| | JIRA | Title | Component/s | Release Note Description | Relevant to Local Health Departments? Yes or No |
|---|---|---|---|---|---|
| 1 | MAEDSS-15146 | Update *Candida auris* Event Date Hierarchy to Include Earliest Positive Test Result | Antibiotic Resistance/Healthcare Acquired Infections | We are updating the CAURIS event date hierarchy so that the earliest positive test result is prioritized over earliest test result, since CAURIS events may have negative screening results preceeding later positive reuslts. | No |
| 2 | MAEDSS-15845 | Reprocess *Candida auris* Events for Event Date Hierarchy Changes | Antibiotic Resistance/Healthcare Acquired Infections | We will reprocess all CAURIS events so that all CAURIS event dates recalculate according to the updated event date hierarchy (MAEDSS-15146) | No |
| 3 | MAEDSS-15147 | Multi Drug Resistance Organism and *Candida auris (*Modeling Changes) | Antibiotic Resistance/Healthcare Acquired Infections | For MDRO and CAURIS events, we are adding a new Healthcare Admissions History table in the Clinical question package and retiring old hospitalization and supervised care questions that the table is replacing. We are also updating the list of carbapenemases recorded for CPOs in the MDRO administrative question package. | No |
| 4 | MAEDSS-15451 | Create New Reference Group for Admissions Facility Type | Antibiotic Resistance/Healthcare Acquired Infections | We are creating a new reference group of healthcare facility types to be used with new QID: ADMISSIONS_FACILITY_TYPE in the new Healthcare Admissions History table in MDRO and CAURIS events (MAEDSS-15147). | No |
| 5 | MAEDSS-15450 | Create New Reference Group for Unit Type | Antibiotic Resistance/Healthcare Acquired Infections | We are creating a new reference group of healthcare unit types to be used with new QID: ADMISSIONS_FACILITY_UNIT_TYPE in the new Healthcare Admissions History table in MDRO and CAURIS events (MAEDSS-15147). | No |
| 6 | MAEDSS-15492 | Infection Prevention wizard Changes for Multi Drug Resistance Organism and *Candida auris* | Antibiotic Resistance/Healthcare Acquired Infections | We are adding the new Healthcare Admissions History table (MAEDSS-15147) to the IP Wizard for MDRO and CAURIS events, and removing CASE_HOSPITALIZED and EMPLOYED_OR_ATTEND_SUPERVISED_CARE. | No |
| 7 | MAEDSS-15395 | Addition of Message Mapping Guide variables to Multi Drug Resistance Organism events | Antibiotic Resistance/Healthcare Acquired Infections | Adding new QIDs in the Clinical and Risk question packages of MDRO/CAURIS events as a part of NMI according to the HAI MMG guide | No |
| 8 | MAEDSS-15145 | Create CRF Needs Review Cluster Events -- Multi Drug Resistance Organism Workflow | Antibiotic Resistance/Healthcare Acquired Infections | We are creating the new CRF Needs Review Cluster Events – MDRO workflow to help the HAI/AR program better track and manage MDRO and CAURIS cluster events. | No |
| 9 | MAEDSS-15403 | Add New Cluster CRF Reviewed Question to Multi Drug Resistance Organism and *Candida auris* Cluster Events | Antibiotic Resistance/Healthcare Acquired Infections | We are adding the variable "Cluster CRF reviewed:" to MDRO and CAURIS cluster events in the Administrative question package. This question will serve as a trigger to move clusters in/out of the new CRF Needs Review Cluster Events – MDRO workflow (MAEDSS-15145) | No |
| 10 | MAEDSS-15558 | Updating the Case Report Form (CRF) needs review- Active Surveillance workflow | Antibiotic Resistance/Healthcare Acquired Infections | Renaming the workflow "Invasive GAS CRF Review" to capture only GAS events to be reviewed by the program once EOD is complete | No |
| 11 | MAEDSS-15390 | Update to the autogenerated linked event report for Grp A Strep dashboard | Antibiotic Resistance/Healthcare Acquired Infections | Updating the report to extend the time frame from 1/1/2015 to current date to match the Event information extract report | No |
| 12 | MAEDSS-15389 | Update to the autogenerated HAI outbreak report for Grp A Strep dashboard | Antibiotic Resistance/Healthcare Acquired Infections | Updating the report to extend the time frame from 1/1/2015 to current date to match the Event information extract report | No |
| 13 | MAEDSS-15388 | Update the autogenerate Event information extract report for Grp A Strep dashboard | Antibiotic Resistance/Healthcare Acquired Infections | Updating the report to include the sequencing question package QIDs | No |
| 14 | MAEDSS-15249 | Grp A Strep Modeling Changes | Antibiotic Resistance/Healthcare Acquired Infections | In the Risk/Exposure/Control & Prevention question package for Group A Strep (GAS) events, we are adding a new exposure period label, retiring out of date incubation period labels, adding new questions related to supervised care, and retiring some variables related to supervised care and school that are no longer needed. | Yes |
| 15 | MAEDSS-15250 | Grp A Strep Cluster Modeling Changes | Antibiotic Resistance/Healthcare Acquired Infections | We are adding "LTC" as an answer choice for two questions about the unit type affected or involved in screening in the HAI question package for cluster events. | Yes |
| 16 | MAEDSS-15382 | Updating the generated Antibiogram report with new the new Antibiotic list | Antibiotic Resistance/Healthcare Acquired Infections | Updating Antibiogram reports to include the updated list of Antibiotics | No |
| 17 | MAEDSS-15379 | Updating Antibiotics list for the Antibiogram Model | Antibiotic Resistance/Healthcare Acquired Infections | Adding/retiring antibiotics required for generating the Antibiogram report | No |
| 18 | MAEDSS-15394 | Addressing HL7 2.5.1 Susceptibility Messages with Duplicate Organism Results | Antibiotic Resistance/Healthcare Acquired Infections | When the XSL processor encounters ELR messages that contain two results of the same organism isolated from the same specimen under the parent test OBR, it should 1)create the appropriate event(s) in MAVEN, 2) add all results in the message to the event(s), and 3) not create an unknown event. If the accession number, test, and result are the same in a message for more than one parent result, default to the first of the matching parent results. | No |
| 19 | MAEDSS-15392 | Create new workflow for WGS Grp A Strep events | Antibiotic Resistance/Healthcare Acquired Infections & Next Genome Sequencing | New Workflow created to include GAS events that are WGS by the lab that have the "Sequencing Needs Review:" Yes in the Sequencing information QP for the HAI program to be able to keep track and review | No |
| 20 | MAEDSS-15482 | Retire ENX (Mass Notify) Question Package | COVID | Retiring question package related to MassNotify, as the program has been sunset. | Yes |
| 21 | MAEDSS-14284 | Move COVID sequencing data into Sequencing Question Package | COVID, Next Genome Sequencing | Building an execution output to automatically pull COVID sequencing data from the investigation fields to the Sequencing Question Package. Fields to be populated in the Sequencing QP include: NGS_SPEC_NUMBER NGS_REPOSITORY_TYPE NGS_STRAIN (result) NGS_STRAIN (result value) NGS_DATE_MODIFIED | No |

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|---|---|---|---|---|---|
| 22 | MAEDSS-15505 | Add additional QIDs to eCR Event Extract | Electronic Case Reporting (eCR) | Adding additional QIDs to eCR Event extract including: REPORTING_SOURCE REPORTING_SOURCE_DATE EICR_FACILITY_HOSPITALIZATION EICR_ADMISSION_DATE EICR_DISCHARGE_DATE EICR_PREGNANT_STATUS_DATE EICR_GESTATIONAL_WEEKS EICR_PREGNANT_OUTCOME EICR_MEDS_ENDDATE EICR_FILENAME | No |
| 23 | MAEDSS-15489 | Change Answer Option for ESPECR_MEDS_STATUS and ESPECR_MEDS_DOSE_UNITS | Electronic Case Reporting (eCR) | Changing the answer option for these two QIDs to string text instead of current answer options for ESP eCR | No |
| 24 | MAEDSS-15485 | Create EICR_BIRTH_SEX | Electronic Case Reporting (eCR) | Creating new QID to capture patient sex assigned at birth as reported by eCR | No |
| 25 | MAEDSS-15484 | Change Answer Option for EICR_MEDS_STATUS and EICR_MEDS_DOSE_UNITS | Electronic Case Reporting (eCR) | Changing the answer option for these two QIDs to string text instead of current answer options for AIMS eCR | No |
| 26 | MAEDSS-15469 | Create EICR_TRIBAL_AFFILIATION and EICR_TRIBAL_AFFILIATION_ENROLLED | Electronic Case Reporting (eCR) | Creating new hidden QIDs to capture the name of the tribe that the patient is affiliated with and the enrollment status of the patient for that tribe (Y/N) | No |
| 27 | MAEDSS-15467 | Change Answer Options for ESPECR_PREGNANT_OUTCOME | Electronic Case Reporting (eCR) | Changing the answer option for a patient's outcome of pregnancy to be consistent with the reference code list established in 15466 (value set consistent with eCR CDA IG). This is for ESP eCR. | No |
| 28 | MAEDSS-15466 | Create new reference code list ECR_PREGNANCY_OUTCOMES | Electronic Case Reporting (eCR) | Creating a new reference code list to capture answer options for a patient's pregnancy outcome consistent with the value set within the eCR CDA IG | No |
| 29 | MAEDSS-15465 | Change Answer Option for EICR_PREGNANT_OUTCOME | Electronic Case Reporting (eCR) | Changing the answer option for a patient's outcome of pregnancy to be consistent with the reference code list established in 15466 (value set consistent with eCR CDA IG). This is for AIMS eCR. | No |
| 30 | MAEDSS-15790 | Fix spelling error of ITLS question in 09. Other External Data Sources | Epidemiology Program - Hepatitis | Fix incorrect spelling of 'Identity' | No |
| 31 | MAEDSS-15231 | Update list of substances available as response options for drug use questions in viral hepatitis events | Epidemiology Program - Hepatitis | Add additional substances to dropdown list in HAV, HBV and HCV drug use questions | Yes |
| 32 | MAEDSS-15441 | Modify Acute HBV Identification workflow to remove age and HCV co-infection criteria | Epidemiology Program - Hepatitis | Removed age and HCV co-infection criteria | No |
| 33 | MAEDSS-15340 | Unable to upload roster into MAVEN QA for "HEPB" product code | Epidemiology Program - Hepatitis | Fix custom code preventing the creation of HEPB events via roster | No |
| 34 | MAEDSS-15567 | Update "Confirmed HCV Teleform Print" workflow | Epidemiology Program - Hepatitis | Boston as an official city was excluded from this workflow prior to the transition from BoSS to MAVEN. Removed exclusion now that Boston is on MAVEN | No |
| 35 | MAEDSS-15536 | Add child question to capture D+R- transplants in Hep C cases | Epidemiology Program - Hepatitis | New question to collect whether transplant was D+R- | Yes |
| 36 | MAEDSS-15299 | Update first.postive.then.negative caseattribute to include detected and reactive results | Epidemiology Program - Hepatitis | Add detected and reactive snomeds into case attribute in case no positive snomeds are present | No |
| 37 | MAEDSS-15294 | Fix INFECTIOUS_PERIOD_START_DATE and INFECTIOUS_PERIOD_STOP_DATE so they appear in HAV, Rubella and Mumps cases again | Epidemiology Program - Hepatitis & Immunication | Fields were inadvertenly removed in 2023 for HAV, mumps and rubella. Adding fields back. | |
| 38 | MAEDSS-15835 | Add new criteria for when Norovirus (caliciviruses) events should enter the LBOH Case Report Forms (CRF) are pending workflow | Epidemiology Program - Foodborne & Waterborne | Norovirus (caliciviruses) events will now automatically enter the LBOH CRF are pending WF when LBOH Acknowledged = Yes and new age criteria are met; otherwise, an epi needs to manually answer Yes to Initiate LBOH Investigation. | Yes |
| 39 | MAEDSS-15834 | Modify LBOH Notification but no follow-up required workflow by adding Norovirus (caliciviruses) events that fit age criteria | Epidemiology Program - Foodborne & Waterborne | LBOHs are no longer recommended to investigate norovirus events for individuals between the ages of 12 years and 80 years old so Norovirus (caliciviruses) events for individuals within this age group will now flow into the LBOH Notification but no follow-up required workflow. | Yes |
| 40 | MAEDSS-15833 | Add age criteria for Norovirus (caliciviruses) events to enter LBOH Notification for Routine Disease workflow | Epidemiology Program - Foodborne & Waterborne | Age criteria was added for when Norovirus (caliciviruses) events should enter the LBOH Notification for Routine Disease WF because investigation of norovirus events is now recommended for only children under 12 years old and adults 80 years and over. | Yes |
| 41 | MAEDSS-15831 | Turn on Question "Initiate LBOH investigation?" for Norovirus (caliciviruses) | Epidemiology Program - Foodborne & Waterborne | Adding new question to Norovirus (caliciviruses) events for when to manually send an event to an LBOH if age criteria are not met (e.g., for a cluster investigation). | Yes |
| 42 | MAEDSS-15823 | Correct spelling of answer choice "Employee" for QID: VS_HC_SET_TYPE_EXPOSURE in EPI/IMM model | Epidemiology Program - Foodborne & Waterborne | Correcting a misspelling of Legionella event variable. | Yes |

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|---|---|---|---|---|---|
| 43 | MAEDSS-15761 | Optimize FBI Event Information Extract | Epidemiology Program - Foodborne & Waterborne | FBI Event Information Extract is optimized to run quicker and to run for the full length of time in MAVEN. | Yes |
| 44 | MAEDSS-15397 | Reference code list for Legionella labs that are sent to LBOHs for investigation without need for FB review | Epidemiology Program - Foodborne & Waterborne | Creating new reference group called REVIEW_LEGIONELLA_EVENTS for Legionella LOINC/SNOMED combinations that require review by an epidemiologist before being sent to an LBOH for investigation. | Yes |
| 45 | MAEDSS-15375 | Yersinia Risk questions updates | Epidemiology Program - Foodborne & Waterborne | Adding new questions to align with risk factors outlined by CSTE including in the Clinical QP (iron overload, blood product transfusion) and Risk QP (drinking water, food, animal, diaper, and large gathering exposures). | Yes |
| 46 | MAEDSS-15374 | Add new question "Probable type:" (QID: PROBABLE_TYPE) to the Admin QP for enteric events | Epidemiology Program - Foodborne & Waterborne | Adding new question to distinguish between probable events based on culture independent diagnostic tests (CIDTs) and epidemiologically linked cases. | Yes |
| 47 | MAEDSS-15368 | Campylobacter Risk Question updates | Epidemiology Program - Foodborne & Waterborne | Adding, modifying, and retiring of risk data elements primarily for Campylobacter events, including adding raw shellfish exposure questions. | Yes |
| 48 | MAEDSS-15229 | Update Legionella labs to bypass Enteric and Waterborne Lab Review workflow, and conduct review of Legionella tests | Epidemiology Program - Foodborne & Waterborne | Updating the Enteric and Waterborne Lab Review WF so Legionella events with labs that don't require review by an epidemiologist for clinical compatiblity will bypass the WF and go directly to the LBOH for investigation. | Yes |
| 49 | MAEDSS-15117 | Add CSTE-recommended data elements to MAVEN for Salmonella events | Epidemiology Program - Foodborne & Waterborne | Adding/removing questions in the Salmonella Risk QP to align with risk factors outlined by CSTE, including shellfish/seafood, unpasteurized milk/cheese, and fruit/leafy green consumption. | Yes |
| 50 | MAEDSS-15493 | Creation of new report for pulling Sal data from MAVEN into S3 bucket for use in Pathogen Genomics Data Viz Tool (Lantern) | Epidemiology Program - Foodborne & Waterborne & Next Genome Sequencing | Creates a new report with variables relevant to Salmonella for Lantern tool to run automatically M-F | No |
| 51 | MAEDSS-15376 | Add new question "Toxin:" (QID: TOXIN) to the Sequencing QP for CHOL events | Epidemiology Program - Foodborne & Waterborne & Next Genome Sequencing | Adding a new question "Toxin" to the Sequencing QP for Vibrio events to capture whole genome sequencing information that shows if Vibrio cholerae species is toxigenic and may cause Cholera disease. | |
| 52 | MAEDSS-14851 | Automation of current reports for pulling Sal data from MAVEN for use in Pathogen Genomics Data Viz Tool (Lantern) | Epidemiology Program - Foodborne & Waterborne & Next Genome Sequencing | Creating copy of two reports (Lab Info Extract and Linked Events Report) with criteria specific to Lantern tool to run automatically M-F | No |
| 53 | MAEDSS-14705 | Adding datestamp variables to the Enteric WGS Question Package | Epidemiology Program - Foodborne & Waterborne & Next Genome Sequencing | Adds new datestamps to the Sequencing QP to identify when genomic data from SEDRIC was first entered (NGS_INPUT_DATE) and when a national cluster ID was first entered (NGS_CLUSTER_FIRSTDATE) and last updated (NGS_CLUSTER_LASTDATE) | No |
| 54 | MAEDSS-14245 | Enteric Sequencing SEDRIC automation | Epidemiology Program - Foodborne & Waterborne & Next Genome Sequencing | Implementing/building an API between CDC's SEDRIC system and MAVEN for the daily pull of genomic data into the WGS Question Package for SAL, SHIG, STEC, CHOL, and LIST product codes | No |
| 55 | MAEDSS-15875 | Move school-related QID in Risk History QP | Epidemiology Program - Immunization | Related to MAEDSS-15193. Rearraging QIDs in the back-end so questions remain in logical order on the front-end. | No |
| 56 | MAEDSS-15663 | Optimized MIIS VPD Workflow and add SP | Epidemiology Program - Immunization | Fixes a bug that prevents events from entering the MIIS VPD review workflow, and adds SP events as part of MAEDSS-15481. | No |
| 57 | MAEDSS-15527 | Update VPD 10-Year Morbidity Report | Epidemiology Program - Immunization | Updates to the VPD 10-Year Morbidity Report to accurately count cases. | No |
| 58 | MAEDSS-15502 | Update Mumps and Pertussis Wizards with new QIDs | Epidemiology Program - Immunization | Adding QIDs to Pertussis and Mumps CRF wizards. These questions were added in the previous release and are a priority for CDC. | Yes |
| 59 | MAEDSS-15499 | Retire FOLLOWUP QIDs in Strep Pneumo events | Epidemiology Program - Immunization | Retiring questions related to supervised care location information and household members in Strep Pneumo events in the Risk/Exposure QP. | Yes |
| 60 | MAEDSS-15481 | MIIS Integration: Strep Pneumo | Epidemiology Program - Immunization | Adding Strep Pneumo events to our MIIS integration, allowing vaccine info to populate SP events automatically. | Yes |
| 61 | MAEDSS-14020 | Add Infection Prevention fields to Admin QP in FLU Events | Epidemiology Program - Immunization | Adding the IP module to flu events, so IPs can directly update high priority flu events. | No |
| 62 | MAEDSS-15604 | Infection Prevention wizard updates related to FLU events | Epidemiology Program - Immunization | Related to MAEDSS-15604. | No |
| 63 | MAEDSS-15345 | Create a new variable to document communication concerns for PHB events | Epidemiology Program - Immunization - Perinatal Hepatis B | Add a new variable to document communication concerns to better inform case contact and management. Triggers concern on splashscreen. | No |
| 64 | MAEDSS-15344 | Add PHB communication concern to Hepatitis B events | Epidemiology Program - Immunization - Perinatal Hepatis B | Concern shows on splashscreen if a communication concern is noted | No |
| 65 | MAEDSS-12927 | Add Quantitative HBsAg to Perinatal Hepatitis B workflows | Epidemiology Program - Immunization - Perinatal Hepatis B | Only qualitative HBsAgs are in workflow logic. Adding quantitative. | No |
| 66 | MAEDSS-15521 | Move FLU sequencing data into Sequencing QP | Epidemiology Program - Immunization & Next Genomic Sequencing | Building an execution output to automatically pull FLU sequencing data from the investigation fields to the Sequencing Question Package. Fields to be populated in the Sequencing QP include: NGS_SPEC_NUMBER NGS_REPOSITORY_TYPE NGS_STRAIN (result) NGS_STRAIN (result value) NGS_CLADE NGS_DATE_MODIFIED | No |
| 67 | MAEDSS-15440 | Modify Zoonotic Additional Lab Results Received workflow to include Hansen's disease and Hantavirus product codes | Epidemiology Program - Zoonotics | Add new product codes to workflow | No |
| 68 | MAEDSS-15439 | Modify Zoonotic CRF Needs Review Workflow to include CJD and Hansen's product codes | Epidemiology Program - Zoonotics | Add new product codes to workflow | No |
| 69 | MAEDSS-15359 | Modify SS_JOINT_ACHES_PAINS symptom variable for multiple product codes | Epidemiology Program - Zoonotics | Renamed to "Joint aches/pains (Arthralgia)" to be more in line with case classification definition. | Yes |

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|---|---|---|---|---|---|
| 70 | MAEDSS-15358 | Retire SS_ARTHRALGIA symptom variable for multiple product codes | Epidemiology Program - Zoonotics | Retired and moved product codes to SS_JOINT_ACHES_PAINS. Joint aches/pains and arthralgia mean the same thing. | |
| 71 | MAEDSS-15895 | Update conditions in the EPI/IMM model for SS_SHOCK | Global | Fixed a modeling issue so this question shows up appropriately in Vibrio and Other product codes. | No |
| 72 | MAEDSS-15893 | Update CASE_INFO_REPORT_QUESTION_SETS reference code | Global | Updated the display order and added a new subgroup used to populate the question package report parameter in a new Cluster report (MAEDSS-15357). | No |
| 73 | MAEDSS-15805 | Delete Unknown events that have ONLY ALT and/or AST laboratory results | Global | Unknown events created between 2019 and 2022 with only ALT and/or AST results included are being deleted from MAVEN | No |
| 74 | MAEDSS-15486 | Tribal Affiliation Case Mapping | Global | A case mapping to copy Tribal affiliation date from Party Tab to Demographic QP. | No |
| 75 | MAEDSS-15480 | Change default collapsing of DRUGS_PATIENT_ON repeatable block | Global | Iterations will appear upon opening the question package instead of being collapsed and requiring expansion. | Yes |
| 76 | MAEDSS-15421 | Update answer selection for Race in all models to include Middle East and North Africa (MENA) | Global | Adding Middle East and North Africa (MENA) to all race variable options within MAVEN. Calculated Race will also be updated to reflect this addition and update the Calculated Race value accordingly. | Yes |
| 77 | MAEDSS-15414 | Update hide expressions for Sexual Orientation Gender Identity (SOGI) variables | Global | To better follow MA DPH Sexual Orientation Gender Identity (SOGI) variable data standards a few variables were updated. For individuals that are less than 5 years of age, Current Gender and Transgender Experience will be hidden. For individuals that are less than 12 years of age, Sexual Orientation will be hidden. | Yes |
| 78 | MAEDSS-15357 | Creating new report: Disease Event Cluster/Outbreak Linelist | Global | A new report that generates all the linked events for a particular cluster event in a linelist allowing to extract the question packages in each linked event rather than QP of the cluster itself. This report would be available for all product codes in the EPI/IMM, STD and TB model and can be selected from the Disease dropdown in the parameters that select the report. | No |
| 79 | MAEDSS-15221 | Create a new report: Model Print by Product Code | Global | Report would produce similar output to existing Model Print report with an additional report parameter for product code. | No |
| 80 | MAEDSS-15193 | Create a new variable CASE_HEALTHCARE_WORKER_NEW with child question CASE_HEALTHCARE_WORKER_VHF_NEW | Global | In October 2024 it was noticed that an existing QID: CASE_HEALTHCARE_WORKER had it's message changed during the pandemic from "Is the case a healthcare worker?" to "Employed at the facility above". Multiple product codes were impacted from this change. The original question will be hidden for impacted product codes (See Jira 15192) and this new question is being added. We will use a roster to add any existing data from the old question. | No |
| 81 | MAEDSS-15192 | Hide CASE_HEALTHCARE_WORKER for multiple product codes and CASE_HEALTHCARE_WORKER_VHF for VHF product code | Global | See MAEDSS-15193. | No |
| 82 | MAEDSS-15093 | MAEDSSDateUtils.getDateDiffInDays Is implemented in such a way to return 0 when there is exception (Not Valid Implementation) | Global | Update back end MAVEN calculation process that ensures date difference calculations treat equal dates (datediff = 0) correctly | No |
| 83 | MAEDSS-14600 | Create Tribal Affiliation Rule Output | Global | A rule that will use the Tribal Affiliation variable to assign jurisdiction to events that are of members of the two federally recognized Tribes in Massachusetts, Mashpee Wampanoag and Wampanoag Tribe of Gay Head (Aquinnah). | No |
| 84 | MAEDSS-14599 | Add Tribal Affiliation variable in Demographic QPs | Global | Variable that will house the Tribal affiliation data from the Party Tab. Will be hidden. | No |
| 85 | MAEDSS-14598 | Add Tribal Affiliation variable in Party Tab | Global | Adding a Tribal affiliation variable to the Party Tab. Will be hidden. | No |
| 86 | MAEDSS-14421 | Update custom code in the XSL processor to move data coming in HL7 OBR13 and/or OBR31 segments to populate product codes beyond Hepatitis B | Global | Allows Relevant Clinical Information and Reason of Study information in an HL7 message to populate new fields for all product codes in the EXTERNAL_INFORMATION question package. | Yes |
| 87 | MAEDSS-15783 | HIV - Reprocess HIV cases | HIV Program | Reprocess HIV cases in QA and Prod. | No |
| 88 | MAEDSS-15660 | Add two new subgroups to existing POSITIVE_STI_RESULTS reference code to inform hiv.confirmatory case attribute [HIVCONFIRM subgroup] and COPHI_STATUS [HIV2 subgroup] | HIV Program | Update the POSITIVE_STI_RESULTS reference code so it can be used for the HIV confirmatory case attribute, HIV COPHI events, and the event date hierarchy. | No |
| 89 | MAEDSS-15590 | Update code logic to set COPHI_STATUS to 1 if someone is HIV-2 positive | HIV Program | Update the logic of the COPHI_STATUS variable so that cases with HIV-2 detected flow into the appropriate HIV Suspect COPHI case workflow for investigation. | No |
| 90 | MAEDSS-15589 | Update existing logic for hiv.confirmatory case attribute | HIV Program | Update the HIV confirmatory case attribute to correct the LOINC/SNOMED combinations used in the criteria for several workflows. | No |
| 91 | MAEDSS-15498 | HIV - Update HIV CRF QP | HIV Program | Add additional questions/ clarify language in the HIV CRF QP which are present in the paper form. | No |
| 92 | MAEDSS-15378 | [HIV report] - Not HIV for Review report | HIV Program | The 'Not HIV' report will help the HIV team identify suspect and contact events that are not true HIV events so they can be revoked and closed. These events must meet all of the following criteria: Positive Ag/Ab test, and Negative or not detected HIV NAT, and Negative or indeterminate type-differentiating immunoassay. | No |
| 93 | MAEDSS-15361 | [HIV Wizard] - Grey out HIV CRF wizard | HIV Program | The HIV CRF wizard has been replaced by the HIV CRF QP. This wizard has been updated to update restriction level 5 to prevent entry into the wizard. | No |
| 94 | MAEDSS-15233 | HIV - Update the HIV - HIV/AIDS needs complete CRF workflow | HIV Program | Update the entry criteria, description, and workflow page so this workflow aligns with 08. HIV Additional CRF and Lab Information QP. | No |
| 95 | MAEDSS-15227 | HIV- update workflow: HIV - Attached CRF Ready for Review | HIV Program | Update the entry criteria, description, and workflow page so this workflow aligns with 08. HIV Additional CRF and Lab Information QP. | No |
| 96 | MAEDSS-14228 | [HIV report] Modify report: Lab Line List - HIV | HIV Program | Adds a new column to the existing "Lab Line List - HIV" report to pull Electronic Laboratory Data Source to distinguish between labs sent via ESP and ELR. | No |
| 97 | MAEDSS-15851 | [HIV Report] - Create a Genotype report for eHARS transfer | HIV Program, Next Genome Sequencing | Create a report to extract events that have recently received genotype sequencing results. These results will then be transferred to eHARS. | No |

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|---|---|---|---|---|---|
| 98 | MAEDSS-15813 | [HIV Genotype] - Changes to XSL processor for HIV sequencing results | HIV Program, Next Genome Sequencing | XSL updates are required to map HIV sequencing LOINCs to a single SNOMED and populate the new Sequencing Information fields in the Labs tab, including the Sequence Test and full Sequence result. | No |
| 99 | MAEDSS-14589 | [HIV - Genotype] - Changes to Labs Tab to add new fields for HIV sequencing | HIV Program, Next Genome Sequencing | Create two new fields ("Sequence Test" and "Sequence") added to the Sequencing Information section of the Labs tab. These will be read-only fields. | No |
| 100 | MAEDSS-15429 | Disable INTERSTATE_REPORTING DNT and associated processing modules | Interstate | This DNT was replaced by another that includes susceptibilities and is no longer needed. | No |
| 101 | MAEDSS-15295 | Exclude SARS from the Cases where official address is out of state Workflow | Interstate | COVID19 was removed from the national reporting list as of 2025. The program as well as senior management agreed to treat out of state notifications for COVID19 in a similar fashion to influenza and clostridium difficile, both of which are not nationally notifiable but still reportable in MA. They are sent out only to states that have been onboarded to HL7. | No |
| 102 | MAEDSS-15343 | Create a Bulk Action Button for LBOH Shared Cases Workflows | Local Boards of Health | Create a bulk action button to allow removal of Shared Cases with me and Shared Cases by me workflows | Yes |
| 103 | MAEDSS-15457 | Create report that capture Refugee events that have a TB co-infection but "IGRA Done" is blank | Refugee Immigrant Health Program | Create report that will capture all Refugee events that have a linked TB event (in cross-product notifier) but the "IGRA Done" variable in the Health Assessment QP is blank. | No |
| 104 | MAEDSS-15456 | Add existing QID to "RIHP Follow-up Wizard" | Refugee Immigrant Health Program | Adding "First scheduled clinic visit date" to RIHP Follow-up Wizard | No |
| 105 | MAEDSS-15455 | Update "Pending Post RHA Home Visit" workflow | Refugee Immigrant Health Program | Removes Status, Assigned To, Assigned To Group columns from Pending Post RHA Home Visit workflow. Adds Post RHA Visit CHW Name and Post RHA assigned date columns. | No |
| 106 | MAEDSS-15755 | STD - Update Patient Summary and OOS Case Report (ESP Treatments table) | Sexual Transmitted Disease Program | Update OOS report to remove "ESP TREATMENTS" as these are treatments that have been ordered and have not been verified. | No |
| 107 | MAEDSS-15497 | STD- Risk QP & Field Investigation Wizard- Update variables | Sexual Transmitted Disease Program | Update answer options in Risk History QP and Field Investigation QP to make language more clear/ add additional answer options. | No |
| 108 | MAEDSS-15477 | STD- Add variables to Interview Record Summary Report | Sexual Transmitted Disease Program | Add columns to existing report to pull additional interview variables. | No |
| 109 | MAEDSS-15422 | Update CALCULATED_RACE reference code to reorganize order and add Middle East and North Africa (MENA) | Sexual Transmitted Disease Program | Adding MENA to dropdown option for HIV CRF QP | No |
| 110 | MAEDSS-15402 | Exception while Importing some of the Rosters | Sexual Transmitted Disease Program | Fix the import rosters for the following rosters: HIVLabNameImport; GC/CT labs; HIV Roster MB; HIV Roster MB2 | No |
| 111 | MAEDSS-15883 | TB - Remove options from UNDERLYING_ILLNESS Ref List | Tuberculosis | Removing Diabetes (Type II), Insulin Dependent Diabetes Melitus, TNF Blocking Agents, Organ Transplant, Chronic/Renal Disease/Hemodialysis, Immunosuppression (not HIV/AIDS) from UNDERLYING_ILLNESS ref list filter for TB. These options are being broken out into their own separate QIDs (see MAEDSS-15494 for details). | Yes |
| 112 | MAEDSS-15801 | TB - Create a reference code list containing the L/S combinations that will set the DISEASE_STATE variable in TB Events | Tuberculosis | There is custom code on the back-end of MAVEN that looks at certain L/S combinations and will update DISEASE_STATE to TB_DISEASE. Instead of these L/S combinations being hard coded, a ref code list is being created instead. | No |
| 113 | MAEDSS-15800 | TB - Update custom code to set DISEASE_STATE variable using a reference code list | Tuberculosis | Updates custom code that sets DISEASE_STATE variable to draw from ref code list created in MAEDSS-15801 instead of using hard-coded combinations | No |
| 114 | MAEDSS-15728 | Please update TB events before L Release to address scenarios where VISIT_DATE was populated but VISIT_CLINIC was not | Tuberculosis | This JIRA is a companion to MAEDSS-15464; it will fix cases where the new parent question (which was previously a child question) will populate with a value if there wasn't one before. | No |
| 115 | MAEDSS-15727 | TB - Substitute VISIT_DATE with VISIT_CLINIC in certain TB Wizards | Tuberculosis | Substitutes in the new parent question "VISIT_CLINIC" in wizards where it was previously "VISIT_DATE" | Yes |
| 116 | MAEDSS-15643 | TB - Modify MDR question for CDC reporting | Tuberculosis | The funding formula for MDR was recently changed by the CDC; this change updates the existing MDR QID to reflect the new definition | Yes |
| 117 | MAEDSS-15507 | Create new QID that calculates date 8 weeks from exposure date in TB model | Tuberculosis | Creates POST_EXPOSURE_END_DATE variable which is an autocalculated/autopopulated variable that will take INFECTIOUS_PERIOD_STOP_DATE as input and calculate 8 weeks from that date. | Yes |
| 118 | MAEDSS-15506 | Create new "Contact to Active Cases" report for TB | Tuberculosis | Report that captures events linked to TB_DISEASE events in a given timeframe. If the linked event is a cluster event, it will populate the event IDs linked to that cluster, instead of the cluster ID itself. | No |
| 119 | MAEDSS-15496 | Modify PREVIOUS_HISTORY_OF_TB_DISEASE and PREVIOUS_HISTORY_OF_LTBI_INFECTION questions in TB model | Tuberculosis | Creates new variables to better capture date of previous history of TB Disease/TB Infection events for a given case. Replaces variable that only captured the previous year of TB Disease (which did not meet CDC reporting requirements). | Yes |
| 120 | MAEDSS-15494 | Modify UNDERLYING_ILLNESS question in TB model | Tuberculosis | Removing Diabetes (Type II), Insulin Dependent Diabetes Melitus, TNF Blocking Agents, Organ Transplant, Chronic/Renal Disease/Hemodialysis, Immunosuppression (not HIV/AIDS) options in UNDERLYING_ILLNESS_SPECIFY dropdown. These questions are being broken out into their own on QIDs for improved CDC reporting. | Yes |
| 121 | MAEDSS-15464 | Make VISIT_DATE a child question to VISIT_CLINIC in TB model | Tuberculosis | Currently, VISIT_CLINIC is a child question for VISIT_DATE. The program would like this flipped for ease of review. For scenarios where VISIT_CLINIC was blank but a VISIT_DATE value is available, 'Historical Clinic' will be populated so that the VISIT_DATE value is visible in the question package. | Yes |
| 122 | MAEDSS-15463 | Create Concerns for TB model | Tuberculosis | Four new Concerns are being added for TB events. The first will trigger if the patient moved to MA from another jurisdiction. The second will trigger if the patient moved out of state but was previously being treated in MA. The third will trigger if the case was not confirmed in MA, but follow-up only on RVCT is still required (this is only relevant to MDPH TB staff). The fourth will trigger if the patient is a recent arrival to the US with any Class A/B designation-- the particular Class A/B designation will be stated in the text of the Concern. | Yes |
| 123 | MAEDSS-15462 | Modify RVCT Wizard to add/remove/edit QIDs | Tuberculosis | Modifies "Report of Verified Case of Tuberculosis" Wizard to reflect CDC reporting requirements | Yes |
| 124 | MAEDSS-15460 | Add child question "IGRA done overseas" to "IGRA done" variable | Tuberculosis | Adds new child question "IGRA done overseas" to the "IGRA done" variable in the Medical Information QP. This question has Yes/No/Unknown answer options. | Yes |

| | JIRA | Title | Component/s | Release Note Description | Relevant to Local Health Departments? Yes or No |
|---|---|---|---|---|---|
| 125 | MAEDSS-15459 | Add new dropdown options for "Reason for Referral?" | Tuberculosis | Adds the following dropdown options to the "Reason for Referral" question: LTBI - Refugee, LTBI - Contact, LTBI - B1, B2<19. | Yes |
| 126 | MAEDSS-15531 | Add variables to the TB extract (L Release) | Tuberculosis | Updates the TB NMI extract to capture model/other changes in L release that impacts the extract and CDC reporting. | No |
| 127 | MAEDSS-15491 | Sort repeatable date fields in ascending order in TB/Refugee models | Tuberculosis, Refugee Immigrant Health Program | Sorts certain date fields in TB/Refugee models in ascending order (oldest to newest). | Yes |