

Date Data Entered: _____

Field Form for Carnivore Backtracks & Collection of Genetics Samples

Backtrack ID¹: _____ **Associated Survey Route²:** _____ **Date:** _____

Observer(s) initials: _____ **Observer(s) Affiliation:** USFS FWP NWC Other:

Species (best guess): Lynx Fisher Wolverine **Confidence:** High Med Low **Distance Backtracked (mi):** ____

Weather³: Sunny P.cloudy Overcast Windy Below freezing Above freezing Lt. snow Heavy snow

Days since snowfall: _____ **Tracking conditions:** Excellent Good Fair Poor **Genetics found:** Yes No

Backtrack Comments:

Genetics sample ID⁴: _____ **Type:** Hair Scat Other (describe):

Location of Genetics sample: Daybed Brush Kill site Other (describe):

Waypoint⁵: _____ E _____ N **Datum/Zone:** _____

Genetics Sample Comments (e.g. number of hairs):

Genetics sample ID⁴: _____ **Type:** Hair Scat Other (describe):

Location of Genetics sample: Daybed Brush Kill site Other (describe):

Waypoint⁵: _____ E _____ N **Datum/Zone:** _____

Genetics Sample Comments:

Notes:

1- Backtrack nomenclature: BT(Backtrack #)_PixelID_Date; for example, the first backtrack you do that day in pixel 1469 on Feb 6 will be called "BT01_1469_020611"

2- This is the survey route you were working on when you came across this track; survey route nomenclature: (PixelID)_Date; for example, if you're surveying pixel 1469 on Feb 6, you'll call the route "1469_020611)

3- Circle all that apply

4- Genetics samples should be named so they're clearly associated with the backtracking effort; nomenclature:

Gen(sample#)_BT(backtrack#)_PixelID_Date; for example, the second hair sample you pick up on the first backtrack in pixel 1469 on Feb 6 would be called "Gen02_BT01_1469_020611"

5- If possible, record waypoints in UTM's (Zone 11 or 12, depending on where you are), NAD 83; if other than this datum, please specify the datum/zone you recorded the waypoint in

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Waypoint⁵: _____ E _____ N **Datum/Zone:** _____

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Genetics Sample Comments: