

Table S1: Model Selection Output – Whole animal analysis

	<i>df</i>	<i>AICc</i>	<i>delta</i>
Evaporative water loss			
<i>Above the Inflection Point</i>			
Sex + T _a	5	263.8	0
Sex + T _a + T _a ×Sex	6	266.2	2.32
T _a	4	266.3	2.48
M _b + Sex+ T _a	6	273.5	9.71
M _b + T _a	5	274.2	10.35
<i>Below the Inflection Point</i>			
T _a	5	43.7	0
<i>null</i>	3	47.2	3.47
Sex + T _a	5	48.7	5.03
Sex	4	51.7	7.96
Sex + T _a + Sex×T _a	6	56.3	12.61
Resting metabolic rate			
<i>Above the Inflection Point</i>			
Sex + T _a	5	77.2	0
T _a	4	78.5	1.26
Sex + T _a + Sex×T _a	6	85.0	7.82
M _b + Sex	5	85.6	8.38
M _b + Sex + T _a	6	88.0	10.79
<i>Below the Inflection Point</i>			
<i>null</i>	3	-5.3	0
Sex	4	-3.8	1.46
M _b	4	3.4	8.61
T _a	4	4.1	9.37
Sex + T _a	5	5.7	10.96
Core Body Temperature			
<i>Above the Inflection Point</i>			
T _a	4	162.6	0
Sex + T _a	5	167.0	4.44
M _b + T _a	5	174.7	12.14
Sex +T _a	6	174.9	12.36
M _b + Sex + T _a	6	178.3	15.76

Below the Inflection point

T _a	4	162.6	0
Sex + T _a	5	167.0	4.44
M _b + Sex	5	174.7	12.14
M _b + Sex + T _a × Sex	6	174.9	12.36
M _b + Sex + T _a	6	178.3	15.76

EHL/MHP

Above the Inflection Point

T _a	4	-19.4	0
Sex + T _a	5	-14.2	5.21
M _b + T _a	5	-6.1	13.27
Sex + T _a + T _a × Sex	6	-4.7	14.70
M _b + Sex + T _a	6	-2.5	16.91

Below the Inflection Point

null	3	-87.3	0
Sex	4	-80.9	6.45
T _a	4	-75.1	12.28
M _b	4	-73.1	14.28
Sex + T _a	5	-68.5	18.88

EHL/MHP vs T_a-T_b

Above the Inflection Point

T _a -T _b	4	-5.6	0
Sex + T _a -T _b	5	-0.4	5.13
Sex + T _a -T _b + T _a -T _b × Sex	6	7.7	13.26
M _b + T _a -T _b	5	7.8	13.37
M _b + Sex + T _a -T _b	6	11.4	16.97

Below the Inflection Point

null	3	-82.7	0
Sex	4	-76.3	6.47
T _a -T _b	4	-71.1	11.62
M _b	4	-68.5	14.29
Sex + T _a -T _b	5	-64.5	18.22

Table S2: Model Selection Output – Mass-specific analysis

Mass-specific evaporative water loss			
	<i>df</i>	<i>AICc</i>	<i>delta</i>
<i>Above the Inflection Point</i>			
T _a	4	-696.3	0
Sex + T _a	5	-683.3	12.98
Sex+ T _a + Sex×T _a	6	-666.6	29.68
<i>Below the Inflection Point</i>			
T _a	4	-494.6	0
<i>null</i>	3	-491.6	3.01
Sex + T _a	5	-497.6	14.95
Mass-specific resting metabolic rate			
<i>Above the Inflection Point</i>			
T _a	4	-833.7	0
Sex + T _a	5	-819.2	14.43
Sex + T _a + Sex×T _a	6	-801.7	32.00
<i>Below the Inflection Point</i>			
<i>null</i>	3	-390.9	0
Sex	4	-375.1	15.88
T _a	4	-373.7	17.30

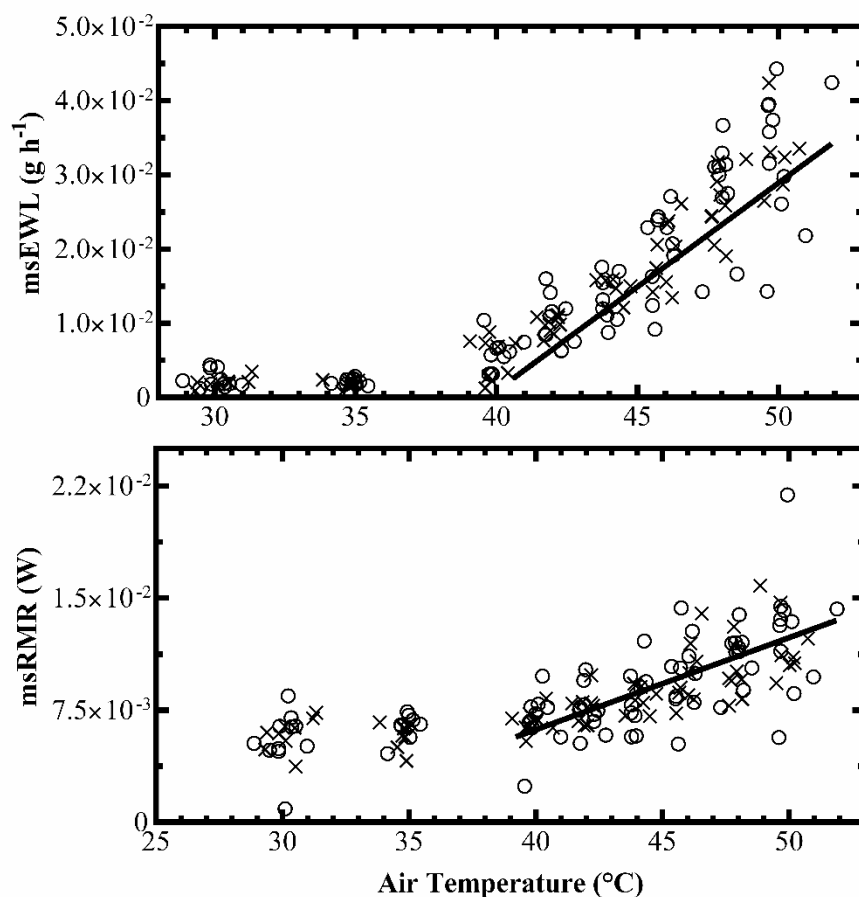


Figure S1. Mass-specific evaporative water loss (msEWL) and mass-specific resting metabolic rate in relation to air temperature (T_a) in southern yellow-billed hornbills (*Tockus leucomelas*). Data were obtained from 10 females (circles) and nine males (crosses) using open flow-through respirometry. Lines indicate linear mixed-effects regression models fitted for $\text{msEWL} = 0.0028T_a - 0.11$ ($T_a > 40.5^\circ\text{C}$) and $\text{msRMR} = 0.000061T_a - 0.18$ ($T_a > 39.2^\circ\text{C}$). Inflections in msEWL occurred at $T_a = 40.5^\circ\text{C}$ for males and $T_a = 39.7^\circ\text{C}$ females, above which EWL increased with increasing T_a . Below the inflection point, sex did not emerge as a significant predictor (see also Table S2). msEWL increased significantly with T_a ($t = 5.71$, $P < 0.001$; Fig. S2). At T_a values above the inflection, msEWL increased linearly and significantly with T_a ($t = 20.45$, $P < 0.001$; Fig. S2), and sex did not emerge as a significant predictor (Table S2). Inflections in msRMR occurred at $T_a = 39.2^\circ\text{C}$ for males and at $T_a = 38.8^\circ\text{C}$ for females. Within the thermoneutral zone the null model provided the best fit (Table S2). At T_a above inflection points, msRMR increased linearly and significantly with T_a ($t = 10.18$, $P < 0.001$; Fig. S2) and sex did not emerge as a significant predictor (Table S2).