## **Supplementary Materials and Methods**

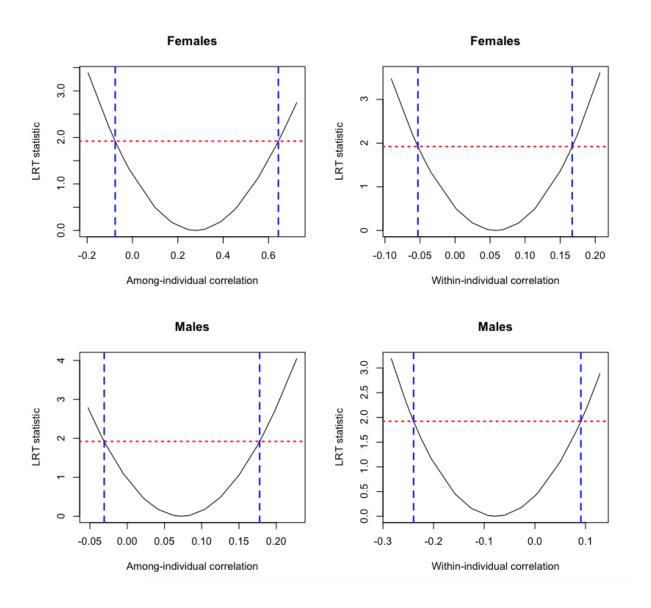
## **Sprint Speed**

Camera views overlapped to ensure that each run attempt was captured in full, from a standing start to maximum speed (which occurred approximately 150 cm along the track). A 1 m long tape measure was fixed to the rubber floor across the central area to calibrate movement speed of each quoll. A reference point was also placed in between the shoulder blades of each quoll - a 4  $\times$  4 cm piece of masking tape with a black dot, visible from the cameras above. Quolls were held 1.25 m from the start of the racetrack, and when released were immediately chased towards the opposite end, 3.25 m from the start line. This method ensured that a minimum of 4 strides of linear acceleration were achieved within the 2 m field of view of the camera.

## **Bite Force**

Bite force was quantified using a custom-built force-transducer (as per Cameron et al., 2013) consisting of a secured unit of two metal plates ( $70 \times 10 \times 2$  mm) separated by a third, larger steel metal pivot plate (3 mm thick). The two smaller plates protruded 12 mm beyond the pivot plate, with a strain gauge (RS Electronics, Sydney, Australia) attached via epoxy resin on the top metal plate. The output from the strain gauge was connected to a custom-made Wheatstone bridge linked to a bridge amplifier (ADInstruments, Sydney Australia). A defined biting point and surface was created on the two protruding plates by covering each in three layers of flesh-like tape (Elastoplast, Beiersdorf, North Ryde, NSW, Australia).

## Results



**Fig. S1.** Plots of profile likelihood tests of significance of among-individual and within-individual correlations between sprint speed and bite force in female and male northern quolls (*D. hallucatus*). Upper and lower 95% confidence intervals are represented by dashed blue lines and the likelihood ratio test (LRT) statistic is marked by the red dashed line. Profile likelihoods were estimated from bivariate mixed models, and calculated using the nadiv package in R. Confidence intervals overlapping 0 indicate non-significant correlations between traits sprint speed and bite force.

**Table S1.** Table of summary statistics from ANOVA conducted on linear mixed effects model of the effect of reproductive season on sprint speed and bite force in male and female northern quolls (*Dasyurus hallucatus*)

	Sum Sq	Mean Sq	NumDF	DenDF	F value	Pr(>F)
Sprint Speed						
Females						
Body Size	0.09	0.09	1	35.34	0.11	0.74
<b>Body Condition</b>	0.44	0.44	1	48.74	0.56	0.46
Season	1.22	0.61	2	40.83	0.77	0.47
Test Number	0.19	0.19	1	47.61	0.24	0.63
<u>Males</u>						
Body Size	0.57	0.57	1	19.29	1.23	0.28
Body Condition	2.15	2.15	1	32.27	4.60	0.04 *
Season	4.67	2.33	2	45.59	5.00	0.01 *
Test Number	<0.001	<0.001	1	55.22	<0.001	1.00
Bite Force						
<u>Females</u>						
Body Size	1.05	1.05	1	35.37	6.82	0.01 *
<b>Body Condition</b>	1.23	1.23	1	42.18	8.00	0.01 *
Season	1.38	0.69	2	37.21	4.49	0.02 *
Test Number	0.07	0.07	1	42.62	0.43	0.52
<u>Males</u>						
Body Size	30.15	30.15	1	23.20	50.02	<0.001 ***
<b>Body Condition</b>	0.13	0.13	1	72.44	0.22	0.64
Season	1.23	0.62	2	70.84	1.02	0.37
Test Number	0.44	0.44	1	47.87	0.72	0.40

**Table S2.** Summary statistics (estimates and 95% CI) from a bivariate mixed model for female northern quolls considering all trials, but retaining only the maximum values for an individual on a given day, of (a) sprint speed and (b) bite force (both standardised to a mean of 0 and a variance of 1).

	Sprint Speed			Bite Force				
	Estimate		95% CI		Estimate		95% CI	
			Lower	Upper			Lower	Uppe
Females								
Fixed Effects								
PC <sub>bodysize</sub>	0.01		-0.12	0.13	0.24	***	0.16	0.32
PCcondition	0.11		-0.08	0.31	0.22	*	0.09	0.35
Age	0.02		-0.36	0.39	0.01		-0.22	0.25
Season		**				**		
Breeding	-0.36		-0.62	-0.10	-0.43		-0.61	-0.24
Post-Breeding	0.04		-0.36	0.44	0.12		-0.15	0.39
Test Number	-0.07		-0.15	0.01	0.02		-0.03	0.07
Variance Components								
Vind	0.30		0.12	0.47	0.04		-0.02	0.10
Ve	0.66		0.51	0.81	0.37		0.29	0.45
Repeatability (R)			0.45	0.40			0.04	0.05
	0.31		0.15	0.46	0.11		-0.04	0.25
Correlation [95% CI]				0.401.6	00 0 571			
Among-individual [r <sub>ind</sub> ] Within-individual [r <sub>e</sub> ]								
within-individual [rej				0.07 [—0	7.00,0.10]			
Males Fixed Effects								
PC <sub>bodysize</sub>	0.07		0.01	0.13	0.23	***	0.18	0.27
PCcondition	0.10		-0.08	0.27	0.06		-0.09	0.20
Age						*		
Season	-0.05		-0.34	0.23	-0.16		-0.45	0.12
Breeding	-0.07		-0.62	0.48	0.37		-0.15	0.88
Post-Breeding	-0.19	*	-0.35	-0.03	-0.06		-0.20	0.08
Test Number	0.10		-0.08	0.27	0.06		-0.09	0.20
Variance Components								
$V_{ind}$	0.56		0.33	0.79	0.03		-0.13	0.18
Ve	0.41		0.28	0.53	0.68		0.48	0.87
Repeatability (R)			0.15	0 = 0			<b>6</b> 4 =	6
	0.58		0.43	0.73	0.04		-0.18	0.25
Correlation [95% CI]								
Among-individual [r <sub>ind</sub> ]					.07;0.15]			
Within-individual [r <sub>e</sub> ]				-0.04 [-0	0.20;0.13]			

Fixed effects included are the same as Table 2 and 3, except for run trial number which was not fitted in this model because only a maximum value per session for each trait was taken. Significant fixed effect estimates are bolded; Signif. codes:  $0^{\text{t****}}$  0.001 '\*\*'. Variance components of the random effects include individual identity ( $V_{\text{ind}}$ ) and residual ( $V_{\text{e}}$ ), as well as among individual ( $r_{\text{ind}}$ ) and within-individual ( $r_{\text{e}}$ ) correlation estimates. Variance components were used to calculate repeatability.

**Table S3.** Repeatability estimates (*R*) and 95% confidence intervals (CI) in (a) body size, (b) body mass, and (c) body condition at each temporal scale extracted from univariate models for wild female and male northern quolls (*Dasyurus hallucatus*)

	Body Size		Body Mas	s		Body Co	Body Condition			
	Estimate	959	95% CI		95%	6 CI	Estimate	95%	95% CI	
		Lower	Upper	-	Lower	Upper	-	Lower	Upper	
Females										
Variance										
Components										
$V_{ind}$	0.66	0.33	0.98	0.10	0.05	0.15	0.16	0.07	0.26	
$V_{ m year}$	0.33	0.10	0.56	0.01	-0.02	0.04	0.04	-0.04	0.13	
$V_{\sf season}$	0.09	-0.01	0.19	0.05	0.03	0.08	0.08	-0.01	0.18	
$V_{e}$	0.10	0.02	0.18	0.03	0.01	0.05	0.11	0.03	0.19	
Repeatability										
R <sub>long-term</sub>	0.56	0.35	0.76	0.53	0.35	0.70	0.41	0.21	0.61	
<b>R</b> year	0.84	0.78	0.90	0.58	0.44	0.71	0.52	0.37	0.66	
R <sub>season</sub>	0.91	0.84	0.99	0.86	0.76	0.97	0.72	0.51	0.93	
R <sub>short-term</sub>	-	-	-	-	-	-	-	-	-	
Males										
Variance										
Components										
$V_{ind}$	6.46	4.78	8.15	1.27	0.91	1.62	0.35	0.21	0.49	
$V_{\sf season}$	0.26	0.13	0.39	0.01	-0.12	0.14	<0.01	<0.01	<0.01	
$V_{e}$	0.09	0.02	0.16	0.20	0.06	0.33	0.29	0.20	0.37	
Repeatability										
Rlong-term	0.95	0.93	0.97	0.86	0.81	0.92	0.55	0.41	0.70	
R <sub>season</sub>	0.99	0.98	0.99	0.87	0.77	0.96	0.55	0.41	0.70	
Rshort-term	-	-	-	-	-	-	-	-	-	

Repeatability of performance traits at each temporal scale were calculated using variance components extracted from each univariate model. Variance components of the random effects included in the models: individual identity ( $V_{ind}$ ), age ( $V_{year}$  – females only), season ( $V_{season}$ ), day of test number ( $V_{day}$ ), and residual ( $V_e$ ).

**Table S4.** Results of linear models testing for significance between females and males in Mass, Body Size (PC1), Body Condition (PC2), Maximum Bite Force (N), and Maximum Sprint Speed (m s<sup>-1</sup>)

Group Average							
	Females	Males	t value	df	Pr (> t )		
Mass (g)	337	486	-19.27	516.05	< 0.001		
Body Size (PC1)	-1.25	2.03	<b>-</b> 24.39	526.34	< 0.001		
<b>Body Condition</b>	0.12	-0.19	5.99	740.85	< 0.001		
(PC2)							
Max Bite (N)	77.6	107.8	3.68	287.47	< 0.001		
Max Sprint (m s <sup>-1</sup> )	4.17	4.35	1.08	279.73	0.28		

**Table S5.** Summary statistics (estimates and 95% CI) from a univariate mixed model of the effects of fixed variables on body condition (PC<sub>condition</sub>) for male and female northern quolls (*D. hallucatus*).

	Body Condition					
	Estimate		95% CI			
			Lower	Upper		
Fixed Effects						
$PC_{bodysize}$	0.17	***	0.14	0.20		
Sex		***				
Sex_male	-0.11		-0.38	0.15		
Season	-0.01	***	-0.05	0.06		
Test Sequence	0.01		-0.01	0.04		
Sex × Season						
Sex_male × Season	-0.36	***	-0.45	-0.26		