Bovine Tuberculosis and Brucellosis Surveillance Results Monthly Reports, Federal Fiscal Year (FY) 2016 September 1-30, 2016

New Information – Bovine Tuberculosis (TB)

- A cow slaughtered was reported as histo compatible and PCR positive for M. tuberculosis
 complex presumably M. bovis. The cow was confirmed as Canadian origin by DNA matching of
 the identification devices and lesion tissue. Canada has been noitified of this case. Culture and
 WGS is pending.
- A 5-year old cow slaughtered was reported with a histo compatible lesion. PCR was negative for M. tuberculosis complex, M. avium and MAP. Culture is pending.

Update of Previously Reported Information

- Trace-outs from the small Michigan menagerie herd (consisting of a few steers, goats, pigs, and chickens) located in the Modified Accredited (MA) zone have been completed. There were no out-of-State traces associated with this herd and only one source trace to a premises that tested negative. The herd was detected in May 2015 on an annual whole-herd test (WHT) and was depopulated in August 2015. This is the 62nd TB-infected herd found in Michigan since 1998.
- The State of Michigan paid indemnity to depopulate the adult animals from the Michigan beef herd located in the MA zone. One additional infected animal was found. The feeders remained on the test and remove protocol until all feeders had moved to slaughter by the end of July 2016. The herd was detected in December on an annual WHT. This is the 63rd TB-infected herd found in Michigan since 1998.
- The infected Michigan beef herd located in the MAZ, which was detected during an annual WHT, is being managed under a test and removal herd plan. The first removal test found an additional animal with lesions and was confirmed to be infected with M. bovis. This is the 64th TB-infected herd found in Michigan since 1998.
- One new TB-infected herd was detected in a Michigan beef herd located in the MA zone through the epidemiological investigation performed on herd #64. This is the 65th TB-infected herd found in Michigan since 1998. Whole genome sequencing found the isolates from herds #64 and #65 were not closely related.
- Depopulation was completed on March 31, 2016, for the first Texas dairy quarantined in October 2014. The initial estimate within herd prevalence was 5.9 percent, with a herd size of over 10,000 cattle. Evidence of TB was found in 171 depopulated cattle over 6 months of age.
- The second Texas dairy, quarantined in October 2014, completed three whole-herd removal tests without evidence of TB. A total of three TB-infected cattle were associated with this herd on initial testing. A quarantine releasing test was completed in December and the dairy was released from quarantine on January 13, 2016. It will be retested on five yearly assurance tests.
- Two Texas organic dairies and the associated Texas feedlot and heifer raiser were quarantined on April 8, 2015. A total of 56 lesioned cattle have been removed from the dairy of approximately 11,000 cattle. In a WHT in March, 168 caudal-fold test positives were found and 134 continue awaiting slaughter and postmortem examination. The owner did not accept indemnity values presented, and steps to resolve perceived value disparity continue.
- A Holstein steer was found through routine slaughter in Arizona in January. The epidemiological investigation had mistakenly identified that the steer as originating from Arizona. It is now clear that the steer was fed in California. Lack of specific identification information on the steer has lead to multiple potential consignors in California. An investigation of its origin continues.
- The TB-infected herd detected in Indiana as a result of the epidemiological investigation of six TB-infected steers identified at slaughter in May 2016 has been depopulated.

Table 1. Bovine TB cases found through routine slaughter inspection, FY 2016.^a

| | New TB September | | Cumulative TB Cases October 1, 2015-September 30, 2016 | | | |
|--|---------------------|-----------------|---|--------------|-------|--|
| Laboratory Status | Fed cattle | Adult cattle | Fed cattle | Adult cattle | Total | |
| M. bovis cases, confirmed ^b | 0 | 1 | 13 | 1 | 14 | |
| PCR pending | 0 | 0 | 1 | 0 | 1 | |
| PCR negative, culture pending | 0 | 1 | 0 | 0 | 1 | |

^a Animals detected through routine slaughter inspection. Animals sent to slaughter with a 1-27 permit,

Table 2. Livestock herds confirmed infected with bovine TB, FY 2016, including test-and-remove managed herds under quarantine and carried forward from previous fiscal years.

| State | Federal Fiscal Year Detected | Method of Detection | Herd Type | Herd Manage- ment Plan | Closest Isolates by WGS (Number of SNP Differences) ^a | Wildlife Surveillance Planned? | Approximate Number of Animal Traces | States With Traced Cattle |
|-----------------|---------------------------------------|------------------------|--------------------|---|--|---|---|--|
| MI ^b | 2008 | Surveillance | Cervid | Permanent quarantine | MI | Ongoing | None | None |
| MI ^b | 2008 | Surveillance | Cervid | Permanent quarantine | MI | Ongoing | None | None |
| TX | 2015 | Slaughter | Dairy | Depopulation | 2004 TX (16), 2007 NM dairies (15), 2011 AZ roping steer from CHI (25) | Small survey completed with no evidence of TB | None | None |
| TX | 2015 | Epidemiology | Dairy | Test and remove | Same as above | Not indicated | 68 | MI |
| TX | 2015 | Slaughter | Dairy ^c | Test and remove | 1997 MX (8) | Small survey completed with no evidence of TB | 4,150 | AZ, CA, CO, MI, MO, MN, NM, IN, TX |
| MI | 2015 | Area Testing | Mixed | Depopulation | Common ancestor with MI free ranging deer (1 SNP) | Ongoing | 1 source | None |
| MI | 2016 | Area Testing | Beef | State funded Depopulation of adults, all feeders have been slaughtered | Similar to isolates from herd #54 (Alpena Dairy) and opossums on that premises. | Ongoing | 2 source, 4 traceouts | None |
| MI | 2016 | Area Testing | Beef | Test and Remove | Common ancestor with MI free- | Ongoing | Pending | Pending IL |

[&]quot;Permit for Movement of Restricted Animals" are not included.

^b Confirmed by PCR testing and/or culture.

| | | | | | ranging deer (3). Does not match any previously identified livestock WGS. | | | |
|----|------|----------------------------------|------|--------------------|---|----------|-----------------------|---------|
| MI | 2016 | Epidemiological Investigation | Beef | Test and Remove | Common ancestor with MI free- ranging deer (6). Same common ancestor with previous infection on this premises in 2003 (#27) | Ongoing | 9 source, 2 traceouts | None |
| IN | 2016 | Slaughter | Beef | Depopulation | 2009 Cervid (1) | Underway | 2 traceouts | None |
| MI | 2016 | Movement Test | Beef | Pending | Pending | Ongoing | Pending | Pending |

^aWGS = whole genome sequencing, SNP = single nucleotide polymorphisms; () = number of SNP differences to the most recent common ancestor compared to other isolates in the National Veterinary Services Laboratories database.

Brucellosis - Herds Remaining under Quarantine and Test-and-Remove Procedures or Awaiting Assurance Test Results

FY 2016

Wyoming

- Affected beef herd: Testing of a portion of a Park County (DSA) commercial cow-calf beef herd on October 28, 2015, revealed one reactor (an 4-year-old cow). The herd of 515 adults was placed under quarantine. Tissues from this cow cultured positive for *B. abortus* on November 19, 2015. Testing of the balance of the herd in early December revealed no nonnegative animals. All the adjacent/contact herds have been identified and have begun to be tested. The cow originated from Montana (non-DSA) and was shipped to this ranch in November 2013. The herd has completed its second negative WHT. An epidemiologically separate management unit of this producer (NorthUnit) will have an assurance test in the spring or next fall. All traces have been completed with no other positive animals found. Updates: The third herd test (post-calving test) conducted on July 11-12, 2016, revealed no test positives and subsequently the quarantine on the herd was lifted on July 18, 2016. An assurance test will be performed in approximately one year, likely during the 2017 fall pregnancy check.
- Affected beef herd: On November 9-11, 2015, testing of a portion of a Sublette County (DSA) commercial cow-calf beef herd revealed five reactor cows. The herd of 717 cattle was placed under quarantine on November 17, 2015. Tissues from several of these reactors cows cultured positive for *B. abortus* as reported on December 9, 2015. Testing of the balance of the herd in early December revealed six more reactor animals. All seven adjacent/contact

^bTwo hunt facility herds detected in previous fiscal years under permanent quarantine are included.

^cThis herd consists of two sister dairies and their associated common heifer raising/feedlot facility.

herds have been identified and have begun to be tested. All traces have been completed with no other positive animals found. The first negative whole herd test was completed in January 2016. The second whole herd test on March 14-15 revealed two more reactors. The whole herd test conducted May 21-22 revealed two more reactors. Testing will resume in the fall of 2016 after the herd returns from summer pasture.

Updates: None

FY 2015

Montana

• Affected beef herd: On November 8, 2014, one reactor (an 8-year-old cow) was detected during testing of a group of 171 head in a Madison County commercial cow-calf beef herd. The herd had a DSA herd plan in place, as a portion of the herd grazes within the DSA. Whole herd testing of 2,338 head was completed December 10, 2015, and no additional seropositive animals were found. An Affected Herd Plan was put in place December 11th. All the DSA-adjacent herds have been identified and have tested negative. Whole-herd test post-removal of the reactor was completed in January 2015 and all animals were negative. Post-calving WHT completed and all animals were brucellosis test negative. Quarantine has been released. All non-DSA adjacent herds (27) have been identified and all tested negative. Assurance testing was completed in January 2016 with all animals testing negative. Traces are pending.

Updates: None

• Affected beef herd: On October 23, 2014, one reactor (a 3-year-old cow) was detected during testing of a group of 262 head in a Park/Carbon County commercial cow-calf beef herd. The reactor animal was part of a cohort that grazes seasonally in the DSA and was tested when leaving the DSA, per State law. The herd was placed under verbal quarantine October 25th. An Affected Herd Plan is in place. On October 31, 2014, the reactor's seropositive status was confirmed, and milk samples submitted to NVSL for culture on November 12th were found negative for *Brucella* spp. On December 4th, tissues from the reactor animal were collected and submitted to NVSL, which reported isolation of *B. abortus* biovar 1 on December 12, 2014. Whole-herd testing was completed December 18, 2014, and no additional seropositive animals were found. All the Park County-adjacent herds have been tested and are negative. Whole-herd test post-removal of the reactor was completed in January 2015 and all animals were negative. WHT was completed March 11, 2015, and all animals were negative. Quarantine was lifted March 16, 2015. All Carbon County adjacent herds (4) have tested negative. Assurance test was completed in October 2015 and all animals tested negative. Traces are pending.

Updates: None

FY 2013

Montana

• Affected beef herd: On September 5, 2013, a single brucellosis reactor (2-year-old female) was detected on a DSA brucellosis pre-slaughter test of 42 head in Madison County. *B. abortus* biovar 1 was confirmed September 25, 2013. A herd brucellosis test (1116 head) detected one reactor and one suspect. The tissues were submitted, but no *B. abortus* isolation was made from either animal. DSA herd and affected herd plans are in place.

Epidemiological work has determined the exposure occurred sometime in the spring of 2013. Nine adjacent herds have tested brucellosis negative. A whole-herd test conducted on January 8, 2014 was negative. No *Brucella* was isolated from tissues submitted from the suspect cow, detected on January test, or its fetus. There were no trace-in animals identified. Twenty of 22 trace-out epidemiological investigations, covering 467 cattle in nine States, have been completed. The post-calving whole-herd test in late March 2014 was negative and the quarantine for the herd was lifted in late March 2014. The fall assurance testing was completed in October 2014; all cattle tested negative. Awaiting disposition of final two traces.

Updates: None

FY 2011

Montana

• Affected privately-owned bison herd: A brucellosis-affected privately owned bison herd was detected in Gallatin County in November 2010. This herd was detected as part of Montana's DSA herd management plan testing. This herd is under quarantine with movement controls and an affected herd management plan in place. All trace-out and trace- in epidemiological investigations associated with this herd have been completed and closed. The fall 2013 test detected 13 seropositive (*B. abortus* biovar 1 isolated) animals out of 4050 head of bison tested. Annual fall testing began October 20, 2014 and was completed November 21, 2014. At the conclusion of the annual fall testing, 16 reactors and three suspects had been identified. Fall testing in 2015 concluded November 6th with a final count of two suspects and six reactors identified out of 3262 head tested. Next test will be fall 2016.

Updates: Fall testing will begin mid-October 2016.

Table 3: Livestock herds confirmed as brucellosis affected, FY 2011-2016.

| State/ FY | Method of Detection | Herd Type | Affected Herd Management Plan | Genotyping Descriptive Results | Wildlife Surveillance Planned | Animals Being Traced | States Receiving Traced Cattle/Bison |
|--------------|---|--------------|-------------------------------------|--|-------------------------------------|----------------------------|---|
| WY 2016 | DSA Herd Plan Test | Beef | Quarantine with test & remove | Most closely related to NER & Franz elk feedgrounds and 2004 Teton County beef herd isolates | Undetermined | Pending | Traces have been completed |
| MT 2015 | DSA Herd Plan Test | Beef | Quarantine with test & remove | Most closely related to 2 elk isolates (2009 – 2010) | Yes | Pending | Traces are pending |
| MT 2015 | DSA Herd Plan Test | Beef | Quarantine with test & remove | Closest common ancestor – wild bison (1985) | Yes | Pending | Traces are pending |
| MT 2013 | DSARequired pre-slaughter testing | Beef | Quarantine with test & remove | Common ancestor with wild elk from the same area | Yes | 424 | 0 trace-in States 9 trace-out States: 20 of 22 traces c ompleted (CA-1, MN-1; open) |