



**Comments on the
Grand Mesa, Uncompahgre and Gunnison
Forest Plan Revision**

Submitted by:

North American Packgoat Association

November 26, 2021

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VIA ELECTRONIC SUBMITTAL

RE: Comments on the Grand Mesa, Uncompahgre and Gunnison Forest Plan Revision

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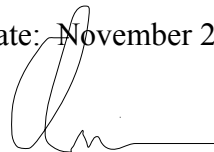
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On behalf of the North American Packgoat Association, I hereby timely submit these Comments on the Grand Mesa, Uncompahgre and Gunnison National Forests (“CMUG”) Draft Environmental Impact Statement (“DEIS”) for the Draft Revised Land Management Plan (“LMP Revision”). If you have any questions concerning these comments or need further information, you may contact NAPgA or Andrew Irvine at the emails and phone numbers indicated above.

Date: November 26, 2021



Andrew A. Irvine
of Andrew A. Irvine, P.C.

I. Introduction to Comments

The North American Packgoat Association (“NAPgA”) timely submits comments on the Grand Mesa, Uncompahgre and Gunnison National Forests (“CMUG” or “Forest”) Draft Environmental Impact Statement (“DEIS”) for the Draft Revised Land Management Plan (“LMP Revision”). Notice of the LMP Revision for the CMUG appeared at 86 Fed. Reg. 44,711-12 (Aug. 13, 2021) and provided the comment deadline of November 12, 2021. The CMUG extended that deadline to November 26, 2021 on its website. *See* <https://www.fs.usda.gov/detail/gmug/landmanagement/planning/?cid=fseprd937839>.

The North American Packgoat Association, Inc. is an organization established specifically for promoting packing with pack goats. The organization was incorporated in March 2001 as a 501(c)(3) non-profit organization. NAPgA seeks to further the pursuit of goatpacking by sharing the knowledge, ideas and experiences of its members, by promoting the use of pack goats to the public as a means of low impact wilderness transportation and recreation, by serving as an advisory group on local and national land use issues, and by engaging in other activities related to educating the public about goatpacking.

NAPgA appreciates this opportunity to comment on the CMUG DEIS for the LMP Revision. NAPgA and its numerous goatpacking-members will be affected by the management direction proposed in the draft LMP Revision. The proposed management direction would result in the curtailment of goatpacking in one of the premier goatpacking areas in the nation, and set a bad precedent for other forests to follow in managing goatpacking as a recreational use. These comments will better inform the DEIS and LMP Revision and further develop the efficacy of the management direction as defined by the LMP Revision.

Goatpacking on the CMUG should not be curtailed as proposed by the LMP Revision.

II. Legal Background for the Comments

A. NEPA Prohibits Uninformed Agency Action

In passing NEPA, Congress “recogniz[ed] the profound impact of man’s activity on the interrelations of all components of the natural environment” and set out “to create and maintain conditions under which man and nature can exist in productive harmony.” 42 U.S.C. § 4331(a). To bring federal action in line with Congress’ goals and to foster environmentally informed decision-making by federal agencies, NEPA “establishes ‘action-forcing’ procedures that require agencies to take a ‘hard look’ at environmental consequences.” *W. Watersheds Project v. Kraayenbrink*, 632 F.3d 472, 486 (9th Cir. 2011) (citing *Metcalf v. Daley*, 214 F.3d 1135, 1141 (9th Cir. 2000)). Foremost among those procedures is the preparation of an environmental impact statement (“EIS”). *Id.*

Agencies considering “major Federal actions significantly affecting the quality of the human environment” are required to prepare an EIS. 42 U.S.C. § 4332(C). The EIS “shall provide full and fair discussion of [the] significant environmental impacts” of the proposed action. 40 C.F.R. § 1502.1. That discussion serves two purposes:

First, it ensures that the agency, in reaching its decision, will have available, and will carefully consider, detailed information concerning significant environmental impacts. Second, it guarantees that the relevant information will be made available to the larger audience that may also play a role in both the decisionmaking process and the implementation of that decision.

W. Watersheds Project, 632 F.3d at 487 (quoting *Dep't of Transp. v. Pub. Citizen*, 541 U.S. 752, 768 (2004)). This process does not mandate particular substantive results, but “NEPA . . . prohibits uninformed . . . agency action.” *Robertson v. Methow Valley Citizens Council*, 490 U.S. 332, 351 (1989). By focusing agency and public attention on the environmental effects of proposed action, “NEPA ensures that the agency will not act on incomplete information, only to regret its decision after it is too late to correct.” *Marsh v. ONRC*, 490 U.S. 360, 371 (1989).

B. Review Under the APA

The Administrative Procedure Act (“APA”), 5 U.S.C. §§ 701-706, provides for judicial review of agency actions, such as those at issue here.¹ Under the APA, a reviewing court shall “hold unlawful and set aside agency action, findings, and conclusions found to be . . . arbitrary, capricious, an abuse of discretion, or otherwise not in accordance with law; . . . [or] without observance of procedures required by law.” 5 U.S.C. § 706(2)(A), (D). Although the arbitrary and capricious standard is a “narrow one,” the court is required to “engage in a substantial inquiry” and a “thorough, probing, in-depth review.” *Native Ecosystems Council v. U.S. Forest Serv.*, 418 F.3d 953, 960 (9th Cir. 2005) (quoting *Citizens to Preserve Overton Park, Inc. v. Volpe*, 401 U.S. 402, 415-16 (1971)).

Under this standard, an agency decision is to be reversed as arbitrary and capricious if the agency has “. . . entirely failed to consider an important aspect of the problem, [or] offered an explanation that runs counter to the evidence before the agency. . . .” *Motor Vehicle Mfrs. Ass’n v. State Farm Mutual Auto. Ins. Co.*, 463 U.S. 29, 43 (1983). “The reviewing court should not attempt itself to make up for such deficiencies.” *Id.* (citation omitted). Most fundamentally, the agency must “examine the relevant data and articulate a satisfactory explanation for its action including a ‘rational connection between the facts found and the choice made.’” *Motor Vehicle*, 463 U.S. at 53 (quotation omitted).

Where, as here, there has been a change in policy from allowing goatpacking on the CMUG to curtailing goatpacking on the Forest, judicial review starts with the presumption that the change in policy is *not* justified by the administrative record. *Motor Vehicle*, 463 U.S. at 42. Additionally, the traditional presumption of agency expertise “may be rebutted if the decisions,

¹ NEPA claims are subject to judicial review under the APA, 5 U.S.C. § 706(2)(A). See *Dep't of Transp. v. Pub. Citizen*, 541 U.S. at 763; *Marsh*, 490 U.S. at 375–76; *League of Wilderness Defenders-Blue Mtns. Biodiversity Project v. U.S.*, 549 F.3d 1211, 1215 (9th Cir. 2008) (the APA provides authority for the court’s review of decisions under NEPA); *W. Watersheds Project v. U.S. Forest Serv.*, 2006 WL 292010, *2 (D. Idaho) (same).

even though based on scientific expertise, are not reasoned.”” *W. Watersheds Project v. Ashe*, No. 11-462, 2013 WL 2433370 at *5 (D. Idaho June 4, 2013) (citations omitted).

In addition to the requirements of the NEPA and the APA, Forest Service regulations require that “best available science” be taken into account in forest planning. 36 C.F.R. § 219.3. In taking “best available science” into account, the Forest Service must “document how the best available science information was used to inform the assessment, the plan decision, and the monitoring program” and such documentation must “[i]dentify what information was determined to be the best available scientific information, explain the basis for that determination, and explain how the information was applied to the issues considered.” *Id.*

III. Background on the LMP Revision

The LMP Revision makes significant changes to the CMUG’s existing management of goatpacking on the Forest. At Guideline FW-GDL-SPEC-14, the Forest further provides: “To maintain long-term population viability for bighorn sheep, the Forest Service should minimize the potential for recreational pack goats to interact with bighorn sheep. The Forest Service should manage recreational pack goats consistently with its management of domestic sheep within the comparable allotment(s) area.” *Id.*

This guideline follows Standard FW-STND-SPEC-13, applicable to domestic sheep, which provides: “On active grazing allotments, maintain effective separation between domestic sheep and bighorn sheep herds. Effective separation is defined as spatial or temporal separation between bighorn sheep and domestic sheep. See associated management approach.” LMP Revision at 29.

Under “Management Approaches” for Big Game Species, the Forest provides: “To implement GDL-SPEC-13, Tier 1 bighorn sheep herds with the greatest potential to contribute to population viability in the plan area should be prioritized. Tier 2 herds, where they interact or have the potential to interact with Tier 1 herds, should also be prioritized. Use the most current version of the Western Association of Fish and Wildlife Agency’s *Recommendations for Domestic Sheep and Goat Management in Wild Sheep Habitat* to inform management.” *Id.* at 30.

The DEIS explains, “Public feedback requested an alternative that would provide full separation of domestic and bighorn sheep within a specified timeframe. The agency’s national policy, however, is to provide for effective separation, and this is reflected in the draft revised forest plan and alternatives.” DEIS Vol. 1 at 25. Neither the “public feedback” nor the “agency’s national policy” is cited or found in the DEIS and associated documents. There is no indication provided of how or why separation of pack goats from bighorn sheep was considered in the alternatives.

The DEIS indicates that “Pneumonia/Respiratory disease” is a potential “risk factor” for Rocky Mountain and Desert bighorn sheep, and references the “*Insects and Disease* section of the 2018 assessment” for further discussion. DEIS Vol. 1 at 147-48; DEIS Vol. 2 at 108 (same). The GMUG Revised Draft Forest Assessments: Identifying and Assessing At-Risk Species at 46

(March 2018) also indicates that “Pneumonia/Respiratory disease” is a potential “risk factor” for Rocky Mountain and Desert bighorn sheep.

The DEIS further discusses “disease epizootics,” indicating that: “[i]n Colorado, the susceptibility of bighorn sheep to pathogens introduced by domestic sheep is regarded as the primary factor limiting bighorn sheep populations (George et al. 2009).” DEIS Vol. 1 at 200. The DEIS further provides:

Effective separation is defined by science-based estimates of bighorn sheep core herd range and movements across the landscape in relationship to domestic sheep areas and managing potential contact rates to an acceptable level to reduce the risk of disease transmission. Management to maintain separation would also address the risk factor for disease epizootics and would address competition with domestic animals. The indicator for this analysis is a qualitative discussion of the effects of plan components.

Id. at 202.

Concerning “Disease Transmission and Effective Separation from Domestic Sheep and Goats” the DEIS provides:

The current forest plan makes little to no recognition of the risk that disease transmission from domestic sheep poses to their wild cousins. However, the current plan does appear to allow a wide enough array of adaptive management flexibility for managers to have options to reduce risk somewhat. Trailing of domestic sheep through bighorn sheep habitat still occurs. All action alternatives contain two Forestwide plan components that address the risk of disease transmission from domestic sheep to bighorn sheep:

- FW-STND-SPEC-13, which addresses “effective separation” of bighorn and domestic sheep as a standard. The plan component does not define effective separation, but the emphasis of the planning rule on “best available science” means that the definition would be based on current science. As our scientific understanding of the species needs changes, so could the definition of “effective separation” as regards the risk of disease transmission from domestic animals to wild animals.
- FW-GDL-SPEC-14 addresses the risk of disease transmission from goat to bighorn sheep, requiring the Forest Service to minimize the potential for interaction and to manage pack goats consistently with the way sheep are managed.

The impact of these two components on the disease transmission issue faced by bighorn sheep would be strong. SPEC-13 for “effective separation” would become a mandatory component of renewed allotment management plans. The guideline SPEC-14 does provide greater flexibility regarding separation between pack goats and bighorn sheep, but still requires justification for any occasion when the Forest Service allows the users or permittees to not comply.

Compared to the existing forest plan, these two components—over time, as incorporated into individual allotments as they are renewed—would likely greatly reduce the risk of disease transmission from domestic animals to bighorn sheep. Any remaining risk would occur from domestic animals escaped from handlers or permitted areas, or from bighorn sheep wandering well outside known herd ranges—stochastic events that can be hard to predict or manage.

Overall, the adoption of the Forestwide components of the action alternatives would reduce the potential for disease transmission from domestic sheep to bighorn sheep. This would be a direct, long-term moderately beneficial impact to bighorn sheep in the GMUG.

Id. at 203.

With regard to “Plan Components” under the “Insects and Disease” section of DEIS Volume 2, the DEIS provides: “Disease transmission to bighorn sheep is addressed in FW-STND-SPEC-13 and FW-GDL-SPEC-14, which manages the potential for interaction between domestic sheep, pack goats, and bighorn sheep.” DEIS Vol. 2 at 109. No further discussion concerning pack goats or the impacts of the alternatives on goatpacking appears in the DEIS or LMP Revision.

IV. Comments on the DEIS and LMP Revision

To assist the CMUG, NAPgA’s comments generally refer to specific pages of the DEIS and LMP Revision that form the basis for each comment; however, some comments may apply more broadly. Comments are intended to apply to all listed pages, or generally, and should be addressed in the context of each of the listed pages or in general.

NAPgA looks forward to the CMUG’s responses to its comments. In addition to its general obligation to respond to public comments under 40 C.F.R. § 1503.4(a), the CMUG must specifically “discuss at appropriate points in the final [EIS] any responsible opposing view which was not adequately discussed in the draft [EIS] and . . . indicate the agency’s response to the issues raised.” *Ctr. for Biological Diversity v. U.S. Forest Serv.*, 349 F.3d 1157, 1167 (9th Cir. 2003) (quoting 40 C.F.R. § 1502.9(b)). A failure to do so is itself a NEPA violation. *Id.* at 1168. The CMUG must also “insure the professional integrity, including scientific integrity, of the discussions and analyses” included in its DEIS. 40 C.F.R. § 1502.24.

1. The DEIS Does Not Present any Science on Disease Transmission from Domestic Goats, Especially Pack Goats. To Ensure the Scientific Integrity of the DEIS and Forest Plan, the CMUG Should Remove Unsupported Statements Concerning Pack Goats from the DEIS and LMP Revision.

In evaluating the environmental impacts of a proposed action, NEPA requires federal agencies to ensure the scientific integrity of an EIS by considering appropriate studies and data. 40 C.F.R. § 1502.24. The CMUG must “insure the professional integrity, including scientific integrity, of the discussions and analyses” included in its DEIS. *Id.* An agency may not rely on conclusory statements unsupported by data, authorities, or explanatory information. *Seattle*

Audubon Soc'y v. Moseley, 798 F. Supp. 1473, 1480-83 (W.D. Wash. 1992), *aff'd*, 998 F.2d 699 (9th Cir. 1993). NEPA requires that an agency candidly disclose in its EIS the risks and effects of its proposed actions, and that it respond to adverse opinions held by respected scientists. *Seattle Audubon*, 798 F. Supp. at 1482 (citing *Friends of the Earth v. Hall*, 693 F. Supp. 904, 937 (W.D. Wash. 1988)). Further, under NEPA, courts have held that agency actions based on unexplained assumptions are arbitrary and capricious. *Ctr. for Biological Diversity v. U.S. Dep't of the Interior*, 623 F.3d 633, 650 (9th Cir. 2010); *see also Dow Agrosciences LLC v. Nat'l Marine Fisheries Serv.*, 707 F.3d 462, 470 (4th Cir. 2013) (agency must explain why lab tests reflect nature).

The CMUG has failed to ensure the professional integrity, including scientific integrity, of the discussions and analyses in the DEIS as required under NEPA. The CMUG appears to be operating on incomplete information concerning disease transmission from domestic goats, including pack goats, to bighorn sheep, and also appears to be ignoring important aspects of the problem of disease transmission as well as offering explanations in the DEIS that run counter to the evidence before the CMUG. Much of the analysis and discussion in the DEIS with regard to pack goats lacks factual or scientific support.

The DEIS indicates: "In Colorado, the susceptibility of bighorn sheep to pathogens introduced by domestic sheep is regarded as the primary factor limiting bighorn sheep populations (George et al. 2009)." DEIS Vol. 1 at 200. This appears to be the only scientific information provided in the DEIS concerning disease transmission.

George et al. 2009 does not appear to establish or discuss disease transmission from domestic goats, and certainly not pack goats, to bighorn sheep. Yet, the CMUG is curtailing goatpacking on the Forest under Guideline FW-GDL-SPEC-14.

What is the scientific justification for curtailing goatpacking on the Forest? None appears to be provided. There are no facts or scientific references provided to support the statements in the DEIS and LMP Revision as they apply to domestic goats, especially pack goats, and the population of bighorn sheep on the CMUG. There are no facts or science presented indicating that a domestic goat, particularly a pack goat, has ever transmitted disease to bighorn sheep in the wild.

If the CMUG is going to implicate pack goats in disease transmission to bighorn sheep and curtail goatpacking on the Forest, it must provide science showing that pack goats carry disease lethal to bighorn sheep, that there is significant risk of disease transmission from pack goats to bighorn sheep and that disease transmission from pack goats would have lasting effects on population performance. Please provide such information to the public for review and comment or otherwise remove the unsupported statements and guideline concerning pack goats from the DEIS and LMP Revision.

2. The CMUG Must Consider Dr. Margaret Highland's Research Concerning the Limited Prevalence of *Mycoplasma ovipneumoniae* in Pack Goats.

The CMUG has failed to consider recent scientific research indicating that pack goats do not commonly carry *Mycoplasma ovipneumoniae*. This research by Dr. Margaret Highland,

Research Veterinarian with the Animal Disease Research Unit-ARS-USDA is presented in Exhibit B. Dr. Highland's research indicates that pack goats do not commonly carry the disease-causing organisms associated with bighorn sheep die-offs. The results of the testing performed for Dr. Highland's research are also included in Exhibit B, so that the CMUG can consider the results and verify the legitimacy and scientific method in the research. Dr. Highland's research is in the process of being published, but has already been presented, *see, e.g.*, <https://pdfs.semanticscholar.org/presentation/4bb7/616fa740f42ceda2c55d275f0a8032fc6ca8.pdf> and has been considered by the Forest Service on other occasions (but, not on the CMUG).

Under the APA and NEPA, the CMUG is required to consider the fundamental aspect of the problem of disease transmission, namely, whether pack goats can actually carry and transmit *M. ovi* to bighorn sheep in the wild. *See Motor Vehicle*, 463 U.S. at 43. The CMUG is also required to examine relevant data, consider opposing viewpoints, ensure the scientific integrity of its discussions, and articulate a satisfactory explanation for its action. *See id.* at 42-43, 53; *Ctr. for Biological Diversity v. U.S. Forest Serv.*, 349 F.3d at 1167 (quoting 40 C.F.R. § 1502.9(b)).

Moreover, and in addition to the requirements of the APA and NEPA, Forest Service regulations require that "best available science" be taken into account in forest planning. 36 C.F.R. § 219.3. In taking "best available science" into account, the Forest Service must "document how the best available science information was used to inform the assessment, the plan decision, and the monitoring program" and such documentation must "[i]dentify what information was determined to be the best available scientific information, explain the basis for that determination, and explain how the information was applied to the issues considered." *Id.* The Forest Service Land Management Planning Handbook, FSH 1909.12, directs the CMUG's use of the best available scientific information and provides that where research is relevant, accurate and reliable, the Forest Service should include it as the best available scientific information. *See* FSH 1909.12, 42.13.

As a result, this science presented by Dr. Highland must be considered in the DEIS under the APA and NEPA, as well as the implications of pack goats not being carriers of *M. ovi*. If pack goats are not carriers of disease-causing pathogens, then they do not pose a risk of disease transmission to bighorn sheep on the CMUG.

In sum, the CMUG must review and consider Dr. Highland's research in the DEIS. Such consideration is required by the APA, NEPA and the Forest Service's own planning regulations. Dr. Highland's research indicates that pack goats are rarely carriers of *M. ovi*. As a result, pack goats do not pose a significant risk of disease transmission to bighorn sheep on the CMUG. Pack goats cannot transmit disease they do not have. These points must be considered in the DEIS and LMP Revision.

3. The DEIS Must Specifically Identify and Discuss the Threat of Disease Transmission from Pack Goats to Bighorn Sheep.

The CMUG should explain in the DEIS what the risk of disease transmission between pack goats and bighorn sheep actually is. Notably, contact between pack goats and bighorn sheep has never occurred before on the Forest, so risk does not mean that contact is more likely

than not, otherwise such contact would have already occurred. There is no scale of risk to inform the reader about the actual likelihood of contact. The CMUG should explain what they mean by “risk,” including the various scales of risk from high to low. Also, the CMUG should explain how contact between pack goats and bighorn sheep on the Forest would actually occur. What does the CMUG mean by “interact?” LMP Revision at 29. Would a bighorn sheep approach a pack goat on a trail, in the presence of the pack goat’s human owner and “interact?” Would a bighorn sheep enter into a camp in a forested area where there is a pack goat, again in the presence of its human owner, and “interact” there? Is this nose-to-nose or sexual contact? When the CMUG considers risk and “interaction” in the DEIS it is unclear what the CMUG is talking about and how such “interaction” would occur. These things should be explained. Likewise, the CMUG should discuss the likelihood of contact in understandable terms and present how such contact would occur based on the behavior of bighorn sheep and use and training of pack goats.

In reality, there is limited use of pack goats on the CMUG, so for disease transmission to occur on the Forest, during one of the few goatpacking trips taken each year, a bighorn sheep would have to (1) leave its herd and its summer habitat in the high country, (2) find a human and pack goat camp, (3) sneak into that camp without causing any disturbance in the pack goats and without being detected by the humans, (4) ask the pack goats to not be alarmed, to remain still and to muffle their bells and collars, (5) find a tethered goat that is infected by and shedding strains of *M. ovi*, (6) make physical contact with that goat sufficient for disease transmission, and (7) sneak back out of camp and return to its herd and infect other bighorn sheep. It is a far-fetched scenario that has never happened before.

In reality, there is almost no overlap in time or space between pack goats and bighorn sheep on the CMUG; bighorn sheep are not prone to leave their herd/habitat and wander into human and pack goat camps; pack goats react noisily when they are alarmed by other wildlife, including bighorn sheep; the vast majority of pack goats do not carry and shed strains of *M. ovi*; and it is unknown whether bighorn sheep can even be infected with strains of *M. ovi* from pack goats resulting in fatal respiratory disease. The facts do not support the CMUG’s assumption that there is a likelihood of disease transmission from pack goats to bighorn sheep on the Forest.

Before undertaking management action concerning the risk of contact and disease transmission between pack goats and bighorn sheep on the CMUG, the Forest should provide an analysis of the current risk posed by pack goats. This could be done with a quantitative risk assessment. Regardless, the CMUG has not presented any scientific information indicating that pack goats pose a significant risk. Rather, pack goats rarely use the CMUG, rarely carry disease and are very unlikely to contact a bighorn sheep, particularly when handled according to established guidelines, so pack goats would appear to pose negligible risk. Why then are they being curtailed on the CMUG? The CMUG should answer this threshold question. The CMUG does not provide any explanation for curtailing pack goat use and the conclusion to curtail goatpacking runs counter to the evidence before the agency. Without establishing significant risk, the CMUG’s curtailment of pack goat use is unjustified.

4. The CMUG Must Consult the Agricultural Research Service, within the United States Department of Agriculture, Before Preparing the Final EIS and Record of Decision.

NEPA imposes on federal agencies conducting environmental review a duty to consult with certain other agencies.” Prior to making any detailed statement, the responsible Federal official shall consult with and obtain the comments of any Federal agency which has jurisdiction by law or special expertise with respect to any environmental impact involved [in the proposed action].” 42 U.S.C. § 4332(2)(C). Further, to promote NEPA’s policies of public participation and informed decisionmaking, copies of the EIS and comments thereon from other agencies “shall accompany the proposal through the existing agency review processes.” *Id.*

The regulations implementing these provisions state that “[a]fter preparing a draft environmental impact statement and before preparing a final environmental impact statement the agency shall . . . [o]btain the comments of any Federal agency which has jurisdiction by law or special expertise with respect to any environmental impact involved” 40 C.F.R. § 1503.1(a)(1); *see also id.* § 1500.1(b) (“Accurate scientific analysis, expert agency comments, and public scrutiny are essential to implementing NEPA.” (emphasis added)). “Special expertise” is defined as “statutory responsibility, agency mission, or related program experience.” *Id.* § 1508.26. Under the statute and its implementing regulations, the CMUG has a duty to consult with the Agriculture Research Service (“ARS”) before issuing the Final EIS. *See Idaho Wool Growers Ass’n v. Vilsack*, 816 F.3d 1095, 1103 (9th Cir. 2016).

ARS has “special expertise” concerning significant aspects of the proposed decision, including the mechanics of pathogen transmission in domestic sheep and goats. For example, 7 C.F.R. § 2.65 delegates to ARS, among other matters, the authority to “[c]onduct research concerning domestic animals and poultry, their protection and use, [and] the causes of contagious, infectious, and communicable diseases.” Also, ARS’s mission statement proclaims: “ARS conducts research to develop and transfer solutions to agricultural problems of high national priority and provide information access and dissemination to . . . enhance the natural resource base and the environment” U.S. Department of Agriculture, Agricultural Research Service, ARS: About US, <http://www.ars.usda.gov/aboutus/aboutus.htm>.

Thus, considering the language establishing NEPA’s consultation requirement is expansive, NEPA mandates consultation with any federal agency that has” special expertise with respect to any environmental impact involved.” 42 U.S.C. § 4332(2)(C) (emphasis added); *see also* 40 C.F.R. § 1503.1(a)(1) (“[T]he agency shall . . . [o]btain the comments of any Federal agency which has jurisdiction by law or special expertise with respect to any environmental impact involved. . . .” (emphasis added)). And, further considering that *Warm Springs Dam Task Force v. Gribble* suggests that for the consultation requirement to apply, the particular expertise of an agency does not have to encompass the proposed project as a whole or the issue the proposed project was designed to address. Rather, the expertise need relate only to one of the project’s anticipated environmental effects. *See* 621 F.2d 1017, 1020-21 (9th Cir. 1980) (per curiam); *see also Idaho Wool Growers Ass’n*, 816 F.3d at 1103. It is a clear requirement that the CMUG MUST consult with ARS on issues of disease transmission, such as those presented in the DEIS and LMP Revision, prior to issuing a Final EIS. As a result, the CMUG MUST consult with ARS and should detail such consultation in the Final EIS.

5. The CMUG Fails to Account for the Important Differences Between Pack Goats and Herd Domestic Goats and Domestic Sheep.

The CMUG fails to acknowledge the important differences between pack goats and herd domestic sheep and goats. These differences must be considered in the DEIS and LMP Revision. NEPA prohibits this type of uninformed agency action. *See Robertson*, 490 U.S. at 352 (“NEPA . . . prohibits uninformed . . . agency action.”); *Marsh*, 490 U.S. at 371 (“NEPA ensures that the agency will not act on incomplete information, only to regret its decision after it is too late to correct.”). These differences are critical to the CMUG’s analysis of disease transmission from pack goats to bighorn sheep and must be considered by the Forest under NEPA.

Pack goats are very different from other domestic goats (and domestic sheep), both by breed and by use. These differences result in far less risk to bighorn sheep than the risk posed by domestic goats (or domestic sheep) on grazing allotments. The CMUG DEIS must account for these differences. To consider pack goats the same as other domestic goats (or domestic sheep) for purposes of analyzing the risk of disease transmission to bighorn sheep on the CMUG would be a critical error.

Pack goat owners go to great lengths and expense to find and train particular goats that will not stray from the security of a finite string of pack goats and their owner. Pack goats are inextricably bonded to their owners, which represent the “alpha goat” in their small herd. This is achieved through the processes of imprinting and socialization of pack goats from birth. As a result, pack goats are not prone to straying and remain in very close proximity to the “alpha goat.” Other domestic goats (and domestic sheep), while often included in herds that number in the hundreds or thousands (compared to a string of pack goats ranging from two to ten goats), are not individually trained and, thus, there may be some risk of individual domestic herd goats (or domestic sheep) straying from the herd. The risk associated with domestic sheep or domestic goats transmitting disease to bighorn sheep requires “physical contact” between the domestic animal and the bighorn sheep, therefore, a pack goat that is less likely to stray and thereby come into contact with a bighorn sheep poses a much lower risk of transmission than any number of herd domestic sheep or goats which can wander and stray.

Domestic goat and sheep herds typical to grazing allotments on public land represent larger populations of animals that are more difficult to maintain, and which may not be in immediate proximity of their caretaker at all times. Pack goats, on the other hand, require their owner or “alpha goat” to be present to monitor the herd at all times, and are always in their owner’s immediate presence and control. The small size of a pack goat string and perpetual control of the owner allows pack goats to be tied in unison while on trails, and tethered or high-lined at night (among other best management practices that can be easily implemented) to reduce the risk of contact between a pack goat and a bighorn sheep. Furthermore, if ever in sight of a bighorn sheep, there is always a human present in close proximity to the pack goats, making it extremely unlikely that a bighorn sheep would approach the string. In the presence of wild animals, such as bighorn sheep, pack goats are also on heightened alert and retreat to a position near the “alpha goat,” i.e., their human caretaker. This and the other defining traits of pack goats, and the nature of their use and training, make pack goats far less of a risk of coming into contact with a bighorn sheep than herd domestic goats and/or domestic sheep.

Further, the lifestyle and care of a pack goat differs greatly from that of a typical herd domestic goat or domestic sheep. This difference in care means that pack goats are healthier and less likely to be the carrier of a disease. Pack goats are seen by their owners as a significant investment in time and resources. A pack goat is not viable for packing purposes until at least the age of three or four, and often pack goats do not reach their packing prime until the age of five or six. Thus, a goatpacker will have had to invest a number of years into a pack goat before it is ready to hit the trail. During this time, and throughout a pack goat's life, pack goats see personalized veterinary care in order to keep the goat healthy and prolong their useful life, a luxury that other free ranging herd domestic goats or domestic sheep do not enjoy.

Because of their overall health and stamina, a trained pack goat can bring a sale price of over \$500. This means that a pack goat owner has a large financial interest in each of his or her pack goats. This high financial interest means that the owner of pack goats is likely to see to their care and protection whether that is protection from disease at home, or from contact with other wildlife when on public lands.

Further, typical herd domestic goats and domestic sheep may be sold and intermixed with goats from other herds. In contrast, pack goats—which are treated more like household pets than livestock—are not likely to change owners. The higher frequency that typical herd domestic goats and domestic sheep may be exposed to other domestic stock, would increase the opportunity for disease to spread between individual animals. On the other hand, pack goats are infrequently transferred between owners because of the nature of their function and required bonding. This greatly reduces the risk of exposure of pack goats to various diseases as compared to herd domestic goats and domestic sheep.

Perhaps most critical to the CMUG's analysis of disease transmission from pack goats to bighorn sheep is the fact that the overwhelming majority of pack goats are not known to carry *M. ovi*. If a pack goat did not carry *M. ovi* it would be impossible for that goat to transmit disease to a bighorn sheep. Thus, the risk of disease transmission from that pack goat to a bighorn sheep would be zero. Further, even if a pack goat were to carry *M. ovi* and directly contact a bighorn sheep, there is no science indicating that the pack goat would transmit this pathogen to the bighorn sheep and that the bighorn would succumb to pneumonia as a result. The CMUG did not consider these important factors in its analysis.

Finally, goatpackers limit their visits to the CMUG, as well as their time on the Forest when they do visit. With only a few pack goats per goatpacker and only a few visits by goatpackers per year, for a limited amount of time, the chance that a pack goat would come into contact with a bighorn sheep is extremely unlikely. This factor was not considered by the CMUG.

Here, the CMUG's analysis in the DEIS is completely silent on the differences between pack goats and herd domestic goats and how those differences affect the risk of disease transmission between pack goats and bighorn sheep. These differences are critical and must be considered by the CMUG. An agency decision is to be reversed as arbitrary and capricious if the agency has "entirely failed to consider an important aspect of the problem." *Motor Vehicle Mfrs. Ass'n v. State Farm Mutual Auto. Ins. Co.*, 463 U.S. 29, 43 (1983). The CMUG's silence on the issue will not suffice. The agency's path must be reasonably discerned. *Id.* A court "cannot

infer an agency's reasoning from mere silence or where the agency failed to address significant objections and alternative proposals." *Beno v. Shalala*, 30 F.3d 1057, 1073 (9th Cir. 1994) (citing *Motor Vehicle*, 463 U.S. at 57); see also, e.g., *SEC v. Chenery Corp.*, 332 U.S. 194, 196-97 (1947) ("[i]t will not do for a court to be compelled to guess at the theory underlying the agency's action.").

In conclusion, pack goats are very different than other herd domestic goats or domestic sheep that are grazed on or near the CMUG, and the use of pack goats on the CMUG is very different than the use of other herd domestic goats and domestic sheep. The CMUG DEIS and LMP Revision fail to account for these differences in the analysis of disease transmission from domestic sheep and domestic goats to bighorn sheep on the CMUG. As a result, the DEIS must be revised to consider (1) pack goats separate from other herd domestic goats and domestic sheep and (2) the unlikelihood that pack goats carry disease and (3) the unlikelihood that pack goats would ever come in close contact with bighorn sheep on the CMUG. Further, the CMUG must consider that the nature and use of pack goats on the Forest already achieves the spatial and/or temporal separation recommended by the CMUG to minimize potential disease transmission. Thus, there is no justification and no need for the curtailments of pack goats on the CMUG.

6. The CMUG Should Consider and Discuss Mitigation Measures that Would Allow the Use of Pack Goats on the Forest.

Under NEPA, the CMUG must consider and discuss mitigation measures that would allow the use of pack goats on the Forest. NAPgA has attached a proposed suite of best management practices ("BMPs") and other minimization and mitigation measures at Exhibit A to prevent contact and possible disease transmission between pack goats and bighorn sheep on the CMUG. These, as well as other available practices and measures must be considered by the CMUG in the DEIS.

For example, the CMUG DEIS fails to consider that separation between pack goats and bighorn sheep is maintained by the presence of a human with pack goats, by nighttime tethering or high-lining of pack goats, and by the nature and training of pack goats. The DEIS also failed to consider the use of GPS tracking collars on pack goats, pathogen testing, permitting for pack goat trips, designation of corridors for pack goats, and a host of other measures. Certainly, if pack goats do not carry disease and do not come into contact with bighorn sheep, there is zero risk of disease transmission from pack goats to bighorn sheep. Neither of these scenarios were considered in the DEIS. Instead of considering any of these measures, in violation of NEPA, the CMUG fails to provide any consideration of these best management practices to maintain separation between pack goats and bighorn sheep on the CMUG. All of the proposed alternatives appear to equally restrict goatpacking on the Forest.

BMPs are mitigation measures that can be employed by goatpackers to prevent contact between pack goats and bighorn sheep. 40 C.F.R. § 1508.20 (defining "mitigation measures" to include "[a]voiding the impact" and "[m]inimizing impacts by limiting the degree or magnitude of the action and its implementation"). For a reasonable range of alternatives, the CMUG DEIS must consider implementation of BMPs and mitigation measures, rather than simply concluding that goatpacking on the CMUG must be curtailed. 40 C.F.R. § 1502.14.

An EIS must discuss “mitigation . . . in sufficient detail to ensure that environmental consequences have been fairly evaluated.” *Robertson*, 490 U.S. at 352. An agency is required to “discuss possible mitigation measures in defining the scope of the EIS, 40 CFR § 1508.25(b), in discussing alternatives to the proposed action, § 1502.14(f), and consequences of that action, § 1502.16(h), and in explaining its ultimate decision, § 1505.2(c).” *Id.*; see also *Okanogan Highlands Alliance v. Williams*, 236 F.3d 468, 473 (9th Cir. 2000) (An EIS must contain a “reasonably complete discussion of possible mitigation measures.” (quoting *Robertson*, 490 U.S. at 352)). To be sure, an agency’s final decision must “[s]tate whether all practicable means to avoid or minimize environmental harm from the alternative selected have been adopted, and if not, why they were not.” 40 C.F.R. § 1505.2(c).

Further, NEPA mandates that federal agencies “provide legitimate consideration to alternatives that fall between the obvious extremes.” *Colorado Envtl. Coalition v. Dombeck*, 185 F.3d 1162, 1175 (10th Cir. 1998). More specifically, NEPA is violated when an agency dismisses the consideration of an alternative “in a conclusory and perfunctory manner that [does] not support a conclusion that it was unreasonable to consider them as viable alternatives.” *Davis v. Mineta*, 302 F.3d 1104, 1122 (10th Cir. 2002). “The existence of reasonable but unexamined alternatives renders an EIS inadequate.” *Ilio’ulaokalani Coalition v. Rumsfeld*, 464 F.3d 1083, 1095, 1101 (9th Cir. 2006).

Without an alternative that describes and analyzes the implementation of mitigation measures to prevent contact between pack goats and bighorn sheep, instead of simply curtailing the use of pack goats on the CMUG, the DEIS contains an inadequate range of alternatives. Alternatives considering BMPs and mitigation measures are both reasonable and feasible under the circumstances, and must be analyzed in the DEIS.

In conclusion, the CMUG has failed to discuss and consider mitigation measures that would allow use of pack goats on the Forest while preventing the risk of disease transmission between pack goats and bighorn sheep. As a result, the CMUG must revise the DEIS and LMP Revision to discuss and consider appropriate mitigation measures to prevent the risk of disease transmission between pack goats and bighorn sheep. Proper consideration of such measures should include consideration and adoption of an alternative to allow the use of pack goats on the CMUG. This alternative should consider maintenance of the separation of pack goats and bighorn sheep on the Forest and, thus, achieve avoidance of any potential for disease transmission between pack goats and bighorn sheep.

7. The CMUG Must Evaluate Alternatives that Consider Strengthening Bighorn Sheep Immunity to Disease.

Established epidemiology shows that disease occurs in bighorn sheep populations in the absence of contact with domestic sheep and other animals, including pack goats. These data indicate that infectious agents and other contributing factors involved in the disease process are present within bighorn sheep populations. It appears that most bighorns are getting pneumonia from other bighorns because most of the herds that have outbreaks of pneumonia, are not in contact with domestic sheep or domestic goats. This indicates that the major problem is the lack of a good immune system in the bighorns. As discussed below, there are inherent risks in choosing a management strategy that attempts to isolate bighorn sheep populations from all

perceived transmission risks (when complete isolation is not possible); instead the focus should be on managing population immunity.

The critical component of managing infectious diseases is population immunity. A decision to isolate a given population of bighorn sheep from contact with potential sources of infection assumes the ability for that population to maintain isolation. The wisdom of this management scheme (maintaining immunological naivety) in animal populations within the United States, when sources of infection are present in nature, is questionable at best. Two methods which provide population immunity are vaccination and/or exposure of populations through natural exposure (transmission). This latter situation is also referred to as premonition (resistance to a disease due to the existence of its causative agent in a state of physiological equilibrium in the host and/or by immunity to a particular infection due to previous presence of the causative agent).

A primary risk associated with incomplete immunologic isolation of animal populations is cycles of disease when isolation is broken as opposed to a continuum of managed population immunity through vaccines and/or natural exposure and premonition. When multiple sources of a given pathogen or group of pathogens exist, the prudent long-term health management dictates that population immunity be the primary tool. As an example of population immunity being the most effective management tool, the Lostine River herd of bighorns experienced a die-off in the 1980s, but is now considered the most viable herd in the Hells Canyon area due to successful population immunity. Since bighorn sheep are infecting each other, building up their immune systems could have a beneficial effect on survival from many forms of disease.

Likewise, bighorn sheep face the risk of infection from domestic sheep and other animals on and off the CMUG. Consequently, the curtailment of pack goats on the CMUG, even if there was evidence that pack goats carried and transmitted disease, would not eliminate the risk of disease transmission to bighorns. This fact is not adequately considered in the DEIS. It will be impossible for the CMUG to eliminate the risk of disease transmission to bighorns because of the numerous variables besides pack goats (which are not even a known carrier or transmitter of disease) on the CMUG. As a result, the CMUG must analyze alternative solutions to maintaining bighorn sheep viability.

The CMUG must also analyze the possibility that without interaction between bighorn sheep and other animals, bighorn sheep tolerance to disease may become worse, leading to more widespread die-offs, instead of fewer die-offs. Instead of considering this likelihood, the DEIS only considers one course of action: total separation. Based on the analysis in the DEIS, the most prudent and most logical management action would be to encourage development of immunity in bighorns because total separation is impossible. This action must be considered by the CMUG in the DEIS.

8. Epidemiological Modeling is Needed to Understand How a Range of Factors Affect the Dynamics of Disease Spread Under Various Management Alternatives.

The very limited disease review in the DEIS is generally based on geographic characteristics of the disease in the context of interaction between domestic sheep and bighorn

sheep. While this is a useful component of much needed research, it is not in itself enough to make well-informed recommendations on policy alternatives. There remains limited knowledge of transmission dynamics. Clinical studies have shown bighorn sheep susceptibility to disease from contact with domestic sheep. However, epidemiologic modeling is needed to understand how contacts with domestic sheep, bighorn sheep, and other disease carriers (elk, deer, wild goats, birds, etc.), forage and climatic conditions, and other factors affect the dynamics of the disease spread under various management alternatives. The CMUG does not appear to apply any sort of modeling for the risk of disease transmission on the Forest.

NEPA's procedures require the presentation of "complete and accurate information to decision makers and to the public to allow an informed comparison of the alternatives considered in the EIS." *NRDC v. U.S. Forest Service*, 421 F.3d at 813. Here, further modeling and additional study is needed to determine the added probability of disease transmission among bighorns and from other animals. The probability that healthy "carrier" bighorns are infecting "non-carrier" bighorns is likely high, since a large number of the bighorns on the CMUG may be disease-carriers. Additionally, more information and study should be undertaken to determine the exact mechanism for developing pneumonia in bighorn sheep following association with domestic sheep or other animals. Further, the CMUG must study the development of immunity to disease in bighorn sheep. All of this information should be considered and addressed by the CMUG in the DEIS.

9. The CMUG Fails to Consider the Most Important Aspects of the Problem in the DEIS.

Under the APA, agency decisions under NEPA and NFMA will be set aside if they are "arbitrary, capricious, an abuse of discretion, or otherwise not in accordance with law." 5 U.S.C. § 706(2)(A). Under this standard, judicial review of agency action seeks to determine whether an agency "has relied on factors which Congress has not intended it to consider, entirely failed to consider an important aspect of the problem, offered an explanation for its decision that runs counter to the evidence before the agency, or is so implausible that it could not be ascribed to a difference in view or the product of agency expertise." *See Motor Vehicle Mfrs. Ass'n v. State Farm Mut. Auto. Ins. Co.*, 463 U.S. 29, 43 (1983); *Utah Environmental Congress v. Bosworth*, 443 F.3d 732, 739 (10th Cir. 2006).

In its DEIS, the CMUG has failed to consider and acknowledge that the proposed alternative is unlikely to control disease transmission and is implausible. Disease could still be a factor for bighorn sheep populations on the CMUG, regardless of the closure of the Forest to pack goats. The DEIS fails to give importance to the fact that bighorn sheep themselves on the CMUG in fact already carry the pathogens that lead to disease. Thus, bighorn sheep are at risk of contacting other bighorn sheep that carry the pathogens that can lead to disease.

Because bighorn sheep are carriers of the pathogens that can lead to disease, contact with other bighorn sheep not only puts bighorn sheep populations at risk, but renders irrelevant pack goats as the vector for transmission of the pathogens (assuming that pack goats on the CMUG are carriers of the pathogens). This can mislead readers to believe that eliminating risk of contact on the CMUG between pack goats and bighorn sheep will eliminate the threat of disease transmission. Under this misleading premise, the DEIS appears to be designed to depict pack

goats as a significant cause of disease transmission or even risk of disease transmission, which is not accurate. The alternatives and the discussion in the DEIS must acknowledge more fully the potential futility of alternatives and explain the need for more comprehensive solutions to the problem of disease transmission, such as the development of a vaccine, or the improvement of bighorn sheep immunity, or the improvement of bighorn sheep habitat.

The DEIS also fails to consider that other animals on the CMUG, like elk, deer, birds, etc., may carry the pathogens that can lead to diseases. Thus, contact between cattle and other animals, besides pack goats, and bighorn sheep may lead to disease transmission on the CMUG. The DEIS does not discuss this possibility. In addition, the DEIS fails to acknowledge that bighorn sheep are at risk of contact with domestic sheep and other animals off the areas controlled by the CMUG, and which risk is not mitigated by the alternatives or the ban on pack goat use.

Because the DEIS wholly fails to consider the risks of disease transmission from other bighorns, the risks of disease transmission off the Forest, and risks of disease transmission from other sources, the DEIS is inadequate under NEPA. As a result, the DEIS must be revised to consider risks of disease transfer from other bighorns, off of the forest and from other sources.

10. The DEIS Does Not Properly Address the Relevance of Unavailable or Incomplete Scientific Information.

The CMUG DEIS lacks complete information to assess the potential effects of disease transmission between domestic sheep and domestic goats and bighorn sheep, let alone pack goats and bighorn sheep. The DEIS does little to address the lack of information with its subsequent conclusions.

In situations such as this, where the relevant information for assessing impacts is incomplete or unavailable, the agency preparing the EIS must take the following steps: first, if the incomplete information relevant to reasonably foreseeable adverse effects is essential to a reasoned choice among alternatives and the overall costs of obtaining the information is not exorbitant, the agency must include that information in the EIS. Next, if the relevant information cannot be obtained because the overall costs are exorbitant or the means of obtaining the information are not known, then an agency must include in an EIS:

- (1) a statement that such information is incomplete or unavailable;
- (2) a statement of the relevance of the incomplete or unavailable information to evaluating reasonably foreseeable significant impacts on the human environment; (3) a summary of existing credible scientific evidence which is relevant to evaluating the reasonably foreseeable significant adverse impacts on the human environment; and (4) the agency's evaluation of such impacts based upon theoretical approaches or research methods generally accepted in the scientific community.

40 C.F.R. § 1502.22(b).

Here, the Forest Service fails to take these required steps to address the incomplete or unavailable information relevant to ascertaining the possibility and consequences of disease transmission between domestic sheep/domestic goats and bighorns, and further fails to do so pertaining to the lesser risks of disease transmission from domestic goats in comparison to domestic sheep. The evidence provided in the DEIS suggests the likelihood or risk of disease transfer is largely specific to domestic sheep and not to domestic goats. The DEIS fails here to include contrasting scientific points of view that have studied the differences in disease transfer risk between domestic sheep and bighorns, and domestic goats and bighorns. Here also, the DEIS fails to distinguish relevant information pertaining to disease transfer between other free ranging animals as comparable to easily managed and controlled animals like pack goats. Likewise, the DEIS fails to contain a clear and direct statement that the required information is incomplete or unavailable. The DEIS also fails to discuss the relevance of incomplete or unavailable information in light of evaluation of a reasonably foreseeable environmental impact. Lastly, the DEIS fails to contain the Forest Service's own evaluation of such impacts "based upon theoretical approaches or research methods generally accepted in the scientific community." *Id.*

Instead of honestly evaluating the range of potential scientific opinion applicable to disease transmission between pack goats and bighorns, the Forest Service impermissibly fails to comply with the requirements of the CEQ regulations to address incomplete or unavailable scientific information. Based on this fundamental flaw in the evaluation of environmental consequences in the DEIS, the DEIS should be revised to provide further analysis.

11. The CMUG must Obtain Additional Information for the DEIS.

When particular information "relevant to reasonably foreseeable significant adverse impacts is essential to a reasoned choice among alternatives," the agency must obtain that information and include it in the EIS, unless the cost is "exorbitant or the means to obtain it are not known." 40 C.F.R. § 1502.22. If obtaining the information is too costly or infeasible, the agency can forego its collection, providing full explanation in the EIS. *Id.* § 1502.22(b). "In that case the agency must include in the EIS: (1) A statement that the information is incomplete or unavailable; (2) a statement of the relevance of the incomplete or unavailable information; (3) a summary of relevant "existing credible scientific evidence;" and (4) the agency's evaluation of impacts based on "theoretical approaches or research methods generally accepted in the scientific community." *Id.*

The CMUG has not included the following relevant information in the DEIS:

- Information indicating the differences between pack goats and other domestic goats;
- Information indicating that pack goats carry disease that can be transmitted to bighorn sheep;
- Information indicating that pack goats may come into contact or have come into contact with bighorn sheep on the CMUG;

- Information indicating that BMPs and/or mitigation measures are not effective to ensure separation between pack goats and bighorn sheep on the CMUG;
- Information indicating that pack goats may transmit or have transmitted disease to bighorn sheep on the CMUG;
- Information indicating that bighorn sheep have contracted disease transmitted by pack goats on the CMUG;
- Information indicating that bighorn sheep that have contracted disease transmitted by pack goats on the CMUG have returned to their herds and infected other bighorn sheep;
- Information indicating that bighorn sheep that have contracted disease transmitted by pack goats on the CMUG have returned to their herds and infected other bighorn sheep, which has led to a die-off;
- Information indicating that there is a risk of disease transmission from pack goats to bighorn sheep on the CMUG;
- Information indicating the risk of disease transmission from other animals on and off of the CMUG to bighorn sheep;
- Information indicating the recreational, social and economic impacts on goatpackers of a closure of all or part of the CMUG to pack goats.

EXHIBIT A

NAPgA Best Management Practices (BMP'S)

The BMP document is a living document which is open to editing and updating as needed.

NAPgA created the BMP's to establish responsible common sense guidelines for goatpacking. They are not intended to be overly restrictive or to discourage packgoat use in any way or in any location.

NAPgA will use best available science as a guide in which to measure and develop the BMP's to address wildlife and other resource concerns.

BMP#1: Individually Identify Your Packgoats

Each packgoat shall be individually identified. Each goat shall have a collar with a tag attached to it containing, at a minimum, the current owner's name and phone number.

Packgoats may be identified with a tattoo or microchip which is specific to each individual goat in conjunction with a collar.

Tattoos containing the individual packgoat's Scrapie Herd Number & ID or an official Scrapie ear tag may be used in conjunction with a collar.

BMP#2: Control

All packgoats shall be under direct human supervision at all times. They shall be on leads or have leads attached to their collar/halter.

In camp all packgoats shall be in direct sight or tethered in some fashion (picketing, high lining, etc.).

All packgoats shall be tethered at night within 30 feet of humans and bells will be attached to their collars.

BMP#3: Separation

Goatpackers shall minimize packgoat contact with wildlife.

BMP#4: Lost Packgoat

If a packgoat becomes lost every effort will be exhausted to locate and recover it.

If the owner is unable to locate and recover the lost packgoat the following agencies shall be contacted by telephone as soon as possible.

Information given should include a detailed description of the packgoat (size; color; ears erect, hanging or none, horned or not), any equipment they are carrying and the last known location. A photograph of the packgoat, if possible.

The local County Sheriff's office. Call 911 or the non-emergency line to dispatch of that county. Most hikers, hunters, land owners or citizens will call the sheriff's office first if they find a lost pack stock animal.

The state's Department of Fish and Game or Fish.

The local land management agency responsible for the area where the packgoat was lost. (Forest Service/BLM/DNR).

Post information, including photos if available, at convenience stores, trail heads and camp grounds with owners contact information, goat and gear descriptions.

Contact the North American Pack Goat Association (NAPgA) to report the loss. NAPgA will maintain a documentation file on all lost pack goats. NAPgA will request an initial report as well as an after-action report from the packgoat's owner/user. The information will be used for documentation as well as continued training and educational awareness training for pack goat users.

Contact the North American Pack Goat Association (NAPgA) to report the loss. NAPgA will maintain a documentation file on all lost pack goats. NAPgA will request an initial report as well as an after-action report from the packgoat's owner/user. The information will be used for documentation as well as continued training and educational awareness training for pack goat users.

BMP#5: Leave No Trace

Leave No Trace principles are strongly encouraged.

Leave No Trace principles are found on this website: <https://Int.org/learn/7-principles>

EXHIBIT B

Packgoats and *Mycoplasma ovipneumoniae* Prevalence Study 2016 North American Packgoat Association Summary of Understanding

Mycoplasma ovipneumoniae, often referred to by the nickname “Movi” (or some variation of that) is the pathogen currently believed to be the most likely primary cause of outbreaks of bighorn sheep pneumonia that have threatened recovery of that species. On November 10, 2015 information was presented at The Technical Packgoat Meeting to NAPgA and the Blue Mountain Forest Plan Revision team in Pendleton, Oregon that goats had a 90% prevalence rate of *M. ovipneumoniae*. In clarifying this information Dr. Tom Besser noted in an email Dec 15, 2015 that this information was obtained from a “report of a large US survey of sheep operations tested for MOVI”. Domestic goats are different than domestic sheep and most certainly packgoats are very different from domestic sheep on public lands grazing allotments.

To consider packgoats the same as sheep for purposes of analyzing the risk of disease (pathogen) transmission to bighorn sheep is in error. Packgoat owners train packgoat prospects from a young age. Packgoats are inextricably bonded to their owner, which represents the “alpha goat” in their small herd. The lifestyle and care of a packgoat in herds of 2 to 10 differs greatly from that of a typical herd of domestic sheep or goats which can range in size of hundreds to thousands. Packgoats are seen by their owners as a significant investment in time and resources for 3 or 4 years before they are viable for packing purposes. Throughout a packgoat’s life, the packgoat receives routine veterinary care in order to keep the goat healthy and prolong their useful life.

Available literature at the time of this 2015 meeting quoted decades-old science in its discussion of evidence for “disease transmission” from domestic goats to BHS. There was no, and to date remains no, scientific support to implicate packgoats in BHS die-offs. Goats and sheep are different species and the scientific data from captive commingling experiments concerning pathogen (*M. ovipneumoniae* or other historically examined pathogens, such as members of the Pasteurellaceae family of bacteria) transmission to bighorn sheep and subsequent disease is vastly different. The types of *M. ovipneumoniae* carried by domestic sheep differ genetically from those carried by domestic goats (Maksimovic, Cassirer, unpublished data). Goat types or “strains” of *M. ovipneumoniae* have resulted in relatively mild (non-fatal) respiratory illness, dramatically different than the nearly 100% fatality reported from captive commingling with domestic sheep. To group sheep and goats together, and even packgoats and other types of domestic goats, in the discussion of pathogen or disease transmission falsely implicates packgoats in BHS die-off’s.

In more recent research by Besser *et al.* (2016), not a single domestic goat or bighorn sheep succumbed to any sort of pneumonia before or after being infected with a “goat type” of *M. ovipneumoniae* and not a single animal died as a result of disease during the study. Domestic goats were not shown to cause deaths of bighorn sheep as a result of pathogen (“disease”) transmission, even when the 3 study goats, were inoculated/infected with a “goat type” of *M. ovipneumoniae* and forced to commingle with bighorn sheep for 100 days. All animals in the study, both the domestic goats and bighorn sheep began showing symptoms of respiratory illness, and all of them recovered prior to being euthanized by the researchers. While the publication would imply that “sub-lethal pneumonia” was

induced in the bighorn sheep in this study, this is not consistent with the histopathology reports from lung tissue that was submitted to the Washington Animal Disease Diagnostic Laboratory in Pullman, WA. Those reports indicated that there were minimal to mild changes that are typically seen in small ruminants that are infected with *M. ovipneumoniae* (bronchiolar associated lymphoid tissue (BALT) hyperplasia and hyperplasia of the bronchial/bronchiolar epithelium); but no diagnosis of pneumonia was reported.

NAPgA is the leading organization in making recommendations on how to safely recreate with packgoats around BHS habitat. The complete lack of relevant research regarding *M. ovipneumoniae* prevalence in packgoats lead NAPgA to contact the USDA - Agricultural Research Unit - Animal Disease Research Unit (ARDU) in December of 2015. ARDU and APHIS (Animal and Plant Health Inspection Service) developed a packgoat *M. ovipneumoniae* surveillance research project.

In the spring of 2016 NAPgA recruited packgoat owners to participate in this research project. Consent was obtained from each packgoat owner. The majority of samples were collected by APHIS personnel and the remainder by Margaret Highland, DVM, PhD, Dipl. ACVP. Duplicate swabs were collected by both APHIS personnel and Dr. Highland. One swab was tested in the ARDU-ARS-USDA laboratory and the other was tested in the Washington Animal Disease Laboratory (except for kids <6 months of age and some of the non-packers that were also tested, which were tested only in the USDA-ARS-ARDU laboratory, as a means to save on research funds, since these animals are not used for packing).

A **packgoat owner survey** was completed. Information obtained was as follows:

- Goat information: Age, Sex, Breed
- Number of goats on premises (packers, non-packers)
- Illness(es) within the last year, including pinkeye/respiratory disease
- Any recent (last month) use of antibiotics
- Vaccination and antiparasitic regimen
- Use of packgoats on public lands? Proximity to bighorn sheep?

Samples collected (spring-fall 2016)

Packgoats

- 3 sets of duplicate nasal swabs collected at 4 week minimum intervals (few premises had only 1 or 2 sample collections)
- 1 blood sample for serum
- Other goats (milkers/breeders/etc) on premises were also tested
- At a minimum, 1 or 2 nasal swabs collected, at 1 to 3 time points
- Not all premises had “non-packer” goats on premises sampled
- All samples processed within 72 hours of collection

Sample Testing

- Nasal Swab samples tested by PCR and/or qPCR; positive samples confirmed by DNA sequencing
- PCR = polymerase chain reaction = technique that amplifies a segment of the bacteria’s genome to determine if it is present
- Duplicate nasal swabs from the first sample collection submitted to the Washington Animal Disease Diagnostic Laboratory (qPCR analysis)
- Serum samples are currently banked frozen

Distribution

State	#premises	#packgoats	# other goats	Total
AZ	3	16	23	39
CA	6	16	42	58
CO	8	29	12	41
ID	25	101	35	136
KS	1	13	51	64
MT	5	21	6	27
NM	1	2	0	2
NV	2	8	0	8
OR	9	32	3	35
UT	5	34	2	36
WA	14	65	17	82
WY	4	40	3	43
Total	83	377	194	571

“Other goats” = milkers, bucks, kids under 4 months of age which would not be out packing or on long hikes

WADDL Test Results

# Goats Tested	Detected	Indeterminate *	Not Detected
485 (83premises)	18 (5 premises)	20 (9 premises, 3 overlapped with the detected premises)	474 (72 premises)
	3.7% (6.0%premises)	4.1% (10.8%premises)	92.2% (86.7%premises)

* Indeterminate indicates that either there was an extremely low number of *M. ovipneumoniae* present in the sample OR the sample is truly negative, and the low detection is a false positive

WADDL Laboratory Test Results

NAPgA believes the large number of samples tested by the AAVLD accredited state diagnostic laboratory (WADDL) provide sufficient and valid evidence as to the very low prevalence of *M. ovipneumoniae* in packgoats.

ADRU-ARS-USDA Laboratory Results

8.2%, or 47, of all goats tested (n=571) had at least 1 sample in which *M. ovipneumoniae* was detected. Twenty-six of the positive animals were ≤4 months old, 35 were ≤12 months, and when considering only the “packers”, 3.3% overall had *M. ovipneumoniae* detected on at least 1 sample collection. 10 of the 14 premises with at least 1 positive detection were premises reported to house kids or were a premises in which the packgoat(s) were in recent contact with a positive packgoat or kids from a positive premises. These results have not yet been published in a peer-reviewed venue. Overall NAPgA will provide the complete report after peer-reviewed publication.

This is a living document and will be updated as new scientific evidence-based information is available.

From: Highland, Margaret
Sent: Friday, May 05, 2017 9:59 AM
To: 'Steve Kilpatrick' <skilpatrick@wyomingwildsheep.org>; 'Ron Smith' <rsagebrushsmith@aol.com>; canyonshadows@wyoming.com; johnmionne@gmail.com; packgoat@icloud.com; ctrulock@fs.fed.us; sschacht@fs.fed.us; brandonjhouck@fs.fed.us; rvandervoet@blm.gov; Lander_WYMail@blm.gov; daryl.lutz@wyo.gov; pat_hnilicka@fws.gov; sara@bighorn.org
Cc: 'Knowles, Don (dknowles@vetmed.wsu.edu)' <dknowles@vetmed.wsu.edu>
Subject: RE: Pack Goat Meeting rescheduled

Since this may not occur before a final decision is made on the Shoshone NF, I would like to share with this group the data from the large scale pack goat study that was performed in 2016. While the ocular swabs are now and finally being tested after developing and validating PCR assays for detecting the 4 most common bacterial agents of pink eye (this process was much slower than anticipated by me), the *Mycoplasma ovipneumoniae* results are completed. The following, in quotes, is an email that I shared with Jim Wilder on 12/16/17. Since then we have retested all of the pack goat nasal swabs a 3 time with a more sensitive standard PCR method, the update on the findings from this follow the email correspondence.

“Over the last year we (ADRU-ARS-USDA), in collaboration with APHIS, were able to complete a fairly large scale surveillance study testing nasal shedding/presence of *Mycoplasma ovipneumoniae* in pack goats. We also tested goats that were housed with or on the same premises as domestic goats that were reported by the owner to be used specifically for packing. We also collected ocular swabs from participating goats to test for the presence of the common agents of small ruminant pink eye (*Chlamydophila* sp and *Mycoplasma conjunctivae*, *Moraxella ovis*, and *Acheloplasma oculi*); the ocular swabs are still being analyzed, with hopes of completing analysis this month. Upon analysis completion of the ocular swabs, the plan is to report the results by publishing in a peer-reviewed scientific journal by the end of winter/early spring.

I would like to share with you the following results from the nasal swab samples that were collected:

Nasal swabs were collected 3 times, at 1 month minimum intervals, from participating goats (aside from the handful of animals that were sold, removed from the study as per the owners discretion, or entered into the study late so had fewer sample time points). A couple of the premises had 4 or 5 samples collected. Duplicate nasal swabs were collected at each time point. 1 swab was tested in our USDA laboratory and samples that tested negative were then submitted to an independent laboratory for confirmation of the results (WADDL in Pullman, WA was the independent laboratory).

We tested a total of 576 domestic goats from 84 premises which included the following states (# of premises in parentheses after each): AZ (3), CA (6), CO (7), ID (26), KS (1), MT (5), NM (1), NV (2), OR (9), UT (5), WA (14), WY (4), VT (1). (I believe I had reported that there were 88 premises in earlier info that I shared with Mark P.I forgot to deduct the 4 premises scattered in 4 eastern states that we didn't get tested).

Of all of the premises tested, we confirmed *M. ovipneumoniae* to be present in nasal

secretions from goats on 2 premises, limited to kids ≤ 2 weeks of age at only one test time. We collected additional swabs from 1 of these premises for 5 times total sample collections and the last 3 collection points had no detected *M. ovipneumoniae* and interestingly, all of the adult goats (9 of them) never had *M. ovipneumoniae* detected....the kids (there were 15 of them total) had 3 positives at time point 1, and 2 different kids positive at time point 2 (1st 3 positive were negative at this 2nd time point) and all goats on the premises were negative the last 3 sample collections.

As for the other premises that had a handful of positive kids: I repeat swabbed several of them 1 or 2 more times and they too were subsequently negative on the repeat samplings. This “kid phenomenon” is interesting.....I’ll leave it at that as to save typing time in this already lengthy email, but am happy to discuss further some time if you are interested. One additional premises that had *M. ovipneumoniae* detected 2 of the 3 sample times had a small group of yearling pack goats that were being housed at fence line with an ‘open’ breeding herd of registered Boer goats that were used for shows and sent out to farms for sire purposes. I instructed that owner to move his packers as soon as possible away from the large group of traveling Boer goats.....I suspect that his pack goats may clear (not shed) *M. ovipneumoniae* without the constant potential exposure, as all of his goats were negative on the 3 sample collection (I’d be happy to discuss why I suspect this may be possible with you too, if you’re interested).

The other 81 premises had no confirmed *M. ovipneumoniae* present on any of the nasal swabs collected. Of interest to your local and nearby area, none of the WY, UT, CO, MT herds had confirmed *M. ovipneumoniae* detection at any of the time points. 1 of the places with “kid detected *M. ovipneumoniae*” was in ID, but these kids are the ones that have sense been negative and the adults never positive.

While nothing is ever 100% risk free in life, I think this data strongly supports that there is a very low prevalence of *M. ovipneumoniae* in goats, at least those raised and kept in closed and typically small groups (however, a few of the premises that I tested had 20+ goats though and still negative....even the premises that tested their milk goats).

I would also like to take the time here to give warning that unless researchers and/or diagnosticians are looking beyond the common published techniques for identifying *M. ovipneumoniae*, there is a chance that false positive results will occur...particularly in goats. For example, we know that the published PCR primers, referred to as “LM primers” and qPCR techniques that have been developed in the past based on these primers can (and do) result in false positive results. By “looking beyond” I mean perform standard PCR to amplify a minimum of 2 regions of the bacterial genome and sequence the products/amplicons.....and making sure that the products/amplicons match well-characterized strains of *M. ovipneumoniae* (ie. strains that are characterized by reputable groups such as ATCC). Mycoplasmas are tricky, to say the least. Again, I’m happy to discuss more should you be interested.

Please feel free to let me know, either by email or phone (listed in signature line), if you have questions, comments, or concerns about the information provided herein or if you have anything that you would like to further discuss with me regarding the bighorn pneumonia phenomenon.”

Update following repeated testing using a more sensitive method of detection:
Five of the 83 premises tested (6%) had *M. ovipneumoniae* identified during the repeat nasal sample collections. Premises that had *M. ovipneumoniae* detected in any the goats

had at least 7 goats housed on the premises. *M. ovipneumoniae* was confirmed to be present on the nasal swabs collected from 30 of the 576 total goats tested, meaning that 94.8% of the goats tested had no *M. ovipneumoniae* detected at any of the sample collection time points. Of the 30 total *M. ovipneumoniae* positive goats, 27 (or 90%) of the were ≤ 1 year of age, and 23 of them were < 5 months of age.

During the 2016 North American Pack goat annual gathering (“the Rendy”) held in Oregon, I sampled in total 27 adults and 2 kid goats whose owners brought them to the sample collection site that I set up. Most of these goats were already part of the large pack goat/domestic goat surveillance study and I asked owners if they minded me taking an extra nasal swab from their animals with the thought that perhaps the stressor of travelling or bringing a large group of goats together may result in shedding of *M. ovipneumoniae* from animals that it hadn’t been detected on during the first round of sample collections and it also gave the opportunity to add a couple more premises to the study. *M. ovipneumoniae* **was not detected** on any of the swab samples collected at the Rendy.

It’s unfortunate how long research takes, particularly with something as time sensitive as this seems to be, as I had truly hoped that this entire study would be out in published in a peer-reviewed form at this point (April was my goal). Hoping now for June with fingers crossed that all of the ocular swab testing goes smoothly....and more importantly accurately with good specificity and sensitivity.

Thank you and I look forward to participating in the Pack Goat meeting whenever the final date is decided upon.

Maggie

Margaret A. Highland, DVM, PhD, Dipl. ACVP
Animal Disease Research Unit-ARS-USDA (VMO Researcher)
Washington Animal Disease Diagnostic Laboratory (Adjunct Pathologist)
School for Global Animal Health (Adjunct Faculty)
Washington State University
Pullman, WA 99164

Office phone: 509-335-6327
Cell phone: 608-213-3025
Fax: 509-335-8328

**ACCESSION FORM FOR GENERAL DIAGNOSTICS
Washington Animal Disease Diagnostic Laboratory**

College of Veterinary Medicine, Washington State University

Web Site: <http://waddl.vetmed.wsu.edu>

US Postal Service mailing address:
PO Box 647034
Pullman, WA. 99164-7034

UPS, FedEx or Courier shipping address:
Bustad Hall, Rm.155-N
Pullman, WA. 99164-7034

Phone: (509) 335-9696
FAX: (509) 335 7424
E-Mail: waddl@vetmed.wsu.edu

2016-6030
Ref Vet: Highland, Margaret
Owner: USDA - ARS - ADRU
Breed: Domestic Goat
Routed: jmd

05/10/16
Form 2 pages

Please type or use black ink and print clearly.

Veterinarian or Case Coordinator: Name: Highland		First Name: Maggie	
Clinic: ADRU-ARS-USDA			
Street address: ADBF-WSU		Mailing Address or PO Box:	
City: Pullman	State: WA	Zip: 99164	
Phone: 509-335-6327	Fax: 509-335-8328	E-mail: mah@vetmed.wsu.edu	
Owner: Last Name first: same as above		Guardian Name: (if owner is under 18)	
Farm Name:		First Time Submitter? <input type="checkbox"/> Yes <input type="checkbox"/> No	
Street address:		Mailing Address or PO Box:	
City:	State:	Zip:	
Phone:	Fax:	E-mail:	

Billing: Owner Clinic 3rd Party (preapproval required) Please note: WADDL policy is to bill the clinic if provided, unless-prepaid.
Reporting Preference: Mail Fax Web access - register on web site at <http://waddl.vetmed.wsu.edu>

Please fill out completely as possible:

Specimen(s) Submitted:		Date Collected: <i>April 2016</i>	
(Please use WADDL Animal ID Sheet for multiple animals.)		Date Shipped:	
nasal swabs			
Tests Requested:	<input type="checkbox"/> Necropsy	<input type="checkbox"/> Virology	<input type="checkbox"/> Bacteriology
	<input type="checkbox"/> Histopathology	<input type="checkbox"/> Serology	<input type="checkbox"/> Mycoplasma culture
	<input type="checkbox"/> Toxicology	<input type="checkbox"/> Fungal Culture	<input type="checkbox"/> Parasitology
			<input type="checkbox"/> IHC
			<input checked="" type="checkbox"/> PCR
			<input type="checkbox"/> Other: _____
Note: WADDL reserves the right to modify the tests requested for more efficient case work-up and / or to send specimens to outside laboratories to perform testing not done at WADDL.			
Animal ID (name/tag#)	Species	Breed	Age
see multiple animal form	goat	multiple	1mo-12yrs
Sex	Animal Weight	Duration of Problem	
N/A	N/A	N/A	

* Was animal euthanized? If so, what method? *N/A*

Additional History: *Vaccinations, signs, stress factors, treatments, post mortem findings, pertinent feed or feed additives, clinical lab results, previous WADDL Case Numbers. (Attach additional sheets as necessary.)*

Please save any remaining DNA isolations and call Maggie for pick up.

Bill to ADRU-ARS-USDA acct #RSA 2540-1080

Samples were maintained on ice then frozen w/in 2 days of collection + kept at -20°C since.

WADDL is an official brucellosis testing laboratory. All serology for brucellosis, including abortion screens, requires identification of animals, date of sample collection, and signature of an accredited veterinarian attesting to the following statement:
"I certify that the specimens submitted with this form were collected by me from the animal(s) described on the date indicated."

Veterinarian's, Clinician's or Owner's Signature:	Condition(s) Suspected:
---------------------------------------------------	-------------------------

IDENTIFICATION SHEET FOR MULTIPLE ANIMALS

(To accompany WADDL Accession form, if needed)

Washington Animal Disease Diagnostic Laboratory
 College of Veterinary Medicine, Washington State University
 Mailing address: P.O. Box 647034
 Pullman, WA. 99164-7034
 Phone: (509) 335-9696
 E-Mail: waddl@vetmed.wsu.edu
 Web Site: http://waddl.vetmed.wsu.edu

Shipping address:
 Bustad Hall, Rm. 155-N
 Pullman, WA. 99164-7034
 FAX: (509) 335-7424

2016-6030
 Ref Vet: Highland, Margaret
 Owner: USDA-ARS-ADRU
 Breed: Domestic Goat
 Routing: .md

Owner: ADRU-ARS-USDA

Veterinarian: Maggie Highland

TEST(S) REQUESTED: Mycoplasma ovipneumoniae qPCR

Tube	Animal # or Name	Tube	Animal # or Name	Tube	Animal # or Name	Tube	Animal # or Name
1	3_A	26	5_F	51	3_D	76	
2	3_B	27	5_G	52	7_A	77	
3	3_C	28	5_H	53	7_B	78	
4	11_A	29	6_A	54	7_C	79	
5	11_B	30	6_B	55	7_D	80	
6	11_C	31	6_C	56	7_E	81	
7	11_D	32	6_D	57	11_A	82	
8	16_A	33	6_E	58	11_B	83	
9	16_B	34	6_F	59	11_C	84	
10	4_A	35	6_G	60	11_D	85	
11	4_B	36	8_A	61	12_A	86	
12	4_C	37	8_B	62	12_B	87	
13	4_D	38	8_C	63	12_C	88	
14	4_E	39	8_D	64	12_D	89	
15	4_F	40	9_A	65	12_E	90	
16	4_G	41	9_B	66	12_F	91	
17	10_A	42	9_C	67	12_G	92	
18	10_B	43	2_A	68	12_H	93	
19	10_C	44	2_B	69	12_I	94	
20	10_D	45	2_C	70	12_J	95	
21	5_A	46	2_D	71	12_K	96	
22	5_B	47	2_E	72	12_L	97	
23	5_C	48	3_A	73		98	
24	5_D	49	3_B	74		99	
25	5_E	50	3_C	75		100*	

05/10/16

* For over 100 samples, please copy this form and continue numbering.

Washington Animal Disease Diagnostic Lab

P.O. Box 647034
Pullman, WA 99164-7034
Telephone : (509) 335-9696
Fax : (509) 335-7424

Dr. Margaret Highland
USDA-ARS-ADRU
WSU - 3003 ADBF

Case#: 2016-6030
Report Date: 05/16/16

Pullman, WA 99164-6630

Submittal Date: 05/10/16
Owner: USDA-ARS-ADRU

Species: Domestic Goat

Age:
Sex:

Final Report:

Molecular Diagnostics- Reported on 05/16/16 Authorized by Daniel Bradway, Lab Manager

PCR-Mycoplasma ovipneumoniae SOP: 501.40RT.2016.03.17

Animal	Specimen	Result
[REDACTED] 3_A	Nasal swab	Not detected
[REDACTED] 3_B	Nasal swab	Not detected
[REDACTED] 3_C	Nasal swab	Not detected
[REDACTED] 11_A	Nasal swab	Not detected
[REDACTED] 11_B	Nasal swab	Not detected
[REDACTED] 11_C	Nasal swab	Not detected
[REDACTED] 11_D	Nasal swab	Not detected
[REDACTED] 16_A	Nasal swab	Not detected
[REDACTED] 16_B	Nasal swab	Not detected
[REDACTED] 4_A	Nasal swab	Detected
[REDACTED] 4_B	Nasal swab	Detected
[REDACTED] 4_C	Nasal swab	Detected
[REDACTED] 4_D	Nasal swab	Detected
[REDACTED] 4_E	Nasal swab	Detected
[REDACTED] 4_F	Nasal swab	Detected
[REDACTED] 4_G	Nasal swab	Detected
[REDACTED] 10_A	Nasal swab	Indeterminate
[REDACTED] 10_B	Nasal swab	Not detected
[REDACTED] 10_C	Nasal swab	Not detected
[REDACTED] 10_D	Nasal swab	Not detected
[REDACTED] 5_A	Nasal swab	Not detected
[REDACTED] 5_B	Nasal swab	Not detected
[REDACTED] 5_C	Nasal swab	Not detected
[REDACTED] 5_D	Nasal swab	Not detected
[REDACTED] 5_E	Nasal swab	Not detected
[REDACTED] 5_F	Nasal swab	Not detected
[REDACTED] 5_G	Nasal swab	Not detected
[REDACTED] 5_H	Nasal swab	Not detected

Washington Animal Disease Diagnostic Lab

PCR-*Mycoplasma ovipneumoniae* SOP: 501.40RT.2016.03.17

Animal	Specimen	Result
V.6.A	Nasal swab	Not detected
V.6.B	Nasal swab	Not detected
V.6.C	Nasal swab	Not detected
V.6.D	Nasal swab	Not detected
V.6.E	Nasal swab	Not detected
V.6.F	Nasal swab	Not detected
V.6.G	Nasal swab	Not detected
V.8.A	Nasal swab	Not detected
V.8.B	Nasal swab	Not detected
V.8.C	Nasal swab	Not detected
V.8.D	Nasal swab	Not detected
V.9.A	Nasal swab	Not detected
V.9.B	Nasal swab	Not detected
V.9.C	Nasal swab	Not detected
V.12.A	Nasal swab	Not detected
V.12.B	Nasal swab	Not detected
V.12.C	Nasal swab	Not detected
V.12.D	Nasal swab	Not detected
V.12.E	Nasal swab	Not detected
V.13.A	Nasal swab	Not detected
V.13.B	Nasal swab	Not detected
V.13.C	Nasal swab	Not detected
V.13.D	Nasal swab	Not detected
V.17.A	Nasal swab	Not detected
V.17.B	Nasal swab	Not detected
V.17.C	Nasal swab	Not detected
V.17.D	Nasal swab	Not detected
V.17.E	Nasal swab	Not detected
V.11.A	Nasal swab	Not detected
V.11.B	Nasal swab	Not detected
V.11.C	Nasal swab	Not detected
V.11.D	Nasal swab	Not detected
V.12.A	Nasal swab	Not detected
V.12.B	Nasal swab	Not detected
V.12.C	Nasal swab	Not detected
V.12.D	Nasal swab	Not detected
V.12.E	Nasal swab	Not detected
V.12.F	Nasal swab	Not detected
V.12.G	Nasal swab	Not detected
V.12.H	Nasal swab	Not detected
V.12.I	Nasal swab	Not detected
V.12.J	Nasal swab	Not detected
V.12.K	Nasal swab	Not detected
V.12.L	Nasal swab	Not detected

PCR-*Mycoplasma ovipneumoniae* test comment: This assay detects only *Mycoplasma ovipneumoniae*. Culture is available at WADDL to detect other species of *Mycoplasma* if desired. Fees for culture are available on our website. Please contact the lab if *Mycoplasma* culture or other testing is desired.

Quantity/Description/Routing of Samples

72 nasal swabs

2016-6030
Ref Vet: Highland, Margaret
Owner: USDA-ARS-ADRU
Breed: Domestic Goat
Router: ,md

Sample Condition:	<input type="checkbox"/> Room Temp.	<input type="checkbox"/> On ice	<input checked="" type="checkbox"/> Frozen	<input type="checkbox"/> Fixed	Contents match forms: <input type="checkbox"/> Yes <input type="checkbox"/> No Explain below: <i>not</i>	Opened by:
	Samples Received Via:					
	<input type="checkbox"/> US Mail	<input type="checkbox"/> FedEx	<input checked="" type="checkbox"/> Drop off			
	<input type="checkbox"/> UPS	<input type="checkbox"/> FedEx-R	<input type="checkbox"/> Other:			

Comments for Case Tracking:

by Margaret Highland



05/10/16
page 1 of 1

Sample Label

not

ACCESSION FORM FOR GENERAL DIAGNOSTICS
Washington Animal Disease Diagnostic Laboratory
 College of Veterinary Medicine, Washington State University
 Web Site: <http://waddl.vetmed.wsu.edu>

US Postal Service mailing address:
 PO Box 647034
 Pullman, WA. 99164-7034

UPS, FedEx or Courier shipping address:
 Bustad Hall, Rm. 155-N
 Pullman, WA. 99164-7034

Phone: (509) 335-9696
 FAX: (509) 335 7424
 E-Mail: waddl@vetmed.wsu.edu

2016-6160
 Ref Vet: Highland, Margaret
 Owner: USDA - ARS - ADRU
 Breed: Domestic Goat
 Routed: .ind

Please type or use black ink and print clearly.

Veterinarian or Last Case Coordinator Name: **Highland** First Name: **Maggie**

Clinic: **ADRU-ARS-USDA**

Street address: **ADBF-WSU** Mailing Address or PO Box:

City: **Pullman** State: **WA** Zip: **99164**

Phone: **509-335-6327** Fax: **509-335-8328** E-mail: **mah@vetmed.wsu.edu**

Owner: Last Name first: **same as above** Guardian Name: (if owner is under 18)

Farm Name: First Time Submitter? Yes No

Street address: Mailing Address or PO Box:

City: State: Zip:

Phone: Fax: E-mail:

Billing: Owner Clinic 3rd Party (preapproval required) Please note: WADDL policy is to bill the clinic if provided, unless prepaid.

Reporting Preference: Mail Fax Web access - register on web site at <http://waddl.vetmed.wsu.edu>

Please fill out completely as possible:

Specimen(s) Submitted: **nasal swabs** Date Collected: **4/16-5/16**
 (Please use WADDL Animal ID Sheet for multiple animals.) Date Shipped:

Tests Requested: Necropsy Virology Bacteriology IHC
 Histopathology Serology Mycoplasma culture PCR
 Toxicology Fungal Culture Parasitology Other:

Note: WADDL reserves the right to modify the tests requested for more efficient case work-up and / or to send specimens to outside laboratories to perform testing not done at WADDL.

Animal ID (name/tag#)	Species	Breed	Age	Sex	Animal Weight
see multiple animal form	domestic goats	multiple	1mo-12yrs		
Location of Lesion	No. in group	No. Dead	No. Sick	No. on Premises	Duration of Problem
N/A					N/A

* Was animal euthanized? If so, what method? **N/A**

Additional History: Vaccinations, signs, stress factors, treatments, post mortem findings, pertinent feed or feed additives, clinical lab results, previous WADDL Case Numbers. (Attach additional sheets as necessary.)

Nasal swabs for *M. ovipneumoniae* qPCR
 Please save any remaining DNA isolations and call Maggie for pick up.
 Bill to ADRU-ARS-USDA acct #RSA 2540-1080

WADDL is an official brucellosis testing laboratory. All serology for brucellosis, including abortion screens, requires identification of animals, date of sample collection, and signature of an accredited veterinarian attesting to the following statement:
 "I certify that the specimens submitted with this form were collected by me from the animal(s) described on the date indicated."

Veterinarian's, Clinician's or Owner's Signature: *Maggie Wood* Condition(s) Suspected: **None/Healthy Animals**

IDENTIFICATION SHEET FOR MULTIPLE ANIMALS

(To accompany WADDL Accession form, if needed)

Washington Animal Disease Diagnostic Laboratory
 College of Veterinary Medicine, Washington State University
 Mailing address: P.O. Box 647034 Pullman, WA. 99164-7034
 Phone: (509) 335-9696 E-Mail: waddl@vetmed.wsu.edu
 Shipping address: Bustad Hall, Rm. 155-N Pullman, WA. 99164-7034
 FAX: (509) 335-7424
 Web Site: <http://waddl.vetmed.wsu.edu>

2016 - 6160
 Ref Vet: Highland, Margaret
 Owner: USDA - ARS - ADRU
 Breed: Domestic Goat
 Routing: .md

05/12/16

Owner: ADRU-ARS-USDA

Veterinarian: Maggie Highland

TEST(S) REQUESTED: M. ovipneumoniae qPCR

Tube	Animal # or Name	Tube	Animal # or Name	Tube	Animal # or Name	Tube	Animal # or Name
1	<u>1_A</u>	26	<u>4_N</u>	51	<u>5_S</u>	76	<u>14_C</u>
2	<u>1_B</u>	27	<u>4_O</u>	52	<u>5_T</u>	77	<u>17_A</u>
3	<u>1_C</u>	28	<u>4_P</u>	53	<u>5_U</u>	78	<u>17_B</u>
4	<u>1_D</u>	29	<u>4_Q</u>	54	<u>5_V</u>	79	<u>17_C</u>
5	<u>1_E</u>	30	<u>4_R</u>	55	<u>5_W</u>	80	<u>17_D</u>
6	<u>1_F</u>	31	<u>4_S</u>	56	<u>5_X</u>	81	<u>17_E</u>
7	<u>1_G</u>	32	<u>4_T</u>	57	<u>5_Y</u>	82	<u>17_F</u>
8	<u>7_A</u>	33	<u>5_A</u>	58	<u>5_Z</u>	83	<u>17_G</u>
9	<u>7_B</u>	34	<u>5_B</u>	59	<u>8_A</u>	84	<u>17_H</u>
10	<u>7_C</u>	35	<u>5_C</u>	60	<u>8_B</u>	85	<u>17_I</u>
11	<u>7_D</u>	36	<u>5_D</u>	61	<u>8_C</u>	86	<u>22_A</u>
12	<u>7_E</u>	37	<u>5_E</u>	62	<u>9_A</u>	87	<u>22_B</u>
13	<u>4_A</u>	38	<u>5_F</u>	63	<u>9_B</u>	88	<u>22_C</u>
14	<u>4_B</u>	39	<u>5_G</u>	64	<u>9_C</u>	89	<u>23_A</u>
15	<u>4_C</u>	40	<u>5_H</u>	65	<u>9_D</u>	90	<u>23_B</u>
16	<u>4_D</u>	41	<u>5_I</u>	66	<u>9_E</u>	91	<u>23_C</u>
17	<u>4_E</u>	42	<u>5_J</u>	67	<u>19_A</u>	92	<u>23_D</u>
18	<u>4_F</u>	43	<u>5_K</u>	68	<u>19_B</u>	93	<u>23_E</u>
19	<u>4_G</u>	44	<u>5_L</u>	69	<u>10_A</u>	94	<u>23_F</u>
20	<u>4_H</u>	45	<u>5_M</u>	70	<u>10_B</u>	95	<u>23_G</u>
21	<u>4_I</u>	46	<u>5_N</u>	71	<u>6_A</u>	96	<u>2_A</u>
22	<u>4_J</u>	47	<u>5_O</u>	72	<u>6_B</u>	97	<u>2_B</u>
23	<u>4_K</u>	48	<u>5_P</u>	73	<u>6_C</u>	98	<u>12_A</u>
24	<u>4_L</u>	49	<u>5_Q</u>	74	<u>14_A</u>	99	<u>12_B</u>
25	<u>4_M</u>	50	<u>5_R</u>	75	<u>14_B</u>	100 *	<u>12_C</u>

* For over 100 samples, please copy this form and continue numbering.

IDENTIFICATION SHEET FOR MULTIPLE ANIMALS

(To accompany WADDL Accession form, if needed)

Washington Animal Disease Diagnostic Laboratory
 College of Veterinary Medicine, Washington State University
 Mailing address: P.O. Box 647034 Pullman, WA. 99164-7034
 Shipping address: Bustad Hall, Rm. 155-N Pullman, WA. 99164-7034
 Phone: (509) 335-9696 FAX: (509) 335-7424
 E-Mail: waddl@vetmed.wsu.edu
 Web Site: http://waddl.vetmed.wsu.edu

2016 - 6160
 Ref Vet: Highland, Margaret
 Owner: USDA - ARS - ADRU
 Breed: Domestic Goat
 Routing: .mid

Owner: ADRU-ARS-USDA

Veterinarian: Maggie Highland

TEST(S) REQUESTED: M. ovipneumoniae qPCR

05/12/16

Tube	Animal # or Name	Tube	Animal # or Name	Tube	Animal # or Name	Tube	Anim
1	<u>12_D</u>	26		51		76	
2	<u>20_A</u>	27		52		77	
3	<u>20_B</u>	28		53		78	
4		29		54		79	
5		30		55		80	
6		31		56		81	
7		32		57		82	
8		33		58		83	
9		34		59		84	
10		35		60		85	
11		36		61		86	
12		37		62		87	
13		38		63		88	
14		39		64		89	
15		40		65		90	
16		41		66		91	
17		42		67		92	
18		43		68		93	
19		44		69		94	
20		45		70		95	
21		46		71		96	
22		47		72		97	
23		48		73		98	
24		49		74		99	
25		50		75		100 *	

* For over 100 samples, please copy this form and continue numbering

Washington Animal Disease Diagnostic Lab

P.O. Box 647034
Pullman, WA 99164-7034
Telephone : (509) 335-9696
Fax : (509) 335-7424

Dr. Margaret Highland
USDA-ARS-ADRU
WSU - 3003 ADBF

Case#: 2016-6160
Report Date: 05/16/16

Pullman, WA 99164-6630

Submittal Date: 05/12/16
Owner: USDA-ARS-ADRU

Species: Domestic Goat

Age:
Sex:

Final Report:

Molecular Diagnostics- Reported on 05/16/16 Authorized by Daniel Bradway, Lab Manager

PCR-Mycoplasma ovipneumoniae SOP: 501.40RT.2016.03.17

Animal	Specimen	Result
.1.A	Nasal swab	Not detected
.1.B	Nasal swab	Not detected
.1.C	Nasal swab	Not detected
.1.D	Nasal swab	Not detected
.1.E	Nasal swab	Not detected
.1.F	Nasal swab	Not detected
.1.G	Nasal swab	Not detected
.7.A	Nasal swab	Not detected
.7.B	Nasal swab	Not detected
.7.C	Nasal swab	Not detected
.7.D	Nasal swab	Not detected
.7.E	Nasal swab	Not detected
.4.A	Nasal swab	Not detected
.4.B	Nasal swab	Not detected
.4.C	Nasal swab	Not detected
.4.D	Nasal swab	Not detected
.4.E	Nasal swab	Not detected
.4.F	Nasal swab	Not detected
.4.G	Nasal swab	Not detected
.4.H	Nasal swab	Not detected
.4.I	Nasal swab	Detected
.4.J	Nasal swab	Indeterminate
.4.K	Nasal swab	Not detected
.4.L	Nasal swab	Not detected
.4.M	Nasal swab	Not detected
.4.N	Nasal swab	Not detected
.4.O	Nasal swab	Indeterminate
.4.P	Nasal swab	Indeterminate

Washington Animal Disease Diagnostic Lab

PCR-Mycoplasma ovipneumoniae SOP: 501.40RT.2016.03.17

Animal	Specimen	Result
4.Q	Nasal swab	Indeterminate
4.R	Nasal swab	Not detected
4.S	Nasal swab	Detected
4.T	Nasal swab	Detected
5.A	Nasal swab	Not detected
5.B	Nasal swab	Not detected
5.C	Nasal swab	Not detected
5.D	Nasal swab	Not detected
5.E	Nasal swab	Not detected
5.F	Nasal swab	Not detected
5.G	Nasal swab	Not detected
5.H	Nasal swab	Not detected
5.I	Nasal swab	Not detected
5.J	Nasal swab	Not detected
5.K	Nasal swab	Not detected
5.L	Nasal swab	Not detected
5.M	Nasal swab	Not detected
5.N	Nasal swab	Not detected
5.O	Nasal swab	Not detected
5.P	Nasal swab	Not detected
5.Q	Nasal swab	Not detected
5.R	Nasal swab	Not detected
5.S	Nasal swab	Not detected
5.T	Nasal swab	Not detected
5.U	Nasal swab	Not detected
5.V	Nasal swab	Not detected
5.W	Nasal swab	Not detected
5.X	Nasal swab	Not detected
5.Y	Nasal swab	Not detected
5.Z	Nasal swab	Not detected
8.A	Nasal swab	Not detected
8.B	Nasal swab	Not detected
8.C	Nasal swab	Not detected
9.A	Nasal swab	Not detected
9.B	Nasal swab	Not detected
9.C	Nasal swab	Not detected
9.D	Nasal swab	Not detected
9.E	Nasal swab	Not detected
19.A	Nasal swab	Not detected
19.B	Nasal swab	Not detected
10.A	Nasal swab	Not detected
10.B	Nasal swab	Not detected
6.A	Nasal swab	Not detected
6.B	Nasal swab	Not detected
6.C	Nasal swab	Not detected
14.A	Nasal swab	Not detected
14.B	Nasal swab	Not detected
14.C	Nasal swab	Not detected
17.A	Nasal swab	Not detected
17.B	Nasal swab	Not detected
17.C	Nasal swab	Not detected
17.D	Nasal swab	Not detected

Washington Animal Disease Diagnostic Lab

PCR-Mycoplasma ovipneumoniae SOP: 501.40RT.2016.03.17

Animal	Specimen	Result
[REDACTED] 17.E	Nasal swab	Not detected
[REDACTED] 17.F	Nasal swab	Not detected
[REDACTED] 17.G	Nasal swab	Not detected
[REDACTED] 17.H	Nasal swab	Not detected
[REDACTED] 17.I	Nasal swab	Not detected
[REDACTED] 22.A	Nasal swab	Not detected
[REDACTED] 22.B	Nasal swab	Not detected
[REDACTED] 22.C	Nasal swab	Not detected
[REDACTED] 23.A	Nasal swab	Not detected
[REDACTED] 23.B	Nasal swab	Not detected
[REDACTED] 23.C	Nasal swab	Not detected
[REDACTED] 23.D	Nasal swab	Not detected
[REDACTED] 23.E	Nasal swab	Not detected
[REDACTED] 23.F	Nasal swab	Not detected
[REDACTED] 23.G	Nasal swab	Not detected
[REDACTED] 2.A	Nasal swab	Not detected
[REDACTED] 2.B	Nasal swab	Not detected
[REDACTED] 12.A	Nasal swab	Not detected
[REDACTED] 12.B	Nasal swab	Not detected
[REDACTED] 12.C	Nasal swab	Not detected
[REDACTED] 12.D	Nasal swab	Not detected
[REDACTED] 20.A	Nasal swab	Not detected
[REDACTED] 20.B	Nasal swab	Not detected

PCR-Mycoplasma ovipneumoniae test comment: This assay detects only Mycoplasma ovipneumoniae. Culture is available at WADDL to detect other species of Mycoplasma if desired. Fees for culture are available on our website. Please contact the lab if Mycoplasma culture or other testing is desired.

Quantity/Description/Routing of Samples

103 nasal swabs

- Dropped off by M. Highland

2016-6160
Ref Vet: Highland, Margaret
Owner: USDA - ARS - ADRU
Breed: Domestic Goat
Routed: .ind

Sample Condition:	<input type="checkbox"/> Room Temp.	<input type="checkbox"/> On ice	<input checked="" type="checkbox"/> Frozen	<input type="checkbox"/> Fixed	Contents match forms: <input checked="" type="checkbox"/> Yes <input type="checkbox"/> No Explain below:	Opened by: <i>WHT</i>
	Samples Received Via:					
	<input type="checkbox"/> US Mail	<input type="checkbox"/> FedEx	<input checked="" type="checkbox"/> Drop off			
	<input type="checkbox"/> UPS	<input type="checkbox"/> FedEx-R	<input type="checkbox"/> Other:			

Comments for Case Tracking:

Sample Label

WHT



**ACCESSION FORM FOR GENERAL DIAGNOSTICS
Washington Animal Disease Diagnostic Laboratory**

College of Veterinary Medicine, Washington State University
Web Site: <http://waddl.vetmed.wsu.edu>

US Postal Service mailing address:
PO Box 647034
Pullman, WA. 99164-7034

UPS, FedEx or Courier shipping address:
Bustad Hall, Rm.155-N
Pullman, WA. 99164-7034

Phone: (509) 335-9696
FAX: (509) 335 7424
E-Mail: waddl@vetmed.wsu.edu

2016 - 7117
Ret Vet: Highland, Margaret
Owner: USDA - ARS - ADRU
Breed: Domestic Goat
Routed: md

Please type or use black ink and print clearly.

Veterinarian or Case Coordinator: Last Name: Highland		First Name: Maggie	
Clinic: ADRU-ARS-USDA			
Street address: ADBF 3033		Mailing Address or PO Box:	
City: Pullman	State: WA	Zip: 99163	
Phone: 5-6327	Fax: 5-8328	E-mail: mah@vetmed.wsu.edu	

Owner: Last Name first: same as above		Guardian Name: (if owner is under 18)	
Farm Name:		First Time Submitter? <input type="checkbox"/> Yes <input type="checkbox"/> No	
Street address:		Mailing Address or PO Box:	
City:	State:	Zip:	
Phone:	Fax:	E-mail:	

Billing: Owner Clinic 3rd Party (preapproval required) Please note: WADDL policy is to bill the clinic if provided, unless prepaid.
Reporting Preference: Mail Fax Web access - register on web site at <http://waddl.vetmed.wsu.edu>

Please fill out completely as possible:

Specimen(s) Submitted:		Date Collected: May 2016	
nasal swabs		Date Shipped:	
<small>(Please use WADDL Animal ID Sheet for multiple animals.)</small>			
Tests Requested:		Mycoplasma ovipneumoniae qPCR	
<input type="checkbox"/> Necropsy	<input type="checkbox"/> Virology	<input type="checkbox"/> Bacteriology	<input type="checkbox"/> IHC
<input type="checkbox"/> Histopathology	<input type="checkbox"/> Serology	<input type="checkbox"/> Mycoplasma culture	<input checked="" type="checkbox"/> PCR
<input type="checkbox"/> Toxicology	<input type="checkbox"/> Fungal Culture	<input type="checkbox"/> Parasitology	<input type="checkbox"/> Other:
<small>Note: WADDL reserves the right to modify the tests requested for more efficient case work-up and / or to send specimens to outside laboratories to perform testing not done at WADDL.</small>			
Animal ID (name/tag#)	Species	Breed	Age
see multiple animal form	domestic goats	-	adult
Sex	Animal Weight		
-	-		
Location of Lesion	No. in group	No. Dead	No. Sick
N/A	N/A	N/A	N/A
No. on Premises			
N/A			
Duration of Problem			
N/A			

* Was animal euthanized? If so, what method?

Additional History: Vaccinations, signs, stress factors, treatments, post mortem findings, pertinent feed or feed additives, clinical lab results, previous WADDL Case Numbers. (Attach additional sheets as necessary.)

M. ovipneumoniae qPCR
Please save remaining DNA isolations and call Maggie for pick up.

Bill to ADRU-ARS-USDA acct #RSA 2540-1080

WADDL is an official brucellosis testing laboratory. All serology for brucellosis, including abortion screens, requires identification of animals, date of sample collection, and signature of an accredited veterinarian attesting to the following statement:
"I certify that the specimens submitted with this form were collected by me from the animal(s) described on the date indicated."

Veterinarian's, Clinician's or Owner's Signature:	Condition(s) Suspected:
---------------------------------------------------	-------------------------

IDENTIFICATION SHEET FOR MULTIPLE ANIMALS

(To accompany WADDL Accession form, if needed)

Washington Animal Disease Diagnostic Laboratory
 College of Veterinary Medicine, Washington State University
 Mailing address: P.O. Box 647034 Pullman, WA. 99164-7034
 Phone: (509) 335-9696 E-Mail: waddl@vetmed.wsu.edu
 Web Site: http://waddl.vetmed.wsu.edu

2016-7117
 Ref Vet: Highland, Margaret
 Owner: USDA - ARS - ADRU
 Breed: Domestic Goat
 Routing: .jmd

06/02/16

Owner: ADRU-ARS-USDA

Veterinarian: Highland

TEST(S) REQUESTED: M. ovipneumoniae qPCR

Tube	Animal # or Name	Tube	Animal # or Name	Tube	Animal # or Name	Tube	e
1	<u>13_A</u>	26		51		76	
2	<u>13_B</u>	27		52		77	
3	<u>13_C</u>	28		53		78	
4	<u>13_D</u>	29		54		79	
5	<u>15_A</u>	30		55		80	
6	<u>15_B</u>	31		56		81	
7	<u>15_C</u>	32		57		82	
8	<u>15_D</u>	33		58		83	
9		34		59		84	
10		35		60		85	
11		36		61		86	
12		37		62		87	
13		38		63		88	
14		39		64		89	
15		40		65		90	
16		41		66		91	
17		42		67		92	
18		43		68		93	
19		44		69		94	
20		45		70		95	
21		46		71		96	
22		47		72		97	
23		48		73		98	
24		49		74		99	
25		50		75		100 *	

* For over 100 samples, please copy this form and continue numbering.

Washington Animal Disease Diagnostic Lab

P.O. Box 647034
Pullman, WA 99164-7034
Telephone : (509) 335-9696
Fax : (509) 335-7424

Dr. Margaret Highland
USDA-ARS-ADRU
WSU - 3003 ADBF

Case#: 2016-7117
Report Date: 06/07/16

Pullman, WA 99164-6630

Submittal Date: 06/02/16
Owner: USDA-ARS-ADRU

Species: Domestic Goat

Age: Adult
Sex:

Final Report:

Molecular Diagnostics- Reported on 06/07/16 Authorized by Daniel Bradway, Lab Manager

PCR-Mycoplasma ovipneumoniae SOP: 501.40RT.2016.03.17

Animal	Specimen	Result
13.A	Nasal swab	Not detected
13.B	Nasal swab	Not detected
13.C	Nasal swab	Not detected
13.D	Nasal swab	Not detected
15.A	Nasal swab	Not detected
15.B	Nasal swab	Not detected
15.C	Nasal swab	Not detected
15.D	Nasal swab	Not detected

PCR-Mycoplasma ovipneumoniae test comment: This assay detects only Mycoplasma ovipneumoniae. Culture is available at WADDL to detect other species of Mycoplasma if desired. Fees for culture are available on our website. Please contact the lab if Mycoplasma culture or other testing is desired.

Quantity/Description/Routing of Samples

8 nasal swabs
- dropped off
by MAH

2016 - 7117
Ref Vet: Highland, Margaret
Owner: USDA - ARS - ADRU
Breed: Domestic Goat
Routed: jmd

Sample Condition:	<input type="checkbox"/> Room Temp.	<input type="checkbox"/> On ice	<input checked="" type="checkbox"/> Frozen	<input type="checkbox"/> Fixed	Contents match forms:	Opened by:
Samples Received Via:	<input type="checkbox"/> US Mail	<input type="checkbox"/> FedEx	<input checked="" type="checkbox"/> Drop off	Explain below: <i>not</i>		
	<input type="checkbox"/> UPS	<input type="checkbox"/> FedEx-R	<input type="checkbox"/> Other:			

Comments for Case Tracking:



06/02/16
index: 1 page

Sample Label <input checked="" type="checkbox"/>
<i>MAH</i>

ACCESSION FORM FOR GENERAL DIAGNOSTICS
Washington Animal Disease Diagnostic Laboratory

College of Veterinary Medicine, Washington State University
 Web Site: <http://waddl.vetmed.wsu.edu>

US Postal Service mailing address:
 PO Box 647034
 Pullman, WA. 99164-7034

UPS, FedEx or Courier shipping address:
 Bustad Hall, Rm.155-N
 Pullman, WA. 99164-7034

Phone: (509) 335-9696
 FAX: (509) 335 7424
 E-Mail: waddl@vetmed.wsu.edu

2016-7913
 Ref Vet: Highland, Margaret
 Owner:
 Breed: Domestic Goat
 Routed: md

06/20/16
 Form: 3 pages

Please type or use black ink and print clearly.

Veterinarian or Last Case Coordinator Name: Highland		First Name: Maggie	
Clinic: ADRU-ARS-USDA			
Street address: ADBF 3033		Mailing Address or PO Box:	
City: Pullman	State: WA	Zip: 99164	
Phone: 509-335-6327	Fax: 509-335-8328	E-mail: mah@vetmed.wsu.edu	
Owner Last Name first: same as above		Guardian Name: (if owner is under 18)	
Farm Name:		First Time Submitter? <input type="checkbox"/> Yes <input type="checkbox"/> No	
Street address:		Mailing Address or PO Box:	
City:	State:	Zip:	
Phone:	Fax:	E-mail:	

Billing: Owner Clinic 3rd Party (preapproval required) Please note: WADDL policy is to bill the clinic if provided, unless prepaid.
Reporting Preference: Mail Fax Web access - register on web site at <http://waddl.vetmed.wsu.edu>

Please fill out completely as possible:

Specimen(s) Submitted:		Date Collected: June 2016
nasal swabs		Date Shipped:
<small>(Please use WADDL Animal ID Sheet for multiple animals.)</small>		
Tests Requested:	<input type="checkbox"/> Necropsy <input type="checkbox"/> Virology <input type="checkbox"/> Bacteriology <input type="checkbox"/> IHC <input type="checkbox"/> Histopathology <input type="checkbox"/> Serology <input type="checkbox"/> Mycoplasma culture <input checked="" type="checkbox"/> PCR <input type="checkbox"/> Toxicology <input type="checkbox"/> Fungal Culture <input type="checkbox"/> Parasitology <input type="checkbox"/> Other:	
<small>Note: WADDL reserves the right to modify the tests requested for more efficient case work-up and / or to send specimens to outside laboratories to perform testing not done at WADDL.</small>		
Animal ID (name/tag#)	Species	Breed
see multiple animal form	domestic goats	multiple
Age	Sex	Animal Weight
multiple		
Location of Lesion	No. in group	No. Dead
N/A		N/A
	No. Sick	No. on Premises
	N/A	

* Was animal euthanized? If so, what method?

Additional History: Vaccinations, signs, stress factors, treatments, post mortem findings, pertinent feed or feed additives, clinical lab results, previous WADDL Case Numbers. (Attach additional sheets as necessary.)

M.ovipneumoniae qPCR on each sample.
 Please save remaining DNA isolation and call Maggie for pick up or may request further testing (sequencing) be performed by WADDL, depending on the results of qPCR analyses.

Bill to ADRU-ARS-USDA acct #RSA 2540-1080

WADDL is an official brucellosis testing laboratory. All serology for brucellosis, including abortion screens, requires identification of animals, date of sample collection, and signature of an accredited veterinarian attesting to the following statement:

"I certify that the specimens submitted with this form were collected by me from the animal(s) described on the date indicated."

Veterinarian's, Clinician's or Owner's Signature: <i>Maggie Highland</i>	Condition(s) Suspected: N/A (surveillance)
--------------------------------------------------------------------------	---------------------------------------------------

IDENTIFICATION SHEET FOR MULTIPLE ANIMALS

(To accompany WADDL Accession form, if needed)

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 College of Veterinary Medicine, Washington State University

Mailing address: P.O. Box 647034
 Pullman, WA. 99164-7034
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 Web Site: http://waddl.vetmed.wsu.edu

Shipping address:
 Bustad Hall, Rm.155-N
 Pullman, WA. 99164-7034
 FAX: (509) 335-7424

2016-7913

06/20/16

Ref Vet: Highland, Margaret
 Owner:
 Breed: Domestic Goat
 Routing: md

Owner: ADRU-ARS-USDA

Veterinarian: Highland

TEST(S) REQUESTED: Mycoplasma ovipneumoniae qPCR

Tube	Animal # or Name	Tube	Animal # or Name	Tube	Animal # or Name	Tube	Animal # or Name
1	24_A	26	1_S	51	[REDACTED]	76	5_B
2	24_B	27	1_T	52	[REDACTED]	77	5_C
3	24_C	28	1_U	53	[REDACTED]	78	5_D
4	3_A	29	1_V	54	[REDACTED]	79	5_E
5	3_B	30	1_W	55	[REDACTED]	80	5_F
6	3_C	31	1_X	56	[REDACTED]	81	5_G
7	3_D	32	1_Y	57	[REDACTED]	82	5_H
8	1_A	33	1_Aa	58	[REDACTED]	83	5_I
9	1_B	34	1_Bb	59	[REDACTED]	84	5_J
10	1_C	35	1_Cc	60	[REDACTED]	85	5_K
11	1_D	36	1_Dd	61	[REDACTED]	86	5_L
12	1_E	37	1_Ee	62	[REDACTED]	87	5_M
13	1_F	38	1_Ff	63	[REDACTED]	88	5_N
14	1_G	39	1_Gg	64	[REDACTED]	89	5_O
15	1_H	40	1_Hh	65	[REDACTED]	90	5_P
16	1_I	41	1_Ii	66	[REDACTED]	91	4_A
17	1_J	42	1_Jj	67	[REDACTED]	92	4_B
18	1_K	43	1_Kk	68	[REDACTED]	93	2_A
19	1_L	44	1_Ll	69	[REDACTED]	94	2_B
20	1_M	45	1_Mm	70	[REDACTED]	95	2_C
21	1_N	46	1_Nn	71	[REDACTED]	96	2_D
22	1_O	47	1_Oo	72	4_A	97	2_E
23	1_P	48	[REDACTED]	73	4_B	98	2_F
24	1_Q	49	[REDACTED]	74	4_C	99	2_G
25	1_R	50	[REDACTED]	75	5_A	100*	2_H

* For over 100 samples, please copy this form and continue numbering.

IDENTIFICATION SHEET FOR MULTIPLE ANIMALS

(To accompany WADDL Accession form, if needed)

Washington Animal Disease Diagnostic Laboratory
 College of Veterinary Medicine, Washington State University
 Mailing address: P.O. Box 647034
 Pullman, WA. 99164-7034
 Phone: (509) 335-9696
 E-Mail: waddl@vetmed.wsu.edu
 Web Site: http://waddl.vetmed.wsu.edu

Shipping address:
 Bustad Hall, Rm. 155-N
 Pullman, WA. 99164-7034
 FAX: (509) 335-7424

2016-7913

06/20/16

Ref Vet: Highland, Margaret
 Owner:
 Breed: Domestic Goat
 Routing: md

Owner: ADRU-ARS-USDA

Veterinarian: HIGHLAND

TEST(S) REQUESTED: M. ovipneumoniae qPCR

Tube	Animal # or Name	Tube	Animal # or Name	Tube	Animal # or Name	Tube	Animal # or Name
1	5_A	26	2_A	51		76	
2	5_B	27	2_B	52	6_G	77	
3	5_C	28	2_C	53	13_A	78	
4	5_D	29	2_D	54	13_B	79	
5	5_E	30	3_A	55	13_C	80	
6	5_F	31	3_B	56	13_D	81	
7	5_G	32	3_C	57	13_E	82	
8	5_H	33	3_D	58	13_F	83	
9	5_I	34	4_A	59	2-b-H	84	
10	5_J	35	4_B	60	2-A	85	
11	5_K	36	4_C	61	2-B	86	
12	5_L	37	1_A	62	2-C	87	
13	5_M	38	1_B	63	2-D	88	
14	5_N	39	21_A	64	2-E	89	
15	5_O	40	5_A	65	2-F	90	
16	5_P	41	5_B	66	2-G	91	
17	4_A	42	5_C	67	2-H	92	
18	4_B	43	5_D	68	2-I	93	
19	1_A	44	5_E	69	2-A	94	
20	1_B	45	6_A	70	2B	95	
21	1_C	46	6_B	71	2-C	96	
22	1_D	47	6_C	72	2-D	97	
23	1_E	48	6_D	73	7-A	98	
24	7_A	49	6_E	74	7-B	99	
25	7_B	50	6_F	75		100*	

* For over 100 samples, please copy this form and continue numbering.

Washington Animal Disease Diagnostic Lab

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Pullman, WA 99164-7034
Telephone : (509) 335-9696
Fax : (509) 335-7424

Dr. Margaret Highland
USDA-ARS-ADRU
WSU - 3003 ADBF

Case#: 2016-7913
Report Date: 07/01/16

Pullman, WA 99164-6630

Submittal Date: 06/20/16
Owner:

Species: Domestic Goat

Age:
Sex:

Final Report:

Molecular Diagnostics- Reported on 07/01/16 Authorized by Daniel Bradway, Lab Manager

PCR-Mycoplasma ovipneumoniae SOP: 501.40RT.2016.03.17

Animal	Specimen	Result
[REDACTED] 24.A	Nasal swab	Not detected
[REDACTED] 24.B	Nasal swab	Not detected
[REDACTED] 24.C	Nasal swab	Not detected
[REDACTED] 3.A	Nasal swab	Not detected
[REDACTED] 3.B	Nasal swab	Not detected
[REDACTED] 3.C	Nasal swab	Not detected
[REDACTED] 3.D	Nasal swab	Not detected
[REDACTED] 1.A	Nasal swab	Not detected
[REDACTED] 1.B	Nasal swab	Not detected
[REDACTED] 1.C	Nasal swab	Indeterminate
[REDACTED] 1.D	Nasal swab	Not detected
[REDACTED] 1.E	Nasal swab	Not detected
[REDACTED] 1.F	Nasal swab	Not detected
[REDACTED] 1.G	Nasal swab	Not detected
[REDACTED] 1.H	Nasal swab	Detected
[REDACTED] 1.I	Nasal swab	Not detected
[REDACTED] 1.J	Nasal swab	Not detected
[REDACTED] 1.K	Nasal swab	Not detected
[REDACTED] 1.L	Nasal swab	Not detected
[REDACTED] 1.M	Nasal swab	Not detected
[REDACTED] 1.N	Nasal swab	Not detected
[REDACTED] 1.O	Nasal swab	Not detected
[REDACTED] 1.P	Nasal swab	Not detected
[REDACTED] 1.Q	Nasal swab	Not detected
[REDACTED] 1.R	Nasal swab	Not detected
[REDACTED] 1.S	Nasal swab	Not detected
[REDACTED] 1.T	Nasal swab	Not detected
[REDACTED] 1.U	Nasal swab	Not detected

Washington Animal Disease Diagnostic Lab

PCR-Mycoplasma ovipneumoniae SOP: 501.40RT.2016.03.17

Animal	Specimen	Result
1.V	Nasal swab	Not detected
1.W	Nasal swab	Not detected
1.X	Nasal swab	Not detected
1.Y	Nasal swab	Not detected
1.Aa	Nasal swab	Not detected
1.Bb	Nasal swab	Detected
1.Cc	Nasal swab	Not detected
1.Dd	Nasal swab	Not detected
1.Ee	Nasal swab	Detected
1.Ff	Nasal swab	Detected
1.Gg	Nasal swab	Indeterminate
1.Hh	Nasal swab	Not detected
1.Ii	Nasal swab	Indeterminate
1.Jj	Nasal swab	Detected
1.Kk	Nasal swab	Not detected
1.Ll	Nasal swab	Indeterminate
1.Mm	Nasal swab	Not detected
1.Nn	Nasal swab	Indeterminate
1.Oo	Nasal swab	Detected
4.A	Nasal swab	Not detected
4.B	Nasal swab	Not detected
4.C	Nasal swab	Not detected
5.A	Nasal swab	Not detected
5.B	Nasal swab	Not detected
5.C	Nasal swab	Not detected
5.D	Nasal swab	Not detected
5.E	Nasal swab	Not detected
5.F	Nasal swab	Not detected
5.G	Nasal swab	Not detected
5.H	Nasal swab	Not detected
5.I	Nasal swab	Not detected
5.J	Nasal swab	Not detected
5.K	Nasal swab	Not detected
5.L	Nasal swab	Not detected
5.M	Nasal swab	Not detected
5.N	Nasal swab	Not detected
5.O	Nasal swab	Not detected
5.P	Nasal swab	Indeterminate
4.A	Nasal swab	Not detected
4.B	Nasal swab	Not detected
2.A	Nasal swab	Not detected
2.B	Nasal swab	Not detected
2.C	Nasal swab	Not detected
2.D	Nasal swab	Not detected
2.E	Nasal swab	Not detected
2.F	Nasal swab	Not detected
2.G	Nasal swab	Not detected
2.H	Nasal swab	Not detected
5.A	Nasal swab	Not detected
5.B	Nasal swab	Not detected
5.C	Nasal swab	Not detected
5.D	Nasal swab	Not detected

Washington Animal Disease Diagnostic Lab

PCR-Mycoplasma ovipneumoniae SOP: 501.40RT.2016.03.17

Animal	Specimen	Result
T.5.E	Nasal swab	Indeterminate
T.5.F	Nasal swab	Not detected
T.5.G	Nasal swab	Not detected
T.5.H	Nasal swab	Not detected
T.5.I	Nasal swab	Not detected
T.5.J	Nasal swab	Not detected
T.5.K	Nasal swab	Not detected
T.5.L	Nasal swab	Not detected
T.5.M	Nasal swab	Not detected
T.5.N	Nasal swab	Not detected
T.5.O	Nasal swab	Detected
T.5.P	Nasal swab	Indeterminate
Y.4.A	Nasal swab	Not detected
Y.4.B	Nasal swab	Not detected
X.1.A	Nasal swab	Not detected
X.1.B	Nasal swab	Not detected
X.1.C	Nasal swab	Not detected
X.1.D	Nasal swab	Not detected
X.1.E	Nasal swab	Not detected
Z.7.A	Nasal swab	Not detected
Z.7.B	Nasal swab	Indeterminate
Z.2.A	Nasal swab	Not detected
Z.2.B	Nasal swab	Not detected
Z.2.C	Nasal swab	Indeterminate
Z.2.D	Nasal swab	Not detected
Z.3.A	Nasal swab	Not detected
Z.3.B	Nasal swab	Not detected
Z.3.C	Nasal swab	Not detected
Z.3.D	Nasal swab	Not detected
Z.4.A	Nasal swab	Not detected
Z.4.B	Nasal swab	Not detected
Z.4.C	Nasal swab	Not detected
Z.1.A	Nasal swab	Not detected
Z.1.B	Nasal swab	Not detected
Z.21.A	Nasal swab	Not detected
Z.R.5.A	Nasal swab	Not detected
Z.R.5.B	Nasal swab	Not detected
Z.R.5.C	Nasal swab	Indeterminate
Z.R.5.D	Nasal swab	Indeterminate
Z.R.5.E	Nasal swab	Not detected
Z.6.A	Nasal swab	Not detected
Z.6.B	Nasal swab	Not detected
Z.6.C	Nasal swab	Not detected
Z.6.D	Nasal swab	Not detected
Z.6.E	Nasal swab	Not detected
Z.6.F	Nasal swab	Not detected
Z.6.G	Nasal swab	Not detected
Z.6.H	Nasal swab	Not detected
Z.13.A	Nasal swab	Not detected
Z.13.B	Nasal swab	Not detected
Z.13.C	Nasal swab	Not detected
Z.13.D	Nasal swab	Not detected

Washington Animal Disease Diagnostic Lab

PCR-Mycoplasma ovipneumoniae SOP: 501.40RT.2016.03.17

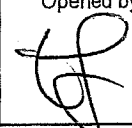
Animal	Specimen	Result
[REDACTED] 13_E	Nasal swab	Not detected
[REDACTED] 13_F	Nasal swab	Not detected
[REDACTED] 2_A	Nasal swab	Not detected
[REDACTED] 2_B	Nasal swab	Not detected
[REDACTED] 2_C	Nasal swab	Not detected
[REDACTED] 2_D	Nasal swab	Not detected
[REDACTED] 2_E	Nasal swab	Not detected
[REDACTED] 2_F	Nasal swab	Not detected
[REDACTED] 2_G	Nasal swab	Not detected
[REDACTED] 2_H	Nasal swab	Detected
[REDACTED] 2_I	Nasal swab	Not detected
[REDACTED] 2_A	Nasal swab	Not detected
[REDACTED] 2_B	Nasal swab	Not detected
[REDACTED] 2_C	Nasal swab	Not detected
[REDACTED] 2_D	Nasal swab	Not detected
[REDACTED] 7_A	Nasal swab	Not detected
[REDACTED] 7_B	Nasal swab	Not detected

PCR-Mycoplasma ovipneumoniae test comment: This assay detects only Mycoplasma ovipneumoniae. Culture is available at WADDL to detect other species of Mycoplasma if desired. Fees for culture are available on our website. Please contact the lab if Mycoplasma culture or other testing is desired.

Quantity/Description/Routing of Samples

149 dry swabs

2016-7913
Ref Vet: Highland, Margaret
Owner:
Breed: Domestic Goat
Routed: md


Sample Condition:	<input type="checkbox"/> Room Temp.	<input type="checkbox"/> On ice	<input checked="" type="checkbox"/> Frozen	<input type="checkbox"/> Fixed	Contents match forms: <input checked="" type="checkbox"/> Yes <input type="checkbox"/> No Explain below:	Opened by: 
	Samples Received Via:					
	<input type="checkbox"/> US Mail	<input type="checkbox"/> FedEx	<input checked="" type="checkbox"/> Drop off			
	<input type="checkbox"/> UPS	<input type="checkbox"/> FedEx-R	<input type="checkbox"/> Other:			



06/20/16
Index: 1 page

Comments for Case Tracking:

Sample Label



**ACCESSION FORM FOR GENERAL DIAGNOSTICS
Washington Animal Disease Diagnostic Laboratory**

College of Veterinary Medicine, Washington State University

Web Site: <http://waddl.vetmed.wsu.edu>

US Postal Service mailing address:
PO Box 647034
Pullman, WA. 99164-7034

UPS, FedEx or Courier shipping address:
Bustad Hall, Rm.155-N
Pullman, WA. 99164-7034

Phone: (509) 335-9696
FAX: (509) 335 7424
E-Mail: waddl@vetmed.wsu.edu

2016 - 10050
Ret Vet: Highland, Margaret
Owner: USDA - ARS - ADRU
Breed: Domestic Goat
Routed: md

08/04/16
form: 3 pages

Please type or use black ink and print clearly.

Veterinarian or Case Coordinator: Name: Highland		First Name: Maggie	
Clinic: ADRU-ARS-USDA			
Street address: ADBF 3033		Mailing Address or PO Box:	
City: Pullman	State: WA	Zip: 99164	
Phone: 509-335-6327	Fax: 509-335-8328	E-mail: mah@vetmed.wsu.edu	
Owner: Last Name first: same as above		Guardian Name: (if owner is under 18)	
Farm Name:		First Time Submitter? <input type="checkbox"/> Yes <input type="checkbox"/> No	
Street address:		Mailing Address or PO Box:	
City:	State:	Zip:	
Phone:	Fax:	E-mail:	

Billing: Owner Clinic 3rd Party (preapproval required) Please note: WADDL policy is to bill the clinic if provided, unless prepaid.

Reporting Preference: Mail Fax Web access - register on web site at <http://waddl.vetmed.wsu.edu>

Please fill out completely as possible:

Specimen(s) Submitted:		Date Collected: July 2016	
(Please use WADDL Animal ID Sheet for multiple animals.)		Date Shipped: n/a	
nasal swabs			
<input type="checkbox"/> Necropsy	<input type="checkbox"/> Virology	<input type="checkbox"/> Bacteriology	<input type="checkbox"/> IHC
<input type="checkbox"/> Histopathology	<input type="checkbox"/> Serology	<input type="checkbox"/> Mycoplasma culture	<input checked="" type="checkbox"/> PCR
<input type="checkbox"/> Toxicology	<input type="checkbox"/> Fungal Culture	<input type="checkbox"/> Parasitology	<input type="checkbox"/> Other:
<i>Note: WADDL reserves the right to modify the tests requested for more efficient case work-up and / or to send specimens to outside laboratories to perform testing not done at WADDL.</i>			
Animal ID (name/tag#)	Species	Breed	Age
see multiple animal form	domestic goats	multiple	multiple
Sex	Animal Weight	Location of Lesion	
		N/A	
No. in group	No. Dead	No. Sick	No. on Premises
	N/A	N/A	
* Was animal euthanized? If so, what method? N/A			

Additional History: Vaccinations, signs, stress factors, treatments, post mortem findings, pertinent feed or feed additives, clinical lab results, previous WADDL Case Numbers. (Attach additional sheets as necessary.)

M. ovipneumoniae qPCR on each sample
Please save remaining DNA isolations and call Maggie for pick up or may request further testing (sequencing) be performed by WADDL, depending on the results of qPCR analysis.

Please bill to ADRU-ARS-USDA account #RSA 2540-1080

WADDL is an official brucellosis testing laboratory. All serology for brucellosis, including abortion screens, requires identification of animals, date of sample collection, and signature of an accredited veterinarian attesting to the following statement:

"I certify that the specimens submitted with this form were collected by me from the animal(s) described on the date indicated."

Veterinarian's, Clinician's or Owner's Signature: *Maggie Highland* Condition(s) Suspected: *N/A screening study*

IDENTIFICATION SHEET FOR MULTIPLE ANIMALS

(To accompany WADDL Accession form, if needed)

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 College of Veterinary Medicine, Washington State University
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 Pullman, WA. 99164-7034 Pullman, WA. 99164-7034
 Phone: (509) 335-9696 FAX: (509) 335-7424
 E-Mail: waddl@vetmed.wsu.edu
 Web Site: http://waddl.vetmed.wsu.edu

2016 - 10050
 Ref Vet: Highland, Margaret
 Owner: USDA - ARS - ADRU
 Breed: Domestic Goat
 Routing: md

08/04/16

Owner: Highland, Maggie

Veterinarian: Highland, Maggie

TEST(S) REQUESTED: Movi qPCR

Tube	Animal # or Name	Tube	Animal # or Name	Tube	Animal # or Name	Tube	Animal # or Name
1	8_A	26	5_O	51	4_B	76	1_E
2	8_B	27	5_P	52	4_C	77	1_F
3	4_A	28	2_A	53	4_D	78	1_G
4	4_B	29	2_B	54	1_A	79	1_A
5	4_C	30	2_C	55	1_B	80	1_B
6	4_D	31	2_D	56	1_A	81	1_C
7	4_E	32	6_A	57	1_B	82	1_D
8	4_F	33	1_A	58	1_C	83	1_E
9	4_G	34	2_B	59	1_D	84	1_F
10	1_A	35	2_C	60	6_A	85	1_G
11	1_B	36	2_A	61	6_B	86	5_A
12	5_A	37	2_B	62	6_C	87	5_B
13	5_B	38	2_C	63	2_A	88	5_C
14	5_C	39	2_D	64	2_B	89	5_D
15	5_D	40	25_A	65	2_C	90	5_E
16	5_E	41	25_B	66	2_D	91	5_F
17	5_F	42	25_C	67	2_E	92	5_G
18	5_G	43	25_D	68	2_F	93	5_H
19	5_H	44	25_E	69	2_G	94	5_I
20	5_I	45	1_A	70	2_H	95	5_J
21	5_J	46	1_B	71	2_I	96	5_K
22	5_K	47	1_C	72	1_A	97	5_L
23	5_L	48	1_D	73	1_B	98	5_M
24	5_M	49	1_E	74	1_C	99	5_N
25	5_N	50	4_A	75	1_D	100 *	5_O

* For over 100 samples, please copy this form and continue numbering.

IDENTIFICATION SHEET FOR MULTIPLE ANIMALS

(To accompany WADDL Accession form, if needed)

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 College of Veterinary Medicine, Washington State University
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 Web Site: http://waddl.vetmed.wsu.edu

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 Pullman, WA. 99164-7034
 FAX: (509) 335-7424

2016-10050
 Ref Vet: Highland, Margaret
 Owner: USDA-ARS-ADRU
 Breed: Domestic Goat
 Routing: md

08/04/16

Owner: Highland, Maggie

Veterinarian: Highland, Maggie

TEST(S) REQUESTED: Movi qPCR

Tube	Animal # or Name	Tube	Animal # or Name	Tube	Animal # or Name	Tube	Animal # or Name
101	5_P	126	1_E	51		76	
102	5_Q	127	7_A	52		77	
103	5_R	128	7_B	53		78	
104	5_S	129	7_C	54		79	
105	5_T	130	7_D	55		80	
106	5_U	131	2_A	56		81	
107	5_V	132	2_B	57		82	
108	5_W	133	2_C	58		83	
109	5_X	34		59		84	
110	5_Y	35		60		85	
111	3_A	36		61		86	
112	3_B	37		62		87	
113	1_A	38		63		88	
114	1_B	39		64		89	
115	1_C	40		65		90	
116	1_D	41		66		91	
117	1_E	42		67		92	
118	1_F	43		68		93	
119	1_G	44		69		94	
120	5_A	45		70		95	
121	5_B	46		71		96	
122	1_A	47		72		97	
123	1_B	48		73		98	
124	1_C	49		74		99	
125	1_D	50		75		100*	

* For over 100 samples, please copy this form and continue numbering.

Washington Animal Disease Diagnostic Lab

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Pullman, WA 99164-7034
Telephone : (509) 335-9696
Fax : (509) 335-7424

Dr. Margaret Highland
USDA-ARS-ADRU
WSU - 3003 ADBF

Case#: 2016-10050
Report Date: 08/19/16

Pullman, WA 99164-6630

Submittal Date: 08/04/16
Owner: USDA-ARS-ADRU

Species: Domestic Goat

Age:
Sex:

Final Report:

Molecular Diagnostics- Reported on 08/19/16 Authorized by Daniel Bradway, Lab Manager

PCR-Mycoplasma ovipneumoniae SOP: 501.40RT.2016.07.18

Animal	Specimen	Result
18.A	Nasal swab	Not detected
18.B	Nasal swab	Not detected
14.A	Nasal swab	Indeterminate
14.B	Nasal swab	Not detected
14.C	Nasal swab	Indeterminate
14.D	Nasal swab	Indeterminate
14.E	Nasal swab	Not detected
14.F	Nasal swab	Indeterminate
14.G	Nasal swab	Not detected
11.A	Nasal swab	Not detected
11.B	Nasal swab	Not detected
15.A	Nasal swab	Not detected
15.B	Nasal swab	Not detected
15.C	Nasal swab	Not detected
15.D	Nasal swab	Not detected
15.E	Nasal swab	Not detected
15.F	Nasal swab	Not detected
15.G	Nasal swab	Not detected
15.H	Nasal swab	Not detected
15.I	Nasal swab	Indeterminate
15.J	Nasal swab	Not detected
15.K	Nasal swab	Not detected
15.L	Nasal swab	Not detected
15.M	Nasal swab	Not detected
15.N	Nasal swab	Indeterminate
15.O	Nasal swab	Not detected
15.P	Nasal swab	Not detected
12.A	Nasal swab	Not detected

Washington Animal Disease Diagnostic Lab

PCR-Mycoplasma ovipneumoniae SOP: 501.40RT.2016.07.18

Animal	Specimen	Result
[REDACTED] 2.B	Nasal swab	Not detected
[REDACTED] 2.C	Nasal swab	Not detected
[REDACTED] 2.D	Nasal swab	Not detected
[REDACTED] 6.A	Nasal swab	Not detected
[REDACTED] 1.A	Nasal swab	Not detected
[REDACTED] 2.B	Nasal swab	Not detected
[REDACTED] 2.C	Nasal swab	Not detected
[REDACTED] 2.A	Nasal swab	Not detected
[REDACTED] 2.B	Nasal swab	Not detected
[REDACTED] 2.C	Nasal swab	Not detected
[REDACTED] 2.D	Nasal swab	Not detected
[REDACTED] 25.A	Nasal swab	Not detected
[REDACTED] 25.B	Nasal swab	Not detected
[REDACTED] 25.C	Nasal swab	Not detected
[REDACTED] 25.D	Nasal swab	Not detected
[REDACTED] 25.E	Nasal swab	Not detected
[REDACTED] 1.A	Nasal swab	Not detected
[REDACTED] 1.B	Nasal swab	Not detected
[REDACTED] 1.C	Nasal swab	Not detected
[REDACTED] 1.D	Nasal swab	Not detected
[REDACTED] 1.E	Nasal swab	Not detected
[REDACTED] 4.A	Nasal swab	Not detected
[REDACTED] 4.B	Nasal swab	Not detected
[REDACTED] 4.C	Nasal swab	Not detected
[REDACTED] 4.D	Nasal swab	Not detected
[REDACTED] 1.1.A	Nasal swab	Not detected
[REDACTED] 1.1.B	Nasal swab	Not detected
[REDACTED] 1.1.A	Nasal swab	Not detected
[REDACTED] 1.1.B	Nasal swab	Not detected
[REDACTED] 1.1.C	Nasal swab	Not detected
[REDACTED] 1.1.D	Nasal swab	Not detected
[REDACTED] 6.A	Nasal swab	Not detected
[REDACTED] 6.B	Nasal swab	Not detected
[REDACTED] 6.C	Nasal swab	Not detected
[REDACTED] 2.A	Nasal swab	Not detected
[REDACTED] 2.B	Nasal swab	Not detected
[REDACTED] 2.C	Nasal swab	Not detected
[REDACTED] 2.D	Nasal swab	Not detected
[REDACTED] 2.E	Nasal swab	Not detected
[REDACTED] 2.F	Nasal swab	Not detected
[REDACTED] 2.G	Nasal swab	Not detected
[REDACTED] 2.H	Nasal swab	Indeterminate
[REDACTED] 2.I	Nasal swab	Not detected
[REDACTED] Y.1.A	Nasal swab	Not detected
[REDACTED] Y.1.B	Nasal swab	Not detected
[REDACTED] Y.1.C	Nasal swab	Not detected
[REDACTED] Y.1.D	Nasal swab	Not detected
[REDACTED] Y.1.E	Nasal swab	Not detected
[REDACTED] Y.1.F	Nasal swab	Not detected
[REDACTED] Y.1.G	Nasal swab	Not detected
[REDACTED] 1.A	Nasal swab	Not detected
[REDACTED] 1.B	Nasal swab	Not detected

Washington Animal Disease Diagnostic Lab

PCR-Mycoplasma ovipneumoniae SOP: 501.40RT.2016.07.18

Animal	Specimen	Result
1.C	Nasal swab	Not detected
1.D	Nasal swab	Not detected
1.E	Nasal swab	Not detected
1.F	Nasal swab	Not detected
1.G	Nasal swab	Not detected
5.A	Nasal swab	Not detected
5.B	Nasal swab	Not detected
5.C	Nasal swab	Not detected
5.D	Nasal swab	Not detected
5.E	Nasal swab	Not detected
5.F	Nasal swab	Not detected
5.G	Nasal swab	Not detected
5.H	Nasal swab	Not detected
5.I	Nasal swab	Not detected
5.J	Nasal swab	Not detected
5.K	Nasal swab	Not detected
5.L	Nasal swab	Not detected
5.M	Nasal swab	Not detected
5.N	Nasal swab	Not detected
5.O	Nasal swab	Not detected
5.P	Nasal swab	Not detected
5.Q	Nasal swab	Not detected
5.R	Nasal swab	Not detected
5.S	Nasal swab	Not detected
5.T	Nasal swab	Not detected
5.U	Nasal swab	Not detected
5.V	Nasal swab	Not detected
5.W	Nasal swab	Not detected
5.X	Nasal swab	Not detected
5.Y	Nasal swab	Not detected
3.A	Nasal swab	Not detected
3.B	Nasal swab	Not detected
1.A	Nasal swab	Not detected
1.B	Nasal swab	Not detected
1.C	Nasal swab	Not detected
1.D	Nasal swab	Not detected
1.E	Nasal swab	Not detected
1.F	Nasal swab	Not detected
1.G	Nasal swab	Not detected
5.A	Nasal swab	Not detected
5.B	Nasal swab	Not detected
1.A	Nasal swab	Not detected
1.B	Nasal swab	Not detected
1.C	Nasal swab	Not detected
1.D	Nasal swab	Not detected
1.E	Nasal swab	Not detected
7.A	Nasal swab	Not detected
7.B	Nasal swab	Not detected
7.C	Nasal swab	Not detected
7.D	Nasal swab	Not detected
2.A	Nasal swab	Not detected
2.B	Nasal swab	Not detected

Washington Animal Disease Diagnostic Lab

PCR-Mycoplasma ovipneumoniae SOP: 501.40RT.2016.07.18

Animal	Specimen	Result
2-C	Nasal swab	Not detected
1.H	Nasal swab	Not detected

PCR-Mycoplasma ovipneumoniae test comment: This assay detects only Mycoplasma ovipneumoniae. Culture is available at WADDL to detect other species of Mycoplasma if desired. Fees for culture are available on our website. Please contact the lab if Mycoplasma culture or other testing is desired.

Quantity/Description/Routing of Samples

133 nasal swabs
- dropped off by
Maggie Highland

2016 - 10050
Ref Vet: Highland, Margaret
Owner: USDA - ARS - ADRI
Breed: Domestic Goat
Routed: md

Sample Condition:	<input type="checkbox"/> Room Temp.	<input type="checkbox"/> On ice	<input checked="" type="checkbox"/> Frozen	<input type="checkbox"/> Fixed	Contents match forms: <input type="checkbox"/> Yes <input checked="" type="checkbox"/> No Explain below:	Opened by: <i>MA</i>
	Samples Received Via:		<input type="checkbox"/> US Mail	<input type="checkbox"/> FedEx		
	<input type="checkbox"/> UPS	<input type="checkbox"/> FedEx-R	<input type="checkbox"/> Other:			

Comments for Case Tracking:



08/04/16
page 1 of 1

Sample Label <input checked="" type="checkbox"/>
<i>MA</i>

ACCESSION FORM FOR GENERAL DIAGNOSTICS
Washington Animal Disease Diagnostic Laboratory

College of Veterinary Medicine, Washington State University

Web Site: <http://waddl.vetmed.wsu.edu>

US Postal Service mailing address:
 PO Box 647034
 Pullman, WA. 99164-7034

UPS, FedEx or Courier shipping address:
 Bustad Hall, Rm.155-N
 Pullman, WA. 99164-7034

Phone: (509) 335-9696
 FAX: (509) 335 7424
 E-Mail: waddl@vetmed.wsu.edu

2016 - 12311
 Ref Vet: Highland, Margaret
 Owner: USDA - ARS - ADRU
 Breed: Domestic Goat
 Routed: md

Please type or use black ink and print clearly.

Veterinarian or Case Coordinator: Name: Highland		First Name: Maggie	
Clinic: ADRU-ARS-USDA			
Street address: ADBF 3033		Mailing Address or PO Box:	
City: Pullman	State: WA	Zip: 99164	
Phone: 509-335-6327	Fax: 509-335-8328	E-mail: mah@vetmed.wsu.edu	
Owner: Last Name first: same as above		Guardian Name: (if owner is under 18)	
Farm Name:		First Time Submitter? <input type="checkbox"/> Yes <input type="checkbox"/> No	
Street address:		Mailing Address or PO Box:	
City:	State:	Zip:	
Phone:	Fax:	E-mail:	



09/21/16
form 2 pages

Billing: Owner Clinic 3rd Party (preapproval required) Please note: WADDL policy is to bill the clinic if provided, unless prepaid.
Reporting Preference: Mail Fax Web access - register on web site at <http://waddl.vetmed.wsu.edu>

Please fill out completely as possible:

Specimen(s) Submitted: (Please use WADDL Animal ID Sheet for multiple animals.)	Date Collected: Aug-Sept 2016
nasal swabs-frozen (-20C)	Date Shipped: n/a

<input type="checkbox"/> Necropsy	<input type="checkbox"/> Virology	<input type="checkbox"/> Bacteriology	<input type="checkbox"/> IHC
<input type="checkbox"/> Histopathology	<input type="checkbox"/> Serology	<input type="checkbox"/> Mycoplasma culture	<input checked="" type="checkbox"/> PCR
<input type="checkbox"/> Toxicology	<input type="checkbox"/> Fungal Culture	<input type="checkbox"/> Parasitology	<input type="checkbox"/> Other:

Note: WADDL reserves the right to modify the tests requested for more efficient case work-up and / or to send specimens to outside laboratories to perform testing not done at WADDL.

Animal ID (name/tag#) see multiple animal form	Species domestic goats	Breed multiple	Age multiple	Sex	Animal Weight
Location of Lesion N/A	No. in group	No. Dead N/A	No. Sick N/A	No. on Premises	Duration of Problem N/A

* Was animal euthanized? If so, what method? N/A

Additional History: Vaccinations, signs, stress factors, treatments, post mortem findings, pertinent feed or feed additives, clinical lab results, previous WADDL Case Numbers. (Attach additional sheets as necessary.)

M. ovipneumoniae qPCR on each sample
 Please save remaining DNA isolations and call Maggie for pick up or may request further testing (sequencing) be performed by WADDL, depending on the results of qPCR analysis.

Please bill to ADRU-ARS-USDA account #RSA 2540-1094

WADDL is an official brucellosis testing laboratory. All serology for brucellosis, including abortion screens, requires identification of animals, date of sample collection, and signature of an accredited veterinarian attesting to the following statement:

"I certify that the specimens submitted with this form were collected by me from the animal(s) described on the date indicated."

Veterinarian's, Clinician's or Owner's Signature:	Condition(s) Suspected:
---------------------------------------------------	-------------------------

IDENTIFICATION SHEET FOR MULTIPLE ANIMALS

(To accompany WADDL Accession form, if needed)

Washington Animal Disease Diagnostic Laboratory
 College of Veterinary Medicine, Washington State University
 Mailing address: P.O. Box 647034
 Pullman, WA. 99164-7034
 Phone: (509) 335-9696
 E-Mail: waddl@vetmed.wsu.edu
 Web Site: http://waddl.vetmed.wsu.edu

Shipping address:
 Bustad Hall, Rm. 155-N
 Pullman, WA. 99164-7034
 FAX: (509) 335-7424

2016 – 12311

09/21/16

Ref Vet: Highland, Margaret
 Owner: USDA – ARS – ADRU
 Breed: Domestic Goat
 Routing: md

Owner: Highland, Maggie

Veterinarian: Highland, Maggie

TEST(S) REQUESTED: Movi qPCR

Tube	Animal # or Name	Tube	Animal # or Name	Tube	Animal # or Name	Tube	Animal # or Name
1	2_A	26	2_F	51	4_B (3)	76	1_J (2)
2	2_B	27	2_G	52	4_C (3)	77	1_L (2)
3	2_C	28	2_H	53	4_D (3)	78	1_N (2)
4	2_D	29	2_I	54	4_E (3)	79	1_O (2)
5	9_A	30	2_J	55	4_F (3)	80	1_P (2)
6	9_B	31	3_A	56	4_G (3)	81	1_Q (2)
7	9_C	32	3_B	57	4_H (3)	82	1_R (2)
8	9_D	33	3_C	58	4_I (3)	83	1_S (2)
9	9_E	34	4_A	59	4_J (3)	84	1_T (2)
10	5_A	35	4_B	60	4_K (3)	85	1_U (2)
11	5_B	36	4_C	61	4_L (3)	86	1_V (2)
12	5_C	37	4_D	62	4_M (3)	87	1_W (2)
13	5_D	38	4_E	63	4_N (3)	88	1_X (2)
14	5_E	39	3_A	64	4_O (3)	89	1_Y (2)
15	5_F	40	3_B	65	4_S (3)	90	1_HH (2)
16	5_G	41	3_C	66	9_F (2)	91	1_II (2)
17	5_H	42	3_D	67	9_G (2)	92	1_KK (2)
18	5_I	43	3_E	68	17_J (2)	93	1_LL (2)
19	5_J	44	3_F	69	17_K (2)	94	1_MM (2)
20	5_K	45	3_G	70	1_A (2)	95	1_NN (2)
21	2_A	46	3_H	71	1_B (2)	96	1_SS (2)
22	2_B	47	A_3_I	72	1_D (2)	97	1_ZZ (2)
23	2_C	48	26_A	73	1_E (2)	98	1_BC (2)
24	2_D	49	26_B	74	1_F (2)	99	2_H (4)
25	2_E	50	4_A (3)	75	1_G (2)	100 *	

* For over 100 samples, please copy this form and continue numbering.

Washington Animal Disease Diagnostic Lab

P.O. Box 647034
Pullman, WA 99164-7034
Telephone : (509) 335-9696
Fax : (509) 335-7424

Dr. Margaret Highland
USDA-ARS-ADRU
WSU - 3003 ADBF

Case#: 2016-12311
Report Date: 10/05/16

Pullman, WA 99164-6630

Submittal Date: 09/21/16
Owner: USDA-ARS-ADRU

Species: Domestic Goat

Age:
Sex:

Final Report:

Molecular Diagnostics- Reported on 10/05/16 Authorized by Daniel Bradway, Lab Manager

PCR-Mycoplasma ovipneumoniae SOP: 501.40RT.2016.07.18

Animal	Specimen	Result
[REDACTED] 2.A	Nasal swab	Not detected
[REDACTED] 2.B	Nasal swab	Not detected
[REDACTED] 2.C	Nasal swab	Not detected
[REDACTED] 2.D	Nasal swab	Not detected
[REDACTED] 9.A	Nasal swab	Not detected
[REDACTED] 9.B	Nasal swab	Not detected
[REDACTED] 9.C	Nasal swab	Not detected
[REDACTED] 9.D	Nasal swab	Not detected
[REDACTED] 9.E	Nasal swab	Not detected
[REDACTED] 5.A	Nasal swab	Not detected
[REDACTED] 5.B	Nasal swab	Not detected
[REDACTED] 5.C	Nasal swab	Not detected
[REDACTED] 5.D	Nasal swab	Not detected
[REDACTED] 5.E	Nasal swab	Not detected
[REDACTED] 5.F	Nasal swab	Not detected
[REDACTED] 5.G	Nasal swab	Not detected
[REDACTED] 5.H	Nasal swab	Not detected
[REDACTED] 5.I	Nasal swab	Not detected
[REDACTED] 5.J	Nasal swab	Not detected
[REDACTED] 5.K	Nasal swab	Not detected
[REDACTED] 2.A	Nasal swab	Not detected
[REDACTED] 2.B	Nasal swab	Not detected
[REDACTED] 2.C	Nasal swab	Not detected
[REDACTED] 2.D	Nasal swab	Not detected
[REDACTED] 2.E	Nasal swab	Not detected
[REDACTED] 2.F	Nasal swab	Not detected
[REDACTED] 2.G	Nasal swab	Not detected
[REDACTED] 2.H	Nasal swab	Not detected

Washington Animal Disease Diagnostic Lab

PCR-Mycoplasma ovipneumoniae SOP: 501.40RT.2016.07.18

Animal	Specimen	Result
2.I	Nasal swab	Not detected
2.J	Nasal swab	Not detected
3.A	Nasal swab	Not detected
3.B	Nasal swab	Not detected
3.C	Nasal swab	Not detected
4.A	Nasal swab	Not detected
4.B	Nasal swab	Not detected
4.C	Nasal swab	Not detected
4.D	Nasal swab	Not detected
4.E	Nasal swab	Not detected
3.A	Nasal swab	Not detected
3.B	Nasal swab	Not detected
3.C	Nasal swab	Indeterminate
3.D	Nasal swab	Indeterminate
3.E	Nasal swab	Not detected
3.F	Nasal swab	Indeterminate
3.G	Nasal swab	Not detected
3.H	Nasal swab	Not detected
3.I	Nasal swab	Not detected
26.A	Nasal swab	Not detected
26.B	Nasal swab	Not detected
4.A (3)	Nasal swab	Not detected
4.B (3)	Nasal swab	Not detected
4.C (3)	Nasal swab	Indeterminate
4.D (3)	Nasal swab	Indeterminate
4.E (3)	Nasal swab	Not detected
4.F (3)	Nasal swab	Not detected
4.G (3)	Nasal swab	Not detected
4.H (3)	Nasal swab	Not detected
4.I (3)	Nasal swab	Not detected
4.J (3)	Nasal swab	Indeterminate
4.K (3)	Nasal swab	Not detected
4.L (3)	Nasal swab	Indeterminate
4.M (3)	Nasal swab	Indeterminate
4.N (3)	Nasal swab	Not detected
4.O (3)	Nasal swab	Not detected
4.S (3)	Nasal swab	Indeterminate
9.F (2)	Nasal swab	Not detected
9.G (2)	Nasal swab	Not detected
17.J (2)	Nasal swab	Indeterminate
17.K (2)	Nasal swab	Not detected
11.A (2)	Nasal swab	Not detected
11.B (2)	Nasal swab	Not detected
11.D (2)	Nasal swab	Not detected
11.E (2)	Nasal swab	Not detected
11.F (2)	Nasal swab	Not detected
11.G (2)	Nasal swab	Not detected
11.J (2)	Nasal swab	Not detected
11.L (2)	Nasal swab	Not detected
11.N (2)	Nasal swab	Not detected
11.O (2)	Nasal swab	Not detected
11.P (2)	Nasal swab	Not detected

Washington Animal Disease Diagnostic Lab

PCR-Mycoplasma ovipneumoniae SOP: 501.40RT.2016.07.18

Animal	Specimen	Result
1-Q (2)	Nasal swab	Not detected
1-R (2)	Nasal swab	Not detected
1-S (2)	Nasal swab	Not detected
1-T (2)	Nasal swab	Not detected
1-U (2)	Nasal swab	Not detected
1-V (2)	Nasal swab	Indeterminate
1-W (2)	Nasal swab	Not detected
1-X (2)	Nasal swab	Not detected
1-Y (2)	Nasal swab	Not detected
1_HH (2)	Nasal swab	Indeterminate
1_II (2)	Nasal swab	Not detected
1_KK (2)	Nasal swab	Not detected
1_LL (2)	Nasal swab	Not detected
1_MM (2)	Nasal swab	Indeterminate
1_NN (2)	Nasal swab	Not detected
1_SS (2)	Nasal swab	Not detected
1_ZZ (2)	Nasal swab	Not detected
1_BC (2)	Nasal swab	Not detected
2_H (4)	Nasal swab	Not detected

PCR-Mycoplasma ovipneumoniae test comment: This assay detects only Mycoplasma ovipneumoniae. Culture is available at WADDL to detect other species of Mycoplasma if desired. Fees for culture are available on our website. Please contact the lab if Mycoplasma culture or other testing is desired.

Quantity/Description/Routing of Samples

99 nasal swabs -> MD
per MATH

by M. Highland
-> 250B

2016-12311
Ref Vet: Highland, Margaret
Owner: USDA - ARS - ADPRU
Breed: Domestic Goat
Routed: md

Sample Condition:	<input type="checkbox"/> Room Temp.	<input type="checkbox"/> On ice	<input checked="" type="checkbox"/> Frozen	<input type="checkbox"/> Fixed	Contents match forms: <input type="checkbox"/> Yes <input type="checkbox"/> No Explain below:	Opened by: <i>not</i>
	Samples Received Via:		<input checked="" type="checkbox"/> Drop off			
	<input type="checkbox"/> US Mail	<input type="checkbox"/> FedEx	<input type="checkbox"/> Other:			
	<input type="checkbox"/> UPS	<input type="checkbox"/> FedEx-R				

Comments for Case Tracking:

MD to verify



09/21/16
pages: 1 page

Sample Label <i>[Signature]</i>
