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BLOOD PARASITES OF BIRDS FROM CHIRIQUI AND PANAMA PROVINCES IN THE REPUBLIC OF PANAMA

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Abstract: Blood smears were examined from 3,715 birds from four areas in the Republic of Panama. Hematozoa were present in at least 142 (50%) of the 281 species examined. An overall prevalence of 18% of the individual birds was noted: Haemoproteus 9%, Plasmodium 5%, Leucocytozoon <1%, Trypanosoma 2%, Atoxoplasma/Lankesterella <1% and microfilariae 3%. Prevalence in each of the four study areas varied from 13% to 28%. Distribution is analyzed by orders, families and individual species of Plasmodium are tabulated. New host records from 170 species of birds are recorded.

INTRODUCTION

A previous report (Galindo and Sousa, 1966) listed the findings of a survey of blood parasites of birds from a single region of Panama. Huff and Wetmore (1967) have added several additional records. Previous records of Hematozoa from birds of the neotropical region were summarized by White, et al. (1978) including the available data from the Republic of Panama. These authors reported on the distribution of blood parasites in the Neotropics on the basis of the avian-host families and the region in which they occur. The present paper summarizes a study of blood parasites of birds from four different areas of the Pacific slope of the Isthmus of Panama.

MATERIALS AND METHODS

Description of the study areas: Specimens for this study were procured from four areas. Two collecting sites, Cuesta de Piedras and La Fortuna, are located in the highlands of Chiriqui Province in western Panama and the other two areas, Juan Mina and Bayano, near sea level in Panama Province just east of the Panama Canal (Fig. 1). The

two locations in western Panama are enriched with species predominantly representative of the Central American avifauna. These locations provided a sample of the Panamanian avifauna similar to that found in the Atlantic regions of western Panama. The collecting sites sampled in eastern Panama Province are more influenced by the Colombian (South America) avifauna.

La Fortuna (Area A) is located in the highlands of Chiriqui Province, some 25 km north of Gualaca. The area has been selected as the site of a new hydroelectric project where the Chiriqui River dam will be constructed. The collection area was located approximately 1,000 m above sea level. It is considered a rainy Premontane Life Zone with an annual precipitation of about 4,000 mm. In this area, a total of 209 bird species have been noted (Adames, 1977) with strong representation of species known from the western Atlantic slope and the Costa Rican avifauna.

Cuesta de Piedra (Area B) is a collecting site located on a small farm approximately 1,300 m in elevation just below the town of El Volcan, on the slopes of the Chiriqui Volcano. The area

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PANAMA A TLANTIC OCEAN Area B PACIFIC OCEAN

FIGURE 1. Map of the Republic of Panama indicating the general location of collecting sites for birds in La Fortuna, Chiriqui Province (Area A), Cuesta de Piedra, Chiriqui Province (Area B), Juan Mina, Panama Province (Area C) and Bayano, Panama Province (Area D).

used to be, some 25 years ago, under a cover of cloud forest, but it is now completely cleared and utilized mostly as grazing pastures with some cultivation of citrus, plantain, sugar cane and vegetables.

Juan Mina (Area C) has been a mosquito collecting field station for entomologists of the Gorgas Memorial Laboratory for many years. It is located in an old citrus plantation on the south bank of the middle Chagres river, at the old boundary of Panama and the Canal Zone. The plantation has been overgrown for many years by second growth, and some parts have been converted into grazing pastures. Nearby are primary forests of the transitional type approaching the Tropical Moist Forest association.

Bayano (Area D) is located about 70 km east of Panama City and the Panama Canal. It is now an island remaining in the Bayano River Hydroelectric dam reservoir after the recent flooding of the basin. The area corresponds to tropical forest transitional to tropical dry forest. The average rainfall is 2,000 mm corresponding with the Pacific regime of precipitation pattern of the Republic of Panama, with a prolonged wet season from April to November.

Methods. The blood smears which constitute the basis of this report were obtained during a period of eight years (1969 through 1976), as a byproduct of an extensive survey for arthropod-borne viruses in Panama. The procedures followed were described previously (Galindo and Sousa, 1966). Indentification of avian hosts was made by Pedro Galindo and Eustorgio Mendez from the Gorgas Memorial Laboratory and questionable identities were confirmed from study skins submitted to the late Alexander Wetmore of the Smithsonian Institution in Washington, D.C. and Eugene Eisenmann of the American Museum of Natural History. Specimens collected in La Fortuna area were identified by Robert S. Ridgely of Yale University and R. Hinds from the Gorgas Memorial Laboratory. Species names were used according to Ridgely (1976).

The blood films were fixed with methyl alcohol in the field and treated with Giemsa stain by conventional methods in the laboratory. Each smear was examined under an oil immersion objective (100×) and 6× ocular for at least 10 minutes before discarding as "no parasite found". Most positive smears were examined for longer periods in search of mixed infections or of additional stages which might clarify diagnosis. Representative blood films have been deposited in the International Reference Center for Avian Haematozoa at Memorial University, St. Johns, Newfoundland (Accession Nos. 62219-62238).

RESULTS AND DISCUSSION

Geographical analysis:

Single smears were obtained from a total of 3,715 individuals, representing 281 species, 189 genera, 36 families, and 15 orders. They included 366 birds, representing 80 species, 68 genera, 23

families, and 9 orders from Area A: 221 birds, 49 species, 39 genera, 19 families and 7 orders from Area B; 891 birds, 101 species, 80 genera, 27 families and 11 orders from Area C; and, 2,237 birds, 194 species, 146 genera, 31 families and 11 orders from Area D. Tabulations of findings from each area are presented in Table 1. A total of 661 birds were found infected with at least one type of Hematozoa from all areas of collection, for an overall infection rate of 18%. Birds collected in Juan Mina (Area C) near the Panama Canal showed the highest rate of infection (28%). The lowest rate of infection (13%) was observed in the avifauna examined from Bayano (Area D) in eastern Panama.

Microfilariae were more frequently detected in La Fortuna (Area A) where a 4% prevalence appeared significantly higher than the 2-3% prevalence observed in the other areas. Leucocytozoon was not found frequently in the blood of resident birds. Only 20 individuals, <1% of the total sample, were found infected with this parasite. Infections with Leucocytozoon were diagnosed mostly (19 of 20 cases) in migrant bird species

TABLE 1. Avian Hematozoa from four areas of the Republic of Panama.

	Collection	Total No.		und itive		% Inf	ecte	d w	ith ^a	l
Area	localities	birds examined	No.	%	P	Н	L	Т	Mf	Other
A	La Fortuna, Chiriqui Province	366	68	18.6	3.6	10.9	0.3	0.8	4.4	0.0
В	Cuesta de Piedra, Chiriqui Province	221	57	25.8	4.5	19.5	1.3	1.3	2.3	0.9
С	Juan Mina, Panama Province	891	250	28.0	13.3	12.9	0.0	1.0	2.2	0.6
D	Bayano, Panama Province	2,237	286	12.8	2.5	6.2	0.7	2.6	2.9	0.0
All areas		3,715	661	17.8	5.3	9.0	0.5	2.0	2.9	0.2

^aP = Plasmodium; H = Haemoproteus; L = Leucocytozoon; T = Trypanosoma; Mf = Microfilariae; other: Atoxoplasma, Lankesterella, etc.

which may have acquired the infections elsewhere. Haemoproteus infections were most frequently detected in birds examined from Cuesta de Piedras (Area B). Hemoproteid parasites were demonstrated in 20% of the birds examined from this area. The prevalence of Plasmodium was 13% in birds from Area C, ranging from three to five times that of the other areas.

Distribution by Avian Host:

Table 2 shows the distribution of birds according to the avian orders and families included in the sample. The largest proportion of the birds were Passeriformes (3,155 individuals), but Apodiformes (280) and Columbiformes (155) were also well represented. Except for the Piciformes (60) all other orders were represented by smaller samples. Haemoproteus was a frequent parasite among Columbiformes, Apodiformes and Passeriformes. Plasmodium was seen almost exclusively in Passeriformes. Only two out of 197 birds found infected with this genus were from other avian orders (Caprimulgiformes and Columbiformes).

A total of 3,155 Passeriform birds (205 species, 17 families) were examined for blood parasites. Only two families, Vireolaniidae and Sylviidae, were represented by less than 30 individuals. The sample size varied from 30 to 597 birds examined in the other 15 families.

The prevalence of infection varied markedly among the families. The Icteridae (64%) and Vireolaniidae (100%) showed the highest prevalence of infection with Hematozoa. However, the latter family was represented by only two birds. Other families, such as Pipridae and Hirudinidae, showed the lowest rates of infection. Less than 1% of 320 birds of the family Pipridae and none of 48 individuals from the Hirudinidae were found infected. Within these two extremes, Cotingidae, Vireonidae, Turdidae and Thraupidae showed prevalences from 27 to 48%. The prevalence

among other passerine families included in this study varied from 10 to 25%. According to the parasites found: Plasmodium was identified in 35% of the positive birds. These malarial parasites were often seen in birds of the Formicariidae, Tyrannidae, Fringillidae, Thraupidae, Turdidae, Vireonidae and Icteridae. Plasmodium was infrequent in Cotingidae. Haemoproteid parasites were often observed in most passerine birds (44% of the positives), but were not found in Hirudinidae, Pipridae, and Troglodytidae. Leucocytozoon detected only among Turdidae (23%), Cotingidae (3%) and Parulidae (50%) usually in migratory species. Atoxoplasma/Lankesterella parasites were seen only in few instances among Thraupidae (2 of 597 birds), Icteridae (1/44) and Fringillidae (3/545) representing less than 1% of the total birds examined.

Haemoproteus columbae Kruse 1890, the principal blood parasite found among Columbiformes, was particularly prevalent in Columbina talpacoti (55%). Haemoproteus archilochus Coatney and West 1938 and H. trochili White, Bennett and Williams 1979 were the only hematozoan parasites seen in hummingbirds (Apodiformes: Trochilidae). Ten of 24 species of hummingbirds were found infected with these haemoproteids. Haemoproteus and Plasmodium were widely distributed among Passeriformes. Out of 24 Haemoproteus infections found among Cotingidae, 83% were recognized in the black and white becard (Pachyramphus albogriseus) and the white-winged becard (P. polychopterus). Plasmodium was often seen in the spotted antbird (Hylophylax naevioides), the social flycatcher (Myiozetetes similis), the gray-colored robin (Turdus grayi), the red-eyed vireo (Vireo olivaceus), the crimson-backed tanager (Ramphocelus dimidiatus), the blue-gray tanager (Thraupis episcopus), the yellowbilled cacique (Amblycercus holosericeus), the black-striped sparrow (Arremonops conirostris) and the vari-

Total Infected	Total	Infected		Tota	birds i	Total birds infected with:	with:		
Host Species	Birds	Birds	Ь	H	1	Ĺ.	Mf	Other	Area
TINAMIFORMES (Sub-Total)	-	С	0	С	С	c	С	0	
Tinamidae		С	0	0	С	0	C	С	
CICONIIFORMES	_	0	0	С	С	C	0	0	
Ardeidae	_	0	0	С	С	С	=	С	
FALCONIFORMES	1	æ	0	-	0	-	3 1	С	
Accipitridae		С	0	С	0	-	0	С	
Falconidaae	**	≈	0	_	С	0	3 1	c	
Herpetotheres cachinnans	-	_	0	0	=	=	*	С	<u>_</u>
Micrastur ruficollis	\$1	\$1	0	-	0	C	*	c	C.D
GALLIFORMES	31	_	0	_	0	0	C	С	
Cracidae	31	_	0	-	-	-	=	С	
Penelope purpurascens	_	_	0	_	c	0	C	c	_
Uninfected species	_	0	0	0	0	0	0	0	
GRUIFORMES	9	_	0	0	С	С	_	С	
Eurypygidae	_	_	С	0	=	_	-	0	
Eurypyga helias	-	-	0	0	0	*_	*_	0	<u> </u>
Rallidae	īc	0	0	0	-	С	0	С	
COLUMBIFORMES	155	09	1	57	0	-	-	0	
Columbidae	155	09	_	57	0	_	_	=	
Claravis pretiosa	98	<u>::</u>	0	13	0	0	0	0	Ω
Columbina talpacoti	\$ 2	44	0	43	С	С	_	С	B.C.D
Geotrygon chiriquensis	-	_	C	0	C	*_	=	С	<
Leptotila verreauxi	% 7	31	-	_	С	c	Э	0	B.C.D
Uninfected species	15	С	C	0	0	=	=	0	
PSITTACIFORMES	2	3 1	0	ઝ	=	-	_		
Psittacidae	[~	3 1	0	-	0	0	_	C	
Brotogeris jugularis	7	_	С	0	0	C	*-	c	<u> </u>
Pionus menstruus	-	-	0	-	С	C	C	C	<u>-</u>
Uninfected species	S)	0	С	C	0	С	0	0	
STRIGIFORMES	\$1	_	0	-	0	С	0	С	
Strigidae	21	_	С	-	С	0	0	c	
Otus choliba	_	_	С	-	0	0	0	c	ĸ
Uninfected species	_	C	0	c	-	0	=	=	

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IABLE 2. (continued)									
	Total	Infected		Tota	birds i	Total birds infected with:	with:		
Host Species	Birds	Birds	P	Н	Г	Т	Mf	Other	Area
CUCULIFORMES	7	С	0	0	=	С	С	0	
CAPRIMULGIFORMES	x	_	_	0	Э	0	С	_	
Caprimulgidae	x	-	-	С	=	0	С		
Nyctidromus albicollis	x	-	*_	0	0	С	C	_	B,C
APODIFORMES	780	50	0	.50	Э	0	0	С	
Trochilidae	280	70	0	50	0	=	0	c	
Amazilia amabilis	31	21	0	<u>.</u> 71	=	=	0	С	_
Amazilia edward	24	77*	0	4	0	=	0	С	B,D
Amazilia tzacatl	œ	7	0	*,	=	0	-	С	B,D
Anthracothorax									
nigricollis	31	-	0	*.	0	0	-	С	<u> </u>
Chalybura buffoni	10	4	С	4	0	0	-	0	<u> </u>
Dmophila julie	9		0	*-	0	0	С	0	_
Glaucis hirsuta	7.4	77	0	27	=	0	0	0	A,U
Phaeochroa cuvierii	**	7	0	*7	0	0	С	С	<u> </u>
Phaethornis anthopilus	23	_	0	*_	0	0	C	0	_
Phaethornis guy	25	_	0	-	0	0	-	С	<
Uninfected species	103	0	0	0	0	0	С	0	
TROGONIFORMES	4	31	0	0	С	0	31	0	
Trogonidae	4	2	0	0	С	0	7	0	
Trogon clathratus	_	-	0	0	0	0	-	0	V
Trogon massena	_	-	0	С	0	0	<u>*</u> _	0	_
CORACIIFORMES	17	23	0	27	0	0	0	С	
Alcedinidae	7	7	0	27	0	0	c	0	<u> </u>
Chloroceryle aenea	4	2	0	7*	0	0	0	С	ပ
Uninfected species	n	С	0	0	0	0	0	С	
Momotidae	10	С	0	0	С	0	C	0	
PICIFORMES	09	4	0	7	=	0	21	0	
Bucconidae	2.7	-	0	-	0	0	С	С	
Notharchus macrorhynchus	-	-	0	-	0	0	=	0	
Uninfected species	23	0	0	0	0	0	0	0	

ABLE 2. (continued)									
Picidae	17.7	_	С	_	С	=	=	0	
Melanerpes rubricapillus	5.	_	=	*_	=	=	c	0	.). Y
Uninfected species	×	С	c	С	=	C	=	0	
Ramphastidae	5.	21	С	0	0	=	21	0	
Aulacorhynchus prassinus	21	÷1	0	С	0	=	÷1	=	<
Uninfected species	-1	С	0	0	0	0	С	С	
ASSERIFORMES	3,155	564	961	250	50	7.1	86.	9	
Dendrocolaptidae			_	_	0	_	_	С	
Campylorhamphus pusillus	_	-	0	С	0	=	*-	0	<
Dendrocincla homochroa	21	÷1	*-	*-	С	=	c	0	=
Dendrocolaptes certhia	ıc	_	С	0	0	*_	0	0	Ϋ́
Uninfected species	96	0	С	С	С	0	=	0	
Furnariidae	35	31	С	-	0	_	0	С	
Automolus ochrolaemus	x	31	С	*_	С	*	0	0	<u></u>
Uninfected species	27	С	С	0	С	С	0	С	
Formicariidae	281	98:	16	9	0	r.	:: :	0	
Cercomacra tyrannina	12	21	С	, ,	С	С	С	С	ਹ ਹ
Cymbilaimus lineatus	_	-	С	С	0	=	*-	0	<u>-</u>
Formicarius analis	=	_	С	*_	C	=	С	0	<u> </u>
Gymnocichla nudiceps	3]	51	*-	c	0	<u>*</u>	С	С	=
Gymnopithys leucaspis	3.1		*∻	*_	C	=	С	c	G S
Hylophylax naevioides	£;	9	* 9	С	0	C	С	C	ਹ ਹ
Myrmeciza immaculata		-	С	c	0	*_	С	-	<u>-</u>
Myrmeeiza longipes	Ξ	~	0	*_	С	C	<u>\$</u> 1	0	<u> </u>
Myrmotherula fulviventris	2	_	*_	c	C	-	0	0	=
Phaenostictus meleannani	61	_	С	c	-	*_	0	c	_
Taraba major	13	-	*_	c	С	C	С	c	ਹ ਹ
Thamnophilus doliatus	£	⋾	-	*_	-	3 1	0	0	<u> </u>
Thamnophilus punctatus	<u>.c.</u>	~ :	*∴	c	0	=	С	-	<u> </u>
Uninfected species	5	С	С	=	=	C	-	=	
Pipridae	350	**	n	0	С	=	0	С	
Manacus vitellinus	99	21	ઝા	Ξ	C	=	С	С	<u> </u>
Pipra erythrocephala	?] ?]	-	*-	=	=	0	С	С	<u> </u>
Uninfected species	2352	С	0	0	0	=	0	=	

ABLE 2. (continued)

TABLE 2. (continued)									
	Total	Infected		Total	birds in	Total birds infected with:	vith:		
Host Species	Birds	Birds	P	Н	Г	Т	Mf	Other	Area
Cotingidae	67	35	-	24	-	=	n	С	
Attila spadiceus	15	≈	0	*_	*_	*-	С	С	A.U
Cotinga nattererii	_	_	0	С	0	*_	С	0	<u>-</u>
Lipaugus unirufus	æ		0	*-	С	*_	С	c	<u>_</u>
Pachyramphus albogriseus	Ξ	31	0	С	0	<u>\$</u> 1	С	C	B,D
Pachyramphus cinnamomeus	19	Ξ	0	10*	0	rC	≈	0	<u> </u>
Pachyramphus polychopterus	14	Ξ	-	10	0	_	С	О	C.D
Platypsaris homochrous	?1	_	0	*_	С	0	C	С	2
Rhytipterna holerythra	_	_	0	*_	0	0	0	0	<u>_</u>
Uninfected species	-	С	0	0	0	0	0	0	
Tyrannidae	513	99	50	36	0	01	- -	0	
Capsiempsis flaveola	7	31	0	-	0	*_	Э	0	C:D
Contopus cinereus	æ	_	0	*_	0	0	0	0	ပ
Contopus sordidulus	9	_	0	*_	0	0	С	0	B,C,D
Contopus sp.	1	-	0	_	0	0	0	·O	ပ
Elaenia chiriquensis	01	23	*2	0	0	0	0	С	B,C
Elaenia flavogaster	42	7	0	*9	0	*_	0	0	B,C,D
Empidonax flaviventris	x	_	*-	0	0	_	0	0	C,D
Empidonax traillii	<u>8</u> 1	≈	0	က	0	0	0	0	C'D
Legatus leucophaius	3	-	0	*.	0	0	С	0	B,C
Mionectes olivaceus	ច	-	0	0	0	*_	0	0	A,C
Myiarchus tuberculifer	6	က	က	0	0	*_	0	0	A,B,C
Myiobius atricaudus	x	1	0	0	0	*_	0	0	A,D
Myiodynastes maculatus	10	4	0	*n	0	0	<u>*</u> _	0	A,C,D
Myiopagis gaimardii	က	-	0	0	С	*_	*	0	<u> </u>
Myiophobus fasciatus	73	-	0	*_	0	0	0	0	ပ
Myiozetetes similis	+	7	*9	-	0	0	С	0	B,C,D
Ornithion brunneicapillum	1	-	0	0	0	*_	С	0	<u> </u>
Phylloscartes flavovirens	ત્ર	_	0	*_	0	0	С	0	ပ
Pipromorpha oleaginea	168	7	*-	*,	С	0	-	С	A,B,C
Platyrinchus coronatus	15	-	0	0	0	0	*_	С	A.C.D
Sublegatus arenarum	ıc	3 1	0	<u>*</u> 71	0	0	0	О	ပ

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Indian 2. (Continued)									
Toronotricens orythrurus	Ξ	-	C	C	0	*	C	0	CD
Tyranniscus villissimus	; c:	-	. —	0	· C	· C	· C	· C	
Tyrannus melancholicus	98	17	9	11	0	0	0	0	A.B.C
Flycatcher (unidentified)		-	0	_	0	-	0	0	D
Uninfected species	94	0	0	0	0	0	0	0	
Hirundinidae	48	0	0	0	0	0	0	0	
Troglodytidae	140	5	2	Ö	0	0	ဢ	0	
Henicorhina leucophrys	27	က	*_	0	0	0	*	0	V
Thryothorus modestus	12	-	0	0	0	0	*_	0	A,C
Troglodytes musculus	45	1	1*	0	0	0	0	0	A,B,D
Uninfected species	26	0	0	0	0	0	0	0	
Turdidade	174	47	24	6	Ξ	0	7	0	
Catharus fuscater	11	7	* 9	0	0	0	* 2	0	V
Catharus minimus	16	2	0	0	2	0	0	0	Ω
Catharus ustulatus	69	13	4	0	6	0	1	0	A,B,C
Myadestes melanops	12	87	5*	0	0	0	0	0	V
Turdus albicollis	1	-	0	-	0	0	0	0	В
Turdus grayi	55	19	12	9	0	0	က	0	A,B,C,I
Turdus obsoletus	က	23	0	*_	0	0	*	0	V
Turdus plebejus	-	1	0	*-	0	0	0	0	V
Uninfected species	9	0	0	0	0	0	0	0	
Sylviidae	11	-	0	-	0	0	0	0	
Rhamphocaenus rufiventris	10	1	0	*.	0	0	0	0	A,D
Uninfected species	1	0	0	0	0	0	0	0	
Vireolaniidae	2	8	0	-	0	0	1	0	
Smaragdolanius pulchellus	2	8	0	* _	0	0	*.	0	C,D
Vireonidae	35	36	12	21	0	9	4	0	
Hylophilus flavipes	22	7	*	*:	0	* 2	*_	0	B,C
Vireo flavoviridis	14	10	5 *	*9	0	* 3	* ი	0	B,C,D
Vireo olivaceus	48	19	6	15	0	7	0	0	B,C,D
Uninfected species	œ	0	0	0	0	0	0	0	
Parulidae	142	16	7	9	œ	က		0	
Dendroica castanea	34	_ග	*5	2*	*	* ი	*_	0	A,D
Dendroica petechia	27 (0	→ (0	0	0	0	Ω
Protonotaria citrea	7	П.	*	0	0	0	0	0	

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TABLE 2. (continued)									
	Total	Infected		Tota	birds i	Total birds infected with:	with:		
Host Species	Birds	Birds	Ь	Н	r	F	Mf	Other	Area
Seiurus motacilla	_	1	*-	0	0	0	0	0	D
Seiurus noveboracensis	18	2	2	0	2	0	0	0	C,D
Vermivora peregrina	43	2	*.	0	*3	0	0	0	A,D
Uninfected species	42	0	0	0	0	0	0	0	
Thraupidae	597	160	48	83	0	50	56	2	
Chlorospingus ophthalmicus	5	က	5*	*_	0	0	0	0	A
Chlorospingus pileatus	-	1	0	*_	0	0	0	0	A
Eucometis penicillata	29	9	0	*	0	* 2	0	0	C,D
Euphonia anneae	10	_	0	0	0	0	*	0	A
Euphonia laniirostris	47	က	0	2	0	*	*	0	A,C,D
Habia fuscicauda	5	2	1*	*_	0	0	0	0	C,D
Piranga olivacea	3	2	0	1	0	*	0	0	D
Piranga rubra	12	6	1	œ	0	-	0	0	\mathbf{A},\mathbf{D}
Ramphocelus dimidiatus	20	24	14*	**	0	* 2	2*	0	C,D
Ramphocelus passerinii	14	12	-	12	0	0	0	*-	В
Ramphocelus icteronotus	47	18	*_	*	0	* 2	*9	0	C,D
Rhodinocichla rosea	4	2	5 *	0	0	0	0	0	ပ
Tachyphonus rufus	12	4	_	*3	0	0	-	0	C,D
Tachyphonus luctuosus	35	20	*	18*	0	*	0	0	C,D
Tangara icterocephala	17	13	0	13*	0	0	* _	0	V
Tangara inornata	17	9	*-	*	0	*	4	0	A,C,D
Tangara larvata	20	5	0	5 *	0	0	*	0	B,C,D
Thraupis episcopus	202	5 6	13*	4	0	9	4	*_	A,B,C
Thraupis palmarum	22	က	5 *	-	0	0	0	0	C,D
Uninfected species	20	0	0	0	0	0	0	0	
Coerebidae	30	4	-	-	0	_	7	0	
Coereba flaveola	24	2	1*	0	0	*	*_	0	B,D
Dacnis cayana	5	2	0	*_	0	0	*_	0	C,D
Uninfected species	1	0	0	0	0	0	0	0	
Icteridae	44	88	10	11	0	4	6	-	
Amblycercus holosericeus	11	10	10*	0	0	0	0	*_	C,D
Cacicus cela	11	4	0	1	0	0	* с	0	Ω

(continued on page 216)

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ADLE 2. (continued)									
Icterus aurocapillus	ıc	1	0	*-	0	*1	0	0	D
Icterus chrysater	10	10	0	*x	0	* ი	**	0	Ω
Icterus mesomelas	4	3	0	0	0	0	*2	0	C_D
Zarhynchus wagleri	2	-	0	*_	0	0	0	0	Ð
Uninfected species	_	0	0	0	0	0	0	0	
Fringillidae	545	128	20	49	0	6	34	က	
Arremon aurantiirostris	<u>&</u> 7	9	9	0	0	0	0	0	C_D
Arremonops conirostris	49	20	14	က	0	*-	9	*2	A,B,C,D
Atlapetes brunneinucha	24	16	0	15	0	0	*	0	V
Cyanocompsa cyanoides	29	18	œ	5*	0	*9	4	0	C,D
Oryzoborus funereus	33	9	4	-	0	0	*.	0	C_D
Pitylus grossus	77	1	0	0	0	0	1	0	Ď
Saltator albicollis	75	31	1	*.	0	0	*_	0	A,B,C,D
Saltator atriceps	9	-	0	*-	0	0	0	0	O
Saltator maximus	æ	10	0	0	0	0	10*	0	A,B,C
Sporophila aurita	164	38 38	12	24	0	*2	5	*_	A,B,C,D
Sporophila schistacea	4	_	0	*	0	0	0	0	Ω
Tiaris olivacea	98	9	*n	0	0	0	**	0	A.B.C.D
Volatinia jacarina	25	က	2	1	0	0	0	0	BD
Uninfected species	7	0	0	0	0	0	0	0	
Total	3,715	661	197	335	20	73	107	2	
Percent		17.8	5.3	9.0	0.5	5.0	5.9	0.2	
1 1 1 11 11 11 11 11 11 11 11 11 11 11	(OEO: 1 '								

*New host records (based on White, et al., 1979). P = Plasmodium: P = Plasmodium

terella types.

NOTE: No parasites were found in the following birds. The number after the species indicates number of birds examined; the

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TINAMIFORMES: Tinamidae, Crypturellus soui 4C. CI. CONIIFORMES: Ardeidae, Florida caerulea 1C. FALCONIFORMES: Accipitridae, Accipiter bicolor 1D, Buteo platypterus 2AD, Leucopternis albicollis 1D. GALLIFORMES: Cracidae, Ortalis cinereiceps 1D. GRUIFORMES: Rallidae, Gallinula chloropus 2C, Laterallus albigularis 1C, Porphyrula letter indicates area where collected.

martinica 2C. COLUMBIFORMES: Columbidae, Columba nigrirostris 2D, Columbina minuta 1D, Leptotila cassinii 9CD. PSITTACIFORMES: Psittacidae, Forpus conspicillatus 1D, Pyrrhura hoffmanni 1D. STRIGIFORMES. Strigidae, Otus guatemalae 1A. CUCULIFORMES: Cuculidae, Crotophaga ani 2CD, Coccysus minor 1D, Piaya cayana 2D, Tapera naevia 1D. APODIFORMES: Trochilidae, Campylopterus hemileucurus 6A, Colibri thalassinus 1A, Chalybura urochrysia 2D, Chlorostilbon

TABLE 2. (continued)

Heliotryx barroti 1D, Lampornis castaneoventris 2A, Phaethornis longuemareus 2D, Phaethornis superciliosus 46D, Threnetes ruckeri 7D, TROGONIFORMES: Trogonidae, Trogon violaceus 1D, CORACIIFORMES: Chloroceryle americana 1A; Momotidae, Momotus momota 10BCD. PICIFORMES: Bucconidae, Malacoptila panamensis 13D, Monasa morphoeus 2D, Nonnula frontalis 8D; Picidae, pylorhamphus trochilirostris 5D, Deconychura longicauda 17AD, Dendrocincla fuliginosa 23CD, Dendrocolaptes picumnus 1D, Glyphorynchus spirurus 14AD, Lepidocolaptes souleyetii griseicapillus 3A, Xiphorhynchus erythropygius 3A, Xiphorhynchus guttatus 17CD, Xiphorhynchus lachrymosus 9D, Furnariidae, Anabacerthia Cranioleuca erythrops 1A, Margarornis assimilis 1D, Elvira chionura 1A, Eutoxeres aquila 6A, Florisuga Alcedinidae, Ceryle torquata 1B, Chloroceryle amazona 1B, Campephilus melanoleucos 8D, Melanerpes formicivorus 1A, Melanerpes pucherani 8D, Picumnus olivaceus 1D; Ramphastidae, Pteroglossus torquatus 6D, Ramphastos sulturatus 1D. PASSERIFORMES: Dendrocolaptidae, Cam rubiginosus 1D, Sclerurus albigularis 1A, Sclerurus guatemalen 6AD, Synallaxis brachyura 1B, Syndactyla subalaris 3A, Xenops minutus 12CD; Formicariidae, Dysithamnus mentalis Chloropipo holochlora 20D, Pipra coronata 2C, Pipra mentalis mellivora 1B, Heliodoxa jacula 26A, Heliomaster longirostris 11) 3A, Dysithamnus puncticeps 1C, Formicarius rufipectus 1C, Myrmotherula surinamensis 11D, Thamnistes anabatinus 1D, Thamnophilus nigriceps 6D; Pipridae, Corapipo altera 3A, .94D. Schiffornis turdinus 13D; Cotingidae. Pachyramphus Myrmotherula axillaris 13D, Sittasomus exsulvariegaticeps 2A, Myrmeciza 848

superciliaris 1A, Lophotriccus pileatus 2A, Mitrephanes phaeocercus 2A, Myiarchus panamensis 2C, Myiobius sulpureipygius 5D, Myiodynastes luteiventris 1B, Myiodynastes hemichrysus 1D, Myiozetetes cayanensis 6CD, Myiozetetes granadensis 10B, Oncostoma olivaceum 1D, Onychorthynchus mexicanus 17CD, Pitangus sulphuratus 1C, Platyrinchus mystaceus 3A, Rhynchocyclus olivaceus 11CD, Sayornis nigricans 6A, Serpophaga cinerea 3A, Todirostrum cinereum 3CD, Todirostrum sylvia 1C, Tolmomyias sulphurescens 2D, obsoletum 3CD, Cnipodectes subbrunneus 2D, Colonia colonus 3D, Contopus virens 1B, Empidonax virescens 3CD, Leptopogon Tyrannulus elatus 3CD; Hirundinidae, Hirundo rustica 3D, Pygochelidon cyanoleuca 12A, Stelgidopteryx ruficollis 33 ABCD; Troglodytidae, Campylorhynchus albobrunneus 1D, Cyphorhinus phaeocephalus 11CD, Henicorhina leucosticta 24D, Microcerculus marginatus 4D, Thryothorus atrogularis 1C, Turdidae, Catharus fuscescens 6D; Sylviidae, Polioptila plumbea 1D; Vireonidae, Hylophilus aurantiifrons 1D, Wilsonia pusilla 1B; Thraupidae, Euphonia fulvicrissa 10D, Habia rubica 9C, Tangara gyrola 1B; Coerebidae, Cyanerpes cyaneus 1C, Icteridae, Icterus galbula 1D; Fringillidae, Atlapetes rufus 1D; Tyrannidae, Aphanothriccus audax 1D, Camptostoma Basileuterus fulvicauda 5AD, Dendroica pensylvanica 1D, Mniotilta varia 2A, Myioborus miniatus 12A, Oporornis for ludovicianus 2D, Sporophila Thryothorus fasciatoventris 11D, Thryothorus rufalbus 4A Hylophilus decurtatus 2BD, Hylophilus ochraceiceps 5D Parulidae, Basileuterus culicivorus 1A, Basileuterus delattrei 1B mosus 6AD, Parula pitiayumi 1B, Wilsonia canadenis 12AD 3A, Pheucticus nigricollis 2B. able seedeater (Sporophila aurita). It is possible that some of the low level infections identified as *Haemoproteus* may actually represent *Plasmodium* cases with only gametocytes circulating in the blood of the naturally infected avian host at the time of sampling.

A total of 170 new host-parasite records are included in the findings of this survey (Table 2).

IDENTIFICATION: Most of the infections produced parasitemias of less than one parasite per 10,000 red blood cells. Specific identity in a single smear survey must be based exclusively on morphological characteristics of a limited number of visible forms. Many overlapping characteristics in the currently accepted classification of these parasites make diagnosis precarious. For example, the finding of very few elongate gametocytes and a single schizont containing four merozoites could be interpreted as an early stage of several species of Plasmodium. All the infections with Leucocytozoon contained gametocytes and were often too few in numbers to justify specific assignment. Only, L. dubreuili of the Turdidae could be recognized. With the exception of Haemoproteus columbae, H. trochili and H. archilochus cited above, which were sufficiently numerous and characteristic to leave little doubt as to their identity, all other haemoproteid infections had such a low parasitemia that species identification was not plausible. Types 2 and 4 of Greiner et al. (1977) were the gametocyte forms usually observed. The trypanosomes, which are characteristically pleomorphic, were scarce when found in the individual blood preparations and no attempt was made to classify them to species.

The identity of the *Plasmodium* species are tabulated in Table 3. The authors are well aware of the fact that with further material and/or isolation and study in experimental hosts these designations might prove in error and that even some

of the infections herein reported as *Haemoproteus* might prove to be *Plasmodium* instead. We have attempted to designate species on the basis of closest attainments (Greiner et al., 1975a) to major characteristics of the species assigned and hope this approach will aid future investigators in the clarification of this problem and toward a more exact interpretation of the distribution of species of *Plasmodium* prevalent in avian hosts.

CONCLUSIONS

The prevalence of Hematozoa in the neotropical avifauna is substantially lower than that of the Nearctic (White et al., 1978). Birds in the nearctic region of the Western Hemisphere have been found to harbor at least one type of hematozoan parasite in 75% of the species examined (Greiner et al., 1975b) in contrast to neotropical birds which have been found infected with hematozoa in only 53% of the species examined (White et al., 1978). The present study of the avian Hematozoa in the birds of the Republic of Panama has shown at least 142 (50%) out of a total of 282 species harbored at least one hematozoan parasite. This is comparable to findings for the Neotropic, but much lower than indicated for the Nearctic. The prevalence of Hematozoa in 3,715 birds of the Republic of Panama examined during the present study was 18%, identical with the 18% rate of infection reported by White et al. (1978) for 20,880 birds previously examined for blood parasites in the neotropical regions. This is, however, much lower than the 27% prevalence recorded for Central American birds by the same authors. A comparison of our findings with previous data is summarized in Table 4.

Present findings on the hematozoan parasites of birds collected in four areas of the pacific slopes of the Republic of Panama have shown that the prevalence of infection varies from 13% in eastern

TABLE 3. Frequency of Plasmodium infections in birds from the Republic of Panama	ium infection	s in birds	from the	Republic	of Pan	ama.					
		ź	Ž					Plas	Plasmodium*		
Host	Area	Exam.	Inf.	Re	Po	ij	Fi	Na	He	Va	88
Leptotila verreauxi	၁	6	-		-						
Nyctidromus albicollis	၁	œ						1			
Dendrocincla homochroa	Ω	12	_							_	
Gymnocichla nudiceps		25	_							_	
Gymnopithys leucaspis	<u> </u>	30	က							≈	
Hylophylax naevioides	C,D	32	9	2						9	
Myrmotherula fulviventris	Ω	10	-							_	
Taraba major	၁	9	_							_	
Thamnophilus doliatus	ပ	15	1				_				
Thamnophilus punctatus	C,D	15	က	2						_	
Manacus vitellinus	ပ	19	2					2			
Pipra erythrocephala	ပ	18	_					1			
Pachyramphus polychopterus	ပ	7	_					_			
Empidonax flaviventris	ပ	1	-					-			
Elaenia chiriquensis	ပ	6	2					2			
Myiarchus tuberculifer	C,D	ī	က					23		-	
Myiozetetes similis	ပ	27	9					9			
Pipromorpha oleaginea	ပ	7	_					_			
Tyranniscus villissimus	D	က	_					_			
Tyrannus melancholicus	ပ	21	9					9			
Henicorhina leucophrys	ď	27	_							_	
Troglodytes musculus	Q	40	-	-							
Catharus fuscater	V	11	9	27						9	
Catharus ustulatus	B,D	65	4							4	
Myadestes melanops	V	12	7			_				-	
Turdus grayi	A,B,C	52	15	7						11	
Hylophilus flavipes	ပ	20	_					_			
Vireo flavoviridis	ပ	7	27					21			
Vireo olivaceus	B,C,D	48	6	-		21		≈		4	
Dendroica castanea	Ω	£	27							27	
Protonotaria citrea	<u> </u>	7	-							-	
	1	-	-		-						

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Vermivora peregrina Chlorospingus ophthalmicus Habia fuscicauda											
Chlorospingus ophthalmicus Habia fuscicauda	_	 	_		_						
Habia fuscicauda	<	ī.	Ç1	_						21	
D_{i} L_{i}	ن	-;	-					_			
riranga ruora	<u>-</u>	9	-							_	
Ramphocelus dimidiatus	C.D	02	Ξ	71				31		6	
Ramphocelus icteronotus	ပ	21	(~	_				≈	_	≈	
Ramphocelus passerinii	ĸ	! :	_	_							
Rodinocichla rosea	ပ	7	ા					?I			
Tachyphonus rufus	ပ	=	-	_							
Tachyphonus luctuosus	<u>_</u>	*	≈	≈							
Tangara mornata	ບ	1	1					-			
Thraupis episcopus	၁	181	<u>::</u>					21		_	
Thraupis palmarum	C'D	7.7	21	_				_			
Coereba flaveola	<u> </u>	55	1							_	
Amblycercus holosericeus	ပ	01	01		21					œ	
Arremon aurantiirostris	C,D	£7.	9	7						7	
Arremonops conirostris	C:D	41	14	**						21	
Cyanocompsa cyanoides	C,D	29	x	≈	_					÷	
Oryzoborus funereus	ပ		7					_		≈	
Saltator albicollis	၁	47	_								-
Sporophila aurita	A.B,C	9 8	21							77	
Tiaris olivacea	_	25	က	က							
Volatinia jacarina	B,D	25	21							_	_
Total birds infected											
with Plasmodium		1,392	197	35	9	÷	_	::: :::	-	106	31
Percent of total birds											
infected with Plasmodium			14.1	2.5	0.4	0.5	0.07	æ. æ.	0.07	9.7	0.14
Percent of Plasmodium											
infections				17.8	3.0	1.5	0.5	56.9	0.5	53.8	0.1

Table 4. Comparison of blood parasites of birds in Panama with findings in Central America.

	Panama avifauna	Central America
	(Present work)	(White et al., 1978)
Total no. birds examined	3,715	5,335
Total no. found infected	661	1,446
Percent infected	17.8	27.1
Prevalence of infection		
(percent) with		
Haemoproteus	9.0	19.9
Plasmodium	5.3	7.5
Leucocytozoon	0.5	0.8
Microfilariae	2.9	5.4
Other	0.2	1.0

Panama (Bayanao, Area D) to 28% in central Panama (Juan Mina, Area C) close to the old Canal Zone. Such differences might be due to sample size, seasonal changes, the species of birds and ecological conditions of the collection site, more than the result of simple geographical separation. Distribution and density of vector species probably also play an important role. The

prevalence of Hematozoa among passerine birds of Almirante, Bocas del Toro Province (Gallindo and Sousa, 1966), La Fortuna and Cuesta de Piedras, Chiriqui Province, and Juan Mina, in Panama Province varied slightly (22 to 28%) in contrast to the much lower rate of infection (13%) among passerines of the Bayano area in eastern Panama Province.

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