

Supplementary Fig. 02 Example of a horizontally oriented extended specimen datasheet for vertebrate capture release study

Site _____ Species _____ Net Date _____ Net Open Time _____ Net Close Time _____ Exit Count _____ Process Date _____ Researchers _____

IDENTITY				TRACK	DIET	MEASUREMENTS					MISC SAMPLES		SWABS	BLOOD				NOTES	
UniqueID #		M/F?	Mom/ baby pair? Y/N	Processing time start	Fed before process- ing? Y/N	Weight (g)	Body (cm)	Fore- arm (mm)	Tibia (mm)	Ear (mm)	Gonads	Hair Y/N	Ectos (# BF, MS, F, M, T)	UR (#)	Amt eppendorf (ul) <i>Target: >400ul</i>	# aliquot serum	Amt RNA-tu be (ul) <i>Target: >40ul</i>	# thin slides	List anomalies, deviations from standard protocol, unreliable equipment. If bat is GPS tagged or necropsied, note here.
		class (M: J./A. F: J/ NL/P/L) * if YOY	ID # of mom/ baby	Processing time stop										# aliquot RBC					
Sample ID	Recap? Y/N	Tooth Assess (new, mild, mod, heavy) Sex:	Amt milk (ul)	Temp							(L)			(W)		Wing Punch in EtOH (#)		THR (#)	
				Start:							L:	Hair:	BF:	UR:		SER:		THN:	
		Class:		Stop:							W:	WP:	MS:	FEC:		PEL:		Filter:	
SampleID:	Recap:	Tooth:		Temp:									F:	THR:		BC:			
		Sex:		Start:							L:	Hair:	BF:	UR:		SER:		THN:	
		Class:		Stop:							W:	WP:	MS:	FEC:		PEL:		Filter:	
SampleID:	Recap:	Tooth:		Temp:									F:	THR:		BC:			
		Sex:		Start:							L:	Hair:	BF:	UR:		SER:		THN:	
		Class:		Stop:							W:	WP:	MS:	FEC:		PEL:		Filter:	
SampleID:	Recap:	Tooth:		Temp:									F:	THR:		BC:			
		Sex:		Start:							L:	Hair:	BF:	UR:		SER:		THN:	
		Class:		Stop:							W:	WP:	MS:	FEC:		PEL:		Filter:	
SampleID:	Recap:	Tooth:		Temp:									F:	THR:		BC:			
		Sex:		Start:							L:	Hair:	BF:	UR:		SER:		THN:	
		Class:		Stop:							W:	WP:	MS:	FEC:		PEL:		Filter:	
SampleID:	Recap:	Tooth:		Temp:									F:	THR:		BC:			
		Sex:		Start:							L:	Hair:	BF:	UR:		SER:		THN:	
		Class:		Stop:							W:	WP:	MS:	FEC:		PEL:		Filter:	
SampleID:	Recap:	Tooth:		Temp:									F:	THR:		BC:			