

# Bird diversity in the production forest management unit in North Seram, Central Maluku Regency, Indonesia

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## ABSTRACT

Bird diversity hotspots are important locations for species conservation. Conservation actions at these hotspot locations will protect important forest areas outside the designated conservation areas. This study aimed to measure the importance of the location of bird diversity hotspots in the area of Production Forest Management Unit (PFMU) in the Wae Sapalewa area in North Seram, Central Maluku. The method used was a territorial search survey to determine hotspot locations. Point Count method was used to collect bird species information. The analysis was performed quantitatively and descriptively so that it is easy to understand. There were 7 locations of bird diversity hotspots with important species status. 121 species of birds were found of 9 endemic species of Seram Island (*Gymnophaps mada*, *Eos semilarvata*, *Lorius domicella*, *Cacatua moluccensis*, *Halcyon lazuli*, *Rhipidura dedemi*, *Basilornis corythaix*, *Philemon subcorniculatus*, *Myzomela blasii*), 4 endangered species (*Halcyon lazuli*, *Cacatua moluccensis*, *Lorius domicella*, *Eulipoa wallacei*), 6 unique species (*Casuarus casuarinus*, *Charmosyna placentis*, *Micropsitta bruijnii*, *Alisterus amboinensis*, *Ninox squamipila*, *Tanysiptera galatea*, *Rhyticeros plicatus*) and 21 limited distributed species. This type of bird is referred to as the "Indicator Bird Species". Production forest areas such as PFMU have potential bird diversity hotspots that can be applied in management blocks for the conservation and ecotourism purposes.

**Key words:** Bird, conservation, diversity hotspot, forest management unit.

## INTRODUCTION

The concentration location of various bird species that are typical with a high level of endemism in an area has been designated as an important location for conservation. This location is considered as a part of the concept of the biodiversity hotspot locations for a smaller scale, such as Endemic Bird Area or EBA (Vale *et al.*, 2018) and Key Biodiversity Area or KBA (Dudley *et al.*, 2014; IUCN, 2016; Tantipisanuh *et al.*, 2016; Donald *et al.*, 2019). With an interest value in the bird species, the concentration location of various bird species with a high level of endemism can be referred to as an important Bird Diversity Hotspot location. Endemic bird species have been proven to be used in many places as an effective indicator for assessing the overall diversity of communities in an area (Mekonen, 2017). Areas where endemic bird species are found also support a variety of other important animal species and various types of plants (Tanalgo *et al.*, 2015; Şekercioglu *et al.*, 2019).

At present, forestry development is directed at forest management in accordance with its main function and designation at the site level, and for this reason the government has issued a Forest Management Unit (FMU) development policy based on Minister of Forestry Decree No.230/Kpts-II/2003. One of them is PFMU Wae Sapalewa which was determined through Minister of Forestry Decree No.336/Menhut-II/2010 dated May 25, 2010, covering an area of 67,057 ha located in North Seram Subdistrict, Central Maluku Regency, Maluku Province.

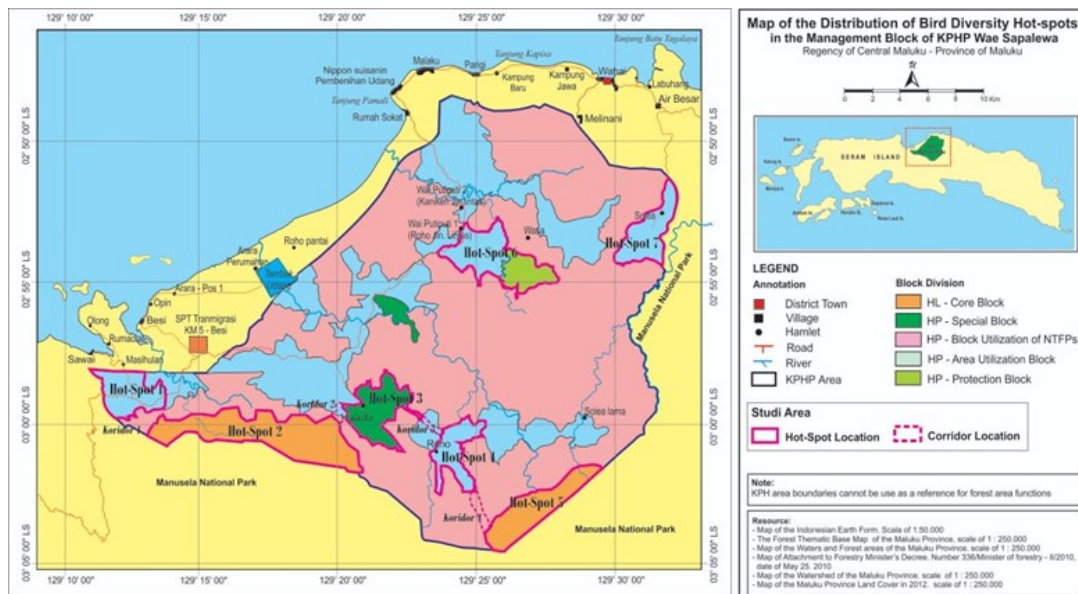
Production Forest Management Unit (PFMU) Wae Sapalewa is a Forest Management Unit intended for timber forest production businesses (Forest Area Consolidation Center Region IX Ambon, 2013). However, this Forest Management Unit (FMU) has a rich biodiversity that is interesting to study. Therefore, this FMU area is believed to be used to identify biodiversity hotspot locations on a small scale, which can be applied to management blocks according to the work plan of FMU, especially in management blocks intended for protection and other uses, so as to produce simple hotspot locations in the form of bird diversity hotspots. This will facilitate management by the FMU organization responsible for limited forest areas to achieve the goals of conservation and utilization of environmental services in the future. For this purpose, seven bird diversity hotspots in the PFMU Wae Sapalewa area have been identified which are believed to provide value and benefits for achieving sustainable forest management at the site level for the North Seram Island forest area (Persulesy *et al.*, 2019). The aims of this study were to examine the richness and abundance of bird species with an important status that exist at the seven bird diversity hotspots identified in the PFMU Wae Sapalewa area, and to study the conservation values of the seven hotspots.

## MATERIALS AND METHODS

### Study area

This study was conducted in the PFMU Model Wae Sapalewa area, North Seram Subdistrict, Central

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**Figure 1.** Distribution of hotspot locations in the Wae Sapalewa PFMU management block.

Maluku Regency, Seram Island, Maluku Province, Indonesia. Geographical location is at 129°10'15"- 129° 31'47" East Longitude and 2°47'39"- 3°4'38" South latitude (Forest Area Consolidation Center Region IX Ambon, 2013). The locations of the Biodiversity Hotspots were selected using groups of bird species as indicators, which were then used as criteria in assessing and determining the distribution of Biodiversity Hotspot locations. The map of the location distribution and forms of the Bird Diversity Hotspots is presented in Figure 1. Since it used bird species as an indicator, the Biodiversity Hotspot can also be referred as the Bird Diversity Hotspot.

The dominant type of land cover in the PFMU Wae Sapalewa is a lowland rainforest <300 m, and some of which has experienced exploitation of timber stands in the 1980s to early 2000s. There are also two other types of land cover which are quite extensive, i.e., freshwater swamp forest that experiences periodic inundation, but a small portion experiences permanent inundation, and lowland forest formation on limestone. Physiography of PFMU Wae Sapalewa area consists mostly of sloping and undulating areas, mainly in the North and East; while the West, South and Southeast are more dominated by hilly physiography with narrow valleys. The highest part in the South and Southeast is Mount Kalapahin; 400 m, asl, and the highest part in the West is Morita Hill; 350 m, asl (CoLUPSIA, 2013.) Based on the classification of climate regions by Schmidt and Ferguson, the PFMU Wae Sapalewa area is included in climate zone B (Schmidt and Ferguson, 1951; Lakitan, 2002).

The locations of Bird Diversity Hotspots and the form of the Wae Sapalewa PFMU management block are presented in Table 1. The Bird Diversity Hotspot has the shape and boundaries that follow the form of the Wae Sapalewa PFMU management blocks and the natural boundaries in the form of ridges and river borders as follows: Hotspots located throughout the Protection Forest (PtF) - Core Block, Production Forest (PdF) - Special Block, and Production Forest (PdF) - Protection Block. Hotspot locations whose boundaries cover only part of the form of management blocks, such as hotspot locations in the Production Forest (PdF) - Area Utilization Block. Furthermore, the shape and boundaries of hotspot locations in more detail utilize the boundaries of sub-block distribution, and also the natural boundaries in the form of ridges, valleys and rivers.

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**Table 1.** Bird diversity hotspot locations and the Wae Sapalewa PFMU management block forms.

Hotspot Location	Forms in the Wae Sapalewa PFMU Management Block
Masihulan	PdF - Area Utilization Block
Kaluala Mountain	PtF - Core Block
Hualu	PdF - Special Block
Roho	PdF - Area Utilization Block
Kalapain Mountain	PtF - Core Block
Wai Putiputi	PdF - Area Utilization Block, and PdF / Protection Block
Solea	PdF - Area Utilization Block, and PdF / HHBK-HA Utilization Block

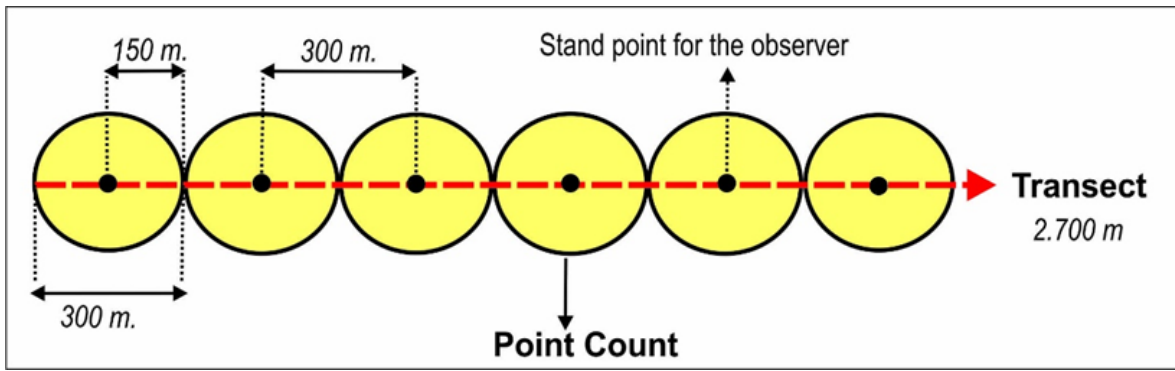


Figure 2. Design of point count on the transects

**Methods of study**

The research was conducted in March 2015 to December 2018. The bird data collection was the Point Count method (Vergara *et al.*, 2010; Sullivan, 2012; Blackwellet *al.*, 2013). Point Count was an imaginary circular area, where the observer stands in the middle while looking in all directions for data collection for 15 minutes. Each Point Count was also called a Data Collection Station. The shape of the Point Count in the field was as follows: the length of the circle area of the Point Count was 300 m and the radius length of the circle was 150 m. Point count was placed on the transect (working line), with a distance between the center of the point count was 300 m. The total length of the transect at the hotspot location was 10.8 Km, divided into 4 transects, each was 2.7 Km long. Each transect had 10 Point Count with a total of 4 transects at each hotspot location, thus there were a total of 40 Point Count at each hotspot location (Figure 2).

**Data collection**

Data collection was performed based on the point count on the transect that was determined according to the representation of the transect location in the studied habitat, in which the observer stood in the center of the circle while observing in all directions for data collection for 15 minutes, then the observer moved to the next Point Count position by walking slowly. Data collection was performed only once, in the morning between 07.00 CET and 10.00 CET. Each transect was traced only once, and this was intended to obtain truly independent field data. The data variables collected were 1). Type information includes: hours of observation; type; number of contacts; number of individuals; form of observation; 2). Habitat information includes: Land cover type; Land cover conditions; Physiographic form; Place height; 3). Other information includes: Season and weather; and human activities. Factors to be considered in placing observation transects were the diversity of land cover types, land cover forms and variations in altitude, and these factors are important ecosystem gradients in the biodiversity distribution.

**Data analysis**

**Species Richness Index (R<sub>1</sub>)**

Species richness index is the total number of species in a community, the value depends on the sample size of the area and the time needed to achieve it, as explained in the description of the data collection method above. The calculation of species richness index used the Margalef equation (Bibi and Ali, 2013).

$$R = \frac{(S - 1)}{\text{Ln. } N}$$

Where, R = Margalef species richness index  
 S = number of species observed  
 N = number of individuals of all types  
 Ln = natural logarithm value

Richness index category:

- R < 2.5 Low species richness
- 2.5 > R < 4 Medium species richness
- R > 4 High species richness

**Abundance Classes Based on Amount of Observation Time**

Analysis of Abundance Classes based on observation time was only intended for indicator bird species groups. This abundance class was based on the length of time or the total time of one bird species observed, then divided the total time into several categories of abundance classes. The classification of abundance classes was made according to the category of abundance classes by (Issa, 2019), which was modified in accordance with the conditions of the field collection data, as presented in Table 2.

Table 2. Categories of abundance classes

Duration of Time Observed	Abundance Value	Abundance Class
< 20 minutes	1	Rare
21 – 50 minutes	2	Uncommon
51 – 100 minutes	3	Often
101 – 200 minutes	4	Common
> 201 minutes	5	Very Common

**Relative Abundance Index (RAI)**

The relative abundance based on the Relative Abundance Index (RAI) referred to the number of individuals of a bird species compared to the total number of individuals of all bird species at all data collection stations. The Relative Abundance Index value is an indication which states the abundance condition of one bird species population to the population abundance of all bird species in the study location. The relative abundance index used the equation adopted from (Issa, 2019) (Table 3).

$$RAI = \frac{\text{Number of individuals of a species } (ni)}{\text{The total number of individuals found } (N)} \times 100 \%$$

**Table 3.** Categories of relative abundance

Relative Abundance Index Value	Abundance Category
< 15 %	Low
15 – 20 %	Medium
> 20 %	High

## RESULTS

### *Bird Species Diversity at Hotspot Locations*

There were 121 species of birds found in 7 Bird Diversity hotspot locations. Based on the species status; there were 9 endemic species of Seram Island, 4 endangered species according to the IUCN list, 21 limited distributed species, and 6 unique species (Table 4 and Figure 3).

**Table 4.** Species of birds found at 7 hotspot locations Expressed by the number of individuals of each species for each hotspot location

Bird Species Family	Scientific Name	Masihulan	Mt. Kaluala	Huau	Roho	Mt. Kalapahin	Wai Putiputi	Solea
CASUARIDAE	<i>Casuarium casuarium</i> <sup>Unique</sup>	1	7	1	6			5
FREGATIDAE	<i>Fregata ariel</i>	3						
PHALACROCOCIDAE	<i>Phalacrocorax melanoleucos</i>						1	2
ARDIDAE	<i>Egretta picata</i>						4	6
	<i>Egretta intermedia</i>						2	3
	<i>Egretta garzetta</i>	4					3	5
	<i>Yxobrychus sinensis</i>						1	1
THRESKIORNITHIDAE	<i>Therkiornis molucca</i>	1		1			1	3
ACCIPITRIDAE	<i>Pandion haliaetus</i>	2	1	1			1	2
	<i>Aviceda subcristata</i>	3	2	2	1	1	1	2
	<i>Haliaastur indus</i>	14	9	7	4	3	8	13
	<i>Haliaeetus leucogaster</i>	1		1			1	1
	<i>Accipiter novaehollandiae</i>		1			1		1
	<i>Accipiter erythrauchen</i> <sup>En.M - BST</sup>	1	2	4	1	3		5
	<i>Accipiter meyerianus</i>	1	1	1				
	<i>Ictinaetus malayensis</i>	6	5	1				
	<i>Aquila gurneyi</i>	2	3	1	1	1		3
FALCONIDAE	<i>Falco moluccensis</i>	7	4	5		1	3	5
	<i>Falco severus</i>		2	1				2
ANATIDAE	<i>Tandorna radjah</i>							1
	<i>Nettapus pulchellus</i>				2		3	
	<i>Anas gracilis</i>				2			4
MEGAPODIDAE	<i>Megapodius reinwardt</i>	2	10	6	8	5	4	4
	<i>Eulipoa wallacei</i> <sup>BST-VU</sup>							3
PHASIANIDAE	<i>Coturnix chinensis</i>	4		6				6
RALLIDAE	<i>Gallirallus philippensis</i>				1		1	2
	<i>Amaurornis olivaceus</i>	2			1		2	2
	<i>Amaurornis phoenicurus</i>						1	1
	<i>Porphyrio porphyrio</i>						1	
SCOLOPACIDAE	<i>Tringa stagnatilis</i>				2		2	5
COLUMBIDAE	<i>Columba livia</i>	12					10	16
	<i>Streptopelia chinensis</i>	8	4	7	3		4	9
	<i>Macropygia amboinensis</i>	8	7	5	5	3	4	5
	<i>Reinwardtoena</i>		8		6	6		
	<i>Gymnophaps mada</i> <sup>En.S,B</sup>	28	17	9	25	31	21	18
	<i>Chalcophaps indica</i>	4	4	2	1	2	2	3
	<i>Geopelia striata</i>	4	1	3	2		3	2
	<i>Ptilinopus superbus</i>	76	60	70	36	30	66	69

### *Species richness*

The Margalef species richness index in Table 5 showed that all hotspot locations had a high species richness index because the resulting richness index was at 7.96515-11.21258, and it means above the high richness index criteria of  $R > 4$ . This condition was caused by the high number of bird species found in all hotspot locations, ranging from 65 to 98 species, the number of individual bird species recorded at each hotspot location was quite good, and the total number of individuals recorded at each hotspot location was also good.

As per location, it was shown that the Solea and Masihulan Hotspot locations had a species richness index that was higher than other hotspot locations, each having a value at 11.01675 in Masihulan and 11.21258 in Solea. This was influenced by the number of species and the total number of individuals of all bird species collected at

Table 4 contd.

	<i>Ptilinopus rivoli</i>	31	26		22	29		
	<i>Ptilinopus viridis</i>	112	96	81	84	57	58	67
	<i>Ptilinopus melanospila</i>		7		5	3		
	<i>Ducula perspicillata</i> <sup>BST</sup>	126	121	110	112	111	90	123
	<i>Ducula concina</i> <sup>BST</sup>	30	16	26				30
	<i>Ducula bicolor</i>	6					8	
PSITTACIDAE	<i>Eos bornea</i> <sup>En.M - BST</sup>	320	302	184	243	190	205	145
	<i>Eos semilarvata</i> <sup>En.S - BST</sup>		7			9		
	<i>Trichoglossus haematodus</i>	223	190	177	135	112	231	257
	<i>Lorius domicella</i> <sup>En.S - BST - VU</sup>	3						
	<i>Chamosyna placentis</i> <i>Unique</i>	57	32	74	37	36	78	81
	<i>Micropsitta bruijnii</i> <sup>Unique</sup>	7	30			12		
	<i>Cacatua moluccensis</i> <i>En.S, KI - BST - VU</i>	26	21	16	22	30	6	14
	<i>Eclectus roratus</i>	28	29	26	18	12	23	32
	<i>Geoffroyus geoffroyi</i>	96	81	88	76	53	77	90
	<i>Tanygnathus megalorhynchus</i>	108	78	95	68	56	81	87
	<i>Alisterus amboinensis</i> <i>Unique</i>	17	47	15	37	10	40	23
CUCULIDAE	<i>Cuculus saturates</i>		1		2			1
	<i>Cacomantis sepulcralis</i>	8	5	6	4	2	8	5
	<i>Cacomantis variolus</i>	2	1	1	1	1	2	3
	<i>Chrysococcyx minutillus</i>	2					1	
	<i>Eudynamys scolopacea</i>	18	6	20	12		10	16
	<i>Eudynamys cyanocephala</i>	12	5	10	6		4	7
	<i>Scythrops novaehollandiae</i>	36	37	22	17	5	28	33
CENTROPODID AE	<i>Centropus bengalensis</i>	48		34			20	28
TYTONIDAE	<i>Tyto sororcula</i>		4		5			
STRIGIDAE	<i>Otus magicus</i>	22	14	18	13	7	13	19
	<i>Ninox squamipila</i> <sup>BST</sup>	6	2	4	4	1	5	3
CAPRIMULGID AE	<i>Caprimulgus macrurus</i>	8		5			3	14
APODIDAE	<i>Collocalia vanikorensis</i>	280	240	310	110	275	255	305
	<i>Collocalia infuscata</i>	45	30	60		25	40	55
	<i>Collocalia esculenta</i>	920	860	990	720	660	825	1060
	<i>Apus pasificus</i>	26		18			12	37
HEMIPROCINID AE	<i>Hemiprocne mystacea</i>	33	8	27	5		21	34
HALCYONIDAE	<i>Tanysiptera galatea</i> <sup>Unique</sup>	5	3		6		4	8
	<i>Halcyon lazuli</i> <sup>End.S, A, H - BST - VU</sup>	2	2		3		2	
	<i>Halcyon chloris</i>	24	24	8	21	2	18	20
	<i>Halcyon sancta</i>	14	13	10	12	5	11	18
ALCEDINIDAE	<i>Ceyx lepidus</i>	2	3	1	4		3	
	<i>Alcedo atthis</i>	1	3		3		2	2
MEROPIDAE	<i>Merops ornatus</i>	44	16	38	8		24	36
CORACIDAE	<i>Eurystomus orientalis</i>	9	2	10	1	2	7	12
BUCEROTIDAE	<i>Rhyticeros plicatus</i> <sup>Unique</sup>	86	60	58	70	56	49	47
PITTIDAE	<i>Pitta erythrogaster</i>		1		1	2		
	<i>Pitta elegans</i>		1			1		
HIRUNDINIDAE	<i>Hirundo rustica</i>	120	30	140			60	80
	<i>Hirundo tahitica</i>	55		40			25	45
MOTACILLIDAE	<i>Motacila flava</i>	6		9			7	11
	<i>Motacila cinerea</i>	12	3	18	2	2	12	9
CAMPEPHAGID AE	<i>Coracina novaehollandiae</i>	7	13	16			12	10
	<i>Coracina atriceps</i> <sup>En.M - BST</sup>	33	21	18	14	17	26	20
	<i>Coracina ceramensis</i> <sup>En.M - BST</sup>	15	16	7	10	11	7	10

Table 4 contd.

PYCNONOTIDAE	<i>Ixos affinis affinis</i>	66	78	71	55	40	53	58
DICRURIDAE	<i>Dicrurus bracteatus</i>	207	201	199	115	92	188	231
CORVIDAE	<i>Corvus enca</i>	23	8	25			17	31
SYLVIIDAE	<i>Locustella fasciolata</i>	13		10				8
	<i>Acrocephalus orientalis</i>	3		1				3
	<i>Phylloscopus borealis</i>	11	5	9		4		9
CISTICOLIDAE	<i>Cisticola exilis</i>	14		8			6	15
MUSCICAPIDAE	<i>Muscicapa griseisticta</i>	28	22	26	13	16	26	23
	<i>Ficedula buruensis</i> <sup>En.M-BST</sup>	4	4	2	1	8		
MONARCHIDAE	<i>Monarcha trivirgatus</i>	26	48	8	21	27	29	21
	<i>Myagra galeata</i> <sup>BST</sup>	31	30	32	21	22	18	18
RHIPIDURIDAE	<i>Rhipidura leucophrys</i>	21		12	8		15	28
	<i>Rhipidura rufiventris</i>	18	28	5	18	14	16	3
	<i>Rhipidura dedemi</i> <sup>En.S-BST</sup>	7	14	2	14	15	5	9
PACHYCEPHALIDAE	<i>Pachycephala pectoralis</i>		5			3		
	<i>Pachycephala griseonota</i> <sup>BST</sup>	11	10	5	14	14	7	15
ARTAMIDAE	<i>Artamus leucorhynchus</i>	14	8	19		10	13	21
STURNIDAE	<i>Aplonis mysolensis</i>	425	365	330	215	175	380	525
	<i>Aplonis metallica</i>	355	315	190	260	180	310	330
	<i>Basilornis corythaix</i> <sup>En.S-BST</sup>	60	48	58	41	45	29	47
MELIPHAGIDAE	<i>Philemon subcorniculatus</i> <sup>En.S-BST</sup>	93	173	187	129	121	128	155
	<i>Myzomela blasii</i> <sup>End.S,B,A-BST</sup>	1	2			3		
	<i>Myzomela sanguinolenta</i>		5			6		
NECTARINIIDAE	<i>Nectarinia aspasia</i>	128	112			1	1	1
	<i>Nectarinia jugularis</i>	128	112	106	99	90	110	130
DICAEIDAE	<i>Dicaeum vulneratum</i> <sup>En.M-BST</sup>	154	123	164	164	110	104	146
ZOSTEROPIDAE	<i>Zosterops montanus</i>		65			47		
	<i>Zosterops atrifrons</i>	110	145	80	120	240		75
ESTRILDINAE	<i>Erythrura trichroa</i>	16		35	18		22	32
	<i>Lonchura molucca</i>	135		155	95		120	155
	<i>Lonchura Malacca</i>	26	7	15	11	10	27	32
<b>Total Number of Individual Species</b>		<b>5.470</b>	<b>4.588</b>	<b>4.721</b>	<b>3452</b>	<b>3.174</b>	<b>4.232</b>	<b>5.293</b>
<b>Number of Species</b>		<b>97</b>	<b>87</b>	<b>85</b>	<b>75</b>	<b>65</b>	<b>90</b>	<b>98</b>

Note: End.S=Endemic Seram, End. S, Br= Endemic Seram and Buru, BST= Limited Distributed Bird, End.S,Kl= Endemic Seram and Lease, End. S, A=Endemic Seram and Ambon, VU= Vulnerable Bird, End.S,A,H= Endemic Seram, Ambon, and Haruku, End. M= Endemic Maluku, U= Unique Bird

the two hotspots which were higher when compared to other hotspots. Then it was followed by Wai Putiputi Hotspot at 10.54128, Mt. Kaluala Hotspot at 10.18233, Huaulu Hotspot at 9.88933, and Roho Hotspot at 9.02990; while Mt. Kalapahin Hotspot had the lowest species richness index, which was 7.96515.

#### Relative Abundance Index of Indicator Bird

The Relative Abundance Index (RAI) of indicator birds at the hotspot locations according to Table 4 are as follows: 4 species with high RAI in all locations, between 10.29% - 27.21%, were *Eos bornea*, *Philemon subcorniculatus*, *Dicaeum vulneratum* and *Ducula perspicillata*. One species with medium RAI in all locations was *Rhyticeros p.*, with RAI value of 5.41% - 7.19%. While 22 other species of indicator birds had low RAI values, between 0.08% - 4.90%. Recapitulation per RAI category at each hotspot location in Table 6 showed that bird species that

had high RAI values were spread evenly in all hotspot locations as many as 4 species, bird species medium RAI were spreading slightly varied between 1-3 species, as well as bird species with low RAI spread varied between 13-20 species.

#### Abundance Classes of Indicator Birds Based on Amount of Observation Time

The results of the abundance class study of the indicator bird species at Hotspot locations are presented in Table 5. In the abundance class column, it appears that the indicator bird species is in the Rare to Very Common abundance class. The explanation was that six species were in the Rare abundance class with the observed amount of time between 3 - 20 minutes, i.e. *Accipiter erythrauchen*, *Halcyon lazuli*, *Myzomela blasii*,



Figure 3. Important bird species found at bird diversity hotspot locations

Table 5. Richness index of bird species at hotspot locations and physiographic conditions

Hotspot Location	Species Number	Individual Number	Richness Index	Land cover	Topography
Masihulan	97	5.470	11.01675	Primary lowland forest Secondary lowland forest Fields, mixed farms	Wavy Hilly < 300 m Hilly > 300 m
Mt. Kaluala	87	4.588	10.18233	Primary lowland forest freshwater swamp forest	Wavy Hilly 300 Hilly > 300 m
Huaulu	85	4.721	9.88933	Primary lowland forest Logged forest Fields, mixed farms	Wavy Hilly 300
Roho	75	3.452	9.02990	Primary lowland forest Logged forest Fields, mixed farms	Wavy Hilly 300
Mt. Kalapahin	65	3.174	7.96515	Primary lowland forest Primary hilly forest	Hilly < 300 m Hilly > 300 m
Wai Putiputi	90	4.233	10.54128	Secondary lowland forest Fields, mixed farms	Wavy Hilly < 300 m
Solea	98	5.293	11.21258	Secondary lowland forest Fields, mixed farms	Wavy Hilly < 300 m

*Eos semilarvata*, *Eulipoa wallacei* and *Lorius domicella*. Seven species were in the Rare to Uncommon abundance class with an observed amount of time between 5 - 50 minutes, i.e. *Gymnophaps mada*, *Rhipidura dedemi*, *Ninoxsquampila s*, *Pachycephala griseonota*, *Casuaris casuaris*, *Ficedula Rushensis* and *Micropsitta bruijnii*. One species was in the Uncommon abundance class with an observed amount of time between 22 - 40 minutes, i.e., *Tanysiptera galatea*. Four species were in the Uncommon to Often abundance class with an observed amount of time between 21 - 95 minutes, i.e., *Ducula concina*, *Alisterus amboinensis*, *Coracina atriceps*, *Coracina ceramensis*. Two species were in the Uncommon to Common abundance class with an observed amount of time between 24-110 minutes, i.e., *Cacatua moluccensis* and *Charmosyna placenthis*.

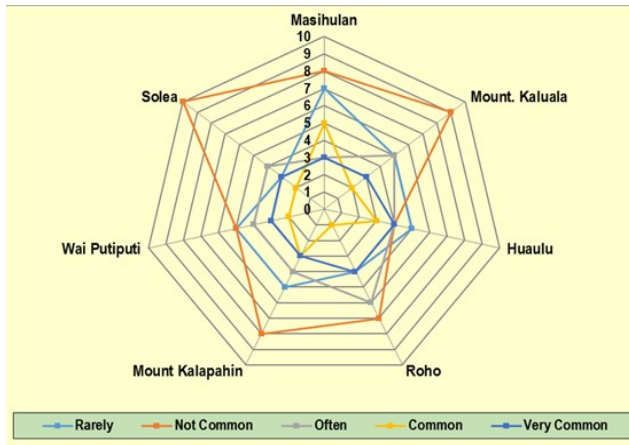
Three species were included in the Often to Common abundance class with an observed amount of

time between 70 - 162 minutes, i.e., *Rhyticeros plicatus*, *Myagra galeata* and *Basilornix corythaix*. Two species were in the Common to Very Common abundance class with an observed amount of time between 117 - 322 minutes, i.e., *Dicaeum vulneratum* and *Eos bornea*. Two species were in the Very Common abundance class with an observed amount of time between 207 - 441 minutes, i.e., *Ducula perspicillata* and *Philemon subcorniculatus*.

Figure 4 shows that the indicator bird species at hotspot locations were mostly in the Uncommon abundance class that spread dominantly at 5 hotspot locations (Masihulan, Mt. Kaluala, Roho, Mt. Kalapahin and Solea). Then it was followed by Rare abundance class that spread a lot at 4 hotspot locations (Masihulan, Huaulu, Mt. Kalapahin and Wai Putiputi), then the Often abundance class was spread at 2 hotspot locations (Roho and Solea), while the distribution of Common and Very Common abundance classes were low in all hotspot locations.

**Table 6.** Relative abundance index value of indicator bird species at hotspot locations

Bird Species	Masihulan	Mt. Kaluala	Huauulu	Roho	Mt. Kalapahin	Wai Putiputi	Solea
<i>Eos bornea</i> <sup>End. M - BST</sup>	26.14	27.21	18.85	24.82	22.22	24.78	15.61
<i>Philemon subcorniculatus</i> <sup>En.S - BST</sup>	15.77	15.59	19.16	13.18	14.15	15.48	16.68
<i>Dicaeum vulneratum</i> <sup>End. M - BST</sup>	12.58	11.08	16.80	16.75	12.87	12.58	15.72
<i>Ducula perspicillata</i> <sup>BST</sup>	10.29	10.90	11.27	11.44	12.98	10.88	13.24
<i>Rhyticeros plicatus</i> <sup>U</sup>	7.03	5.4	5.94	7.15	6.55	5.93	5.06
<i>Charmosyna placentis</i> <sup>U</sup>	4.66	2.88	7.58	3.78	4.21	9.43	8.72
<i>Basilornis corythaix</i> <sup>End.S - BST</sup>	4.90	4.32	5.94	4.19	5.26	3.50	5.06
<i>Myagra galeata</i> <sup>BST</sup>	2.53	2.70	3.28	2.15	2.57	2.18	1.94
<i>Gymnophaps mada</i> <sup>End.S,B - BST</sup>	2.29	1.53	0.92	2.55	3.63	2.54	1.92
<i>Cacatua moluccensis</i> <sup>End.S,KI - BST - VU</sup>	2.12	1.89	1.64	2.25	3.51	0.72	1.51
<i>Alisterus amboinensis</i> <sup>U</sup>	1.39	4.23	1.54	3.78	1.17	4.84	2.47
<i>Coracina atriceps</i> <sup>End. M - BST</sup>	2.70	1.89	1.84	1.43	1.99	3.14	2.15
<i>Ducula concina</i> <sup>BST</sup>	2.45	1.44	2.66	0.00	0.00	0.00	3.23
<i>Pachycephala griseonota</i> <sup>BST</sup>	0.90	0.90	0.51	1.43	1.64	0.85	1.61
<i>Coracina ceramensis</i> <sup>End. M - BST</sup>	1.23	1.44	0.72	1.02	1.29	0.85	1.08
<i>Rhipidura dedemi</i> <sup>End.S - BST</sup>	0.57	1.26	0.20	1.43	1.75	0.60	0.97
<i>Micropsitta bruijnii</i> <sup>U</sup>	0.57	2.70	0.00	0.00	1.40	0.00	0.00
<i>Ficedula buruensis</i> <sup>End. M - BST</sup>	0.33	0.36	0.20	0.10	0.94	0.00	0.00
<i>Casuaris casuaris</i> <sup>U</sup>	0.08	0.63	0.10	0.61	0.00	0.00	0.54
<i>Tanysiptera galatea</i> <sup>U</sup>	0.41	0.27	0.00	0.61	0.00	0.48	0.86
<i>Ninox squamipila</i> <sup>BST</sup>	0.49	0.18	0.41	0.41	0.12	0.60	0.32
<i>Accipiter erythrauchen</i> <sup>End. M - BST</sup>	0.08	0.18	0.41	0.10	0.35	0.00	0.54
<i>Myzomela blasii</i> <sup>End.S,B,A - BST</sup>	0.08	0.18	0.00	0.51	0.35	0.36	0.43
<i>Halcyon lazuli</i> <sup>End.S,A,H - BST - VU</sup>	0.16	0.18	0.00	0.31	0.00	0.24	0.00
<i>Eos semilarvata</i> <sup>End.S - BST</sup>	0.00	0.63	0.00	0.00	1.05	0.00	0.00
<i>Eulipoa wallacei</i> <sup>BST - VU</sup>	0.00	0.00	0.00	0.00	0.00	0.00	0.32
<i>Lorius domicella</i> <sup>End.S - BST - VU</sup>	0.25	0.00	0.00	0.00	0.00	0.00	0.00
Total	100	100	100	100	100	100	100
High RAI category > 10,1 %	4	4	4	4	4	4	4
Medium RAI category 5,1 % – 10 %	1	1	3	1	2	2	3
Low RAI category < 5 %	20	20	13	17	15	13	15
Total Species	25	25	20	22	21	19	22



**Figure 4.** Condition of abundance class distribution of indicator bird species at Hotspot locations

**DISCUSSION**

A total of 121 bird species were found at 7 hotspot locations (Table 4), with a total encounter of bird species at hotspot locations between 65 to 98 species, and important indicator bird species were between 19 to 25 species (Figure 3). The number of species is sufficient for a forest area, especially forests in an archipelago. This showed that the hotspot locations studied in the PFMU Wae Sapalewa area have a good level of species

Richness and diversity, and have a high species conservation value. This condition was confirmed by the high species richness index value (R1) of bird species at 7 hotspot locations with a species richness index value exceeding  $R > 4$ , which was between 7.96515 and 11.21258 (Table 5).

This species richness index value showed that forest areas in bird hotspot locations are forest areas with good ecological conditions as shown by the data in Table 5, and are a suitable habitat for the lives of various species of birds. (Wilcoxon *et al.*, 2015) stated that habitat that is still in good condition and contains a variety of food sources has the possibility of many bird species. Habitat found at hotspot locations was quite diverse and consisted of variations of land cover ranging from open areas in the form of cultivation land (Li *et al.*, 2011), grasslands with high bushes (Lee and Martin, 2017), then secondary forests to primary forests (Ayat & Tata, 2015; Prabowo *et al.*, 2016; Duraisamy *et al.*, 2018). In addition, hotspot locations also had physiographic variations from lowland wavy areas, low hilly areas to high hilly areas. There were also mixed productive community farms and old farms that are already in the form of dusung (heritage traditional farms). This condition provides a good opportunity for bird activities according to their niche in the variety of habitat, which is associated with foraging, shelter and reproduction activities (Tryjanowski *et al.*, 2015; Wiedarti *et al.*, 2016; Baxter, 2019).

**Table 7.** Abundance classes of indicator bird species at Hotspot locations expressed by the observed amount of time

Bird Species Scientific name	Total Observed Time (/Minute)							Abundance Class Category
	Masihulan	Mt. Kaluala	Huau lu	Roho	Mt. Kalapahin	Wai Putiputi	Solea	
<i>Ducula perspicillata</i> <sup>BST</sup>	441	345	355	322	283	220	345	Very Common
<i>Philemon subcorniculatus</i> <sup>En.S - BST</sup>	330	222	270	259	207	212	245	Very Common
<i>Dicaeum vulneratum</i> <sup>En.M - BST</sup>	322	205	274	286	150	210	227	Common - Very Common
<i>Eos bornea bornea</i> <sup>En.M - BST</sup>	157	187	202	205	151	117	141	Common - Very Common
<i>Rhyticeros plicatus</i> <sup>U</sup>	125	100	170	125	90	99	125	Often - Common
<i>Myagra galeata</i> <sup>BST</sup>	162	125	108	99	90	90	80	Often - Common
<i>Basilornis corythaix</i> <sup>En.S-BST</sup>	130	86	125	70	78	80	95	Often - Common
<i>Cacatua moluccensis</i> <sup>En.S,KI-BST - VU</sup>	110	101	80	90	85	24	50	Uncommon - Common
<i>Chamosyna placensis</i> <sup>U</sup>	55	40	90	60	33	110	100	Uncommon - Common
<i>Alisterus amboinensis</i> <sup>U</sup>	50	86	54	80	28	95	40	Uncommon - Often
<i>Coracina atriceps</i> <sup>En.M - BST</sup>	85	52	45	35	45	40	25	Uncommon - Often
<i>Gymnophaps mada</i> <sup>En.S,B</sup>	40	20	30	39	45	30	30	Uncommon - Often
<i>Coracina ceramensis</i> <sup>En.M - BST</sup>	55	34	22	25	25	23	24	Uncommon - Often
<i>Ducula concina</i> <sup>BST</sup>	45	30	54	0	0	0	54	Uncommon - Often
<i>Rhipidura dedemi</i> <sup>En.S - BST</sup>	30	42	10	50	48	24	36	Rare - Uncommon
<i>Pachycephala griseonota</i> <sup>BST</sup>	30	32	17	40	35	18	25	Rare - Uncommon
<i>Ninox squamipila</i> <sup>BST</sup>	25	25	24	25	5	15	25	Rare - Uncommon
<i>Tanyptera galatea</i> <sup>U</sup>	22	25	0	30	0	25	40	Rare - Uncommon
<i>Casuaris casuaris</i> <sup>U</sup>	10	25	5	24	0	0	35	Rare - Uncommon
<i>Ficedula buruensis</i> <sup>En.M - BST</sup>	25	18	15	6	25	0	0	Rare - Uncommon
<i>Micropsitta bruijnii</i> <sup>U</sup>	15	0	0	0	15	0	0	Rare - Uncommon
<i>Accipiter erythrauchen</i> <sup>En.M - BST</sup>	5	7	18	3	20	0	14	Rare
<i>Halcyon lazuli</i> <sup>End.S,A,H - BST - VU</sup>	18	15	0	10	0	16	0	Rare
<i>Myzomela blasii</i> <sup>End.S,B,A - BST</sup>	3	7	0	0	8	0	0	Rare
<i>Eos semilarvata</i> <sup>En.S - BST</sup>	0	15	0	0	17	0	0	Rare
<i>Eulipoa wallacei</i> <sup>BST - VU</sup>	0	0	0	0	0	0	12	Rare
<i>Lorius domicella</i> <sup>En.S - BST - VU</sup>	12	0	0	0	0	0	0	Rare
Total Observed Time	2.302	1.876	1.968	1.883	1.483	1.448	1.768	
S Very Common Abundance Class	3	3	4	4	1	3	3	
S Common Abundance Class	5	3	3	1	3	2	2	
S Often Abundance Class	3	4	4	5	4	4	4	
S Uncommon Abundance Class	8	10	4	8	8	5	10	
S Rare Abundance Class	6	5	5	4	5	5	3	

In Table 6, there are 4 known bird species with high Relative Abundance Index (RAI) at 7 hotspot locations, i.e., *Eos bornea*, *Philemon subcorniculatus*, *Dicaeum vulneratum*, and *Ducula perspicillata*. One species with medium RAI at all hotspot locations was *Rhyticeros plicatus*, and another species with medium RAI, but only recorded at 4 hotspot locations was *Basilornix corythaix*. The other 21 species of bird indicators had a low RAI. Recapitulation per RAI category at each hotspot location shows that 4 species of birds with high RAI were spread evenly in all hotspot locations, while species of birds with medium RAI were spread between 1 to 3 species, and species of birds with low RAI were spread between 13 to 20 species. This condition shows that the abundance of indicator bird species in hotspot locations is very varied. There were species with high abundance, and there were those with low abundance, and even very low (Marchese, 2015; Araneda *et al.*, 2018).

The results of the Abundance Class analysis based on the total observed time in Table 5 shows that there were 4 species of birds in the Common to Very Common abundance class, i.e., *Ducula perspicillata*, *Philemon subcorniculatus*, *Dicaeum vulneratum*, and *Eos bornea*. There were 3 species in the Often to Common abundance class, i.e., *Rhyticeros plicatus*, *Myagra galeata*, and *Basilornix corythaix*. While 20 other species were in the Rare to Uncommon class abundance. This condition shows that some species of indicator birds were easily found, some other species

were rather difficult to find, while most species were difficult to find and some were very difficult to find in hotspot locations (Sussman *et al.*, 2019; Bellanthudawa *et al.*, 2019).

Based on the value of Relative Abundance Index (RAI) and Abundance Class, it was shown that 4 species of birds with high RAI value were in the Common to Very Common abundance class, whereas 2 species with the medium RAI value were in the Often to Common abundance class. From these two study methods, it is found that there were 6 species of birds, i.e., *Ducula perspicillata*, *Philemon subcorniculatus*, *Dicaeum vulneratum*, *Eos bornea*, *Rhyticeros plicatus*, and *Basilornix corythaix* that can be found in all hotspot locations with a significant level of encounter certainty. Relative Abundance Index values and Abundance Class categories can be indicators of the status and presence of important indicator bird species at hotspot locations (Mueller *et al.*, 2014; Bellanthudawa *et al.*, 2019; Persulesy *et al.*, 2019). The combination of these two forms of abundance studies provides reasonably good information about interesting indicator bird species as tourist attraction objects and ecotourism at bird diversity hotspot locations (Nicolaidis, 2014; Hakim, 2017).

Endemic bird species need attention because they are often the basis for consideration in conservation issues (Indrawan *et al.*, 2007; Mittermeier & Rylands, 2017). Endemic bird species such as those in the PFMU Wae Sapalewa area have become an attractive tourist

attraction in Seram Island, especially at the Manusela National Park area (Manusela National Park Center, 2016). This promising condition is an indication about the conservation value and the ecotourism selling value of bird diversity hotspots in the PFMU Wae Sapalewa area. Overall, 9 endemic bird species of Seram Island were found in 7 hotspot locations (Table 4), endemic birds found in these hotspot locations ranged from 5 to 8 species of 14 endemic bird species in Seram Island and surrounding islands. Eight species were found at hotspot locations of the Masihulan and Gunung Kaluala; 7 species were found at hotspot locations of Roho, Mount Kalapahin, and Wai Putiputi; 6 species were found at the hotspot location of Solea; and 5 species were found at hotspot locations of Huaulu. This finding revealed that the identified locations of bird diversity hotspots in the PFMU Wae Sapalewa area can be defined as the endemic bird hotspots. A research in Taiwan stated that an observed hotspot can be defined as an 'endemic hotspot' if there are at least 5 out of 17 species of endemic birds were found (Ko *et al.*, 2014).

## CONCLUSION

This research showed that overall 121 species of birds were found in 7 hotspot locations, with the total number of bird encounters in hotspot locations between 65 to 98 species, and important indicator bird species numbered between 19 to 25 species. This number of species is sufficient for a forest area, especially forests in an archipelago.

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