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# How to study a predator that only eats a few meals a year: high-frequency accelerometry to quantify feeding behaviours of rattlesnakes (*Crotalus* spp.)

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## Abstract

**Background** Many snakes are low-energy predators that use crypsis to ambush their prey. Most of these species feed very infrequently, are sensitive to the presence of larger vertebrates, such as humans, and spend large portions of their lifetime hidden. This makes direct observation of feeding behaviour challenging, and previous methodologies developed for documenting predation behaviours of free-ranging snakes have critical limitations. Animal-borne accelerometers have been increasingly used by ecologists to quantify activity and moment-to-moment behaviour of free ranging animals, but their application in snakes has been limited to documenting basic behavioural states (e.g., active vs. non-active). High-frequency accelerometry can provide new insight into the behaviour of this important group of predators, and here we propose a new method to quantify key aspects of the feeding behaviour of three species of viperid snakes (*Crotalus* spp.) and assess the transferability of classification models across those species.

**Results** We used open-source software to create species-specific models that classified locomotion, stillness, predatory striking, and prey swallowing with high precision, accuracy, and recall. In addition, we identified a low cost, reliable, non-invasive attachment method for accelerometry devices to be placed anteriorly on snakes, as is likely necessary for accurately classifying distinct behaviours in these species. However, species-specific models had low transferability in our cross-species comparison.

**Conclusions** Overall, our study demonstrates the strong potential for using accelerometry to document critical feeding behaviours in snakes that are difficult to observe directly. Furthermore, we provide an 'end-to-end' template for identifying important behaviours involved in the foraging ecology of viperids using high-frequency accelerometry. We highlight a method of attachment of accelerometers, a technique to simulate feeding events in captivity, and a model selection procedure using biologically relevant window sizes in an open-access software for analyzing acceleration data (AcceleRater). Although we were unable to obtain a generalized model across species, if more data are incorporated from snakes across different body sizes and different contexts (i.e., moving through natural habitat), general models could potentially be developed that have higher transferability.

**Keywords** Accelerometry, Animal behaviour, Foraging ecology, *Crotalus oreganus*, *Crotalus horridus*, *Crotalus viridis*, Feeding

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## Background

A basic understanding of the natural history and ecological role of a species often involves monitoring and measuring the behaviours of individuals and establishing an activity budget [1, 2]. Activity budgets and behavioural profiles are particularly critical for understanding how individuals may respond to a changing environment, such as human-induced rapid shifts in temperature [3, 4] and habitat quality [5, 6]. Traditionally, quantifying activity budgets requires numerous hours of direct observation of individuals through methods such as scan sampling or focal sampling [7]. These direct observation methods have been applied to both free-ranging and captive animals across a wide variety of taxa [e.g., 8, 9, 10, 11, 12, 13, 14, 15]. However, a number of constraints make direct observation infeasible for many species. Examples include species that are difficult to observe in their habitat (e.g., dense forest, underground, turbid water, etc.), are only active at night, travel long distances in a short period of time, or are solitary predators, where simultaneous observation across a sample of individuals is impractical. In addition, human presence can drastically alter animal behaviour [16], and individuals may, therefore, not exhibit species-typical behaviour in the presence of human observers.

Recent innovations in animal-borne biologging technologies have provided novel approaches that help circumvent the problems associated with using direct observation to quantify behaviour [17]. The attachment of devices that continuously log acceleration values has been particularly useful for quantifying metrics of movement and behavioral states, as many behaviors result in unique acceleration profiles that can be identified with high accuracy and precision through machine learning methods [19]. The miniaturization and affordability of animal-borne accelerometers has made this approach a key tool for characterizing behaviour of free-ranging animals. Ecologists initially began attaching accelerometers to marine mammals because of the challenges of direct observations on these organisms [18, 19]. Subsequently, terrestrial and freshwater ecologists have adopted this novel method and have been able to use animal-borne accelerometers to quantify activity budgets [e.g., 20, 21], frequencies of key behaviours [e.g., 22, 23], and energy expenditure [e.g., 21, 24]. Accelerometers are now being paired with other biologging devices such as acoustic recorders [25, 26] and global positioning systems [27] to provide even further insight into the activity budgets and behaviour of free-ranging animals and how they may differ across environmentally relevant conditions (e.g., photoperiod, moon phase, seasonality, etc.).

Although accelerometry is becoming an integral and important tool to further understand behaviour,

taxonomic representation in these studies is uneven. The majority of research has focused on larger-bodied mammals, birds, and marine taxa [19]. Because logistical details of how to best attach and configure devices are likely to differ greatly across groups, the widespread adoption of animal-borne accelerometry across different groups of organisms is facilitated by studies that document best practices and provide details on overcoming methodological hurdles. For example, we know comparatively little concerning the detailed behavioural ecology of most snake species, despite the fact that this globally distributed group are often abundant predators that play vital roles in ecological communities. Some snake species also have major impacts on both human health and medicine (snakebite envenoming has been recognized by the World Health Organization as a Neglected Tropical Disease), and others have invaded native ecosystems with devastating effects on local fauna [28–30]. Viperid snakes in particular represent an ideal group for investigating animal behaviour using accelerometry. Detailed studies of the behavioural ecology (specifically foraging ecology) of this group are difficult, because these snakes are cryptic, rarely encountered, sensitive to human presence, and spend long periods of time hidden from view. Although some methodological approaches have been developed to study specific aspects of their foraging ecology (e.g., dietary analysis, fixed videography coupled with radio telemetry), none of these methods [31–35] allow for the quantification of key aspects of hunting behaviour and outcomes of predatory encounters.

Quantifying behavioural data via attaching animal-borne accelerometers to free-ranging individuals is the next step in advancing our knowledge of foraging ecology in viperids. Most species within this group are characterized as ambush (i.e., sit-and-wait) predators and progress through a series of distinctive behavioural stages when capturing prey. A typical feeding event involves: (1) an initial search for an ambush site, (2) a prolonged wait while remaining cryptic, (3) the targeting, striking, and envenomation of any suitable prey that comes within range, (4) the release of envenomated prey to avoid retaliatory attacks, (5) strike-induced chemosensory searching to locate the carcass of envenomated prey, and (6) swallowing and digesting the prey item [reviewed 36]. Many of the behaviours exhibited by viperids during a feeding event are distinct, and could result in unique acceleration signatures that would enable us to quantify the frequency and outcome of predation events across a large sample of free-ranging snakes. Such data would represent the first comprehensive analysis of feeding behaviour within an ecologically critical and globally distributed group of predators. Previous work pioneering the use of low-frequency accelerometers implanted into the body cavities

of Western Diamond-backed Rattlesnakes (*Crotalus atrox*), Burmese Pythons (*Python bivittatus*), and Timber Rattlesnakes (*Crotalus horridus*) illustrates the general utility of this approach [37, 58, 64] and underscores the need to develop new attachment and analysis techniques to quantify detailed behavioural profiles for these predators.

The computational burden imposed by high-resolution accelerometer data remains one of the most significant hurdles to effective use, particularly in validation studies on novel focal taxa [38–40]. Acceleration signatures for unique behaviours can be impacted by multiple factors, such as body size and the duration of the behaviour. Thus, it may not be appropriate to use behavioural classification models developed for one species (or even one population) on another similar species. However, it is plausible that a model developed on one species could have high transferability to another similar species if that species is comparable in body size and if they perform analogous behaviours. Accordingly, accelerometer studies must provide detailed step-wise procedures and, when possible, evaluate the potential for transferability of specific methods across similar focal taxa to streamline future applications. If using a single classification model across multiple species is possible, it should be prioritized, as it could simplify classification and increase the adoption of accelerometers to quantify behaviour of the taxa involved [41, 42]. Viperids are an ideal taxon for applying a single classification model across multiple species, as many are similar in body size and go through the same distinct behavioural stages to capture prey. Thus, we hypothesized that computational models would show high transferability across congeneric viperid species (i.e., high recall, precision, and accuracy), and that future studies could forgo the labor-intensive task of simulating feeding events in captivity to develop adequate classification models.

In this study, we used animal-borne accelerometry to develop behavioural classification models for key feeding behaviours of three species of viperids (Timber Rattlesnakes (*Crotalus horridus*), Western Rattlesnakes (*C. oreganus*), and Prairie Rattlesnakes (*C. viridis*) and assessed the transferability of these models across different species. First, we developed a simple and cost-effective methodology to attach accelerometry devices anteriorly to a venomous snake. Our next objective was to develop and validate machine learning models that would allow us to accurately classify key feeding behaviours in all three species. Finally, we assessed the transferability of species-specific classification models via a cross-species comparison to evaluate the feasibility of building general classification models that would apply across a range of closely related species with similar behavioural profiles.

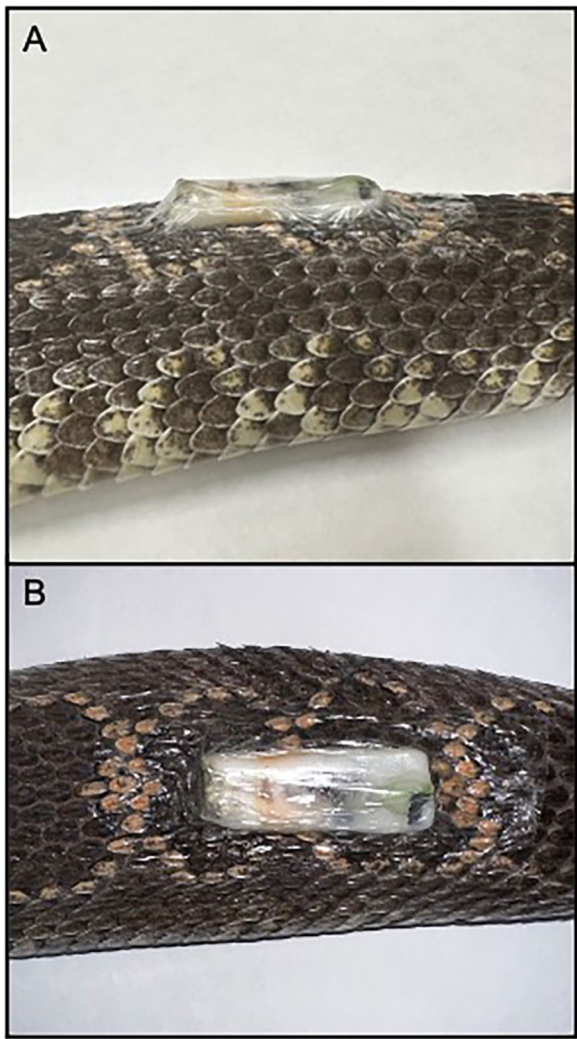
## Methods

### Captive validation

All rattlesnake species were collected from the wild using visual encounter surveys at different locations (*C. horridus* ( $n=17$ ) from Georgia; *C. oreganus* ( $n=14$ ) from Southern California; and *C. viridis* ( $n=7$ ) from Western Texas and Southwestern New Mexico) and transported to either Georgia College and State University (*C. horridus*) or San Diego State University (*C. oreganus* and *C. viridis*).

For each trial, an accelerometer (AXY-5, Technosmart Europe Srl., Rome, Italy) recording tri-axial acceleration values at 25 Hz was attached to the dorsum of individual rattlesnakes (*C. horridus*, *C. oreganus*, or *C. viridis*). Accelerometers were placed anteriorly on each individual at a distance of 25% of the snout vent length of the snake, so that the relative body placement was consistent across individuals. This allowed the three acceleration channels of the device to be placed so they represented the animal's dorso-ventral axis (heave), the anterior–posterior axis (surge), and the lateral axis (sway). The devices were adhered to the snake via a transparent bandage material (3 M™ Tegaderm™; see Fig. 1) that covered the accelerometry device, but only adhered to a few of the scale rows lateral to the dorsal vertebrae. This was important to ensure that sufficient dermal elasticity was retained as prey items were ingested [59]. To stage a feeding trial, each study animal was placed into a custom arena (2×2 m) made of corrugated plastic or glass and video recorded with a stationary camera (PatrolMaster 1296P UHD Body Camera; Amcrest UltraHD Video Security System) recording at 60 frames per second placed above the arena. Immediately after a rattlesnake was placed into an arena, a defensive assay was performed, where personnel agitated the individual with a plush toy attached to a long pole to prompt defensive strike behaviours. After the defensive assay, snakes were allowed to acclimate to the arena for approximately 12 h before a feeding assay was conducted. For the feeding assay, rattlesnakes were video recorded as they were fed a mouse that was allowed to move freely into the arena (*Mus musculus*, their typical food in captivity). It was necessary to use live mice to replicate the feeding behaviors of free-ranging snakes as closely as possible, as past studies have shown that rattlesnakes respond differently to live vs. euthanized prey items [65].

Following exploratory analyses of videos, we chose to group behaviours into four classes that appeared to result in distinct acceleration signatures and corresponded to critical phases of the feeding cycle: still, locomotion, strike, and swallow (Table 1; Fig. 2). Behavioural scores were time-matched to accelerometer readings (to within 1 s) to generate annotated acceleration data sets



**Fig. 1** A lateral (A) and dorsal (B) view of an accelerometer placed posteriorly from the snout at a distance of 25% of the snout-vent length of the snake and adhered via a transparent bandage material (3 M™ Tegaderm™)

by recording a video of the exact time (using the Exact Time™ application by ©Neurovat, 2023) as the accelerometry device began logging data (Technosmart devices emit a visible signal when powering on). We then also recorded the Exact Time™ (©Neurovat, 2023) at the

outset of all videos recording animals wearing accelerometry devices.

**Behavioural classification algorithms**

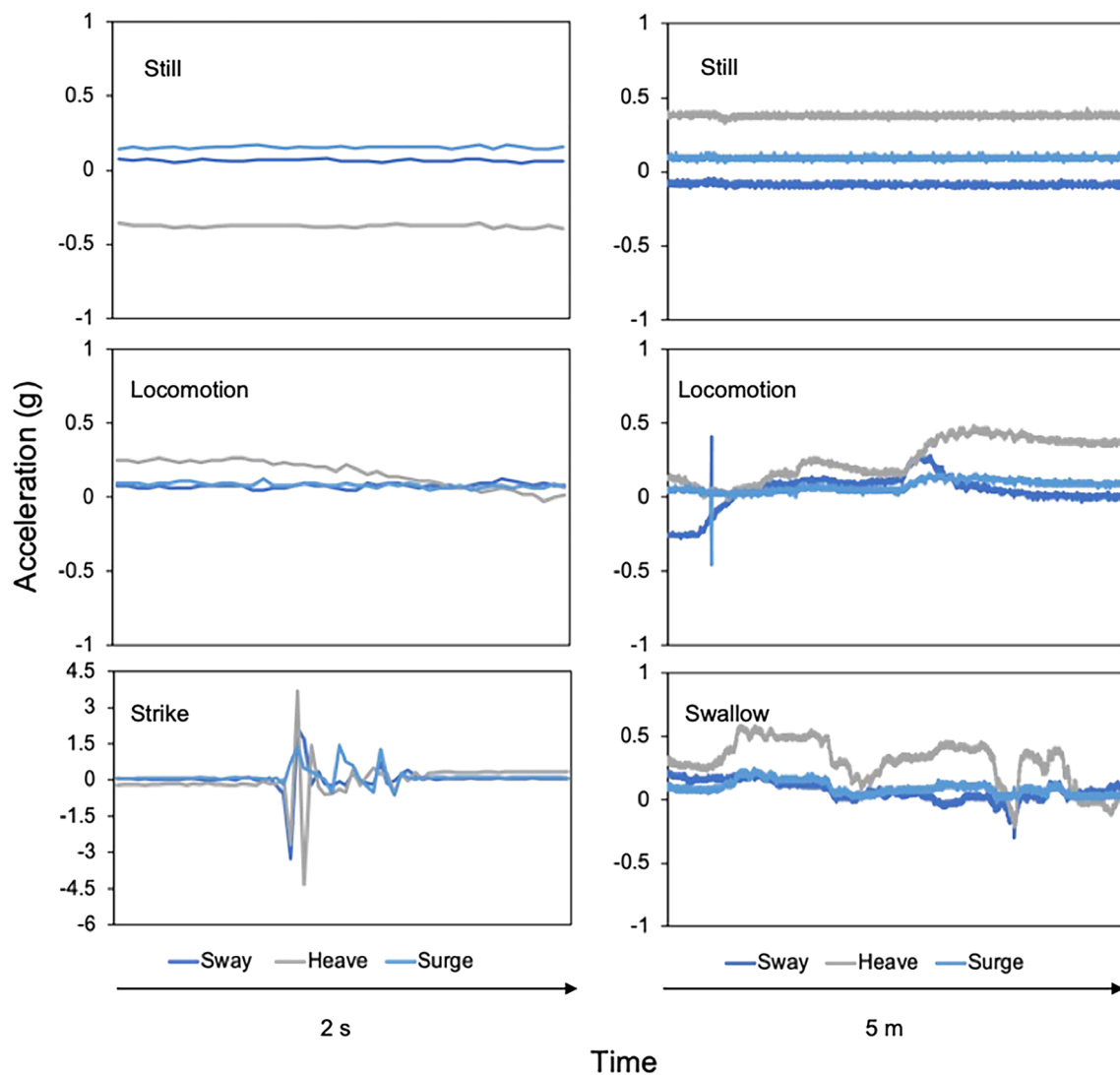
We used the open-access software AcceleRater to train classification algorithms based on our captive training data sets [38]. Because of the extreme differences in the durations of the strike and swallow behaviors (e.g., a typical strike is less than 1 s, while a swallowing event can last over 5 m) leading to complications with window size selection, we analyzed the data separately to identify striking and swallowing behaviors. Thus, we attempted to independently obtain the best model for (1) ‘strike’-focused models, with strike, still, and locomotion as behavior classes; and (2) ‘swallow’-focused models, with swallow, still, and locomotion as behavior classes. In addition, because class imbalance can lead to inference issues when using machine-learning methods [60], all data sets were manually balanced, so that each behavioural class had an equal number of samples relative to the behaviour that had the fewest samples from our captive trials (in all cases, this was either the behaviour swallow or strike). We then ran a linear support vector machine (SVM) [61], a decision tree [62], and a random forest [63] algorithm that either included all summary statistics available in AcceleRater (mean, standard deviation, skewness, kurtosis, maximum, minimum, vector norm, covariance, Pearson correlation, dynamic body acceleration, overall dynamic body acceleration (ODBA), mean-diff, std-diff, wave amplitude, line crossings, 25 percentile, 50 percentile, 75 percentile) or just the summary statistics of mean, standard deviation, and ODBA at window sizes of 3, 6, 12, 24, and 40 s for ‘swallow’ models, and window sizes of 0.4, 0.8, 1.2, and 1.6 s for ‘strike’ models. For each unique model and window size, we trained the model via a (50/50) train-test split, where a random 50% of the data were used to train each model and the remaining 50% of the data set was used to test the model.

We used the same methods as Clermont et al. [43] to identify which algorithm optimally classified behaviours. For each algorithm, a confusion matrix was built in AcceleRater that counted true positives (TP), true negatives (TN), false positives (FP), and false negatives (FN).

**Table 1** Description and function of four rattlesnake behaviours used for accelerometry classification

Behaviour	Description	Function
Still	Not moving while coiled or stretched out	Either in ambush or resting
Locomotion	Any type of body locomotion other than strike or swallow	Movement
Strike	Fast extension of the anterior third of the body toward a target	Envenomate a prey item or defense
Swallow	Prey item passes through gullet underneath accelerometer	Consumption





**Fig. 2** Acceleration signatures of the sway (dark blue), heave (gray), and surge (light blue) axes for the behaviours still, locomotion, and strike at a 2 s time burst, and for the behaviours still, locomotion, and swallow over a period of 5 min

Accuracy, precision, and recall for each behavioural class were calculated using these counts and are defined as follows (redrawn from Clermont et al. [43]). Accuracy is the proportion of correct classifications either into or out of a given behaviour category:

$$\text{Accuracy} = \frac{\text{TP} + \text{TN}}{\text{TP} + \text{TN} + \text{FP} + \text{FN}}$$

Precision is the proportion of classifications into a given behaviour category that were correct:

$$\text{Precision} = \frac{\text{TP}}{\text{TP} + \text{FP}}$$

Recall is the proportion of instances of a behaviour classification into the correct category:

$$\text{Recall} = \frac{\text{TP}}{\text{TP} + \text{FN}}$$

Higher precision denotes fewer false positives and higher recall denotes fewer false negatives.

#### Cross-species comparison: statistical analyses

To assess the transferability of algorithms trained on one rattlesnake species when tested on a different rattlesnake species exhibiting the same behaviour, we used the top performing models from each species to annotate and label already known behaviours. If there was more than one

top model, we used the model that most consistently performed well across each species. Thus, we were able to test how well the best performing ‘strike’ and ‘swallow’ models were able to correctly classify the same behaviours of other species using a factorial experimental design, where we tested (1) *C. horridus* on *C. oreganus* and *C. viridis*; (2) *C. oreganus* on *C. horridus* and *C. viridis*; and (3) *C. viridis* on *C. horridus* and *C. oreganus*. Subsequently, for each test, we calculated overall accuracy (or percent correctness) and the precision, recall, and F1-score of each behavioural class. The F1-score was calculated as follows:

$$F1 \text{ score} = \frac{2 \times \text{Precision} \times \text{Recall}}{\text{Precision} + \text{recall}}.$$

## Results

### Behavioural classification using accelerometry data

The window sizes and classification algorithms that predicted the top model(s) for both ‘strike’ and ‘swallow’ models varied across species (Tables 2, 3; Additional file 1). For *C. horridus* the top performing algorithm for ‘swallow’ models was a random forest that included all summary statistics at a window size of 3 s (all three behaviours with accuracies > 81%) and there were multiple top performing algorithms for ‘strike’ models, where all three behaviours had an accuracy, precision, and recall of 100% (Tables 2, 3). For *C. oreganus*, the top performing algorithm for ‘swallow’ models was a random forest that included all summary statistics at a window size of 6 s (all three behaviours with accuracies > 90%) and the top performing algorithm for ‘strike’ model was a random forest that included all summary statistics at a window size of 0.4 s (all three behaviours with accuracies > 98%;

**Table 2** Accuracy, precision, and recall for the top algorithm, window size, and summary statistics for *Crotalus horridus*, *C. oreganus*, and *C. viridis*. Behaviours included in these models were swallow, locomotion, and still

Species	Window size (s)	Summary statistics	Algorithm	Classification performance	Swallow	Still	Locomotion	Weighted average
<i>C. horridus</i>	3	All	RF	Accuracy	90.83	83.11	81.54	85.16
				Precision	87.12	81.61	66.57	78.43
				Recall	85.38	67.41	81.23	78.00
<i>C. oreganus</i>	6	All	RF	Accuracy	93.62	99.57	94.04	95.74
				Precision	90.06	98.69	92.31	93.69
				Recall	91.19	100.00	90.00	93.73
<i>C. viridis</i>	3	All	RF	Accuracy	88.2	97.35	89.79	91.78
				Precision	88.32	94.3	81.12	87.91
				Recall	74.2	97.95	90.79	87.65

The weighted average across each behaviour is also denoted

RF random forest

**Table 3** Accuracy, precision, and recall for the top algorithm, window size, and summary statistics for *Crotalus horridus*, *C. oreganus*, and *C. viridis*

Species	Window size (s)	Summary statistics	Algorithm	Classification performance	Strike	Still	Locomotion	Weighted average
<i>C. horridus</i>	0.4	All	RF	Accuracy	100.00	100.00	100.00	100.00
				Precision	100.00	100.00	100.00	100.00
				Recall	100.00	100.00	100.00	100.00
<i>C. oreganus</i>	0.4	All	RF	Accuracy	99.45	100.00	99.45	99.64
				Precision	100.00	100.00	98.25	99.42
				Recall	98.39	100.00	100.00	99.46
<i>C. viridis</i>	0.4	All	RF	Accuracy	97.87	100.00	97.87	98.58
				Precision	100.00	100.00	94.12	98.04
				Recall	93.33	100.00	100.00	97.78

Behaviours included in these models were strike, locomotion, and still. The weighted average across each behaviour is also denoted

RF random forest

Tables 2, 3). For *C. viridis*, the top performing algorithm for ‘swallow’ models was a random forest that included all summary statistics at a window size of 3 s (all three behaviours with accuracies > 88%) and there were two top performing algorithms for ‘strike’ models, where all three behaviours had an accuracy, precision, and recall of 100% (Tables 2, 3). Thus, for each species we were able to develop models that performed well and correctly classified our behaviours of interest.

### Cross-species validation

The overall accuracy, and the precision, recall, and *F1*-scores of each behavioural class varied across species tests for ‘swallow’ and ‘strike’ models (Tables 4, 5). The top cross-species ‘swallow’ model (random forest model developed using *C. viridis* data and tested on *C. oregonus*) had an overall accuracy of only 58%. Across all ‘swallow’ models, locomotion behaviour was classified correctly most frequently, but was still often confused with swallow (Table 4; Additional file 2). The top cross-species ‘strike’ model (random forest model created using *C. viridis* data and tested on *C. oregonus*) had an overall accuracy of 73%. In general, cross-species ‘strike’ models were able to classify strikes correctly, but performed poorly at discerning still and locomotion from strike (Table 5; Additional file 2).

### Discussion

Our results indicate that high-frequency acceleration data logged from an anteriorly adhered device has definite potential for quantifying key feeding behaviours in taxa of predators that have proven difficult to study using direct behavioural observation. Using accelerometry data to characterize species-typical hunting behaviours in viperid snakes, we were able to create species-specific models that classified the behaviours strike, swallow, locomotion, and remaining still with high precision, accuracy, and recall. The high success rates of our preliminary models show that using this technique could have strong ecological relevance in future studies and open up new opportunities for quantifying behaviour across viperids and other snakes. In addition, we identified a low cost, reliable, non-invasive attachment method for accelerometry devices to be placed anteriorly on snakes, as is likely necessary for accurately classifying distinct behaviours in these species. However, species-specific models had low transferability in our cross-species comparison. Thus, it may be necessary to develop and validate classification models within the same species to avoid any species-specific differences in behaviour or body size variance that could affect acceleration data. Our study also took place in a captive environment, and additional field

**Table 4** Overall accuracy (i.e., percent correctness) of each test and precision, recall, and *F1*-scores for each behavioural classification when applying top swallow algorithm for each species of *Crotalus* to the two other species

Species model	Tested species	Window size (s)	Summary statistics	Algorithm	Classification performance	Swallow	Still	Locomotion	Weighted average	Overall accuracy
<i>C. horridus</i>	<i>C. oregonus</i>	3	All	RF	Precision	0.51	0.42	0.73	0.55	0.47
					Recall	0.50	0.76	0.16	0.47	
					<i>F1</i> -score	0.51	0.54	0.27	0.44	
<i>C. horridus</i>	<i>C. viridis</i>	3	All	RF	Precision	0.50	0.53	0.66	0.56	0.54
					Recall	0.56	0.68	0.39	0.54	
					<i>F1</i> -score	0.53	0.6	0.49	0.54	
<i>C. oregonus</i>	<i>C. horridus</i>	6	All	RF	Precision	0.64	0.87	0.46	0.66	0.53
					Recall	0.91	0.07	0.62	0.53	
					<i>F1</i> -score	0.61	0.12	0.63	0.46	
<i>C. oregonus</i>	<i>C. viridis</i>	6	All	RF	Precision	0.45	1.00	0.57	0.67	0.51
					Recall	0.73	0.00	0.79	0.51	
					<i>F1</i> -score	0.56	0.01	0.67	0.41	
<i>C. viridis</i>	<i>C. horridus</i>	3	All	RF	Precision	0.37	0.50	0.37	0.49	0.42
					Recall	0.83	0.18	0.25	0.42	
					<i>F1</i> -score	0.51	0.26	0.36	0.38	
<i>C. viridis</i>	<i>C. oregonus</i>	3	All	RF	Precision	0.47	0.71	0.66	0.61	0.58
					Recall	0.46	0.61	0.68	0.58	
					<i>F1</i> -score	0.55	0.66	0.54	0.58	

The top algorithm, window size, summary statistics, and weighted averages are also denoted

RF random forest

**Table 5** Overall accuracy (i.e., percent correctness) of each test and the precision, recall, and *F1*-scores for each behavioural classification when applying top strike algorithm for each species of *Crotalus* to the two other species

Species model	Tested species	Window size (s)	Summary statistics	Algorithm	Classification performance	Strike	Still	Locomotion	Weighted average	Overall accuracy
<i>C. horridus</i>	<i>C. oregonus</i>	0.4	All	RF	Precision	1.00	0.38	0.49	0.62	0.63
					Recall	0.88	0.02	1.00	0.63	
					<i>F1</i> -score	0.93	0.05	0.65	0.54	
<i>C. horridus</i>	<i>C. viridis</i>	0.4	All	RF	Precision	1.00	0.00	0.45	0.48	0.55
					Recall	0.73	0.00	0.91	0.55	
					<i>F1</i> -score	0.85	0.00	0.60	0.48	
<i>C. oregonus</i>	<i>C. horridus</i>	0.4	All	RF	Precision	0.4	0.00	0.78	0.39	0.46
					Recall	1.00	0.00	0.39	0.46	
					<i>F1</i> -score	0.57	0.00	0.52	0.36	
<i>C. oregonus</i>	<i>C. viridis</i>	0.4	All	RF	Precision	0.34	0.00	0.89	0.41	0.36
					Recall	0.99	0.00	0.09	0.36	
					<i>F1</i> -score	0.51	0.00	0.16	0.22	
<i>C. viridis</i>	<i>C. horridus</i>	0.4	All	RF	Precision	0.75	0.00	0.60	0.45	0.67
					Recall	1.00	0.00	1.00	0.67	
					<i>F1</i> -score	0.86	0.00	0.75	0.54	
<i>C. viridis</i>	<i>C. oregonus</i>	0.4	All	RF	Precision	1.00	0.56	0.93	0.83	0.73
					Recall	0.98	1.00	0.21	0.73	
					<i>F1</i> -score	0.99	0.72	0.35	0.68	

The top algorithm, window size, summary statistics, and weighted averages are also denoted

RF random forest

validation sampling may be necessary to build models applicable to free-ranging snakes.

### Species-specific models

For all three species (*C. horridus*, *C. oregonus*, and *C. viridis*), we created a model selection framework to determine the top performing models that could classify swallow and strike behaviours, alongside both still and locomotion. We used what we deemed as biologically relevant window sizes for each separate ‘swallow’ and ‘strike’ model to determine the top performing model. Furthermore, we tested three different supervised machine-learning algorithms across varying biologically relevant window sizes, providing a general framework within AcceleRater to find a top performing model. For our results, the best performing algorithms were almost all random forests, which was unsurprising, as it is typically the approach with the highest success rate in other studies using supervised machine-learning algorithms to classify animal behaviour [27, 43–45]. However, some studies are beginning to explore unsupervised machine-learning algorithms (e.g., ‘deep learning’; [39, 46–48]) which could benefit the classification of animal behavior using acceleration data in the future. Nonetheless, our

top performing ‘strike’ and ‘swallow’ models had high accuracy, precision, and recall within all three species.

For all three species, the behaviour class strike had very high accuracy (> 97%), precision (all models = 100%), and recall (> 93%). This was expected, as rattlesnake strikes are extremely fast and have a short duration, thus it was predictable that a machine-learning algorithm could develop a model able to classify the behaviour strike from still and locomotion. Furthermore, the behaviour class swallow also had high accuracy (> 88%), precision (> 87%), and recall (> 74%) across all three species. This behaviour, which represents the movement of the body when a prey item passes down the esophagus towards the stomach of the animal and underneath the dorsally attached accelerometer, represents a key step in studying feeding ecology in free-ranging snakes. Past studies of viperid feeding ecology have quantified feeding behaviors in free-ranging snakes using a combination of radio tracking and fixed videography to assemble large data sets on ambush behaviour, quantification of site residence times, prey encounter rates, and strike success rates [31–35]. However, because viperid snakes almost always release struck prey after envenomation, videography studies have not been able to estimate the actual frequency of successful feeding events, given that the swallowing of prey or failure to locate prey items almost always occurs once the



animals have left the video frame. Our study indicates that validated machine-learning models applied to accelerometry data sets from free-ranging snakes should be able to accurately estimate the true number of prey items ingested by snakes. We suspect that swallowing is amenable to accurate classification, because it involves a pattern of upward movement on the accelerometer that is likely rare and should most often occur in temporal proximity to a strike. Nevertheless, additional validation may be necessary before applying our models to free-ranging snakes, as we only tested individuals feeding on a narrow size range of prey. Although the mice we used in our study are representative of the 'typical' prey size taken by these individuals in nature, free-ranging viperids swallow a range of carcass sizes [49–51] and a more accurate validation set might need to include an equivalent range of prey size.

The behaviour classes locomotion and still were included in each 'strike' and 'swallow' model. In all cases, our top performing models classified both behavioural classes locomotion and still with high accuracy, precision, and recall (Tables 2, 3). Thus, our accurate classification of behaviours involved in feeding events across these three viperid species is similar to the performance reached with other predator species [25, 27, 43, 52].

#### Accelerometer attachment

In addition to developing models to predict behaviours involved with foraging ecology in viperids, we describe a non-invasive procedure for attaching accelerometers anteriorly to viperids and potentially other small vertebrates as well (e.g., other snakes, lizards, etc.). Many feeding behaviour movements involve only the head and upper body, so attaching the accelerometer close to the head is probably necessary for the success of this technique. For example, feeding strikes may involve a rapid forward surge of the head and neck region, with the lower body remaining more or less still [53], and prey passing through the gullet are then held in the stomach for a prolonged digestion period, making it unlikely that lower regions of the body exhibit unique acceleration patterns for either striking or swallowing behaviours.

We investigated several ways to attach accelerometers, but found that attaching the accelerometer via a transparent bandage material (3 M™ Tegaderm™) on the dorsum with the leading edge at 25% SVL was the simplest and most effective technique. Although being close to the head is desirable for accelerometer attachment, the degree to which devices can be attached anteriorly is also limited by the size of the device relative to the snake. Snakes ingest comparatively large prey items relative to their head and body size, and the elastic skin of their anterior body must be able to expand during ingestion. A

device attached to the dorsum with adhesive tape should only adhere to the few scale rows lateral to the vertebrae, lest it interfere with the snake's ability to ingest large prey. All three of our species tested were broadly similar in body size, with adults ~600–1100 mm in SVL, and we found that attaching devices at 25% SVL did not interfere with normal movement or feeding behaviours. However, we suspect this attachment methodology may need to be modified for smaller-bodied snakes.

Although we recognize the use of an adhesive bandage is a short-term attachment method (snakes generally shed the device during each ecdysis cycle), high-frequency accelerometry on relatively small-bodied vertebrates is inherently limited in duration. Devices small enough for external attachment (our customized AXY-5; Technosmart Europe Srl., Rome, Italy) were configured as flat rectangular packages (32 L × 10 W × 8 H mm) and could collect ~30 days of data at a sampling frequency of 25 Hz. Preliminary field tests of our technique attaching devices to free-ranging *C. viridis* have found that devices generally remain attached for the entire 30-day battery life of the device, although detachments are frequent enough that we adhere a 0.5 g micro-VHF transmitter to the accelerometer, so that we can recover any devices that detach (RJH, JLH, and RWC, pers. observation). We also caution that the success of this technique may be somewhat species-specific; in the current study, we qualitatively noted significant variation in adherence of devices among species, with *C. viridis* and *C. oreganus* retaining devices for long periods of time, whereas devices attached to *C. horridus* detached much more readily during both captive and field tests (DLD, AFT, and MLT, pers. observation).

#### Cross-species comparison

Although accelerometry models are typically developed and tested within a species, several other research groups have also noted the potential utility of cross-species modeling [42, 54–57]. Previous studies have shown both high and low transferability of models across similar species, but when more complicated behaviours are included, the models typically have low transferability. For example, Dickinson et al. [57] examined model transferability of a captive phylogenetically similar species and captive conspecifics (surrogate species) in Caprids for calibrating behavioural classification of 11 different behaviours and found low model transferability (<55%). However, the possibility of cross-species models is still important to assess, because the use of a previously developed and validated model could save researchers extensive time and effort in creating a unique validation data set when a suitable one already exists. For instance, Auge et al. [42] found high model transferability across two species

of freshwater turtles, where accelerometry was coupled with water sensing technology to classify activity states. Therefore, a general model could be useful in many different scenarios, including instances when behavioural categories are broad (e.g., moving vs. still), free-ranging animals are difficult to record, or the target species is especially cryptic and difficult to find.

Despite highly accurate species-specific models, we found low transferability of all of our top performing models across species. Overall accuracy ranged from 42–58% to 36–73% for ‘swallow’ and ‘strike’ models, respectively. Across all ‘swallow’ models, we found that locomotion was classified correctly most frequently, but still was often confused with swallow (Table 4; Additional file 2). This may be because the behavioural class swallow is similar to still, where the accelerometer remains motionless until the prey item passes down the esophagus and underneath the accelerometer, presumably forcing the accelerometer upwards. Broadly, cross-species ‘strike’ models were able to classify strikes correctly, but performed poorly at discerning still and locomotion from strikes (Table 5; Additional file 2). Although it is unclear why the transferability of ‘strike’ models was low, it is possible that there was too much variation in body size across the three species, or that snake strikes are more variable in duration and acceleration than is apparent from direct observation. However, it is important to note that the *C. viridis* ‘strike’ model had a reasonably accurate transferability of 67% and 73% overall accuracy on *C. horridus* and *C. oreganus*, respectively. Consequently, a data set that includes a larger sample of individuals across more variable body sizes may increase overall transferability among species. Although we are still uncertain why models were not generalizable across species, we think it is unlikely that different rattlesnake species differ fundamentally from one another in the kinematics of their various feeding behaviors, and that generalizable models may be achieved by building larger validation data sets that incorporate a broader range of snake sizes and movement contexts (i.e., including movements of free-ranging individuals through natural habitats, rather than relying solely upon captive individuals). Such data sets should be able to train models that can recognize the broader range of variation present in behavioral classifications. We also suggest that future approaches should experiment with machine-learning algorithms that use more complex and involved training methods (such as over-sampling, under-sampling, moving/sliding window sizes, etc.).

## Conclusions

Overall, our study demonstrates the strong potential for using accelerometry to document critical feeding behaviours in snakes that are difficult to observe directly. Many

snakes are extreme low-energy specialists, and may only feed once every few months—a pattern that has generally stymied attempts to fully document predation behaviours for most species. Furthermore, we provide an ‘end-to-end’ template for identifying important behaviours involved in the foraging ecology of viperids using tri-axial accelerometry. We highlight a method of attachment of accelerometers, a technique to simulate feeding events in captivity, and a model selection procedure using biologically relevant window sizes in an open-access software for analyzing acceleration data. Although we were unable to obtain a generalized model across congeneric species, we believe this goal is still feasible if more data are incorporated from a broader range of movement and feeding contexts, including snakes with more varied body sizes moving in more naturalistic settings. This goal is important for driving more widespread adoption of accelerometry tools within particular taxonomic groups, as researchers starting with a general viperid feeding model could spend much less effort in validation, and more effort in characterizing movement and behavior under varied ecological contexts that would lead to rich comparative data sets.

## Abbreviations

ODBA	Overall dynamic body acceleration
TP	True positive
TN	True negative
FP	False positive
FN	False negative

## Supplementary Information

The online version contains supplementary material available at <https://doi.org/10.1186/s40317-023-00332-3>.

**Additional file 1.** Model results for species-specific algorithms at varying window sizes.

**Additional file 2.** Cross-species model performance statistics.

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## Author contributions

RJH, DLD, and RWC conceived and designed the study. RJH, DLD, JLH, AFT, MT and RWC collected the data. RJH, TM, and JS analyzed the data. RJH wrote the original draft of the manuscript. RWC, DLD, RJH, TEH, and JS provided supervision and mentorship. All authors critically contributed to the drafts and gave final approval for publication. All authors read and approved the final manuscript.

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## Availability of data and materials

The full data sets used and/or analyzed during the current study are available from the corresponding author.

## Declarations

### Ethics approval and consent to participate

All procedures on vertebrates involved in this study were approved by the San Diego State University Institutional Animal Care and Use Committee (22-07-008C) and the Georgia College and State University Institutional Animal Care and Use Committee (2020-D).

### Consent for publication

Not applicable.

### Competing interests

The authors declare that they have no competing interests.

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