

Erratum

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Sex ratio determination and worker reproduction
in the slave-making ant *Harpagoxenus sublaevis*A.F.G. Bourke¹, T.M. van der Have², and N.R. Franks¹¹ School of Biological Sciences, University of Bath, Claverton Down, Bath, Avon BA2 7AY, Great Britain² Department of Population Biology and Evolution, University of Utrecht, Padualaan 8, 3584 CH Utrecht, The Netherlands

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Printing errors were unfortunately introduced in Table 3 that were not in the page proofs; these errors were in *line 6 from the bottom* and resulted in the numbers in that line being printed in the wrong columns. Table 3 is reprinted below in the correct form.

Table 3. Genotype distributions of ants in 49 *H. sublaevis* colonies. + = 1985 colony subjected to electrophoresis in year of collection (used for investigating male parentage); * = genotype of colony queen, in the 10 colonies where known (in these colonies genotype numbers do not include the colony queen). Collecting area: The woods were divided into 6 adjoining collecting areas. These were arbitrary except that areas 1, 2 and 4 lay on the left of the road through the woods (see Methods), whereas areas 3A, 3B and 5 lay on its right. This road constituted the only obvious potential gene flow barrier (to queens, not males) in an otherwise uniform open woodland habitat. Area 6, where colony S 85 1 was collected, lay slightly away from the main areas. Colonies designated 1/2 came from either areas 1 or 2

Host class	Colony no	Me						Mdh-2				Area		
		Females				Males		Females		Males				
		96/96	96/100	100/100	100/104	96	100	104	97/100	100/100	97		100	
HS+	S83	22		8	8				9	2			1	
LA		54	6	7						6			1	
n=33	S84	38	*	10						10*			1/2	
		73		60					3	57			2	
		95			13					13			1/2	
		99			9				4	5			1/2	
		114		5	7					12			1/2	
	S85	1	6	5			2			11		2		6
		4		6	9					15				3B
		9+		44	*		10			44*		10		3A
		10			7					7				3B
		15			9		7		7	2	4	3		3B
		17+		6	6		1			12		3		3B
		18			21		8		9	12	5	3		3B
		22	10	9			10	7		19		17		3A
		23+		19*	20		10	3		18*	21	4	9	3A
		26	1	6						7				3A
		27+		27*	20		3	3		23*	11	4	2	3A
		37			8						8			2
		40			19						19			2
		42		6							6			2
48			10						10			4		
70+	46	37*			4	3			83*		7	4		
71		11						4	7			4		

Table 3 (continued)

Host class	Colony no	Me						Mdh-2				Area	
		Females				Males		Females		Males			
		96/96	96/100	100/100	100/104	96	100	104	97/100	100/100	97		100
		74		9					9			4	
		75+		26*			36		26*		36	5	
		83		7					7			5	
		84		17					17			5	
		96+		1*			48		72*		48	5	
		97	6	7					13			5	
		101+		4			7	16	4		23	5	
		106+			10			20	10		20	1	
	S86	19		25					25			5	
HS+	S85	24		8					3	5	2	2	3A
LA+		54		14	15		6	4		29		10	4
LM		60			11	15		1	6			7	4
<i>n=7</i>		62			10					10			4
		68			25			9		25		9	4
		77+		44*	32		15	4		76*		19	5
		107+			14*			3		-		-	1
HS+	S84	25		5	8					13			1/2
LM	S85	12	7	10			1			17		1	3B
		13			13			1		13		1	3B
<i>n=9</i>		16			24					24			3B
		38		3	2					5			2
		53			19			2		19		2	4
		61		8						8			4
	S86	5		20						20			5
		17		3*	5					8*			5