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# **LAMPIRAN**

Lampiran 1. Konstanta biomassa a dan b untuk setiap jenis/spesies ikan karang herbivora yang ditemukan pada penelitian ini (sumber: datamermaid.org)

Family	Name	Biomass Constant A	Biomass Constant B	Average Length
Acanthuridae	<i>Acanthurus auranticavus</i>	0.02291	2.96	10 - 30 cm
	<i>Acanthurus nigricans</i>	0.0263	2.93	12 cm
	<i>Acanthurus pyroferus</i>	0.02344	2.96	10 - 20 cm
	<i>Acanthurus thompsoni</i>	0.01698	2.99	10 - 20 cm
	<i>Ctenochaetus striatus</i>	0.01569	3.058599	5 - 15 cm
	<i>Ctenochaetus tominiensis</i>	0.02344	2.97	12 cm
	<i>Naso annulatus</i>	0.05103	2.71537	12 - 20 cm
	<i>Naso lituratus</i>	0.0324	2.94	15 - 25 cm
	<i>Naso unicornis</i>	0.029529	2.923551	30 - 35 cm
	<i>Naso vlamingii</i>	0.03104	2.843	10 - 25 cm
	<i>Zebrasoma scopas</i>	0.034129	2.939876	10 - 15 cm
	<i>Zebrasoma veliferum</i>	0.034107	2.861415	15 - 35 cm
Scaridae	<i>Cetoscarus bicolor</i>	0.0276	2.92	20 - 25 cm
	<i>Chlorurus bleekeri</i>	0.0415	2.946	8 - 35 cm
	<i>Chlorurus microrhinos</i>	0.021734	3.012728	35 cm
	<i>Chlorurus spilurus</i>	0.020118	3.059482	10 - 35 cm
	<i>Hipposcarus longiceps</i>	0.0161	3.05	25 - 30 cm
	<i>Scarus chameleon</i>	0.01445	3.04	15 - 20 cm
	<i>Scarus dimidiatus</i>	0.0278	3.049	8 - 25 cm
	<i>Scarus flavipectoralis</i>	0.01995	3.01	6 - 35 cm
	<i>Scarus forsteni</i>	0.0142	3.13	15 cm
	<i>Scarus frenatus</i>	0.01889	3.06	20 - 30 cm
	<i>Scarus ghobban</i>	0.015696	3.016738	12 - 15 cm
	<i>Scarus hypselopterus</i>	0.00794	3.11	15 - 30 cm
	<i>Scarus niger</i>	0.024136	3.147753	10 - 30 cm
	<i>Scarus oviceps</i>	0.0144	3.14	12 - 35 cm
	<i>Scarus quoyi</i>	0.0565	2.818	10 - 35 cm
	<i>Scarus rivulatus</i>	0.0184	3.058	12 - 35 cm
	<i>Scarus scaber</i>	0.0278	2.857	6 - 25 cm
	<i>Scarus schlelegeli</i>	0.020801	3.00244	12 cm
	<i>Scarus spinus</i>	0.00794	3.11	15 cm
	<i>Scarus tricolor</i>	0.0229	3.106	15 - 30 cm
Siganidae	<i>Siganus canaliculatus</i>	0.0232	2.8	20 cm
	<i>Siganus corallinus</i>	0.00234	3.82079	10 - 35 cm
	<i>Siganus guttatus</i>	0.054842	2.662305	30 cm
	<i>Siganus puelloides</i>	0.01761	3.02839	12 - 35 cm
	<i>Siganus punctatissimus</i>	0.01585	3.07	10 - 25 cm
	<i>Siganus punctatus</i>	0.019708	3.075761	12 - 20 cm
	<i>Siganus virgatus</i>	0.0204	3.236	5 - 25 cm
	<i>Siganus vulpinus</i>	0.01585	3.07	10 - 25 cm

## Lampiran 2. Hasil transformasi data

```
> with(species.comp, shapiro.test(sum.sp[Year == "2021"]))  
Shapiro-Wilk normality test  
data: sum.sp[Year == "2021"]  
W = 0.55842, p-value = 2.353e-08  
> with(species.comp, shapiro.test(sum.sp[Year == "2022"]))  
Shapiro-Wilk normality test  
data: sum.sp[Year == "2022"]  
W = 0.62454, p-value = 2.229e-08
```

a

```
F test to compare two variances  
data: sum.sp by Year  
F = 0.23147, num df = 29, denom df = 35, p-value = 0.000124  
alternative hypothesis: true ratio of variances is not equal to 1  
95 percent confidence interval:  
0.1153013 0.4758351  
sample estimates:  
ratio of variances  
0.2314682
```

b

1. Untuk melakukan uji-t, dilakukan uji *Shapiro-test* dan *F-test* untuk memenuhi asumsi uji *t-test*.
  - (a) *Shapiro-test* menunjukkan data komposisi jenis ikan karang herbivora  $p<0,05$ , sehingga data tidak memenuhi asumsi normalitas uji-t.
  - (b) *F-test* menunjukkan data komposisi jenis ikan karang herbivora  $p>0,05$ , sehingga data memenuhi asumsi homogenitas uji-t.

```
> with(species.comp, shapiro.test(sp.logg2[Year == "2021"])) #p>0.05  
Shapiro-Wilk normality test  
data: sp.logg2[Year == "2021"]  
W = 0.96229, p-value = 0.354  
> with(species.comp, shapiro.test(sp.logg2[Year == "2022"])) #p>0.05  
Shapiro-Wilk normality test  
data: sp.logg2[Year == "2022"]  
W = 0.96666, p-value = 0.3414
```

a

```
F test to compare two variances  
data: sp.logg2 by Year  
F = 0.63518, num df = 29, denom df = 35, p-value = 0.2139  
alternative hypothesis: true ratio of variances is not equal to 1  
95 percent confidence interval:  
0.3164028 1.3057577  
sample estimates:  
ratio of variances  
0.635181
```

b

2. Data komposisi jenis ikan karang herbivora tidak memenuhi asumsi normalitas dan homogenitas untuk uji-t, sehingga dilakukan *treatment* data dengan mentransformasi data menjadi log2.
  - (a) *Shapiro-test* menunjukkan data komposisi jenis ikan karang herbivora  $p>0,05$ , sehingga data memenuhi asumsi normalitas uji-t.
  - (b) *F-test* menunjukkan data komposisi jenis ikan karang herbivora  $p>0,05$ , sehingga data memenuhi asumsi homogenitas uji-t. Sehingga uji-t dilanjutkan dengan menggunakan data transformasi log2.

```
> with(species.comp, shapiro.test(sp.logg10[Year == "2021"]))  
Shapiro-Wilk normality test  
data: sp.logg10[Year == "2021"]  
W = 0.96229, p-value = 0.354  
> with(species.comp, shapiro.test(sp.logg10[Year == "2022"]))  
Shapiro-Wilk normality test  
data: sp.logg10[Year == "2022"]  
W = 0.96666, p-value = 0.3414
```

a

```
F test to compare two variances  
data: sp.logg10 by Year  
F = 0.63518, num df = 29, denom df = 35, p-value = 0.2139  
alternative hypothesis: true ratio of variances is not equal to 1  
95 percent confidence interval:  
0.3164028 1.3057577  
sample estimates:  
ratio of variances  
0.635181
```

b

3. Data komposisi jenis ikan karang herbivora tidak memenuhi asumsi normalitas dan homogenitas untuk uji-t, sehingga dilakukan *treatment* data dengan mentransformasi data menjadi log10.

## Lampiran 2. Lanjutan

- (a) *Shapiro-test* menunjukkan data komposisi jenis ikan karang herbivora  $p>0,05$ , sehingga data memenuhi asumsi normalitas uji-t.
- (b) *F-test* menunjukkan data komposisi jenis ikan karang herbivora  $p>0,05$ , sehingga data memenuhi asumsi homogenitas uji-t.

```
> with(species.comp, shapiro.test(sp.sqrtt[Year == "2021"]))  
Shapiro-Wilk normality test  
data: sp.sqrtt[Year == "2021"]  
W = 0.81216, p-value = 0.0001101  
> with(species.comp, shapiro.test(sp.sqrtt[Year == "2022"]))  
Shapiro-Wilk normality test  
data: sp.sqrtt[Year == "2022"]  
W = 0.82498, p-value = 5.316e-05
```

a

F test to compare two variances

data: sp.sqrtt by Year

F = 0.39359, num df = 29, denom df = 35, p-value = 0.01191

alternative hypothesis: true ratio of variances is not equal to 1

95 percent confidence interval:

0.1960570 0.8091047

sample estimates:

ratio of variances

0.393586

b

- 4. Data komposisi jenis ikan karang herbivora tidak memenuhi asumsi normalitas dan homogenitas untuk uji-t, sehingga dilakukan *treatment* data dengan mentransformasi data menjadi *square root*.
- (a) *Shapiro-test* menunjukkan data komposisi jenis ikan karang herbivora  $p<0,05$ , sehingga data tidak memenuhi asumsi normalitas uji-t.
- (b) *F-test* menunjukkan data komposisi jenis ikan karang herbivora  $p<0,05$ , sehingga data tidak memenuhi asumsi homogenitas uji-t.

```
> with(species.comp, shapiro.test(sp.fourth[Year == "2021"]))  
Shapiro-Wilk normality test  
data: sp.fourth[Year == "2021"]  
W = 0.91702, p-value = 0.02246  
> with(species.comp, shapiro.test(sp.fourth[Year == "2022"]))  
Shapiro-Wilk normality test  
data: sp.fourth[Year == "2022"]  
W = 0.91665, p-value = 0.01005
```

a

F test to compare two variances

data: sp.fourth by Year

F = 0.50855, num df = 29, denom df = 35, p-value = 0.0655

alternative hypothesis: true ratio of variances is not equal to 1

95 percent confidence interval:

0.2533251 1.0454435

sample estimates:

ratio of variances

0.5085521

b

- 5. Data komposisi jenis ikan karang herbivora tidak memenuhi asumsi normalitas dan homogenitas untuk uji-t, sehingga dilakukan *treatment* data dengan mentransformasi data menjadi *fourth root*.
- (a) *Shapiro-test* menunjukkan data komposisi jenis ikan karang herbivora  $p<0,05$ , sehingga data tidak memenuhi asumsi normalitas uji-t.
- (b) *F-test* menunjukkan data komposisi jenis ikan karang herbivora  $p>0,05$ , sehingga data memenuhi asumsi homogenitas uji-t. Namun uji-t tidak dapat dilanjutkan karena data tidak memenuhi asumsi normalitas.

## Lampiran 2. Lanjutan

```
> with(fish.trans, shapiro.test(Abundance[Year == "2021"]))  
Shapiro-Wilk normality test  
data: Abundance[Year == "2021"]  
W = 0.87782, p-value = 0.01098  
> with(fish.trans, shapiro.test(Abundance[Year == "2022"]))  
Shapiro-Wilk normality test  
data: Abundance[year == "2022"]  
W = 0.72516, p-value = 3.022e-05
```

ab

```
F test to compare two variances  
data: Abundance by Year  
F = 0.13376, num df = 21, denom df = 22, p-value = 2.018e-05  
alternative hypothesis: true ratio of variances is not equal to 1  
95 percent confidence interval:  
0.0563725 0.3201883  
sample estimates:  
ratio of variances  
0.1337587
```

b

6. Untuk melakukan uji-t, dilakukan uji *Shapiro-test* dan *F-test* untuk memenuhi asumsi uji-t.
- (a) *Shapiro-test* menunjukkan data kelimpahan ikan karang herbivora  $p<0,05$ , sehingga data tidak memenuhi asumsi normalitas uji-t.
- (b) *F-test* menunjukkan data kelimpahan ikan karang herbivora  $p>0,05$ , sehingga data memenuhi asumsi homogenitas uji-t. Namun uji-t tidak dapat dilanjutkan karena data tidak memenuhi asumsi normalitas.

```
> with(fish.trans, shapiro.test(log.2[Year == "2021"]))  
Shapiro-Wilk normality test  
data: log.2[Year == "2021"]  
W = 0.91031, p-value = 0.04803  
> with(fish.trans, shapiro.test(log.2[Year == "2022"]))  
Shapiro-Wilk normality test  
data: log.2[Year == "2022"]  
W = 0.85799, p-value = 0.003811
```

ab

```
F test to compare two variances  
data: log.2 by Year  
F = 1.1402, num df = 21, denom df = 22, p-value = 0.7611  
alternative hypothesis: true ratio of variances is not equal to 1  
95 percent confidence interval:  
0.4805472 2.7294440  
sample estimates:  
ratio of variances  
1.140226
```

b

7. Data kelimpahan ikan karang herbivora tidak memenuhi asumsi normalitas dan homogenitas untuk uji-t, sehingga dilakukan *treatment* data dengan mentransformasi data menjadi log2.
- (a) *Shapiro-test* menunjukkan data kelimpahan ikan karang herbivora  $p>0,05$ , sehingga data tidak memenuhi asumsi normalitas uji-t.
- (b) *F-test* menunjukkan data kelimpahan ikan karang herbivora  $p>0,05$ , sehingga data memenuhi asumsi homogenitas uji-t. Namun uji-t tidak dapat dilanjutkan karena data tidak memenuhi asumsi normalitas.

## Lampiran 2. Lanjutan

```
> with(fish.trans, shapiro.test(log.10[Year == "2021"]))
Shapiro-Wilk normality test
data: log.10[Year == "2021"]
W = 0.91031, p-value = 0.04803
> with(fish.trans, shapiro.test(log.10[Year == "2022"]))
Shapiro-Wilk normality test
data: log.10[Year == "2022"]
W = 0.85799, p-value = 0.003811
```

a

**F test to compare two variances**  
 data: log.10 by Year  
 $F = 1.1402$ , num df = 21, denom df = 22, p-value = 0.7611  
 alternative hypothesis: true ratio of variances is not equal to 1  
 95 percent confidence interval:  
 $0.4805472 \text{--} 2.7294440$   
 sample estimates:  
 ratio of variances  
 $1.140226$

b

8. Data kelimpahan ikan karang herbivora tidak memenuhi asumsi normalitas dan homogenitas untuk uji-t, sehingga dilakukan *treatment* data dengan mentransformasi data menjadi log10.
- (a) *Shapiro-test* menunjukkan data kelimpahan ikan karang herbivora  $p<0,05$ , sehingga data tidak memenuhi asumsi normalitas uji-t.
  - (b) *F-test* menunjukkan data kelimpahan ikan karang herbivora  $p>0,05$ , sehingga data memenuhi asumsi homogenitas uji-t. Namun uji-t tidak dapat dilanjutkan karena data tidak memenuhi asumsi normalitas.

```
> with(fish.trans, shapiro.test(square.root[Year == "2021"]))
Shapiro-Wilk normality test
data: square.root[Year == "2021"]
W = 0.96938, p-value = 0.6968
> with(fish.trans, shapiro.test(square.root[Year == "2022"]))
Shapiro-Wilk normality test
data: square.root[Year == "2022"]
W = 0.90601, p-value = 0.03369
```

a

**F test to compare two variances**  
 data: square.root by Year  
 $F = 0.43175$ , num df = 21, denom df = 22, p-value = 0.05901  
 alternative hypothesis: true ratio of variances is not equal to 1  
 95 percent confidence interval:  
 $0.181961 \text{--} 1.033514$   
 sample estimates:  
 ratio of variances  
 $0.4317506$

b

9. Data kelimpahan ikan karang herbivora tidak memenuhi asumsi normalitas dan homogenitas untuk uji-t, sehingga dilakukan *treatment* data dengan mentransformasi data menjadi *square root*.
- (a) *Shapiro-test* menunjukkan data kelimpahan ikan karang herbivora  $p<0,05$ , sehingga data tidak memenuhi asumsi normalitas uji-t.
  - (b) *F-test* menunjukkan data kelimpahan ikan karang herbivora  $p>0,05$ , sehingga data memenuhi asumsi homogenitas uji-t. Namun uji-t tidak dapat dilanjutkan karena data tidak memenuhi asumsi normalitas.

```
> with(fish.trans, shapiro.test(fourth.abundance[Year == "2021"]))
Shapiro-Wilk normality test
data: fourth.abundance[Year == "2021"]
W = 0.97329, p-value = 0.7858
> with(fish.trans, shapiro.test(fourth.abundance[Year == "2022"]))
Shapiro-Wilk normality test
data: fourth.abundance[Year == "2022"]
W = 0.93749, p-value = 0.1587
```

a

**F test to compare two variances**  
 data: fourth.abundance by Year  
 $F = 0.73024$ , num df = 21, denom df = 22, p-value = 0.4749  
 alternative hypothesis: true ratio of variances is not equal to 1  
 95 percent confidence interval:  
 $0.3077579 \text{--} 1.7480238$   
 sample estimates:  
 ratio of variances  
 $0.7302372$

b

## Lampiran 2. Lanjutan

10. Data kelimpahan ikan karang herbivora tidak memenuhi asumsi normalitas dan homogenitas untuk uji-t, sehingga dilakukan *treatment* data dengan mentransformasi data menjadi *fourth root*.
- (a) *Shapiro-test* menunjukkan data kelimpahan ikan karang herbivora  $p>0,05$ , sehingga data memenuhi asumsi normalitas uji-t.
- (b) *F-test* menunjukkan data kelimpahan ikan karang herbivora  $p>0,05$ , sehingga data memenuhi asumsi homogenitas uji-t. Selanjutnya uji-t kelimpahan ikan karang herbivora akan dilakukan menggunakan transformasi data *fourth root*.

```
> with(fish.trans, shapiro.test(Biomass[Year == "2021"]))  
Shapiro-Wilk normality test  
data: Biomass[Year == "2021"]  
W = 0.91309, p-value = 0.05475  
> with(fish.trans, shapiro.test(Biomass[Year == "2022"]))  
Shapiro-Wilk normality test  
data: Biomass[Year == "2022"]  
W = 0.91718, p-value = 0.05799
```

ab

F test to compare two variances

data: Biomass by Year

F = 0.10639, num df = 21, denom df = 22, p-value = 2.878e-06

alternative hypothesis: true ratio of variances is not equal to 1

95 percent confidence interval:

0.04483863 0.25467745

sample estimates:

ratio of variances

0.1063915

11. Untuk melakukan uji-t, dilakukan uji *Shapiro-test* dan *F-test* untuk memenuhi asumsi uji-t.
- (a) *Shapiro-test* menunjukkan data biomassa ikan karang herbivora  $p>0,05$ , sehingga data memenuhi asumsi normalitas uji-t.
- (b) *F-test* menunjukkan data biomassa ikan karang herbivora  $p<0,05$ , sehingga data tidak memenuhi asumsi homogenitas uji-t.

```
> with(fish.trans, shapiro.test(log.2bio[Year == "2021"]))  
Shapiro-Wilk normality test  
data: log.2bio[Year == "2021"]  
W = 0.8881, p-value = 0.01731  
> with(fish.trans, shapiro.test(log.2bio[Year == "2022"]))  
Shapiro-Wilk normality test  
data: log.2bio[Year == "2022"]  
W = 0.91226, p-value = 0.04559
```

ab

F test to compare two variances

data: log.2bio by Year

F = 1.2488, num df = 21, denom df = 22, p-value = 0.6085

alternative hypothesis: true ratio of variances is not equal to 1

95 percent confidence interval:

0.5262889 2.9892505

sample estimates:

ratio of variances

1.24876

12. Data biomassa ikan karang herbivora tidak memenuhi asumsi homogenitas untuk uji-t, sehingga dilakukan *treatment* data dengan mentransformasi data menjadi log2.
- (a) *Shapiro-test* menunjukkan data biomassa ikan karang herbivora  $p<0,05$ , sehingga data tidak memenuhi asumsi normalitas uji-t.

## Lampiran 2. Lanjutan

- (b) *F-test* menunjukkan data biomassa ikan karang herbivora  $p>0,05$ , sehingga data memenuhi asumsi homogenitas uji-t. Namun uji-t tidak dapat dilanjutkan karena data tidak memenuhi asumsi normalitas.

```
> with(fish.trans, shapiro.test(log.10.bio[Year == "2021"]))  
Shapiro-Wilk normality test  
data: log.10.bio[Year == "2021"]  
W = 0.8881, p-value = 0.01731  
> with(fish.trans, shapiro.test(log.10.bio[Year == "2022"]))  
Shapiro-Wilk normality test  
data: log.10.bio[Year == "2022"]  
W = 0.91226, p-value = 0.04559
```

a

```
F test to compare two variances  
data: log.10.bio by Year  
F = 1.2488, num df = 21, denom df = 22, p-value = 0.6085  
alternative hypothesis: true ratio of variances is not equal to 1  
95 percent confidence interval:  
0.5262889 2.9892505  
sample estimates:  
ratio of variances  
1.24876
```

b

13. Data biomassa ikan karang herbivora tidak memenuhi asumsi homogenitas untuk uji-t, sehingga dilakukan *treatment* data dengan mentransformasi data menjadi  $\log_{10}$ .
- (a) *Shapiro-test* menunjukkan data biomassa ikan karang herbivora  $p<0,05$ , sehingga data tidak memenuhi asumsi normalitas uji-t.
- (b) *F-test* menunjukkan data biomassa ikan karang herbivora  $p>0,05$ , sehingga data memenuhi asumsi homogenitas uji-t. Namun uji-t tidak dapat dilanjutkan karena data tidak memenuhi asumsi normalitas.

```
> with(fish.trans, shapiro.test(square.root.bio[Year == "2021"]))  
Shapiro-Wilk normality test  
data: square.root.bio[Year == "2021"]  
W = 0.98997, p-value = 0.9973  
> with(fish.trans, shapiro.test(square.root.bio[Year == "2022"]))  
Shapiro-Wilk normality test  
data: square.root.bio[Year == "2022"]  
W = 0.98263, p-value = 0.9457
```

a

```
F test to compare two variances  
data: square.root.bio by Year  
F = 0.34156, num df = 21, denom df = 22, p-value = 0.01669  
alternative hypothesis: true ratio of variances is not equal to 1  
95 percent confidence interval:  
0.1439491 0.8176117  
sample estimates:  
ratio of variances  
0.3415574
```

b

14. Data biomassa ikan karang herbivora tidak memenuhi asumsi homogenitas untuk uji-t, sehingga dilakukan *treatment* data dengan mentransformasi data menjadi *square root*.
- (a) *Shapiro-test* menunjukkan data biomassa ikan karang herbivora  $p>0,05$ , sehingga data memenuhi asumsi normalitas uji-t.
- (b) *F-test* menunjukkan data biomassa ikan karang herbivora  $p<0,05$ , sehingga data tidak memenuhi asumsi homogenitas uji-t.

## Lampiran 2. Lanjutan

```
> with(fish.trans, shapiro.test(fourth.bio[Year == "2021"]))  
Shapiro-Wilk normality test  
data: fourth.bio[Year == "2021"]  
W = 0.97222, p-value = 0.7619  
> with(fish.trans, shapiro.test(fourth.bio[Year == "2022"]))  
Shapiro-Wilk normality test  
data: fourth.bio[Year == "2022"]  
W = 0.97362, p-value = 0.7747
```

a

```
F test to compare two variances  
data: fourth.bio by Year  
F = 0.63831, num df = 21, denom df = 22, p-value = 0.3081  
alternative hypothesis: true ratio of variances is not equal to 1  
95 percent confidence interval:  
0.2690171 1.5279812  
sample estimates:  
ratio of variances  
0.6383144
```

b

15. Data biomassa ikan karang herbivora tidak memenuhi asumsi homogenitas untuk uji-t, sehingga dilakukan treatment data dengan mentransformasi data menjadi *fourth root*.
- (a) *Shapiro-test* menunjukkan data biomassa ikan karang herbivora  $p>0,05$ , sehingga data memenuhi asumsi normalitas uji-t.
- (b) *F-test* menunjukkan data biomassa ikan karang herbivora  $p>0,05$ , sehingga data memenuhi asumsi homogenitas uji-t. Selanjutnya uji-t dilakukan dengan menggunakan transformasi data *fourth root*.

Lampiran 3. Hasil uji-t komposisi jenis, kelimpahan, dan biomassa ikan karang herbivora

1. Hasil uji-t komposisi jenis ikan karang herbivora.

```
> species.comp %>%
+   group_by(Year) %>%
+   get_summary_stats(sum.sp, type = "mean_sd")
# A tibble: 2 x 5
  Year variable     n   mean    sd
  <fct> <fct> <dbl> <dbl> <dbl>
1 2021  sum.sp     30  29.6  49.7
2 2022  sum.sp     36  61.8 103.

Two Sample t-test

data: sp.logg2 by Year
t = -1.0646, df = 64, p-value = 0.291
alternative hypothesis: true difference in means between group 2021 and group 2022 is not equal to 0
95 percent confidence interval:
-1.6246647  0.4950306
sample estimates:
mean in group 2021 mean in group 2022
            3.676143           4.240960
```

2. Hasil uji-t kelimpahan ikan karang herbivora.

```
> fish.trans %>%
+   group_by(Year) %>%
+   get_summary_stats(fourth.abundance, type = "mean_sd")
# A tibble: 2 x 5
  Year variable     n   mean    sd
  <fct> <fct> <dbl> <dbl> <dbl>
1 2021  fourth.abundance 22  0.59  0.144
2 2022  fourth.abundance 23  0.737 0.169

Two Sample t-test

data: fourth.abundance by Year
t = -3.1562, df = 43, p-value = 0.002918
alternative hypothesis: true difference in means between group 2021 and group 2022 is not equal to 0
95 percent confidence interval:
-0.24223531 -0.05336194
sample estimates:
mean in group 2021 mean in group 2022
          0.5895138          0.7373124
```

3. Hasil uji-t biomassa ikan karang herbivora.

```
> fish.trans %>%
+   group_by(Year) %>%
+   get_summary_stats(fourth.bio, type = "mean_sd")
# A tibble: 2 x 5
  Year variable     n   mean    sd
  <fct> <fct> <dbl> <dbl> <dbl>
1 2021  fourth.bio   22  3.21  0.738
2 2022  fourth.bio   23  4.34  0.924
```

### Lampiran 3. Lanjutan

```
Two Sample t-test
data: fourth.bio by Year
t = -4.4863, df = 43, p-value = 5.335e-05
alternative hypothesis: true difference in means between group 2021 and group 2022 is not equal to 0
95 percent confidence interval:
-1.6259854 -0.6174984
sample estimates:
mean in group 2021 mean in group 2022
3.213002      4.334744
```

Lampiran 4. Kelimpahan ikan karang herbivora berdasarkan site pengamatan

Famili	Spesies	LL		SA		BL		BO		BA		LU		KS		KP		Kelimpahan individu
		Tahun		Tahun		Tahun		Tahun		Tahun		Tahun		Tahun		Tahun		
		2021	2022	2021	2022	2021	2022	2021	2022	2021	2022	2021	2022	2021	2022	2021	2022	
Acanthuridae	<i>Acanthurus auranticavus</i>	1				5	7	4	3			4			3	7	2	36
	<i>Acanthurus nigricans</i>							2							2			4
	<i>Acanthurus pyroferus</i>														2		14	129
	<i>Acanthurus thompsoni</i>														2		3	5
	<i>Ctenochaetus striatus</i>	14	55	22	14	24	67	7	22	17	37	26	27	140	242		714	
	<i>Ctenochaetus toominiensis</i>																3	3
	<i>Naso annulatus</i>			2		1	5			4	2	5	2	1				22
	<i>Naso lituratus</i>					2	2	12				16			8	7		47
	<i>Naso unicornis</i>										6							6
	<i>Naso vlamingii</i>	3															23	26
Scaridae	<i>Zebrasoma scopas</i>					5	4	7		7	2	3			2	60	311	401
	<i>Zebrasoma velifer</i>						1			1		4			2		8	
	<i>Zebrasoma veliferum</i>			2							1		62		3		68	
	<i>Cetoscarus bicolor</i>					1				3			1		2		7	
	<i>Chlorurus bleekeri</i>	15	30	23	78	18	32	4		23	27	15	33	16	10	75		399
	<i>Chlorurus microrhinos</i>								6								6	
	<i>Chlorurus spilurus</i>	3	1	3	46	5	20			1	18	14	3	21	12		147	
	<i>Hippocaricus longiceps</i>										3						3	
	<i>Scarus chameleon</i>						1		3					2			6	
	<i>Scarus dimidiatus</i>	1		5	6	1	3	1	8	1	6	2	8		29		71	
	<i>Scarus flavipectoralis</i>	4	37	4	36	13	62		22	3	24	11	17	12	6		251	
	<i>Scarus forsteni</i>	2				1							2				5	
	<i>Scarus frenatus</i>		3												2		5	
	<i>Scarus ghobban</i>			2	25	1						5					33	
	<i>Scarus hypselopterus</i>							2					2		2		6	
	<i>Scarus niger</i>	2	13	14	3	2	13		6	4	21	4	37	8	10		137	
Siganidae	<i>Scarus oviceps</i>					3						1		2			6	
	<i>Scarus quoyi</i>	4	8	32	6	15	1	12				3		2	3		86	
	<i>Scarus rivulatus</i>	2	34	2	30		3	6									77	
	<i>Scarus scaber</i>		1		1				1		4			9			16	
	<i>Scarus schlegeli</i>					2					2						2	
	<i>Scarus spinus</i>									2			3				5	
	<i>Scarus tricolor</i>	1	2					1	1		2	1					8	
Siganidae	<i>Siganus canaliculatus</i>	12										2					14	
	<i>Siganus corallinus</i>	2				2				3	12	9	17		2		47	
	<i>Siganus guttatus</i>					1											1	
	<i>Siganus puelloides</i>										6	3	11	3	3	3		29
	<i>Siganus punctatissimus</i>	2	4		13	4			8	1	6	4	7		3		52	
	<i>Siganus punctatus</i>				6		2										8	
	<i>Siganus virgatus</i>	5	10	8	2	70	7	10		2	7	10	2	5	1	2		141
	<i>Siganus vulpinus</i>	2	2			3	2		2	14	13	6	8	2	5			59
	Jumlah jenis	1	2	12	14	12	18	20	17	8	18	17	20	22	20	17	21	3112

Lampiran 5. Komposisi jenis ikan karang herbivora

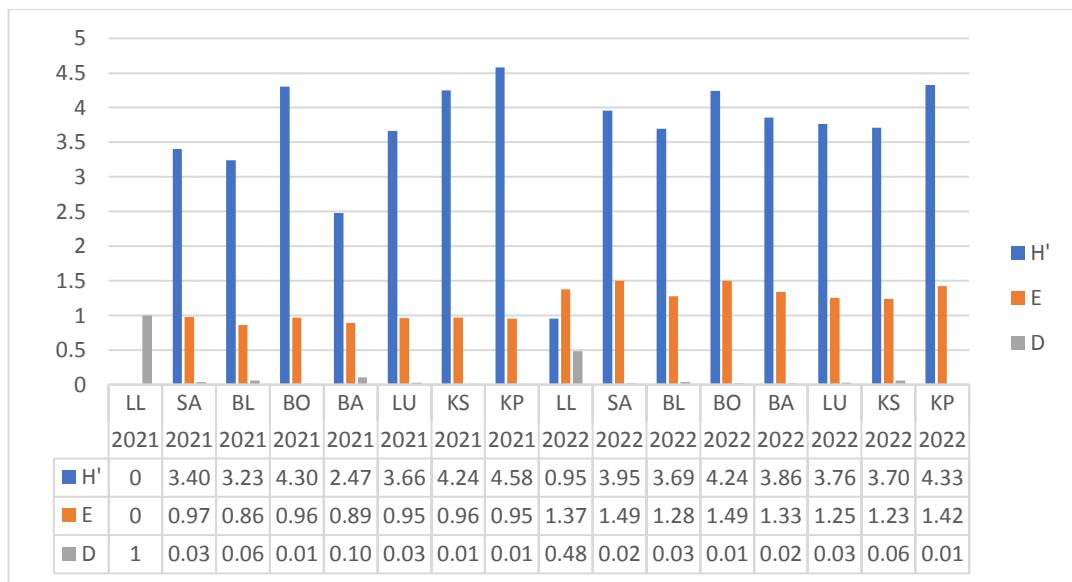
Famili	Jumlah jenis		Total jenis
	2021	2022	
Acanthuridae	9	11	12
Scaridae	16	17	20
Siganidae	5	8	8
<b>Total</b>	<b>30</b>	<b>36</b>	<b>40</b>

Lampiran 6. Nilai standar error (SE) kelimpahan dan biomassa per famili dan tahun ikan karang herbivora

	Year	Fish_family	Av.Biomass	Av.BiomassSE	Av.Abandance	Av.AbandanceSE
1	2021	Acanthuridae	36.82834	18.791013	0.06575000	0.05594439
2	2021	Scaridae	86.40641	32.909238	0.07638095	0.02392050
3	2021	Siganidae	21.61637	7.439914	0.02444444	0.00841398
4	2022	Acanthuridae	146.30069	111.522367	0.20990476	0.19183731
5	2022	Scaridae	274.86492	84.073727	0.16666667	0.04054931
6	2022	Siganidae	68.51517	21.687877	0.05241667	0.03108407

	Year	Biomass	BiomassSE	Abundance	AbundanceSE
1	2021	48.28371	19.56064	0.05552513	0.01584047
2	2022	163.22692	60.16624	0.14299603	0.04697818

## Lampiran 7. Indeks ekologi per site



Lampiran 8. Dokumentasi kegiatan di lapangan: a. persiapan menuju lokasi penelitian; b. persiapan peralatan pengambilan data; c;d. pengambilan data ikan karang herbivora; d. pengambilan peralatan bawah air setelah pendataan; e. tim pengambilan data tahun 2021; f;g. tim pengambilan data tahun 2022

