M2 Direct Care Laboratory Results for the MHS Mart (M2)

(Version 1.00.09)

Future Specification

Revision History

| Version | Date | Originator | Para/Tbl/Fig | Description of Change |
|---------|--------------|------------|-------------------|--|
| 1.00.00 | 04/28/2021 | S. Keane | | Original. |
| 1.00.01 | 06/10/2021 | S. Keane | Entire Document | Update based on functional feedback. |
| | | | | Remove Certifying Provider SSN (Pseudo) |
| | | | | and Ordering Provider SSN (Pseudo). |
| 1.00.02 | 06/30/2021 | S. Keane | Table 3 and | Add descriptions for HIPAA Taxonomy |
| | | | Section VII | Codes, Enrollment Groups, and Eligibility |
| | | | | Groups. |
| | | | | Add Market Transition Date. |
| 1.00.03 | 06/03/2024 | S. Keane | Table 3 | Update M2 Location. Remove Market Transition Date |
| 1.00.03 | | S. Keane | Entire Document | |
| 1.00.04 | 8/01/2024 | S. Keane | Entire Document | Remove Table 2. Update the layout being sent from MDR to M2. |
| 1.00.05 | 8/14/2024 | S. Keane | Table 3 | Removed: MTF Service Area Base Market |
| 1.00.03 | 0,14,2024 | J. Realie | Table 5 | Code, MTF Service Area Expanded Market |
| | | | | Code, MTF Service Area Reporting Market |
| | | | | Code, MTF Service Area Type of Market |
| | | | | Code. |
| | | | | |
| 1.00.06 | 9/10/2024 | S. Keane | Table 1 and Table | Add Sponsor Service and Prime Service |
| | | | 2. | Area. Specify Integer for Line Number. |
| | | | | Convert Genesis Flag to 1/0. Rename Age |
| | | | | Group Common. Remove Single quote |
| | | | | from Result Text and Result Numeric. |
| | | | | Remove Ordering Site Parent. Remove |
| | | | | text "M2 Derived". Rename Table 3 to 2. |
| 1.00.07 | 9/26/2024 | S. Keane | Table 1 and Table | Remove PowerInsight Encounter ID GENESIS. |
| | | | 2 | Update business rules for Line Number, Number of Tests and Number of Results |
| | | | | Maximum. |
| 1.00.08 | 10/2/2024 | S. Keane | Table 1 | Update derivation of Result Abnormal and |
| 2.00.00 | 30, 2, 202 : | | | Result Abnormal Flag. |
| 1.00.09 | 10/15/2024 | S. Keane | Table 1 | Move Number of Results Maximum.to end of |
| | | | | MDR Feed. |

M2 Direct Care Laboratory Results

- **Source:** The most recent MDR CADRE Laboratory Enhanced Results and MDR Genesis Laboratory Results files are the sources. The MDR will merge the two source files into a single feed for M2.
- II. File Format: Regardless of the length specified, all fields are variable length and delimited by "|".
- III. Record Selection: Only FY2019 and forward will be sent to M2. Apply the following filter: include records where LABVALUE (CHCS CADRE) or RESULT_VALUE (MHS GENESIS) is populated. The record key is the Record ID (M2KEY) which is a sequentially assigned number, encoded to base 62 to save space, and continues across Fiscal Years.

For lab test with more than one result, the combination of the following fields will be used to distinguish individual results. Results are ordered by:

(CHCS CADRE): CHCSDMIS ACCESSNO ORDERID LABTEST RECNO

(MHS GENESIS): ENCOUNTER_SK ORDER_SK ACCESSION ORDERABLE_DISP EVENT_SK

MDR will create a Line Number (ROWNUM) for every record with this combination.

Each update will consist of full file refresh. As such, M2 shall purge/remove existing records PRIOR to loading the current/new full file refresh. The load file contains records to be added to the M2 Direct Care Laboratory Results Table. See Table 1 in Section V below.

IV. Refresh Batches: Batches are by Fiscal Year. The Fiscal Year is included in the feed and can be determined by the Date of Service (SERVDATE) field.

Frequency of updates:

- Current Fiscal Year: Weekly.
- Prior Fiscal Year: Semi-annually.
- All Years Prior to Prior Fiscal Year: Annually (October).

Any year refreshed in the MDR should automatically trigger a refresh of the same year's table in the MHS Mart (M2).

V. Table 1. Full File Refresh Layout of MDR Feed to M2: Table 1 includes all fields received or derived in the MDR that are fed to M2. Fields are listed alphabetically by M2 Object Name (after Record ID)

Obtain from M2 CADRE Lab Orders when feasible. Otherwise, obtain from specified source. Sources: CLABE = CADRE LAB Enhanced, CLABR = CADRE Lab Results, GLABO = GENESIS LAB Orders, GLABR = GENESIS LAB Results.

Table 1. MDR Feed to M2

| M2 Object | Format | MDR CADRE LAB SAS Name | MDR GENESIS SAS Name | Business Rule | In M2 CADRE LAB Orders |
|---------------------------------------|----------|---------------------------|-------------------------|--|---------------------------------|
| Record ID | Char(6) | M2KEY | M2KEY | CLABE: none. GLABO: none. | YES |
| Accession Number | Char(20) | ACCESSNO | ACCESSION | CLABE: Set length to 20. GLABR: Set Length to 20. | YES |
| Age | Num(8,0) | PATAGE | PATAGE | CLABE: none. GLABR: none. | YES |
| Age Group Common | Char(1) | AGEGROUP | AGEGRP | CLABE: none. GLABR: none. | YES |
| Appointment Record ID | Char(10) | APPTNO | ENCOUNTER_SK | CHCS: Remove leading and trailing blanks. GENESIS: First 10 digits of ENCOUNTER_SK where RECORD_TYPE_FLAG = "ENCOUNTER RAD". | YES |
| Ben Cat Common | Char(1) | COMBEN | COMBEN_D | CLABE: none. GLABR: none. | YES |
| Beneficiary Category | Char(3) | BENCAT | BENCAT_D | CLABE: none. GLABR: none. | YES |
| Beneficiary First Name | Char(20) | EDIPN | EDIPN | CHCS and GENESIS: Join to MPI Table on UPID. Only visible in the restricted universe. | YES |
| Beneficiary Last Name | Char(26) | EDIPN | EDIPN | CHCS and GENESIS: Join to MPI Table on UPID. Only visible in the restricted universe. | YES |
| Beneficiary Name | Char(70) | EDIPN | EDIPN | CHCS and GENESIS: Join to MPI Table on UPID. Display as LAST, FIRST Middle Initial. Only visible in the restricted universe. | YES |
| Beneficiary Zip Code | Char(5) | PATZIP | ZIP_D | CLABE: none. GLABR: none. | YES |
| Certifying Provider EDIPN | Char(10) | CPEDIPN | PRSNL_COMPL_E DIPN | CLABE: none. GLABR: none. Only visible in the restricted universe. | YES |
| Certifying Provider HIPAA Taxonomy | Char(10) | СРНІРАА | PRSNL_COMPL_H IPAA | CLABE: none. GLABR: none. | YES |
| Certifying Provider Name | Char(61) | CPNAME | PRSNL_COMPL_N AME | CLABE: none. GLABR: none. | YES |
| Certifying Provider NPI | Char(10) | CPNPI | PRSNL_COMPL_N PI | CLABE: none. GLABR: none. | YES |
| Critical High GENESIS | Char(20) | N/A | CRITICAL_HIGH | CLABR: Set to Null. GLABR: none. | NO |
| Critical Low GENESIS | Char(20) | N/A | CRITICAL_LOW | CLABR: Set to Null. GLABR: none. | NO |
| Date of Collection/Exam | yyyymmdd | EXAMDATE | DATE_COMPLETE | CLABE: none. GLABR: Convert to yymmddn8. | YES |
| Date of Order | yyyymmdd | ORDDATE | ORDER_DT_TM_L OCAL | CLABE: none. GLABR: Convert to yymmddn8. | YES |

| M2 Object | Format | MDR CADRE LAB SAS Name | MDR GENESIS SAS Name | Business Rule | In M2 CADRE LAB Orders |
|---------------------------------|-----------|---------------------------|-------------------------|---|---------------------------------|
| Diagnosis Code 1 | Char(10) | N/A | DX_CCE_1 | CLABE: not applicable. GLABR: Use the CCE Diagnosis Code 1 (DX_CCE_1). Remove any decimal points. | NO |
| Diagnosis Code 2 | Char(10) | N/A | DX_CCE_2 | CLABE: not applicable. GLABR: Use the CCE Diagnosis Code 2 (DX_CCE_2). Remove any decimal points. | NO |
| Diagnosis Code 3 | Char(10) | N/A | DX_CCE_3 | CLABE: not applicable. GLABR: Use the CCE Diagnosis Code 3 (DX_CCE_3). Remove any decimal points. | NO |
| Eligibility Group | Char(1) | ELG_GRP | ELG_GRP | CLABE: none. GLABR: none. | YES |
| Enrollment Group | Char(1) | ENR_GRP | ENR_GRP | CLABE: none. GLABR: none. | YES |
| FIN GENESIS | Char(40) | N/A | FIN_E | CLABR: Set to Null. GLABR: Set length to 40. | YES |
| Fiscal Month | Char(2) | FM | FM | CLABE: none. GLABR: none. | YES |
| Fiscal Year | Char(4) | FY | FY | CLABE: none. GLABR: none. | YES |
| Inpatient Record ID | Char(10) | PRN | n/a | CHCS: Remove leading and trailing blanks. Length= 7. GENESIS: First 10 digits of ENCOUNTER_SK where INPT = 'Y'. Length=10. | YES |
| Lab Test Name | Char(100) | LABTEST | ORDERABLE_DISP | CLABR: Fill Down. GLABR: ORDERABLE_DISP. | YES |
| Line Number | Integer | Derive: ROWNUM | Derive: ROWNUM | Group By: chcs clabe: chcsdmis, orderid, accessno, labtest mhs genesis glabr: encounter_sk, order_sk, accession, orderable_disp Each unique combination forms a group. Order By: CHCS CLABE: chcsdmis, orderid, accessno, labtest, recno MHS GENESIS GLABR: encounter_sk, order_sk, accession, orderable_disp, event_sk Line_Number Assignment: Sequentially based on recno for CHCS Sequentially based on event_sk for GENESIS | NO |
| MHS GENESIS Flag | Char(1) | N/A | N/A | CLABR: "0". GLABR: "1". M2 should change 1/0 to Yes/No | YES |
| Millennium Encounter ID GENESIS | Char(22) | N/A | ENCOUNTER_SK | CLABR: Set to Null. GLABR: Set length to 22. | YES |
| MRN GENESIS | Char(40) | N/A | MRN | CLABR: Set to Null. GLABR: Set length to 40. Only visible in the restricted universe. | YES |
| Normal High GENESIS | Char(20) | N/A | NORMAL_HIGH | CLABR: Set to Null. GLABR: none. | NO |
| Normal Low GENESIS | Char(20) | N/A | NORMAL_LOW | CLABR: Set to Null. GLABR: none. | NO |

| M2 Object | Format | MDR CADRE LAB SAS Name | MDR GENESIS SAS Name | Business Rule | In M2 CADRE LAB Orders |
|-------------------------------------|----------|------------------------------|------------------------------|---|---------------------------------|
| Order ID | Char(12) | ORDERID | ORDER_SK | CLABE: none. GLABR: none. | YES |
| Ordering Provider EDIPN | Char(10) | OPEDIPN | PROV_ORD_EDIP N | CLABE: none. GLABR: none. Only visible in the restricted universe. | YES |
| Ordering Provider HIPAA Taxonomy | Char(10) | ОРНІРАА | PROV_ORD_HIPA A | CLABE: none. GLABR: none. | YES |
| Ordering Provider Name | Char(61) | OPNAME | PROV_ORD_NAM E | CLABE: none. GLABO: none. | YES |
| Ordering Provider NPI | Char(10) | OPNPI | PROV_ORD_NPI | CLABE: none. GLABR: none. | YES |
| Ordering Site of Record | Char(4) | ORDDMIS | MTF | CLABE: none. GLABR: none. | YES |
| PATCAT Subcategory | Char(1) | PATCAT2 | N/A | CLABE: none. GLABR: none. | YES |
| Patient Code Category | Char(3) | PATCAT | PATCAT | CLABE: none. GLABR: none. | YES |
| Patient SSN | Char(9) | PATSSN | SSN | CLABE: none. GLABR: none. Only visible in the restricted universe. | YES |
| Person ID | Char(10) | EDIPN | EDIPN | CLABE: none. GLABR: none. Only visible in the restricted universe. | YES |
| Pseudo MRN GENESIS | Char(12) | N/A | MRN | CLABR: Set to Null. GLABR: Encryption algorithm applied to Medical Record Number (MRN). | YES |
| Result Abnormal | CHAR(80) | Derive: ABNORMAL | Derive: ABNORMAL | CLABR: none. GLABR: IF ABNORMAL_FLAG in ('Abnormal', 'Critical', 'High', 'Low') then ABNORMAL = substr(ABNORMAL_FLAG, 1, 1) "" "" trim(NRANGE) "" "" trim(CRANGE) "" "" trim(RESULTS_UNITS); else ABNORMAL = "; | NO |
| Result Abnormal Flag | Char(14) | Derive: ABNORMAL_FLA G | Derive: ABNORMAL_FLA G | CLABR: if substr(abnormal, 1, 1) = "L" then abnormal_flag= "LOW"; else if substr(abnormal, 1, 1) = "H" then abnormal_flag = "HIGH"; else abnormal_flag = ""; GLABR: trim(upper(EVENT_NORMACY)); | NO |

| M2 Object | Format | MDR CADRE LAB SAS Name | MDR GENESIS SAS Name | Business Rule | In M2 CADRE LAB Orders |
|------------------------------------|-----------|---------------------------|--------------------------|--|---------------------------------|
| Result LOINC Code | Char(10) | LOINC | LOINC_CODE | CLABR: Set length to 10. GLABR: Set length to 10. | NO |
| Result LOINC Name GENESIS | Char(100) | N/A | LOINC_DESCRIPTI ON | CLABR: Set to Null. GLABR: Set length to 100. | NO |
| Result Name | Char(60) | LABTESTR | EVENT_CODE | CLABR: obtain LABTESTR when labtestr not missing, merge on chcsdmis, accessno, recno. If missing(labtestr) then delete. GLABR: Obtain EVENT_CODE as labtestr, merge on encounter_key, accession, order_sk, orderable_disp. | YES |
| Result Numeric | 17.4 | LABVALUE | RESULT_VALUE | CLABR: If LABVALUE is numeric, else missing. GLABR: if RESULT_VALUE is numeric, else blank. Remove single quote, if applicable. Use value as received from MDR. For example, 1.5000 should display as 1.5000 and 1.5 should display as 1.5 | YES |
| Result Text | Char(30) | LABVALUE | RESULT_VALUE | CLABR: if LABVALUE is text, else blank. GLABR: if RESULT_VALUE is text, else blank. Remove single quote. | YES |
| Result Units of Measure GENESIS | Char(60) | N/A | RESULTS_UNITS | CLABR: Set to Null. GLABR: none. | NO |
| Result Value | Char(30) | LABVALUE | RESULT_VALUE | CLABR: none. GLABR: Set length to 30 | NO |
| Service Date | yyyymmdd | SERVDATE | SERVICE_DT_TM_ LOCAL | CLABE: none. GLABO: Convert to yymmddn8. | YES |
| Specimen GENESIS | Char(38) | N/A | SPECIMEN | CLABR: Set to Null. GLABR: none. | NO |
| Sponsor ID | Char(9) | SPONSSN | SPONSSN | CLABR: none. GLABR: none. Only visible in the restricted universe. | YES |
| Sponsor Service | Char(1) | SVC | SPONSVC | CLABE: none. GLABO: none. | YES |
| Tmt DMIS ID | Char(4) | TMTDMIS | MTF_RESULTED | CLABE: none. GLABO: none. | YES |
| Tmt Parent DMIS ID | Char(4) | PDMIS | MTF_PARENT_RE C | CLABE: none. GLABR: none. | YES |
| Verified Date | yyyymmdd | CERTDATE | VERIFIED_DT_TM _LOCAL | CLABR: None. GLABR: Convert to yymmddn8. | NO |
| Number of Results Maximum | Num(9,0) | ACCESSNO | ACCESSION | Count number of rows per Accession Number. In other words, return maximum Line Number for each Accession Number. | NO |

M2 Fields: Table 2 includes M2 fields; these are fields that are not in the MDR feed but are visible in M2. These data fields are created in M2 by table joins or calculations made during loading of the data. Table 2 describes these elements and the rules to create them.

Table 2. M2 Fields

| M2 Object | Format | Business Rule |
|--|----------|---|
| Ben Cat Common Des | Char(28) | Based on Ben Cat Common: |
| | | 1 = AD/Guard Family Member |
| | | 2 = Retirees |
| | | 3 = Other |
| | | 4 = AD/Guard |
| Beneficiary Category Desc | Char(30) | ACT=Active Duty |
| | | DA=Active Duty Family |
| | | DCO=Direct Care Only |
| | | DGR=Dep of Guard/Res on AD |
| | | DR=Retiree Family |
| | | DS=Survivor |
| | | GRD=Guard/Res on AD |
| | | IDG=Inactive Guard/Res Family |
| | | IGR=Inactive Guard/Res |
| | | NAT=NATO |
| | | OTH=Other |
| | | RET=Retirees |
| | | UNK=Unknown |
| | | Z=Unknown |
| Certifying Provider HIPAA Taxonomy | Char(50) | Lookup Certifying Provider HIPAA Taxonomy with the Health Care |
| Desc | | Provider Taxonomy Reference Table and return the Description, |
| | | Short. |
| | | C = Continued Health Care Benefit Program (CHCBP) |
| | | D = Direct Care Only |
| | | E = TRICARE Eligible |
| | CI (45) | L = TRICARE for Life (TFL) |
| Eligibility Group Description | Char(45) | R = TRICARE Retired Reserve (TRR) |
| | | S = TRICARE Reserve Select (TRS) |
| | | Y = TRICARE Young Adult (TYA) |
| | | Z = None |
| | | P = TRICARE Prime |
| | | D = Direct Care Only |
| | | L = TRICARE Plus |
| Enrollment Group Description | Char(27) | S = TRICARE Select |
| · | | U = Designated Provider (USFHP) |
| | | Z = Not Enrolled |
| | | Blank = pre-2018 |
| Fiscal Quarter | Char(1) | If FM in (1,2,3) then Fiscal Quarter=1; else if FM in (4,5,6) then Fiscal |
| | | Quarter=2; else if FM in (7,8,9) then Fiscal Quarter = 3; else Fiscal |
| | | Quarter = 4. 1 = FM 1-3 (Oct-Dec) 2 = FM 4-6 (Jan-Mar) 3 = FM 7- |
| | | 9 (Apr-Jun) 4 = FM 10-12 (Jul-Sep) |
| Market Area ID | Char(4) | Join to the Market Table by FY, FM, Beneficiary Zip Code, and |
| | | Sponsor Service. |
| MTF Service Area | Char(4) | Join to the Market Table by FY, FM, Beneficiary Zip Code, and |
| | (- / | Sponsor Service. |
| MTF Service Area Name | Char(40) | Join to the Market Table by FY, FM, Beneficiary Zip Code, and |
| The second secon | (10) | Sponsor Service. |
| | 1 | 1 -6 |

| M2 Object | Format | Business Rule |
|--|----------|---|
| Number of Records | Num(9,0) | Count numbers requested by query. This will be a measure. |
| Number of Tests | Num(9,0) | If Line Number equals 1, then 1, else 0. This will be a measure. |
| Ordering Provider HIPAA Taxonomy Desc | Char(50) | Lookup Ordering Provider HIPAA Taxonomy with the Health Care Provider Taxonomy Reference Table and return the Description, Short. |
| Ordering Site Financial Parent | Char(4) | See DMIS ID specification for rule. |
| Ordering Site Financial Parent Name | Char(40) | See DMIS ID specification for rule. |
| Ordering Site ID/Appoint Record ID | Char(14) | M2 User Defined Object (UDO) that is the concatenation of Ordering Site and Appointment Record ID. |
| Ordering Site ID/InPatient Record ID | Char(14) | M2 User Defined Object (UDO) that is the concatenation of Ordering Site and Inpatient Record ID. |
| Ordering Site of Record Base Market Code | Char(10) | Join to the DMIS ID Table by Ordering Site DMIS ID and FY. Populated FY18+. |
| Ordering Site of Record Command | Char(6) | Join to the DMIS Table by FY and Ordering Site. |
| Ordering Site of Record DHP Code | Char(1) | Join to the DMIS Table by FY and Ordering Site of Record. |
| Ordering Site of Record Expanded Market Code | Char(10) | Join to the DMIS ID Table by Ordering Site DMIS ID and FY. Populated FY18+. |
| Ordering Site of Record Market Name | Char(40) | Join to the DMIS ID Table by Ordering Site DMIS ID and FY. Populated FY18+. |
| Ordering Site of Record Mil Svc | Char(1) | Join to the DMIS Table by FY and Ordering Site of Record. |
| Ordering Site of Record MSMA | Char(3) | Join to the DMIS Table by FY and Ordering Site of Record. |
| Ordering Site of Record Name | Char(40) | Join to the DMIS Table by FY and Ordering Site of Record. |
| Ordering Site of Record Readiness CMD | Char(8) | See DMIS ID specification for rule. |
| Ordering Site of Record Readiness Service | Char(1) | Join to the DMIS ID Table by Ordering Site DMIS ID and FY. Populated FY18+. |
| Ordering Site of Record Reporting Market Code | Char(10) | Join to the DMIS ID Table by Ordering Site DMIS ID and FY. Populated FY18+. |
| Ordering Site of Record Reporting Market Name | Char(40) | Join to the DMIS ID Table by Ordering Site DMIS ID and FY. Populated FY18+. |
| Ordering Site of Record Type of Market Code | Char(10) | Join to the DMIS ID Table by Ordering Site DMIS ID and FY. Populated FY18+. |
| Ordering Site of Record Type of Market Name | Char(40) | Join to the DMIS ID Table by Ordering Site DMIS ID and FY. Populated FY18+. |
| Prime Service Area | Char(1) | Populated as part of the ETL process. |
| Pseudo Patient SSN | Char(9) | Encryption algorithm applied to Patient SSN by the ETL process. |
| Pseudo Person ID | Char(10) | Encryption algorithm applied to Person ID by the ETL process. |
| Pseudo Sponsor ID | Char(9) | Encryption algorithm applied to Sponsor ID by the ETL process. |

| M2 Object | Format | Business Rule | | |
|-----------------------------------|----------|---|--|--|
| Tmt DMIS ID Base Market Code | Char(10) | Join to DMIS ID Table by TMT DMIS ID and FY. Populated FY18+. | | |
| Tmt DMIS ID Command | Char(6) | Join to the DMIS Table by FY and Tmt DMIS ID. | | |
| Tmt DMIS ID DHP Code | Char(1) | Join to the DMIS Table by FY and Tmt DMIS ID. | | |
| Tmt DMIS ID Expanded Market Code | Char(10) | Join to DMIS ID Table by TMT DMIS ID and FY. Populated FY18+. | | |
| Tmt DMIS ID Financial Parent | Char(4) | See DMIS ID specification for rule. | | |
| Tmt DMIS ID Financial Parent Name | Char(40) | See DMIS ID specification for rule. | | |
| Tmt DMIS ID Market Name | Char(40) | Join to DMIS ID Table by TMT DMIS ID and FY. Populated FY18+. | | |
| Tmt DMIS ID MSMA | Char(3) | Join to the DMIS Table by FY and Tmt DMIS ID. | | |
| Tmt DMIS ID Name | Char(40) | Join to the DMIS Table by FY and Tmt DMIS ID. | | |
| Tmt DMIS ID Readiness CMD | Char(8) | See DMIS ID specification for rule. | | |
| Tmt DMIS ID Readiness Parent | Char (4) | Join to DMIS ID Table by TMT DMIS ID and FY. Populated FY18+. | | |
| Tmt DMIS ID Readiness Service | Char (1) | Join to DMIS ID Table by TMT DMIS ID and FY. Populated FY18+. | | |
| Tmt DMIS ID Reporting Market Code | Char(10) | Join to DMIS ID Table by TMT DMIS ID and FY. Populated FY18+. | | |
| Tmt DMIS ID Reporting Market Name | Char(40) | Join to DMIS ID Table by TMT DMIS ID and FY. Populated FY18+. | | |
| Tmt DMIS ID Type of Market Code | Char(10) | Join to DMIS ID Table by TMT DMIS ID and FY. Populated FY18+. | | |
| Tmt DMIS ID Type of Market Name | Char(40) | Join to DMIS ID Table by TMT DMIS ID and FY. Populated FY18+. | | |
| Tmt Parent DMIS ID Name | Char(40) | Join to the DMIS Table by FY and Tmt DMIS Parent. | | |

VI. M2 Location:

Health Care Services / Direct Care / Laboratory and Radiology / Laboratory Results.