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LAMPIRAN

Nama Sapi	Parameter		
	Fragmentasi DNA (%)	**TAU (%)**	**MPU (%)**
Singo	3.17±1.63	92.81±2.18	95.21±1.47
Sinyo	4.27±2.20	91.74±2.51	93.74±2.75
Hercules	3.89±0.85	87.10±3.62	88.85±2.24
Bima	4.21±2.02	88.57±1.90	90.26±4.62
Arjuna	3.68±2.20	89.33±2.36	90.36±1.46
Rewa	3.45±1.41	91.00±1.54	89.60±2.84
Lewa	3.56±1.68	89.81±1.27	90.02±1.86
Rowa	2.07±1.22	91.34±3.63	93.88±2.62
Dewa	3.62±1.36	93.65±2.62	91.55±3.24
Maiwa	4.03±1.75	92.04±3.34	91.75±4.36
Rata-rata	**3.59±1.67**	**90.74±3.09**	**91.52±3.37**
P-*Value*	**0.566**	**0.002**	**0.005**

	Nama sapi									
Parameter	Singo	Sinyo	Hercules	Bima	Arjuna	Rewa	Lewa	Rowa	Dewa	Maiwa
Fragmentasi DNA (%)	3.17±1.63	4.27±2.20	3.89±0.85	4.21±2.02	3.68±2.20	3.45±1.41	3.56±1.68	2.07±1.22	3.62±1.36	4.03±1.75
TAU (%)	92.81±2.18	91.74±2.51	87.10±3.62	88.57±1.90	89.33±2.36	91.00±1.54	89.81±1.27	91.34±3.63	93.65±2.62	92.04±3.34
MPU(%)	95.21±1.	93.74±2.75	88.85±2.24	90.26±4.62	90.36±1.46	89.60±2.84	90.02±1.86	93.88±2.62	91.55±3.24	91.75±36

OUTPUT SPSS
FRAGMENTASI DNA

Deskriptif

Descriptives

Fragmentasi DNA

	N	Mean	Std. Deviation	Std. Error	95% Confidence Interval for Mean		Minimum	Maximum
					Lower Bound	Upper Bound		
Singo	6	3.1667	1.63299	0.66667	1.4529	4.8804	1.00	5.00
Sinyo	6	4.2667	2.19697	0.89691	1.9611	6.5722	2.00	7.00
Hercules	6	3.8867	0.84946	0.34679	2.9952	4.7781	2.40	5.00
Bima	6	4.2117	2.02404	0.82631	2.0876	6.3358	2.00	7.30
Arjuna	6	3.6783	2.19749	0.89712	1.3722	5.9845	1.00	7.50
Rewa	6	3.4500	1.41138	0.57619	1.9688	4.9312	1.00	4.65
Lewa	6	3.5550	1.68296	0.68707	1.7888	5.3212	1.00	5.00
Rowa	6	2.0667	1.21596	0.49641	0.7906	3.3427	0.98	3.90
Dewa	6	3.6167	1.35597	0.55357	2.1937	5.0397	2.00	5.65
Maiwa	6	4.0317	1.75317	0.71573	2.1918	5.8715	1.50	6.54
Total	60	3.5930	1.66580	0.21505	3.1627	4.0233	0.98	7.50

Analysis of Variance

ANOVA

Fragmentasi DNA

	Sum of Squares	df	Mean Square	F	Sig.
Between Groups	21.939	9	2.438	0.860	0.566
Within Groups	141.779	50	2.836		
Total	163.718	59			

TUDUNG AKROSOM UTUH

Deskriptif

Descriptives

TAU

	N	Mean	Std. Deviation	Std. Error	95% Confidence Interval for Mean		Minimum	Maximum
					Lower Bound	Upper Bound		
Singo	6	92.8050	2.17673	0.88865	90.5207	95.0893	90.14	95.21
Sinyo	6	91.7367	2.51224	1.02562	89.1002	94.3731	87.34	94.32
Hercules	6	87.0983	3.62123	1.47836	83.2981	90.8986	82.30	92.32
Bima	6	88.5667	1.90053	0.77589	86.5722	90.5612	85.43	90.75
Arjuna	6	89.3333	2.36083	0.96380	86.8558	91.8109	84.78	91.65
Rewa	6	90.9983	1.54327	0.63004	89.3788	92.6179	89.98	94.02
Lewa	6	89.8100	1.27317	0.51977	88.4739	91.1461	87.34	91.00
Rowa	6	91.3467	3.62630	1.48043	87.5411	95.1522	84.78	94.32
Dewa	6	93.6450	2.61612	1.06803	90.8995	96.3905	89.65	96.08
Maiwa	6	92.0367	3.33712	1.36237	88.5346	95.5388	86.43	95.45
Total	60	90.7377	3.08703	0.39853	89.9402	91.5351	82.30	96.08

Analysis of Variance

ANOVA

TAU

	Sum of Squares	df	Mean Square	F	Sig.
Between Groups	219.848	9	24.428	3.567	0.002
Within Groups	342.408	50	6.848		
Total	562.256	59			

Duncan's Test

TAU

Duncan[a]

		Subset for alpha = 0.05			
Sapi Bali	N	1	2	3	4
Hercules	6	87.0983			
Bima	6	88.5667	88.5667		
Arjuna	6	89.3333	89.3333	89.3333	
Lewa	6	89.8100	89.8100	89.8100	
Rewa	6		90.9983	90.9983	90.9983
Rowa	6		91.3467	91.3467	91.3467
[illegible]	6		91.7367	91.7367	91.7367
[illegible]	6		92.0367	92.0367	92.0367
[illegible]	6			92.8050	92.8050
[illegible]	6				93.6450
		0.107	0.050	0.050	0.131

Means for groups in homogeneous subsets are displayed.
a. Uses Harmonic Mean Sample Size = 6.000.

MEMBRAN PLASMA UTUH

Deskriftif

MPU								
	N	Mean	Std. Deviation	Std. Error	95% Confidence Interval for Mean		Minimum	Maximum
					Lower Bound	Upper Bound		
Singo	6	95.2050	1.46603	0.59850	93.6665	96.7435	92.76	96.75
Sinyo	6	93.7400	2.75433	1.12445	90.8495	96.6305	89.98	96.65
Hercules	6	88.8533	2.24343	0.91588	86.4990	91.2077	86.43	92.13
Bima	6	90.2550	4.61364	1.88351	85.4133	95.0967	85.48	96.70
Arjuna	6	90.3550	1.46231	0.59699	88.8204	91.8896	87.90	92.13
Rewa	6	89.5917	2.84477	1.16137	86.6063	92.5771	86.43	93.87
Lewa	6	90.0217	1.86394	0.76095	88.0656	91.9777	87.00	92.35
Rowa	6	93.8750	2.61593	1.06795	91.1297	96.6203	90.34	96.54
Dewa	6	91.5483	3.23715	1.32156	88.1512	94.9455	86.54	96.50
Maiwa	6	91.7450	4.36452	1.78181	87.1647	96.3253	87.24	97.87
Total	60	91.5190	3.37357	0.43553	90.6475	92.3905	85.48	97.87

Analysis of Variance

ANOVA					
MPU					
	Sum of Squares	df	Mean Square	F	Sig.
Between Groups	240.823	9	26.758	3.107	0.005
Within Groups	430.654	50	8.613		
Total	671.477	59			

Duncan's Test

MPU				
Duncan[a]				
Sapi Bali	N	Subset for alpha = 0.05		
		1	2	3
Hercules	6	88.8533		
Rewa	6	89.5917		
Lewa	6	90.0217	90.0217	
Bima	6	90.2550	90.2550	
Arjuna	6	90.3550	90.3550	
	6	91.5483	91.5483	91.5483
	6	91.7450	91.7450	91.7450
	6		93.7400	93.7400
	6		93.8750	93.8750

Singo	6			95.2050
Sig.		0.147	0.053	0.058
Means for groups in homogeneous subsets are displayed.				
a. Uses Harmonic Mean Sample Size = 6.000.				

DOKUMENTASI

Ket: *Thawing* Sampel Semen Beku

Ket: Pengujian MPU, TAU, dan Fragmentasi DNA

Ket: Pengujian MPU, TAU, Dan Fragmentasi DNA Menggunakan Mikroskop Fluoresen

Ket: Larutan AO