillus_pantothenticus | | 0 |
| organism_bacillus_polymyxa | | 0 |
| organism_bacillus_pumilus | | 0 |
| organism_bacillus_sphaericus | | 0 |
| organism_bacillus_spp | | 0 |
| organism_bacillus_stearothermophilus | | 0 |
| organism_bacillus_subtilis | | 0 |

| Optional: Data values of variable "organism" in AMASS, which are mainly used for the annex | Data values of the variable recorded for "organism" in your microbiology data file | Number of observations |
|--|--|------------------------|
| organism_bacillus_thuringiensis | | 0 |
| organism_brevibacillus_spp | | 0 |
| organism_brevibacterium_acetylicum | | 0 |
| organism_brevibacterium_casei | | 0 |
| organism_brevibacterium_epidermidis | | 0 |
| organism_brevibacterium_mcbrellneri | | 0 |
| organism_brevibacterium_spp | | 0 |
| organism_brucella_abortus | | 0 |
| organism_brucella_canis | | 0 |
| organism_brucella_melitensis | | 0 |
| organism_brucella_spp | | 0 |
| organism_brucella_suis | | 0 |
| organism_burkholderia_cepacia_complex | | 0 |
| organism_burkholderia_pseudomallei | Burkholderia pseudomallei | 432 |
| organism_burkholderia_spp | | 0 |
| organism_campylobacter_spp | | 0 |
| organism_cellulomonas_cellulans | | 0 |
| organism_cellulomonas_spp | | 0 |
| organism_cellulomonas_turbata | | 0 |
| organism_cellulosimicrobium_spp | | 0 |
| organism_chryseobacterium_spp | | 0 |
| organism_citrobacter_amalonaticus | | 0 |
| organism_citrobacter_freundii | | 0 |
| organism_citrobacter_koseri | | 0 |
| organism_citrobacter_spp | | 0 |
| organism_corynebacterium_accolens | | 0 |
| organism_corynebacterium_afermentans | | 0 |
| organism_corynebacterium_amycolatum | | 0 |
| organism_corynebacterium_aquaticum | | 0 |

| Optional: Data values of variable "organism" in AMASS, which are mainly used for the annex | Data values of the variable recorded for "organism" in your microbiology data file | Number of observations |
|--|--|------------------------|
| organism_corynebacterium_bovis | | 0 |
| organism_corynebacterium_cystitidis | | 0 |
| organism_corynebacterium_diphtheriae | | 0 |
| organism_corynebacterium_equi | | 0 |
| organism_corynebacterium_glutamicum | | 0 |
| organism_corynebacterium_haemolyticum | | 0 |
| organism_corynebacterium_jeikeium | | 0 |
| organism_corynebacterium_kutscheri | | 0 |
| organism_corynebacterium_macginleyi | | 0 |
| organism_corynebacterium_matruchotii | | 0 |
| organism_corynebacterium_michiganense | | 0 |
| organism_corynebacterium_minutissimum | | 0 |
| organism_corynebacterium_mycetoides | | 0 |
| organism_corynebacterium_pilosum | | 0 |
| organism_corynebacterium_propinquum | | 0 |
| organism_corynebacterium_pseudodiphtheriticum | | 0 |
| organism_corynebacterium_pseudotuberculosis | | 0 |
| organism_corynebacterium_pyogenes | | 0 |
| organism_corynebacterium_renale | | 0 |
| organism_corynebacterium_spp | | 0 |
| organism_corynebacterium_striatum | | 0 |
| organism_corynebacterium_ulcerans | | 0 |
| organism_corynebacterium_urealyticum | | 0 |
| organism_corynebacterium_xerosis | | 0 |
| organism_cryptococcus_spp | | 0 |
| organism_cutibacterium_spp | | 0 |
| organism_dermabacter_hominis | | 0 |
| organism_dermabacter_spp | | 0 |
| organism_dermacoccus_spp | | 0 |

| Optional: Data values of variable "organism" in AMASS, which are mainly used for the annex | Data values of the variable recorded for "organism" in your microbiology data file | Number of observations |
|--|--|------------------------|
| organism_diphtheroids_spp | | 0 |
| organism_elizabethkingia_anophelis | | 0 |
| organism_elizabethkingia_meningoseptica | | 0 |
| organism_enterobacter_cloacae_complex | | 0 |
| organism_enterobacter_spp | | 0 |
| organism_escherichia_hermannii | | 0 |
| organism_escherichia_spp | | 0 |
| organism_exiguobacterium_acetylicum | | 0 |
| organism_exiguobacterium_spp | | 0 |
| organism_geobacillus_spp | | 0 |
| organism_haemophilus_influenzae | | 0 |
| organism_haemophilus_spp | | 0 |
| organism_hafnia_alvei | | 0 |
| organism_helcobacillus_spp | | 0 |
| organism_helicobacter_pylori | | 0 |
| organism_janibacter_spp | | 0 |
| organism_klebsiella_aerogenes | | 0 |
| organism_klebsiella_oxytoca | Klebsiella oxytoca | 100 |
| organism_klebsiella_spp | | 0 |
| organism_klebsiella_variicola | | 0 |
| organism_knoellia_spp | | 0 |
| organism_kocuria_rosea | | 0 |
| organism_kocuria_spp | | 0 |
| organism_kocuria_varians | | 0 |
| organism_kytococcus_spp | | 0 |
| organism_leclercia_adecarboxylata | | 0 |
| organism_leifsonia_spp | | 0 |
| organism_listeria_monocytogenes | | 0 |
| organism_listeria_spp | | 0 |

| Optional: Data values of variable "organism" in AMASS, which are mainly used for the annex | Data values of the variable recorded for "organism" in your microbiology data file | Number of observations |
|--|--|------------------------|
| organism_microbacterium_arborescens | | 0 |
| organism_microbacterium_imperiale | | 0 |
| organism_microbacterium_spp | | 0 |
| organism_micrococcus_kristinae | | 0 |
| organism_micrococcus_luteus | | 0 |
| organism_micrococcus_roseus | | 0 |
| organism_micrococcus_spp | | 0 |
| organism_micrococcus_varians | | 0 |
| organism_moraxella_spp | | 0 |
| organism_morganella_morganii | | 0 |
| organism_morganella_spp | | 0 |
| organism_neisseria_gonorrhoeae | | 0 |
| organism_neisseria_meningitidis | | 0 |
| organism_nesterenkonia_spp | | 0 |
| organism_ochrobactrum_anthropi | | 0 |
| organism_other_contaminants | | 0 |
| organism_other_enterobacteriaceae | | 0 |
| organism_paenibacillus_alvei | | 0 |
| organism_paenibacillus_macerans | | 0 |
| organism_paenibacillus_polymyxa | | 0 |
| organism_paenibacillus_spp | | 0 |
| organism_pasteurella_spp | | 0 |
| organism_propionibacterium_acnes | | 0 |
| organism_propionibacterium_avidum | | 0 |
| organism_propionibacterium_freudenreichii | | 0 |
| organism_propionibacterium_granulosum | | 0 |
| organism_propionibacterium_jensenii | | 0 |
| organism_propionibacterium_lymphophilum | | 0 |
| organism_propionibacterium_propionicus | | 0 |

| Optional: Data values of variable "organism" in AMASS, which are mainly used for the annex | Data values of the variable recorded for "organism" in your microbiology data file | Number of observations |
|--|--|------------------------|
| organism_propionibacterium_spp | | 0 |
| organism_proteus_mirabilis | | 0 |
| organism_proteus_penneri | | 0 |
| organism_proteus_rettgeri | | 0 |
| organism_proteus_spp | | 0 |
| organism_proteus_stuartii | | 0 |
| organism_proteus_vulgaris | | 0 |
| organism_providencia_spp | | 0 |
| organism_pseudoclavibacter_spp | | 0 |
| organism_pseudomonas_spp | | 0 |
| organism_raoultella_ornitholytica | | 0 |
| organism_raoultella_planticola | | 0 |
| organism_raoultella_spp | | 0 |
| organism_raoultella_terrigena | | 0 |
| organism_serratia_marcescens | | 0 |
| organism_serratia_spp | | 0 |
| organism_shewanella_spp | | 0 |
| organism_shigella_boydii | | 0 |
| organism_shigella_dysenteriae | | 0 |
| organism_shigella_flexneri | | 0 |
| organism_shigella_sonnei | | 0 |
| organism_shigella_spp | | 0 |
| organism_sphingomonas_spp | | 0 |
| organism_staphylococcus_albus | | 0 |
| organism_staphylococcus_arlettae | | 0 |
| organism_staphylococcus_auricularis | | 0 |
| organism_staphylococcus_capitis | | 0 |
| organism_staphylococcus_caprae | | 0 |
| organism_staphylococcus_carnosus | | 0 |

| Optional: Data values of variable "organism" in AMASS, which are mainly used for the annex | Data values of the variable recorded for "organism" in your microbiology data file | Number of observations |
|--|--|------------------------|
| organism_staphylococcus_caseolyticus | | 0 |
| organism_staphylococcus_chromogenes | | 0 |
| organism_staphylococcus_coagulans | | 0 |
| organism_staphylococcus_coagulase_negative | | 0 |
| organism_staphylococcus_cohnii | | 0 |
| organism_staphylococcus_delphini | | 0 |
| organism_staphylococcus_epidermidis | Staphylococcus epidermidis | 742 |
| organism_staphylococcus_equorum | | 0 |
| organism_staphylococcus_felis | | 0 |
| organism_staphylococcus_gallinarum | | 0 |
| organism_staphylococcus_haemolyticus | | 0 |
| organism_staphylococcus_hominis | | 0 |
| organism_staphylococcus_hyicus | | 0 |
| organism_staphylococcus_intermedius | | 0 |
| organism_staphylococcus_kloosii | | 0 |
| organism_staphylococcus_lentus | | 0 |
| organism_staphylococcus_lugdunensis | | 0 |
| organism_staphylococcus_muscae | | 0 |
| organism_staphylococcus_pasteuri | | 0 |
| organism_staphylococcus_piscifermentans | | 0 |
| organism_staphylococcus_pulvereri | | 0 |
| organism_staphylococcus_saccharolyticus | | 0 |
| organism_staphylococcus_saprophyticus | | 0 |
| organism_staphylococcus_schleiferi | | 0 |
| organism_staphylococcus_sciuri | | 0 |
| organism_staphylococcus_simulans | | 0 |
| organism_staphylococcus_spp | | 0 |
| organism_staphylococcus_warneri | | 0 |
| organism_staphylococcus_xylosus | | 0 |

| Optional: Data values of variable "organism" in AMASS, which are mainly used for the annex | Data values of the variable recorded for "organism" in your microbiology data file | Number of observations |
|--|--|------------------------|
| organism_stenotrophomonas_maltophilia | | 0 |
| organism_stenotrophomonas_spp | | 0 |
| organism_streptococcus_agalactiae | | 0 |
| organism_streptococcus_alactolyticus | | 0 |
| organism_streptococcus_anginosus | | 0 |
| organism_streptococcus_bovis | | 0 |
| organism_streptococcus_canis | | 0 |
| organism_streptococcus_constellatus | | 0 |
| organism_streptococcus_cristatus | | 0 |
| organism_streptococcus_dysgalactiae | | 0 |
| organism_streptococcus_gallolyticus | | 0 |
| organism_streptococcus_gordonii | | 0 |
| organism_streptococcus_hyointestinalis | | 0 |
| organism_streptococcus_infantis | | 0 |
| organism_streptococcus_intermedius | | 0 |
| organism_streptococcus_lutetiensis | | 0 |
| organism_streptococcus_mitis | | 0 |
| organism_streptococcus_mutans | | 0 |
| organism_streptococcus_oralis | | 0 |
| organism_streptococcus_parasanguinis | | 0 |
| organism_streptococcus_pluranimalium | | 0 |
| organism_streptococcus_porcinus | | 0 |
| organism_streptococcus_pseudoporcinus | | 0 |
| organism_streptococcus_pyogenes | | 0 |
| organism_streptococcus_salivarius | | 0 |
| organism_streptococcus_sanguinis | | 0 |
| organism_streptococcus_spp | | 0 |
| organism_streptococcus_suis | | 0 |
| organism_streptococcus_thoraltensis | | 0 |

| Optional: Data values of variable "organism" in AMASS, which are mainly used for the annex | Data values of the variable recorded for "organism" in your microbiology data file | Number of observations |
|--|--|------------------------|
| organism_streptococcus_thoraltensis | | 0 |
| organism_streptococcus_uberis | | 0 |
| organism_streptococcus_vestibularis | | 0 |
| organism_trueperella_spp | | 0 |
| organism_vibrio_alginolyticus | | 0 |
| organism_vibrio_carchariae | | 0 |
| organism_vibrio_cholerae | | 0 |
| organism_vibrio_cincinnatiensis | | 0 |
| organism_vibrio_damsela | | 0 |
| organism_vibrio_fluvialis | | 0 |
| organism_vibrio_furnissii | | 0 |
| organism_vibrio_hollisae | | 0 |
| organism_vibrio_metschnikovii | | 0 |
| organism_vibrio_mimicus | | 0 |
| organism_vibrio_parahaemolyticus | | 0 |
| organism_vibrio_spp | Vibrio spp | 47 |
| organism_vibrio_vulnificus | | 0 |
| organism_virgibacillus_spp | | 0 |
| organism_viridans_group_streptococci | | 0 |
| organism_yersinia_enterocolitica | | 0 |
| organism_yersinia_pseudotuberculosis | | 0 |
| organism_yersinia_spp | | 0 |
| | others | 2272 |

Table S2 (Summary): List of number of records per AMASS's "specimen type" in your microbiology data file

| Data values of variable name recorded for "specimen type" in your hospital admission data file | Number of observations |
|--|------------------------|
| specimen_blood | 15878 |
| specimen_cerebrospinal_fluid | 503 |
| specimen_genital_swab | 478 |
| specimen_others | 3454 |
| specimen_respiratory_tract | 9819 |
| specimen_stool | 623 |
| specimen_urine | 19659 |

Table S2: List of data values of the variable recorded for "specimen type" in your microbiology data file

| Data values of variable used for "specimen type" in AMASS | Data values of variable recorded for "specimen type" in your microbiology data file | Number of observations |
|---|---|------------------------|
| specimen_blood | blood | 15878 |
| specimen_cerebrospinal_fluid | csf | 503 |
| specimen_genital_swab | genital swab | 478 |
| specimen_others | pus from wound | 2011 |
| specimen_others | other | 1443 |
| specimen_respiratory_tract | sputum | 9819 |
| specimen_stool | stool | 623 |
| specimen_urine | urine | 19659 |

Table S3: List of variables recorded for "antibiotics" in your microbiology data file

| Variable names used for "antibiotics" described in AMASS | Variable names recorded for "antibiotics" in your microbiology data file | Number of observations containing S, I, or R for each antibiotic | Number of observations containing some data (i.e. not blank) which were not translated into S, I or R* |
|---|--|--|--|
| Amikacin | AK | 100 | 0 |
| Amoxicillin | AMX | 100 | 0 |
| Ampicillin | AP | 846 | 0 |
| Cefepime | FEP | 224 | 0 |
| Cefotaxime | CTX | 986 | 0 |
| Cefoxitin | FOX | 415 | 0 |
| Cefpodoxime | CPD | 0 | 0 |
| Ceftazidime | CAZ | 1292 | 0 |
| Ceftriaxone | CRO | 1439 | 0 |
| Ciprofloxacin | CFX | 1542 | 0 |
| Clindamycin | DA | 486 | 0 |
| Colistin | CL | 0 | 0 |
| Daptomycin | DAP | 0 | 0 |
| Doripenem | DOR | 3 | 0 |
| Ertapenem | ETP | 1303 | 0 |
| Erythromycin | Е | 71 | 0 |
| Gentamicin | GM | 2018 | 0 |
| Imipenem | IPM | 1918 | 0 |
| Levofloxacin | LVX | 0 | 0 |
| Linezolid | LNZ | 21 | 0 |
| Meropenem | MEM | 1419 | 0 |
| Minocycline | MI | 0 | 0 |
| Oxacillin | OX | 0 | 0 |
| Penicillin_G | PG | 60 | 0 |
| Piperacillin_and_tazobactam | TZP | 29 | 0 |
| Sulfamethoxazole_and_trimethoprim | SXT | 1372 | 0 |
| Teicoplanin | TCO | 0 | 0 |
| Tigecycline | TGC | 440 | 0 |
| Vancomycin | VAN | 578 | 0 |
| | department | 4124 | |

^{*} For antibiotics used for the AST, the numbers in this column should be 0. This is because there should be no data that is not translated into S, I or R. This may occur when laboratories record some number into the same column (e.g. a mix of S I R and zone size) or when dictionary is still incomplete (e.g. recording "R" and "R - no zone" but having only "R" in the dictionary); NA=Not applicable. This could occur when the variable names for antibiotics are not used in AMASS.

Table S3 (continue): List of variables recorded for "antibiotics" in your microbiology data file

| Variable names used for "antibiotics" described in AMASS | Variable names recorded for "antibiotics" in your microbiology data file | Number of observations containing S, I, or R for each antibiotic | Number of observations containing some data (i.e. not blank) which were not translated into S, I or R* |
|---|--|--|--|
| | department | 4124 | |

^{*} For antibiotics used for the AST, the numbers in this column should be 0. This is because there should be no data that is not translated into S, I or R. This may occur when laboratories record some number into the same column (e.g. a mix of S I R and zone size) or when dictionary is still incomplete (e.g. recording "R" and "R - no zone" but having only "R" in the dictionary); NA=Not applicable. This could occur when the variable names for antibiotics are not used in AMASS.

Table S4: List of data values of variable recorded for "gender" in your hospital admission data file

| Data values of variable used for "gender" described in AMASS | Data values of variable recorded for "gender" in your hospital admission data file | Number of observations |
|--|--|------------------------|
| male | M | 28570 |
| female | F | 28548 |
| | | 2999 |

Table S5: List of data values of variable recorded for "age" in your hospital admission data file

| Data values of variable used for "age" described in AMASS | Data values of variable recorded for "age" in your hospital admission data file | Number of observations |
|---|---|------------------------|
| Less than 1 year | 0 | 327 |
| 1 to 4 years | 1 | 746 |
| 1 to 4 years | 2 | 706 |
| 1 to 4 years | 3 | 735 |
| 1 to 4 years | 4 | 687 |
| 5 to 14 years | 5 | 740 |
| 5 to 14 years | 6 | 648 |
| 5 to 14 years | 7 | 727 |
| 5 to 14 years | 8 | 676 |
| 5 to 14 years | 9 | 744 |
| 5 to 14 years | 10 | 693 |
| 5 to 14 years | 11 | 705 |
| 5 to 14 years | 12 | 737 |
| 5 to 14 years | 13 | 662 |
| 5 to 14 years | 14 | 691 |
| 15 to 24 years | 15 | 711 |
| 15 to 24 years | 16 | 759 |
| 15 to 24 years | 17 | 718 |
| 15 to 24 years | 18 | 704 |
| 15 to 24 years | 19 | 684 |
| 15 to 24 years | 20 | 655 |
| 15 to 24 years | 21 | 693 |
| 15 to 24 years | 22 | 792 |
| 15 to 24 years | 23 | 693 |
| 15 to 24 years | 24 | 682 |
| 25 to 34 years | 25 | 680 |
| 25 to 34 years | 26 | 665 |
| 25 to 34 years | 27 | 744 |
| 25 to 34 years | 28 | 707 |
| 25 to 34 years | 29 | 655 |

Table S5 (continue): List of data values of variable recorded for "age" in your hospital admission data file

| Data values of variable used for "age" described in AMASS | Data values of variable recorded for "age" in your hospital admission data file | Number of observations |
|---|---|------------------------|
| 25 to 34 years | 30 | 683 |
| 25 to 34 years | 31 | 733 |
| 25 to 34 years | 32 | 689 |
| 25 to 34 years | 33 | 685 |
| 25 to 34 years | 34 | 710 |
| 35 to 44 years | 35 | 725 |
| 35 to 44 years | 36 | 680 |
| 35 to 44 years | 37 | 647 |
| 35 to 44 years | 38 | 682 |
| 35 to 44 years | 39 | 751 |
| 35 to 44 years | 40 | 738 |
| 35 to 44 years | 41 | 703 |
| 35 to 44 years | 42 | 707 |
| 35 to 44 years | 43 | 734 |
| 35 to 44 years | 44 | 750 |
| 45 to 54 years | 45 | 704 |
| 45 to 54 years | 46 | 663 |
| 45 to 54 years | 47 | 716 |
| 45 to 54 years | 48 | 722 |
| 45 to 54 years | 49 | 658 |
| 45 to 54 years | 50 | 686 |
| 45 to 54 years | 51 | 693 |
| 45 to 54 years | 52 | 722 |
| 45 to 54 years | 53 | 689 |
| 45 to 54 years | 54 | 704 |
| 55 to 64 years | 55 | 705 |
| 55 to 64 years | 56 | 747 |
| 55 to 64 years | 57 | 639 |
| 55 to 64 years | 58 | 768 |

Table S5 (continue): List of data values of variable recorded for "age" in your hospital admission data file

| Data values of variable used for "age" described in AMASS | Data values of variable recorded for "age" in your hospital admission data file | Number of observations |
|---|---|------------------------|
| 55 to 64 years | 58 | 768 |
| 55 to 64 years | 59 | 763 |
| 55 to 64 years | 60 | 672 |
| 55 to 64 years | 61 | 709 |
| 55 to 64 years | 62 | 733 |
| 55 to 64 years | 63 | 774 |
| 55 to 64 years | 64 | 715 |
| 65 to 80 years | 65 | 699 |
| 65 to 80 years | 66 | 680 |
| 65 to 80 years | 67 | 742 |
| 65 to 80 years | 68 | 702 |
| 65 to 80 years | 69 | 720 |
| 65 to 80 years | 70 | 791 |
| 65 to 80 years | 71 | 742 |
| 65 to 80 years | 72 | 705 |
| 65 to 80 years | 73 | 674 |
| 65 to 80 years | 74 | 691 |
| 65 to 80 years | 75 | 724 |
| 65 to 80 years | 76 | 725 |
| 65 to 80 years | 77 | 653 |
| 65 to 80 years | 78 | 708 |
| 65 to 80 years | 79 | 730 |
| 65 to 80 years | 80 | 683 |
| More than 80 years | 81 | 677 |
| More than 80 years | 82 | 724 |
| More than 80 years | 83 | 684 |
| More than 80 years | 84 | 725 |
| More than 80 years | 85 | 373 |

Table S6: List of data values of variable recorded for "discharge status" in your hospital admission data file

| Data values of variable name used for "discharge status" described in AMASS | Data values of variable name recorded for "discharge status" in your hospital admission data file | Number of observations |
|---|---|------------------------|
| died | discharge to hospice | 2209 |
| died | dead | 5290 |
| | discharged | 52618 |

Table S7: List of variable names in your microbiology data file

| Variable names used in your microbiology data file |
|--|
| hn |
| spcdate |
| spctype |
| organism |
| department |
| FOX |
| VAN |
| DA |
| AMX |
| тсо |
| LNZ |
| DAP |
| PG |
| OX |
| SXT |
| CRO |
| CAZ |
| E |
| LVX |
| CFX |
| CTX |
| IPM |
| MEM |
| ETP |
| DOR |
| GM |
| AK |
| AP |
| CPD |
| FEP |

Table S7 (continue): List of variable names in your microbiology data file

| Variable names used in your microbiology data file |
|--|
| FEP |
| CL |
| TZP |
| TGC |
| MI |

Table S8: List of variable names in your hospital admission data file

| Variable names used in your hospital admission data file |
|--|
| hn |
| admdate |
| disdate |
| gender |
| age |
| dischargestatus |

| Table S9: Duplicate mapping of data value of variable in your with data values of variable describe in AMASS |
|--|
| There are no duplicate records in dictionary files that need your revision. |
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| Table S10A: List of data values of variable recorded for "ward" in your microbiology data file |
|--|
| No ward column in microbiology data file defined in dictionary for wards. |
| |
| |
| |
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| |
| |

Table S10B: List of data values of variable recorded for "ward" in your hospital admission data file. (Data values will be used if missing ward data in microbiology data file for that merged data record.)

No ward column in hospital admission data file defined in dictionary for wards.