

**Earth HoloGenome Initiative - Metadata Sheet B**

Animal metadata

Sample metadata

Researcher: \_\_\_\_\_

<b>Specimen ID</b>	Own code of field researcher	<b>Tube code</b>	<b>Type</b>	<b>Sample origin</b>	<b>Freshness</b>	<b>Time to freezing</b>	<p><b>Animal metadata</b></p> <p><b>Trapping method</b> Handnet, Mistnet, Pitfall, Sherman, Shot, Other</p> <p><b>Trapped time</b> Maximum time animal has been trapped before sampling</p> <p><b>Length</b> Bats (forearm length), Birds (tarsus length), Mammals (length of hind foot/ greatest length of ear), Reptiles (snout-tail)</p> <p><b>Sex</b> F=Female, M=Male, U=Unknown</p> <p><b>Development &amp; Reproductive status</b> L=Larva, J=Juvenile, A=Adult, P=Pregnant</p> <p><b>Sample metadata</b></p> <p><b>Type</b> F=Faecal, A=Anal/cloacal swab, O=oral swab, S=Skin swab, B=Blood, T=Tissue</p> <p><b>Origin</b> D=Directly from animal, G=From ground, T=From Trap, H=From isolation housing.</p> <p><b>Freshness</b> Minutes or hours before the sample was stored in a preservative. E.g. 10m or 2h</p> <p><b>Time to freezing</b> Time (hours or days) until the tube with the sample was stored minimum at -20 °C. E.g. 4h, 1d, 7d</p> 
<b>Sampling event ID</b>	Linked to Metadata Sheet A	ABC45					
<b>Species</b>	Scientific name						
<b>Trapping method</b>							
<b>Trapped time</b>							
<b>Length</b>	cm						
<b>Weight</b>	grams						
<b>Sex</b>		Notes:					
<b>Dev &amp; Repro</b>							
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<b>Dev &amp; Repro</b>							

**Please, transfer this information to the electronic metadata sheet**

More information: [www.earthhologenome.org/sampling](http://www.earthhologenome.org/sampling)