

**Hypothesis test results for analysis of estrous cyclicity using the continuous-time Markov model**

Stage <sup>a</sup>	Comparison <sup>b</sup>	p-value <sup>c</sup>	Significance <sup>d</sup>	Stage Length Difference <sup>e</sup> (days)
Diestrus	Low-Control	0.840	None	0.8
Diestrus	Mid-Control	1.000	None	0.8
Diestrus	High-Control	0.446	None	1.9
Proestrus	Low-Control	1.000	None	0.0
Proestrus	Mid-Control	0.486	None	0.2
Proestrus	High-Control	1.000	None	-0.0
Estrus	Low-Control	0.123	None	-0.3
Estrus	Mid-Control	0.974	None	-0.1
Estrus	High-Control	1.000	None	0.0

a: Insufficient data to evaluate metestrus stage.

b: Sample sizes for the Control, Low, Mid, and High dose groups respectively were  $n = 10, 10, 10, 10$ .  
Dose levels were 0, 1, 3, 6 mg/mL respectively.

c: The p-values shown were calculated using a permutation null hypothesis testing method and have been adjusted for multiple comparisons using a Hommel correction within each stage.

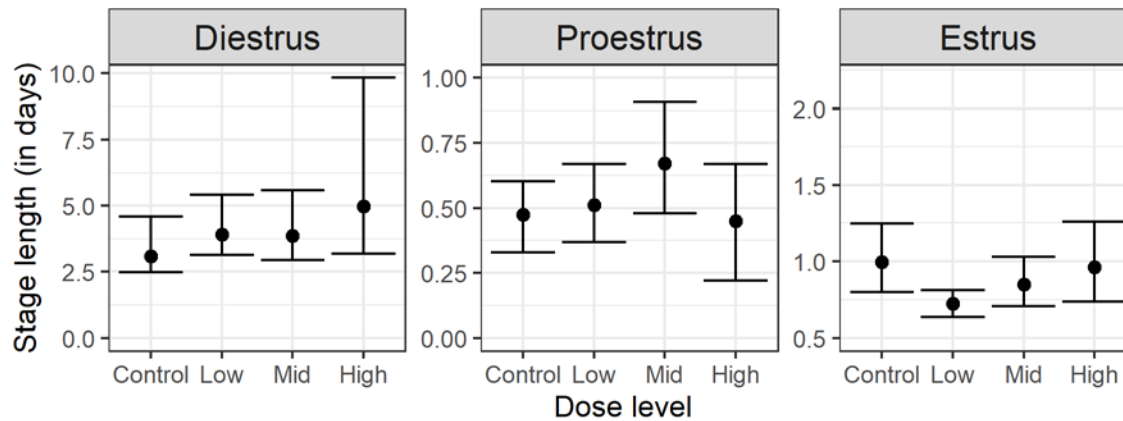
d: Significance is based on the adjusted p-value with a significance level of  $\alpha = 0.05$ .

e: A positive number indicates the estimated stage length in the treated group is longer than in the control group.

**Markov model estimates of stage length and 95% confidence intervals**

	Control (0 mg/mL)		Low dose (1 mg/mL)		Mid dose (3 mg/mL)		High dose (6 mg/mL)	
	Stage Length (days)	95% CI	Stage Length (days)	95% CI	Stage Length (days)	95% CI	Stage Length (days)	95% CI
Diestrus	3.1	(2.5, 4.6)	3.9	(3.1, 5.4)	3.9	(2.9, 5.6)	5.0	(3.2, 9.8)
Proestrus	0.5	(0.3, 0.6)	0.5	(0.4, 0.7)	0.7	(0.5, 0.9)	0.4	(0.2, 0.7)
Estrus	1.0	(0.8, 1.2)	0.7	(0.6, 0.8)	0.8	(0.7, 1.0)	1.0	(0.7, 1.3)
Metestrus <sup>a</sup>	0.1	--	0.1	--	0.1	--	0.1	--

a: Due to a very low number of observations of metestrus, stage lengths were estimated using a profile likelihood approach. As a result, confidence intervals are not available for the metestrus stage length estimate.



Estimates of stage length shown as dots, with bars indicating 95% confidence intervals. Estimates for lengths of metestrus are not shown here due to very low numbers of observations of this stage.