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### **Androgens predict parasitism in female meerkats: a new perspective on a classic trade-off**

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- Results: Relationships between social and reproductive status and FAM concentrations
- Table S1: Factors associated with infection status in female meerkats
- Table S2: Factors associated with parasite abundance in female meerkats
- References

#### **Results: Relationships between social and reproductive status and FAM concentrations**

For all females, the mean FAM concentration was  $73.28 \text{ ng/g} \pm 7.40$ . In line with our previous study (1), FAM concentrations were greater in dominant (mean  $\pm$  SE:  $104.91 \text{ ng/g} \pm 16.46$ ) compared to subordinate (mean  $\pm$  SE:  $59.13 \text{ ng/g} \pm 6.77$ ) females ( $t_{53} = 2.71$ ,  $P = 0.009$ ); however, FAM concentrations varied widely between individual females of the same status (dominant T range: 19.55 - 279.43 ng/g; subordinate T range: 4.43 - 191.55 ng/g). In contrast to our previous findings (1), female FAM concentrations did not vary by reproductive state ( $P > 0.05$ ) and this likely owes to the smaller sample size of the current, versus the previous, study.

**Table S1.** Factors associated with infection status in female meerkats (as determined by GLMs and GLMMs).

Parasite	Predictor <sup>a</sup>	Estimate (SE)	$\chi^2$ <sup>b</sup>	<i>p</i>
<i>Toxocara suricattae</i> <sup>c</sup>	FAM		0.106	0.744
	<b>Social status</b>	<b>1.052 (0.618)</b>	<b>2.909</b>	<b>0.088†</b>
	FAM:Status		0.207	0.65
Spirurida nematode <sup>d</sup>	<b>FAM</b>	<b>3.62 (1.39)</b>	<b>9.6745</b>	<b>0.002*</b>
	Social status		2.791	0.095
	FAM:Status		2.298	0.125
<i>Oxynema suricattae</i> <sup>d</sup>	<b>FAM</b>	<b>2.556 (1.114)</b>	<b>6.646</b>	<b>0.01*</b>
	<b>Social status</b>	<b>1.71 (0.861)</b>	<b>4.784</b>	<b>0.029*</b>
	FAM:Status		0.261	0.609
<i>Pseudandrya suricattae</i> <sup>d,e</sup>	<b>FAM</b>	<b>2.74 (1.176)</b>	<b>7.081</b>	<b>0.008**</b>
	Social status		0.351	0.55
Coccidia <sup>d</sup>	<b>FAM</b>	<b>2.836 (1.241)</b>	<b>8.44</b>	<b>0.004**</b>
	Social status		0.007	0.935
	FAM:Status		0.674	0.412

Note. Factors in bold were included in the final model. Variance in infection with Strongylate nematode was low and precluded statistical analysis. Significance is marked as follows: \*\* $p < 0.01$ , \* $p < 0.05$ , † $p < 0.1$ .

<sup>a</sup> Social status comparisons are made against dominant females.

<sup>b</sup>  $\chi^2$  = likelihood ratio test statistic; d.f. = 1

<sup>c</sup> Results of a GLMM with Individual as a random factor.

<sup>d</sup> Results of a GLM.

<sup>e</sup> Low prevalence precluded our analysis of the FAM and status interaction.

**Table S2.** Factors associated with parasite abundance in female meerkats (as determined by GLMs and GLMMs).

Parasite	Predictor <sup>a</sup>	Estimate (SE)	$\chi^2$ <sup>b</sup>	p
Strongylate nematode <sup>c</sup>	FAM		2.211	0.137
	Social status		0.419	.517
	FAM:Status		0.807	0.369
<i>Toxocara suricattae</i> <sup>d</sup>	FAM		0.478	0.49
	Social status		1.357	0.244
	FAM:Status		1.506	0.471
Spirurida nematode <sup>d</sup>	<b>FAM</b>	<b>1.722 (0.993)</b>	<b>3.101</b>	<b>0.078†</b>
	Social status		0.705	0.401
	FAM:Status		1.391	0.238
<i>Oxyntema suricattae</i> <sup>d</sup>	<b>FAM</b>	<b>1.274 (0.433)</b>	<b>8.918</b>	<b>0.003**</b>
	<b>Social status</b>	<b>0.675 (0.33)</b>	<b>4.08</b>	<b>0.043*</b>
	FAM:Status		0.295	0.587
	<b>Cov: Weight</b>	<b>-0.002 (0.001)</b>	<b>5.383</b>	<b>0.02</b>
<i>Pseudandrya suricattae</i> <sup>d</sup>	<b>FAM</b>	<b>1.614 (0.927)</b>	<b>2.77</b>	<b>0.096†</b>
	Social status		0.056	0.813
	FAM:Status		0.151	0.698
Coccidia <sup>d</sup>	<b>FAM</b>	<b>1.068 (0.346)</b>	<b>8.902</b>	<b>0.003**</b>
	Social status		0.077	0.782
	FAM:Status		0.093	0.76

Note. Factors in bold were included in the final model. Significance is marked as follows: \*\* $p < 0.01$ , \* $p < 0.05$ , † $p < 0.1$ .

<sup>a</sup> Social status comparisons are made against dominant females.

<sup>b</sup>  $\chi^2$  = likelihood ratio test statistic; d.f. = 1

<sup>c</sup> Results of a GLMM with Individual as a random factor.

<sup>d</sup> Results of a GLM.

## References

1. Davies C, Smyth KN, Greene LK, Walsh D, Mitchell J, Clutton-Brock TH, Drea CM. (In Revision for Scientific Reports) Exceptional endocrine profiles characterize the meerkat: sex, status, and reproductive patterns.