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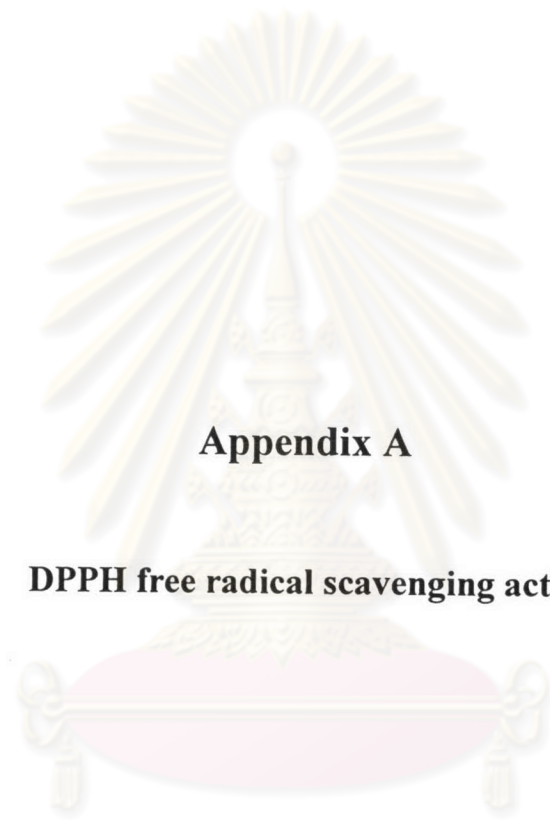
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ศูนย์วิทยทรัพยากร  
จุฬาลงกรณ์มหาวิทยาลัย



**Appendix A**

**DPPH free radical scavenging activity**

ศูนย์วิจัยทรัพยากร  
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Table A13. One-way analysis of variance on the IC<sub>50</sub> values of DPPH inhibition

ANOVA					
	Sum of squares	df	Mean square	F	Sig.
Between groups	130.852	5	26.170	145.463	0.000
Within groups	2.159	12	0.180		
Total	133.011	17			

Table A14. Multiple comparisons on the IC<sub>50</sub> values of DPPH inhibition

Tukey HSD <sup>a</sup>					
Sample	N	Subset for alpha = 0.05			
		1	2	3	4
5	3	1.747			
6	3	2.427			
4	3		3.827		
3	3			5.480	
1	3				8.253
2	3				8.777
Sig.		0.414	1.000	1.000	0.664

Means for groups in homogenous subsets are displayed.

a. Uses Harmonic Mean Sample Size = 3.000.

Where; 1 = Puag-Haad, 2 = oxyresveratrol, 3 = Trolox<sup>®</sup>, 4 = l-ascorbic acid, 5 = EGCG and 6 = pine bark extract

Table A15. One-way analysis of variance on the % inhibition at 0.5 µg/ml of DPPH test

ANOVA					
	Sum of squares	df	Mean square	F	Sig.
Between groups	366.384	5	73.277	16.121	0.000
Within groups	54.545	12	4.545		
Total	420.929	17			

Table A16. Multiple comparisons % inhibition at 0.5 µg/ml of DPPH test

Tukey HSD <sup>a</sup>					
Sample	N	Subset for alpha = 0.05			
		1	2	3	4
3	3	3.232			
2	3	5.853	5.853		
1	3	7.713	7.173		
6	3		9.957	9.957	
4	3			13.004	13.004
5	3				16.757
Sig.		0.300	0.245	0.516	0.333

Means for groups in homogenous subsets are displayed.

a. Uses Harmonic Mean Sample Size = 3.000.

Where; 1 = Puag-Haad, 2 = oxyresveratrol, 3 = Trolox<sup>®</sup>, 4 = l-ascorbic acid, 5 = EGCG and 6 = pine bark extract

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Table A17. One-way analysis of variance on the % inhibition at 1.0 µg/ml of DPPH test

ANOVA					
	Sum of squares	df	Mean square	F	Sig.
Between groups	1215.310	5	243.062	35.225	0.000
Within groups	82.803	12	6.9000		
Total	1298.113	17			

Table A18. Multiple comparisons % inhibition at 1.0 µg/ml of DPPH test

Tukey HSD <sup>a</sup>				
Sample	N	Subset for alpha = 0.05		
		1	2	3
3	3	8.163		
2	3	8.973		
1	3	10.170		
6	3		20.833	
4	3		22.653	
5	3			30.020
Sig.		0.929	0.952	1.000

Means for groups in homogenous subsets are displayed.

a. Uses Harmonic Mean Sample Size = 3.000.

Where; 1 = Puag-Haad, 2 = oxyresveratrol, 3 = Trolox<sup>®</sup>, 4 = l-ascorbic acid, 5 = EGCG and 6 = pine bark extract

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Table A19. One-way analysis of variance on the % inhibition at 2.5 µg/ml of DPPH test

ANOVA					
	Sum of squares	df	Mean square	F	Sig.
Between groups	6055.474	5	1211.095	132.806	0.000
Within groups	109.431	12	9.119		
Total	6164.905	17			

Table A20. Multiple comparisons % inhibition at 2.5 µg/ml of DPPH test

Tukey HSD <sup>a</sup>					
Sample	N	Subset for alpha = 0.05			
		1	2	3	4
2	3	17.423			
1	3	20.057			
3	3	20.337			
4	3		30.19		
6	3			52.317	
5	3				65.833
Sig.		0.837	1.000	1.000	1.000

Means for groups in homogenous subsets are displayed.

a. Uses Harmonic Mean Sample Size = 3.000.

Where; 1 = Puag-Haad, 2 = oxyresveratrol, 3 = Trolox<sup>®</sup>, 4 = l-ascorbic acid, 5 = EGCG and 6 = pine bark extract

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Table A21. One-way analysis of variance on the % inhibition at 5.0 µg/ml of DPPH test

ANOVA					
	Sum of squares	df	Mean square	F	Sig.
Between groups	9135.005	5	1827.001	373.549	0.000
Within groups	58.691	12	4.891		
Total	9193.696	17			

Table A22. Multiple comparisons % inhibition at 5.0 µg/ml of DPPH test

Tukey HSD <sup>a</sup>					
Sample	N	Subset for alpha = 0.05			
		1	2	3	4
2	3	30.223			
1	3	34.800			
3	3		45.250		
4	3			63.533	
6	3				85.387
5	3				86.163
Sig.		0.188	1.000	1.000	0.998

Means for groups in homogenous subsets are displayed.

a. Uses Harmonic Mean Sample Size = 3.000.

Where; 1 = Puag-Haad, 2 = oxyresveratrol, 3 = Trolox<sup>®</sup>, 4 = l-ascorbic acid, 5 = EGCG and 6 = pine bark extract

Table A23. One-way analysis of variance on the % inhibition at 10.0 µg/ml of DPPH test

ANOVA					
	Sum of squares	df	Mean square	F	Sig.
Between groups	3809.288	5	761.858	116.571	0.000
Within groups	78.427	12	6.536		
Total	3887.715	17			

Table A24. Multiple comparisons % inhibition at 10.0 µg/ml of DPPH test

Tukey HSD <sup>a</sup>			
Sample	N	Subset for alpha = 0.05	
		1	2
1	3	55.793	
2	3	56.163	
4	3		85.716
3	3		87.140
6	3		87.167
5	3		87.247
Sigs.		1.000	0.974

Means for groups in homogenous subsets are displayed.

a. Uses Harmonic Mean Sample Size = 3.000.

Where; 1 = Puag-Haad, 2 = oxyresveratrol, 3 = Trolox<sup>®</sup>, 4 = l-ascorbic acid, 5 = EGCG and 6 = pine bark extract

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Table A25. One-way analysis of variance on the % inhibition at 20.0 µg/ml of DPPH test

ANOVA					
	Sum of squares	df	Mean square	F	Sig.
Between groups	946.912	5	189.382	74.184	0.000
Within groups	30.634	12	2.553		
Total	977.546	17			

Table A26. Multiple comparisons % inhibition at 20.0 µg/ml of DPPH test

Tukey HSD <sup>a</sup>					
Sample	N	Subset for alpha = 0.05			
		1	2	3	4
1	3	73.330			
2	3		82.880		
5	3			87.317	
6	3			88.257	
4	3				94.057
3	3				94.787
Sig.		1.000	1.000	0.976	0.992

Means for groups in homogenous subsets are displayed.

a. Uses Harmonic Mean Sample Size = 3.000.

Where; 1 = Puag-Haad, 2 = oxyresveratrol, 3 = Trolox<sup>®</sup>, 4 = l-ascorbic acid, 5 = EGCG and 6 = pine bark extract

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Table A27. One-way analysis of variance on the % inhibition at 50.0 µg/ml of DPPH test

ANOVA					
	Sum of squares	df	Mean square	F	Sig.
Between groups	930.973	5	186.195	52.807	0.000
Within groups	42.311	12	3.526		
Total	973.284	17			

Table A28. Multiple comparisons % inhibition at 50.0 µg/ml of DPPH test

Tukey HSD <sup>a</sup>				
Sample	N	Subset for alpha = 0.05		
		1	2	3
1	3	74.717		
2	3		84.467	
5	3		87.670	
6	3		88.807	
4	3			95.443
3	3			96.083
Sig.		1.000	0.119	0.998

Means for groups in homogenous subsets are displayed.

a. Uses Harmonic Mean Sample Size = 3.000.

Where; 1 = Puag-Haad, 2 = oxyresveratrol, 3 = Trolox<sup>®</sup>, 4 = l-ascorbic acid, 5 = EGCG and 6 = pine bark extract

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Table A29. One-way analysis of variance on the % inhibition at 100.0  $\mu\text{g/ml}$  of DPPH test

ANOVA					
	Sum of squares	df	Mean square	F	Sig.
Between groups	984.667	5	196.933	61.564	0.000
Within groups	38.386	12	3.199		
Total	1023.053	17			

Table A30. Multiple comparisons % inhibition at 100.0  $\mu\text{g/ml}$  of DPPH test

Tukey HSD <sup>a</sup>					
Sample	N	Subset for alpha = 0.05			
		1	2	3	
1	3	74.167			
2	3		84.547		
6	3		87.517		
5	3		88.17		
4	3			95.527	
3	3			96.280	
Sig.		1.000	0.204	0.9948	

Means for groups in homogenous subsets are displayed.

a. Uses Harmonic Mean Sample Size = 3.000.

Where; 1 = Puag-Haad, 2 = oxyresveratrol, 3 = Trolox<sup>®</sup>, 4 = l-ascorbic acid, 5 = EGCG and 6 = pine bark extract

ศูนย์วิจัยทรัพยากร  
จุฬาลงกรณ์มหาวิทยาลัย



**Appendix B**

**Data of Superoxide Radical Scavenging Activity**

ศูนย์วิจัยทรัพยากร  
จุฬาลงกรณ์มหาวิทยาลัย

Table B1. The raw data for the absorbance and superoxide radical inhibition percentages of Puag-Haad

No.	final conc ( $\mu\text{g/ml}$ )	N1						N2						N3											
		Abs at 560 nm			% inh			Abs at 560 nm			% inh			Abs at 560 nm			% inh								
		n	Abs	Blank	Diff	$\bar{x}$	n	% inh	n	Abs	Blank	Diff	$\bar{x}$	n	% inh	n	Abs	Blank	Diff	$\bar{x}$	n	% inh			
1	0.00	1	0.507	0.006	0.497	0.516	0.00	1	0.489	0.006	0.483	0.510	0.00	1	0.494	0.007	0.487	0.491	0.00	1	0.489	0.006	0.483	0.491	0.00
		2	0.538	0.004	0.529	0.516	0.00	2	0.542	0.006	0.536	0.510	0.00	2	0.489	0.006	0.483	0.491	0.00	2	0.489	0.006	0.483	0.491	0.00
		3	0.531	0.006	0.522	0.516	0.00	3	0.517	0.005	0.512	0.510	0.00	3	0.507	0.005	0.502	0.491	0.00	3	0.507	0.005	0.502	0.491	0.00
2	5.00	1	0.507	0.009	0.498	0.504	2.33	1	0.482	0.008	0.474	0.485	4.96	1	0.481	0.008	0.473	0.473	3.60	1	0.481	0.008	0.473	0.473	3.60
		2	0.504	0.008	0.496	0.504	2.33	2	0.491	0.009	0.482	0.485	4.96	2	0.466	0.008	0.458	0.458	3.60	2	0.466	0.008	0.458	0.458	3.60
		3	0.527	0.009	0.518	0.504	2.33	3	0.508	0.009	0.499	0.485	4.96	3	0.496	0.008	0.488	0.488	3.60	3	0.496	0.008	0.488	0.488	3.60
3	10.00	1	0.461	0.010	0.451	0.454	12.02	1	0.466	0.011	0.455	0.442	13.39	1	0.443	0.010	0.433	0.433	9.51	1	0.443	0.010	0.433	0.433	9.51
		2	0.458	0.009	0.449	0.454	12.02	2	0.450	0.010	0.440	0.442	13.39	2	0.470	0.009	0.461	0.461	9.51	2	0.470	0.009	0.461	0.461	9.51
		3	0.471	0.009	0.462	0.454	12.02	3	0.440	0.009	0.431	0.442	13.39	3	0.447	0.009	0.438	0.438	9.51	3	0.447	0.009	0.438	0.438	9.51
4	25.00	1	0.369	0.021	0.348	0.365	29.20	1	0.380	0.021	0.359	0.383	25.02	1	0.368	0.022	0.346	0.346	27.11	1	0.368	0.022	0.346	0.346	27.11
		2	0.392	0.021	0.371	0.365	29.20	2	0.416	0.021	0.395	0.383	25.02	2	0.375	0.021	0.354	0.354	27.11	2	0.375	0.021	0.354	0.354	27.11
		3	0.400	0.023	0.377	0.365	29.20	3	0.417	0.023	0.394	0.383	25.02	3	0.394	0.021	0.373	0.373	27.11	3	0.394	0.021	0.373	0.373	27.11
5	50.00	1	0.297	0.084	0.213	0.232	55.06	1	0.324	0.088	0.236	0.240	53.04	1	0.274	0.078	0.196	0.196	85.60	1	0.274	0.078	0.196	0.196	85.60
		2	0.319	0.074	0.245	0.232	55.06	2	0.313	0.082	0.231	0.240	53.04	2	0.323	0.075	0.248	0.248	85.60	2	0.323	0.075	0.248	0.248	85.60
		3	0.312	0.074	0.238	0.232	55.06	3	0.328	0.076	0.252	0.240	53.04	3	0.337	0.075	0.262	0.262	85.60	3	0.337	0.075	0.262	0.262	85.60
6	100.00	1	0.201	0.167	0.034	0.050	90.31	1	0.243	0.168	0.075	0.054	89.35	1	0.230	0.164	0.066	0.066	85.60	1	0.230	0.164	0.066	0.066	85.60
		2	0.199	0.153	0.046	0.050	90.31	2	0.201	0.159	0.042	0.054	89.35	2	0.213	0.141	0.072	0.072	85.60	2	0.213	0.141	0.072	0.072	85.60
		3	0.224	0.154	0.070	0.050	90.31	3	0.215	0.169	0.046	0.054	89.35	3	0.225	0.151	0.074	0.074	85.60	3	0.225	0.151	0.074	0.074	85.60

Table B2. The raw data for the absorbance and superoxide radical inhibition percentages of oxyresveratrol

No.	final conc (µg/ml)	N1						N2						N3								
		Abs at 560 nm			% inh	$\bar{x}$	n	Abs at 560 nm			% inh	$\bar{x}$	n	Abs at 560 nm			% inh	$\bar{x}$	n			
		Abs	Blank	Diff				Abs	Blank	Diff				Abs	Blank	Diff						
1	0.00	0.568	0.066	0.559	0.00	0.540	1	0.563	0.008	0.555	0.00	0.537	2	0.545	0.007	0.538	0.00	0.537	2			
		0.550	0.007	0.540				0.508	0.006	0.502				0.545	0.008	0.537				0.545	0.008	0.537
		0.528	0.007	0.520				0.562	0.007	0.555				0.542	0.007	0.535				0.542	0.007	0.535
2	5.00	0.505	0.009	0.496	8.34	0.495	1	0.495	0.008	0.487	9.24	0.488	2	0.508	0.008	0.500	3.98	0.515	2			
		0.503	0.009	0.494				0.482	0.008	0.474				0.507	0.009	0.498				0.507	0.009	0.498
		0.502	0.008	0.494				0.510	0.008	0.502				0.557	0.009	0.548				0.557	0.009	0.548
3	10.00	0.484	0.009	0.475	13.71	0.466	1	0.447	0.008	0.439	12.64	0.469	2	0.507	0.010	0.497	10.62	0.480	2			
		0.452	0.010	0.442				0.484	0.009	0.475				0.484	0.009	0.475				0.484	0.009	0.475
		0.488	0.008	0.480				0.504	0.010	0.494				0.476	0.009	0.467				0.476	0.009	0.467
4	25.00	0.451	0.044	0.407	26.25	0.398	1	0.431	0.048	0.383	28.72	0.383	2	0.442	0.043	0.399	25.96	0.397	2			
		0.431	0.047	0.384				0.423	0.041	0.382				0.452	0.047	0.405				0.452	0.047	0.405
		0.444	0.041	0.403				0.435	0.051	0.384				0.437	0.049	0.388				0.437	0.049	0.388
5	50.00	0.407	0.199	0.208	56.52	0.235	1	0.403	0.185	0.218	57.13	0.230	2	0.401	0.182	0.219	56.77	0.232	2			
		0.434	0.189	0.245				0.431	0.179	0.252				0.422	0.199	0.223				0.422	0.199	0.223
		0.435	0.184	0.251				0.414	0.193	0.221				0.431	0.177	0.254				0.431	0.177	0.254
6	100.00	0.216	0.213	0.003	94.32	0.031	1	0.277	0.225	0.052	93.73	0.034	2	0.231	0.224	0.007	95.47	0.024	2			
		0.251	0.190	0.061				0.214	0.205	0.009				0.206	0.195	0.011				0.206	0.195	0.011
		0.233	0.205	0.028				0.268	0.228	0.040				0.263	0.208	0.055				0.263	0.208	0.055

Table B3. The raw data for the absorbance and superoxide radical inhibition percentages of Trolox®

No.	final conc (µg/ml)	N1						N2						N3					
		Abs at 560 nm			% inh	$\bar{x}$	n	Abs at 560 nm			% inh	$\bar{x}$	n	Abs at 560 nm			% inh	$\bar{x}$	n
		Abs	Blank	Diff				Abs	Blank	Diff				Abs	Blank	Diff			
1	0.00	1	0.523	0.008	0.516	0.00	0.525	1	1	0.523	0.008	0.515	0.00	0.518	2	1	0.523	0.007	0.516
		2	0.535	0.008	0.527				2	0.522	0.007	0.515				2	0.539	0.007	0.532
		3	0.538	0.007	0.531				3	0.533	0.008	0.525				3	0.537	0.008	0.529
2	5.00	1	0.410	0.007	0.403	16.96	0.436	1	1	0.468	0.006	0.462	15.11	0.440	2	1	0.414	0.009	0.405
		2	0.451	0.006	0.445				2	0.431	0.006	0.425				2	0.415	0.006	0.409
		3	0.466	0.007	0.459				3	0.441	0.008	0.433				3	0.434	0.006	0.428
3	10.00	1	0.396	0.007	0.389	30.05	0.367	1	1	0.353	0.007	0.346	32.93	0.348	2	1	0.367	0.007	0.360
		2	0.395	0.008	0.387				2	0.356	0.007	0.349				2	0.380	0.007	0.373
		3	0.332	0.007	0.325				3	0.357	0.009	0.348				3	0.359	0.007	0.352
4	25.00	1	0.278	0.008	0.270	47.20	0.277	1	1	0.285	0.006	0.279	45.68	0.282	2	1	0.274	0.006	0.268
		2	0.276	0.007	0.269				2	0.258	0.009	0.285				2	0.309	0.009	0.300
		3	0.300	0.008	0.292				3	0.290	0.009	0.281				3	0.306	0.006	0.300
5	50.00	1	0.168	0.008	0.160	69.00	0.163	1	1	0.152	0.006	0.146	68.10	0.165	2	1	0.164	0.008	0.156
		2	0.162	0.007	0.155				2	0.184	0.008	0.176				2	0.195	0.007	0.188
		3	0.180	0.007	0.173				3	0.181	0.007	0.174				3	0.195	0.007	0.188
6	100.00	1	0.041	0.007	0.034	92.69	0.038	1	1	0.039	0.008	0.031	92.93	0.037	2	1	0.044	0.007	0.037
		2	0.049	0.007	0.042				2	0.049	0.007	0.042				2	0.048	0.008	0.040
		3	0.046	0.007	0.039				3	0.045	0.008	0.037				3	0.045	0.007	0.038

Table B4. The raw data for the absorbance and superoxide radical inhibition percentages of l-ascorbic acid

No.	final conc ( $\mu\text{g/ml}$ )	N1						N2						N3					
		Abs at 560 nm			% inh	$\bar{x}$	n	Abs at 560 nm			% inh	$\bar{x}$	n	Abs at 560 nm			% inh	$\bar{x}$	n
		Abs	Blank	Diff				Abs	Blank	Diff				Abs	Blank	Diff			
1	0.00	0.494	0.005	0.487	0.00	0.491	1	0.519	0.006	0.513	0.00	0.529	2	0.490	0.004	0.486	0.00	0.494	2
		0.489	0.006	0.481				0.545	0.007	0.538				0.490	0.004	0.486			
		0.512	0.006	0.504				0.542	0.007	0.535				0.487	0.006	0.481			
2	5.00	0.477	0.007	0.470	4.76	0.467	1	0.507	0.007	0.500	4.80	0.503	2	0.460	0.006	0.454	4.93	0.469	2
		0.479	0.006	0.473				0.509	0.007	0.502				0.460	0.007	0.453			
		0.466	0.007	0.459				0.516	0.008	0.508				0.507	0.006	0.501			
3	10.00	0.431	0.007	0.424	13.38	0.425	1	0.479	0.008	0.471	14.67	0.451	1	0.421	0.009	0.412	16.00	0.415	2
		0.433	0.008	0.425				0.435	0.008	0.427				0.391	0.008	0.383			
		0.434	0.008	0.426				0.464	0.009	0.455				0.456	0.007	0.449			
4	25.00	0.326	0.008	0.318	30.91	0.339	1	0.350	0.009	0.341	34.74	0.345	1	0.328	0.008	0.320	34.03	0.326	2
		0.366	0.008	0.358				0.349	0.008	0.341				0.326	0.008	0.318			
		0.350	0.009	0.341				0.362	0.009	0.353				0.348	0.009	0.339			
5	50.00	0.220	0.011	0.209	54.96	0.221	1	0.215	0.010	0.205	54.43	0.241	2	0.181	0.011	0.170	62.53	0.185	2
		0.255	0.010	0.245				0.266	0.011	0.255				0.227	0.011	0.216			
		0.220	0.011	0.209				0.273	0.010	0.263				0.181	0.012	0.169			
6	100.00	0.253	0.015	0.238	50.20	0.244	1	0.307	0.013	0.294	43.39	0.299	1	0.267	0.014	0.253	50.91	0.242	2
		0.268	0.014	0.254				0.344	0.014	0.330				0.265	0.015	0.250			
		0.256	0.015	0.241				0.286	0.012	0.274				0.241	0.017	0.224			

Table B5. The raw data for the absorbance and superoxide radical inhibition percentages of EGCG

No.	final conc (µg/ml)	N1						N2						N3					
		Abs at 560 nm			% inh	$\bar{x}$	n	Abs at 560 nm			% inh	$\bar{x}$	n	Abs at 560 nm			% inh	$\bar{x}$	n
		Abs	Blank	Diff				Abs	Blank	Diff				Abs	Blank	Diff			
1	0.00	0.424	0.008	0.401	0.00	0.413	1	0.458	0.010	0.448	0.00	0.482	2	0.533	0.010	0.523	0.00	0.542	2
		0.410	0.009	0.387				0.507	0.010	0.497				0.537	0.008	0.529			
		0.472	0.009	0.451				0.511	0.009	0.502				0.129	0.016	0.113			
2	5.00	0.095	0.019	0.076	77.89	0.091	1	0.153	0.018	0.135	70.84	0.141	2	0.159	0.018	0.141	74.37	0.139	2
		0.107	0.020	0.087				0.161	0.024	0.137				0.181	0.018	0.163			
		0.131	0.020	0.111				0.168	0.018	0.150				0.083	0.025	0.058			
3	10.00	0.076	0.023	0.053	83.13	0.070	1	0.092	0.028	0.064	83.41	0.080	2	0.092	0.025	0.067	87.89	0.066	2
		0.081	0.023	0.058				0.112	0.024	0.088				0.095	0.023	0.072			
		0.119	0.021	0.098				0.112	0.024	0.088				0.106	0.049	0.057			
4	25.00	0.078	0.047	0.031	88.62	0.047	1	0.104	0.052	0.052	88.39	0.056	2	0.078	0.044	0.034	91.40	0.047	2
		0.089	0.045	0.044				0.111	0.042	0.069				0.092	0.042	0.050			
		0.108	0.042	0.066				0.092	0.045	0.047				0.123	0.085	0.038			
5	50.00	0.112	0.081	0.031	92.74	0.030	1	0.124	0.082	0.042	92.54	0.036	2	0.115	0.081	0.034	93.07	0.038	2
		0.114	0.086	0.028				0.114	0.078	0.036				0.123	0.083	0.040			
		0.115	0.084	0.031				0.112	0.082	0.030				0.125	0.085	0.040			
6	100.00	0.117	0.088	0.029	92.82	0.030	1	0.132	0.086	0.046	90.53	0.046	2	0.137	0.089	0.048	91.19	0.048	2
		0.118	0.090	0.028				0.121	0.081	0.040				0.137	0.089	0.048			
		0.120	0.088	0.032				0.133	0.082	0.051				0.137	0.081	0.056			



Table B6. The raw data for the absorbance and superoxide radical inhibition percentages of pine bark extract

No.	final conc ( $\mu\text{g/ml}$ )	N1						N2						N3							
		Abs at 560 nm			% inh	$\bar{x}$	n	Abs at 560 nm			% inh	$\bar{x}$	n	Abs at 560 nm			% inh	$\bar{x}$	n		
		Abs	Blank	Diff				Abs	Blank	Diff				Abs	Blank	Diff					
1	0.00	0.485	0.006	0.472	0.00	0.484	1	0.513	0.009	0.504	0.00	0.500	1	0.560	0.006	0.554	0.00	0.482	1		
		0.495	0.006	0.482				0.461	0.008	0.453				0.457	0.007	0.450				0.450	0.443
		0.510	0.006	0.498				0.553	0.009	0.544				0.450	0.007	0.443					
2	5.00	0.203	0.009	0.194	61.16	0.188	1	0.193	0.012	0.181	62.29	0.189	1	0.212	0.008	0.204	58.67	0.199	1		
		0.189	0.008	0.181				0.207	0.011	0.196				0.204	0.010	0.194					
		0.199	0.010	0.189				0.199	0.010	0.189				0.209	0.009	0.200					
3	10.00	0.070	0.013	0.057	85.33	0.071	1	0.097	0.013	0.084	82.48	0.088	1	0.108	0.014	0.094	82.38	0.085	1		
		0.080	0.013	0.067				0.101	0.013	0.088				0.097	0.014	0.083					
		0.101	0.012	0.089				0.105	0.014	0.091				0.091	0.013	0.078					
4	25.00	0.059	0.026	0.033	92.98	0.034	1	0.064	0.031	0.033	92.27	0.039	1	0.069	0.024	0.045	91.71	0.040	1		
		0.055	0.024	0.031				0.070	0.026	0.044				0.068	0.029	0.039					
		0.066	0.028	0.038				0.065	0.026	0.039				0.062	0.026	0.036					
5	50.00	0.082	0.057	0.025	94.36	0.027	1	0.095	0.052	0.043	92.74	0.036	1	0.087	0.049	0.038	93.23	0.033	1		
		0.073	0.046	0.027				0.085	0.048	0.037				0.076	0.053	0.023					
		0.073	0.043	0.030				0.077	0.048	0.029				0.085	0.048	0.037					
6	100.00	0.160	0.102	0.058	88.22	0.057	1	0.157	0.101	0.056	90.27	0.049	1	0.153	0.108	0.045	88.80	0.054	1		
		0.135	0.088	0.047				0.139	0.094	0.045				0.160	0.098	0.062					
		0.155	0.089	0.066				0.157	0.112	0.045				0.157	0.102	0.055					

Table B7. The average percentage of superoxide inhibition of Puag-Haad (Mean  $\pm$  SD)

No	conc ( $\mu\text{g/ml}$ )	% inhibition			$\bar{x}$	SD
		N1	N2	N3		
1	0.00	0.00	0.00	0.00	0.00	0.00
2	5.00	2.33	4.96	3.60	3.63	1.32
3	10.00	12.02	13.39	9.51	11.64	1.97
4	25.00	29.20	25.02	27.11	27.11	2.09
5	50.00	55.06	53.04	52.04	53.38	1.54
6	100.00	90.31	89.35	85.60	88.42	2.49

Table B8. The average percentage of superoxide inhibition of oxyresveratrol (Mean  $\pm$  SD)

No	conc ( $\mu\text{g/ml}$ )	% inhibition			$\bar{x}$	SD
		N1	N2	N3		
1	0.00	0.00	0.00	0.00	0.00	0.00
2	5.00	8.34	9.24	3.98	7.19	2.81
3	10.00	13.71	12.64	10.62	12.32	1.57
4	25.00	26.25	28.72	25.96	26.98	1.52
5	50.00	56.52	57.13	56.77	56.81	0.31
6	100.00	94.32	93.73	95.47	94.51	0.88

Table B9. The average percentage of superoxide inhibition of Trolox<sup>®</sup> (Mean  $\pm$  SD)

No	conc ( $\mu\text{g/ml}$ )	% inhibition			$\bar{x}$	SD
		N1	N2	N3		
1	0.00	0.00	0.00	0.00	0.00	0.00
2	5.00	16.96	15.11	21.24	17.77	3.14
3	10.00	30.05	32.93	31.20	31.39	1.45
4	25.00	47.20	45.68	44.96	45.95	1.14
5	50.00	69.00	68.10	66.27	67.79	1.39
6	100.00	92.69	92.93	92.71	92.78	0.13

Table B10. The average percentage of superoxide inhibition of l-ascorbic acid  
(Mean  $\pm$  SD)

No	conc ( $\mu\text{g/ml}$ )	% inhibition			$\bar{x}$	SD
		N1	N2	N3		
1	0.00	0.00	0.00	0.00	0.00	0.00
2	5.00	4.76	4.80	4.93	4.83	0.09
3	10.00	13.38	14.67	16.00	14.68	1.31
4	25.00	30.97	34.74	34.03	33.25	2.00
5	50.00	54.96	54.43	62.53	57.31	4.53
6	100.00	50.20	43.39	50.91	48.17	4.15

Table B11. The average percentage of superoxide inhibition of EGCG (Mean  $\pm$  SD)

No	conc ( $\mu\text{g/ml}$ )	% inhibition			$\bar{x}$	SD
		N1	N2	N3		
1	0.00	0.00	0.00	0.00	0.00	0.00
2	5.00	77.89	70.84	74.37	74.37	3.53
3	10.00	83.13	83.41	87.89	84.81	2.67
4	25.00	88.62	88.39	91.40	89.47	1.68
5	50.00	92.74	92.54	93.07	92.78	0.27
6	100.00	92.82	90.53	91.19	91.51	1.18

Table B12. The average percentage of superoxide inhibition of pine bark extract  
(Mean  $\pm$  SD)

No	conc ( $\mu\text{g/ml}$ )	% inhibition			$\bar{x}$	SD
		N1	N2	N3		
1	0.00	0.00	0.00	0.00	0.00	0.00
2	5.00	61.16	62.29	58.67	60.71	1.85
3	10.00	85.33	84.48	82.38	84.06	1.52
4	25.00	92.98	92.27	91.71	92.32	0.64
5	50.00	94.36	92.74	93.23	93.44	0.83
6	100.00	88.22	90.27	88.80	89.10	1.06

Table B13. One-way analysis of variance on the IC<sub>50</sub> values of superoxide radical inhibition

ANOVA					
	Sum of squares	df	Mean square	F	Sig.
Between groups	5893.474	5	1178.695	536.909	0.000
Within groups	26.344	12	2.195		
Total	5919.818	17			

Table B14. Multiple comparisons on the IC<sub>50</sub> values of superoxide radical inhibition

Tukey HSD <sup>a</sup>					
Sample	N	Subset for alpha = 0.05			
		1	2	3	4
5	3	2.843			
6	3	3.847			
3	3		24.897		
4	3			40.243	
2	3				44.313
1	3				44.113
Sig.		0.956	1.000	1.000	0.678

Means for groups in homogenous subsets are displayed.

a. Uses Harmonic Mean Sample Size = 3.000.

Where; 1 = Puag-Haad, 2 = oxyresveratrol, 3 = Trolox<sup>®</sup>, 4 = l-ascorbic acid, 5 = EGCG and 6 = pine bark extract

Table B15. One-way analysis of variance on the % inhibition at 5.0  $\mu\text{g/ml}$  of superoxide radical test

ANOVA					
	Sum of squares	df	Mean square	F	Sig.
Between groups	14664.44	5	2932.888	497.177	0.000
Within groups	70.789	12	5.899		
Total	1435.23	17			

Table B16. Multiple comparisons % inhibition at 5.0  $\mu\text{g/ml}$  of superoxide radical test

Tukey HSD <sup>a</sup>					
Sample	N	Subset for alpha = 0.05			
		1	2	3	4
1	3	3.630			
4	3	4.830			
2	3	7.187			
3	3		17.770		
6	3			60.707	
5	3				74.367
Sig.		0.504	1.000	1.000	1.000

Means for groups in homogenous subsets are displayed.

a. Uses Harmonic Mean Sample Size = 3.000.

Where; 1 = Puag-Haad, 2 = oxyresveratrol, 3 = Trolox<sup>®</sup>, 4 = l-ascorbic acid, 5 = EGCG and 6 = pine bark extract

Table B17. One-way analysis of variance on the % inhibition at 10.0 µg/ml of superoxide radical test

ANOVA					
	Sum of squares	df	Mean square	F	Sig.
Between groups	18703.84	5	3740.767	1145.587	0.000
Within groups	39.184	12	3.265		
Total	18743.02	17			

Table B18. Multiple comparisons % inhibition at 10.0 µg/ml of superoxide radical test

Tukey HSD <sup>a</sup>				
Sample	N	Subset for alpha = 0.05		
		1	2	3
1	3	11.640		
2	3	12.323		
4	3	14.683		
3	3		31.393	
6	3			84.063
5	3			84.810
Sig.		0.365	1.000	0.995

Means for groups in homogenous subsets are displayed.

a. Uses Harmonic Mean Sample Size = 3.000.

Where; 1 = Puag-Haad, 2 = oxyresveratrol, 3 = Trolox<sup>®</sup>, 4 = l-ascorbic acid, 5 = EGCG and 6 = pine bark extract

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Table B19. One-way analysis of variance on the % inhibition at 25.0  $\mu\text{g/ml}$  of superoxide radical test

ANOVA					
	Sum of squares	df	Mean square	F	Sig.
Between groups	13986.43	5	2797.285	1104.068	0.000
Within groups	30.403	12	2.534		
Total	14016.83	17			

Table B20. Multiple comparisons % inhibition at 25.0  $\mu\text{g/ml}$  of superoxide radical test

Tukey HSD <sup>a</sup>					
Sample	N	Subset for alpha = 0.05			
		1	2	3	4
2	3	26.977			
1	3	27.110			
4	3		33.247		
3	3			45.947	
5	3				89.470
6	3				92.320
Sig.		1.000	1.000	1.000	0.308

Means for groups in homogenous subsets are displayed.

a. Uses Harmonic Mean Sample Size = 3.000.

Where; 1 = Puag-Haad, 2 = oxyresveratrol, 3 = Trolox<sup>®</sup>, 4 = l-ascorbic acid, 5 = EGCG and 6 = pine bark extract

Table B21. One-way analysis of variance on the % inhibition at 50.0 µg/ml of superoxide radical test

ANOVA					
	Sum of squares	df	Mean square	F	Sig.
Between groups	5053.750	5	1010.750	236.059	0.000
Within groups	51.381	12	4.282		
Total	5105.131	17			

Table B22. Multiple comparisons % inhibition at 50.0 µg/ml of superoxide radical test

Tukey HSD <sup>a</sup>				
Sample	N	Subset for alpha = 0.05		
		1	2	3
1	3	53.380		
2	3	56.807		
4	3	57.307		
3	3		67.790	
5	3			92.783
6	3			93.443
Sig.		0.257	1.000	0.996

Means for groups in homogenous subsets are displayed.

a. Uses Harmonic Mean Sample Size = 3.000.

Where; 1 = Puag-Haad, 2 = oxyresveratrol, 3 = Trolox<sup>®</sup>, 4 = l-ascorbic acid, 5 = EGCG and 6 = pine bark extract



Table B23. One-way analysis of variance on the % inhibition at 100.0 µg/ml of superoxide radical test

ANOVA					
	Sum of squares	df	Mean square	F	Sig.
Between groups	4720.116	5	944.023	211.823	0.000
Within groups	53.480	12	4.457		
Total	4773.596	17			

Table B24. Multiple comparisons % inhibition at 100.0 µg/ml of superoxide radical test

Tukey HSD <sup>a</sup>				
Sample	N	Subset for alpha = 0.05		
		1	2	3
4	3	48.167		
1	3		88.420	
6	3		89.097	89.097
5	3		91.513	91.513
3	3		92.777	92.777
2	3			94.507
Sig.		1.000	0.190	0.072

Means for groups in homogenous subsets are displayed.

a. Uses Harmonic Mean Sample Size = 3.000.

Where; 1 = Puag-Haad, 2 = oxyresveratrol, 3 = Trolox<sup>®</sup>, 4 = l-ascorbic acid, 5 = EGCG and 6 = pine bark extract



**Appendix C**

**Data of Hydroxyl Radical Scavenging Activity**

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Table C1. The raw data for the absorbance and hydroxyl radical inhibition percentages of Puag-Haad

No	final conc (µg/ml)	N1						N2						N3											
		Abs at 416 nm			% inh			Abs at 416 nm			% inh			Abs at 416 nm			% inh								
		n	Abs	Blank	Diff	Av	n	Abs	Blank	Diff	Av	n	Abs	Blank	Diff	Av	n	Abs	Blank	Diff	Av				
1	0.00	1	0.490	0.000	0.490		1	0.491	0.000	0.491		1	0.596	0.000	0.596		1	0.596	0.000	0.596		1	0.596	0.000	0.596
		2	0.504	0.000	0.504	0.492	0.00	2	0.463	0.000	0.463	0.475	0.00	2	0.553	0.000	0.553	0.567	0.00	2	0.553	0.000	0.553	0.567	0.00
		3	0.481	0.000	0.481			3	0.472	0.000	0.472			3	0.551	0.000	0.551			3	0.551	0.000	0.551		
2	5.00	1	0.352	0.007	0.345		1	0.368	0.007	0.361		1	0.393	0.007	0.386		1	0.393	0.007	0.386		1	0.393	0.007	0.386
		2	0.341	0.007	0.334	0.344	29.97	2	0.343	0.007	0.336	0.345	27.42	2	0.378	0.007	0.371	0.385	32.00	2	0.378	0.007	0.371	0.385	32.00
		3	0.361	0.007	0.354			3	0.345	0.007	0.338			3	0.406	0.007	0.399			3	0.406	0.007	0.399		
3	10.00	1	0.252	0.008	0.244		1	0.233	0.008	0.225		1	0.263	0.009	0.254		1	0.263	0.009	0.254		1	0.263	0.009	0.254
		2	0.244	0.008	0.236	0.238	51.66	2	0.249	0.008	0.241	0.234	50.70	2	0.264	0.009	0.255	0.259	54.35	2	0.264	0.009	0.255	0.259	54.35
		3	0.241	0.008	0.233			3	0.245	0.008	0.237			3	0.276	0.009	0.267			3	0.276	0.009	0.267		
4	20.00	1	0.153	0.012	0.141		1	0.155	0.012	0.143		1	0.165	0.012	0.153		1	0.165	0.012	0.153		1	0.165	0.012	0.153
		2	0.156	0.012	0.144	0.145	70.44	2	0.163	0.012	0.151	0.149	68.72	2	0.168	0.012	0.156	0.152	73.12	2	0.168	0.012	0.156	0.152	73.12
		3	0.163	0.012	0.151			3	0.164	0.012	0.152			3	0.160	0.012	0.148			3	0.160	0.012	0.148		
5	50.00	1	0.108	0.018	0.090		1	0.068	0.018	0.050		1	0.098	0.018	0.080		1	0.098	0.018	0.080		1	0.098	0.018	0.080
		2	0.090	0.018	0.072	0.078	84.20	2	0.072	0.018	0.054	0.053	84.64	2	0.103	0.018	0.085	0.080	85.82	2	0.103	0.018	0.085	0.080	85.82
		3	0.089	0.018	0.071			3	0.073	0.018	0.055			3	0.094	0.018	0.076			3	0.094	0.018	0.076		
6	100.00	1	0.089	0.033	0.056		1	0.074	0.033	0.041		1	0.085	0.033	0.052		1	0.085	0.033	0.052		1	0.085	0.033	0.052
		2	0.091	0.033	0.058	0.058	88.20	2	0.075	0.033	0.042	0.040	91.58	2	0.089	0.033	0.056	0.054	90.41	2	0.089	0.033	0.056	0.054	90.41
		3	0.093	0.033	0.060			3	0.070	0.033	0.037			3	0.088	0.033	0.055			3	0.088	0.033	0.055		

Table C2. The raw data for the absorbance and hydroxyl radical inhibition percentages of oxyresveratrol

No	final conc (µg/ml)	N1						N2						N3					
		Abs at 416 nm			% inh			Abs at 416 nm			% inh			Abs at 416nm			% inh		
		Abs	Blank	Diff	AV	n	% inh	Abs	Blank	Diff	AV	n	% inh	Abs	Blank	Diff	AV	n	% inh
1	0.00	0.463	0.000	0.459	0.450	1	0.00	0.466	0.000	0.466	0.463	1	0.00	0.595	0.000	0.595	0.601	1	0.00
		0.445	0.000	0.441		2		0.458	0.000	0.458		2		0.585	0.000	0.585		2	
		0.455	0.000	0.451		3		0.465	0.000	0.465		3		0.622	0.000	0.622		3	
2	5.00	0.271	0.004	0.267	0.270	1	40.04	0.308	0.004	0.304	0.296	1	36.00	0.417	0.004	0.413	0.389	1	35.29
		0.282	0.004	0.278		2		0.288	0.004	0.284		2		0.378	0.004	0.374		2	
		0.269	0.004	0.265		3		0.305	0.004	0.301		3		0.383	0.004	0.379		3	
3	10.00	0.168	0.004	0.164	0.163	1	63.88	0.186	0.004	0.182	0.178	1	61.56	0.264	0.004	0.260	0.251	1	58.21
		0.163	0.004	0.159		2		0.178	0.004	0.174		2		0.258	0.004	0.254		2	
		0.169	0.004	0.165		3		0.182	0.004	0.178		3		0.243	0.004	0.239		3	
4	20.00	0.103	0.005	0.098	0.095	1	78.90	0.108	0.005	0.103	0.108	1	76.60	0.152	0.005	0.147	0.142	1	76.36
		0.104	0.005	0.099		2		0.114	0.005	0.109		2		0.146	0.005	0.141		2	
		0.093	0.005	0.088		3		0.118	0.005	0.113		3		0.143	0.005	0.138		3	
5	50.00	0.054	0.007	0.047	0.047	1	89.56	0.064	0.006	0.058	0.058	1	87.54	0.073	0.006	0.067	0.070	1	88.35
		0.054	0.007	0.047		2		0.063	0.006	0.057		2		0.080	0.006	0.074		2	
		0.054	0.007	0.047		3		0.064	0.006	0.058		3		0.075	0.006	0.069		3	
6	100.00	0.052	0.010	0.042	0.037	1	91.78	0.058	0.010	0.048	0.046	1	90.06	0.067	0.009	0.058	0.059	1	90.18
		0.044	0.010	0.034		2		0.060	0.010	0.050		2		0.068	0.009	0.059		2	
		0.045	0.010	0.035		3		0.050	0.010	0.040		3		0.069	0.009	0.060		3	

Table C3. The raw data for the absorbance and hydroxyl radical inhibition percentages of Trolox®

No	final conc (µg/ml)	N1						N2						N3					
		Abs at 416 nm			% inh			Abs at 416 nm			% inh			Abs at 416 nm			% inh		
		Abs	Blank	Diff	Avg	n	% inh	Abs	Blank	Diff	Avg	n	% inh	Abs	Blank	Diff	Avg	n	% inh
1	0.00	0.440	0.000	0.440	0.454	1	0.00	0.403	0.000	0.403	0.405	1	0.00	0.424	0.000	0.424	0.441	1	0.00
		0.471	0.000	0.471		2	0.00	0.400	0.000	0.400		2	0.00	0.455	0.000	0.455		2	0.00
		0.451	0.000	0.451		3		0.413	0.000	0.413		3		0.445	0.000	0.445		3	
2	0.50	0.386	0.000	0.386	0.383	1	15.57	0.353	0.000	0.353	0.349	1	13.90	0.378	0.000	0.378	0.371	1	16.01
		0.375	0.000	0.375		2		0.349	0.000	0.349		2		0.364	0.000	0.364		2	
		0.389	0.000	0.389		3		0.345	0.000	0.345		3		0.370	0.000	0.370		3	
3	1.00	0.339	0.000	0.339	0.330	1	27.24	0.320	0.000	0.320	0.316	1	22.12	0.339	0.000	0.339	0.331	1	24.92
		0.330	0.000	0.330		2		0.319	0.000	0.319		2		0.316	0.000	0.316		2	
		0.322	0.000	0.322		3		0.308	0.000	0.308		3		0.339	0.000	0.339		3	
4	2.00	0.284	0.000	0.284	0.270	1	40.60	0.247	0.001	0.246	0.237	1	41.45	0.241	0.000	0.241	0.249	1	43.66
		0.272	0.000	0.272		2		0.249	0.001	0.248		2		0.257	0.000	0.257		2	
		0.253	0.000	0.253		3		0.219	0.001	0.218		3		0.248	0.000	0.248		3	
5	5.00	0.090	0.002	0.088	0.094	1	79.22	0.054	0.003	0.051	0.053	1	86.92	0.044	0.003	0.041	0.050	1	88.75
		0.107	0.002	0.105		2		0.059	0.003	0.056		2		0.054	0.003	0.051		2	
		0.092	0.002	0.090		3		0.055	0.003	0.052		3		0.060	0.003	0.057		3	
6	10.00	0.023	0.003	0.020	0.020	1	95.59	0.020	0.004	0.016	0.018	1	95.48	0.015	0.003	0.012	0.012	1	97.28
		0.022	0.003	0.019		2		0.025	0.004	0.021		2		0.014	0.003	0.011		2	
		0.024	0.003	0.021		3		0.022	0.004	0.018		3		0.016	0.003	0.013		3	
7	20.00	0.017	0.005	0.012	0.012	1	97.36	0.019	0.005	0.014	0.014	1	96.46	0.013	0.005	0.008	0.008	1	98.26
		0.018	0.005	0.013		2		0.019	0.005	0.014		2		0.012	0.005	0.007		2	
		0.016	0.005	0.011		3		0.020	0.005	0.015		3		0.013	0.005	0.008		3	

Table C4. The raw data for the absorbance and hydroxyl radical inhibition percentages of l-ascorbic acid

No	final conc (µg/ml)	N1						N2						N3							
		Abs at 416 nm			% inh			Abs at 416 nm			% inh			Abs at 416 nm			% inh				
		Abs	Blank	Diff	Av	n	Abs	Blank	Diff	Av	n	Abs	Blank	Diff	Av	n	Abs	Blank	Diff	Av	n
1	0.00	0.520	0.000	0.519	0.524	0.00	0.509	0.000	0.509	0.506	0.00	0.416	0.000	0.416	0.410	0.00	0.423	0.000	0.423	0.410	0.00
		0.538	0.000	0.537			0.489	0.000	0.489			0.392	0.000	0.392			0.319	0.000	0.319		
		0.455	0.000	0.455	0.449	14.37	0.376	0.000	0.376	0.371	26.75	0.326	0.000	0.326	0.319	22.34	0.297	0.001	0.296	0.269	34.36
2	0.50	0.448	0.000	0.448			0.384	0.000	0.384	0.371	26.75	0.326	0.000	0.326	0.319	22.34	0.297	0.001	0.296	0.269	34.36
		0.444	0.000	0.444			0.352	0.000	0.352			0.311	0.000	0.311			0.265	0.001	0.264		
		0.336	0.001	0.335	0.318	39.35	0.333	0.001	0.332	0.331	34.65	0.249	0.001	0.248	0.269	34.36	0.191	0.002	0.189	0.172	58.16
3	1.00	0.317	0.001	0.316			0.341	0.001	0.340	0.331	34.65	0.340	0.001	0.340	0.269	34.36	0.265	0.001	0.264	0.269	34.36
		0.304	0.001	0.303			0.321	0.001	0.320			0.320	0.001	0.320			0.163	0.002	0.161		
		0.216	0.002	0.214	0.212	59.50	0.228	0.002	0.226	0.215	57.44	0.167	0.002	0.165	0.172	58.16	0.023	0.002	0.021	0.022	94.64
4	2.00	0.210	0.002	0.208			0.222	0.002	0.220	0.215	57.44	0.220	0.002	0.220	0.172	58.16	0.163	0.002	0.161	0.172	58.16
		0.217	0.002	0.215			0.202	0.002	0.200			0.200	0.002	0.200			0.022	0.002	0.020	0.022	94.64
		0.023	0.002	0.021	0.021	96.06	0.012	0.002	0.010	0.010	98.02	0.023	0.002	0.021	0.022	94.64	0.027	0.002	0.025	0.025	95.86
5	5.00	0.025	0.002	0.023			0.012	0.002	0.010	0.010	98.02	0.010	0.002	0.008	0.017	95.86	0.022	0.002	0.020	0.022	94.64
		0.020	0.002	0.018			0.012	0.002	0.010			0.010	0.002	0.008			0.018	0.002	0.016		
		0.013	0.002	0.011	0.011	97.84	0.009	0.002	0.007	0.007	98.62	0.018	0.002	0.016	0.017	95.86	0.027	0.002	0.025	0.025	95.86
6	10.00	0.014	0.002	0.012			0.009	0.002	0.007	0.007	98.62	0.007	0.002	0.005	0.017	95.86	0.020	0.002	0.018	0.017	95.86
		0.013	0.002	0.011			0.009	0.002	0.007			0.007	0.002	0.005			0.018	0.002	0.016		
		0.013	0.002	0.011	0.011	97.84	0.009	0.002	0.007	0.007	98.62	0.018	0.002	0.016	0.017	95.86	0.027	0.002	0.025	0.025	95.86

Table C5. The raw data for the absorbance and hydroxyl radical inhibition percentages of EGCG

No	final conc (µg/ml)	N1						N2						N3																								
		Abs at 416 nm			% inh			Abs at 416 nm			% inh			Abs at 416 nm			% inh																					
		Abs	Blank	Diff	Av	n	% inh	Abs	Blank	Diff	Av	n	% inh	Abs	Blank	Diff	Av	n	% inh																			
1	0.00	0.478	0.000	0.471		1	0.420	0.000	0.420		1	0.606	0.000	0.606		1	0.606	0.000	0.606		1	0.00	0.00	0.00		1	0.606	0.000	0.606		1	0.606	0.000	0.606		1	0.00	0.00
		0.461	0.000	0.454	0.465	2	0.00	0.489	0.000	0.489	0.460	2	0.590	0.000	0.590		2	0.590	0.000	0.590		2	0.00	0.00		2	0.590	0.000	0.590		2	0.590	0.000	0.590		2	0.00	0.00
		0.478	0.000	0.471		3		0.472	0.000	0.472		3	0.606	0.000	0.606		3	0.606	0.000	0.606		3				3	0.606	0.000	0.606		3	0.606	0.000	0.606		3		
2	5.00	0.297	0.007	0.290		1	0.255	0.007	0.248		1	0.320	0.006	0.314		1	0.320	0.006	0.314		1	46.20	46.20		1	0.320	0.006	0.314		1	0.320	0.006	0.314		1	49.17	49.17	
		0.279	0.007	0.272	0.272	2	41.62	0.261	0.007	0.254	0.248	2	0.305	0.006	0.299		2	0.305	0.006	0.299		2				2	0.305	0.006	0.299		2	0.305	0.006	0.299		2		
		0.260	0.007	0.253		3		0.248	0.007	0.241		3	0.309	0.006	0.303		3	0.309	0.006	0.303		3				3	0.309	0.006	0.303		3	0.309	0.006	0.303		3		
3	10.00	0.162	0.007	0.155		1	0.182	0.007	0.175		1	0.211	0.007	0.204		1	0.211	0.007	0.204		1	63.43	63.43		1	0.211	0.007	0.204		1	0.211	0.007	0.204		1	67.20	67.20	
		0.162	0.007	0.155	0.155	2	66.69	0.175	0.007	0.168	0.168	2	0.206	0.007	0.199		2	0.206	0.007	0.199		2				2	0.206	0.007	0.199		2	0.206	0.007	0.199		2		
		0.162	0.007	0.155		3		0.169	0.007	0.162		3	0.195	0.007	0.188		3	0.195	0.007	0.188		3				3	0.195	0.007	0.188		3	0.195	0.007	0.188		3		
4	20.00	0.131	0.007	0.124		1	0.126	0.007	0.119		1	0.145	0.007	0.138		1	0.145	0.007	0.138		1	72.27	72.27		1	0.145	0.007	0.138		1	0.145	0.007	0.138		1	76.25	76.25	
		0.132	0.007	0.125	0.124	2	73.35	0.147	0.007	0.140	0.128	2	0.146	0.007	0.139		2	0.146	0.007	0.139		2				2	0.146	0.007	0.139		2	0.146	0.007	0.139		2		
		0.130	0.007	0.123		3		0.131	0.007	0.124		3	0.158	0.007	0.151		3	0.158	0.007	0.151		3				3	0.158	0.007	0.151		3	0.158	0.007	0.151		3		
5	50.00	0.092	0.007	0.085		1	0.103	0.007	0.096		1	0.102	0.007	0.095		1	0.102	0.007	0.095		1	79.87	79.87		1	0.102	0.007	0.095		1	0.102	0.007	0.095		1	83.41	83.41	
		0.096	0.007	0.089	0.088	2	81.02	0.098	0.007	0.091	0.093	2	0.106	0.007	0.099		2	0.106	0.007	0.099		2				2	0.106	0.007	0.099		2	0.106	0.007	0.099		2		
		0.098	0.007	0.091		3		0.098	0.007	0.091		3	0.112	0.007	0.105		3	0.112	0.007	0.105		3				3	0.112	0.007	0.105		3	0.112	0.007	0.105		3		
6	100.00	0.079	0.007	0.072		1	0.089	0.007	0.082		1	0.088	0.007	0.081		1	0.088	0.007	0.081		1	82.69	82.69		1	0.088	0.007	0.081		1	0.088	0.007	0.081		1	85.29	85.29	
		0.078	0.007	0.071	0.073	2	84.38	0.087	0.007	0.080	0.080	2	0.088	0.007	0.081		2	0.088	0.007	0.081		2				2	0.088	0.007	0.081		2	0.088	0.007	0.081		2		
		0.082	0.007	0.075		3		0.084	0.007	0.077		3	0.110	0.007	0.103		3	0.110	0.007	0.103		3				3	0.110	0.007	0.103		3	0.110	0.007	0.103		3		

Table C6. The raw data for the absorbance and hydroxyl radical inhibition percentages of pine bark extract

No	final conc (µg/ml)	N1						N2						N3					
		Abs at 416 nm		% inh		n	Av	Abs at 416 nm		% inh		n	Av	Abs at 416 nm		% inh		n	Av
		Abs	Blank	Diff	Abs			Blank	Diff	Abs	Blank			Diff	Abs	Blank	Diff		
1	0.00	0.463	0.000	0.458	0.00	1	0.500	0.000	0.500	0.00	1	0.545	0.000	0.545	0.00	1	0.545	0.000	0.545
		0.471	0.000	0.466	0.00	2	0.497	0.000	0.497	0.00	2	0.536	0.000	0.536	0.00	2	0.536	0.000	0.536
		0.497	0.000	0.492	0.00	3	0.453	0.000	0.453	0.00	3	0.540	0.000	0.540	0.00	3	0.540	0.000	0.540
2	5.00	0.453	0.003	0.450	5.16	1	0.453	0.003	0.450	7.21	1	0.502	0.003	0.499	9.66	2	0.481	0.003	0.478
		0.443	0.003	0.440	5.16	2	0.450	0.003	0.447	7.21	2	0.481	0.003	0.478	9.66	3	0.490	0.003	0.487
		0.456	0.003	0.453	5.16	3	0.452	0.003	0.449	7.21	3	0.490	0.003	0.487	9.66	1	0.459	0.005	0.454
3	10.00	0.389	0.005	0.384	20.48	1	0.383	0.004	0.379	21.19	1	0.459	0.005	0.454	17.49	2	0.453	0.005	0.448
		0.377	0.005	0.372	20.48	2	0.382	0.004	0.378	21.19	2	0.453	0.005	0.448	17.49	3	0.440	0.005	0.435
		0.376	0.005	0.371	20.48	3	0.390	0.004	0.386	21.19	3	0.440	0.005	0.435	17.49	1	0.392	0.008	0.384
4	20.00	0.330	0.008	0.318	32.43	1	0.363	0.008	0.355	29.84	1	0.392	0.008	0.384	30.54	2	0.389	0.008	0.381
		0.324	0.008	0.316	32.43	2	0.352	0.008	0.344	29.84	2	0.389	0.008	0.381	30.54	3	0.369	0.008	0.361
		0.326	0.008	0.318	32.43	3	0.363	0.008	0.355	29.84	3	0.369	0.008	0.361	30.54	1	0.329	0.018	0.311
5	50.00	0.262	0.018	0.244	49.15	1	0.279	0.018	0.261	45.89	1	0.329	0.018	0.311	46.88	2	0.293	0.018	0.275
		0.255	0.018	0.237	49.15	2	0.287	0.018	0.269	45.89	2	0.293	0.018	0.275	46.88	3	0.293	0.018	0.275
		0.257	0.018	0.239	49.15	3	0.272	0.018	0.254	45.89	3	0.293	0.018	0.275	46.88	1	0.256	0.032	0.224
6	100.00	0.208	0.033	0.175	61.99	1	0.225	0.032	0.193	60.01	1	0.256	0.032	0.224	59.56	2	0.250	0.032	0.218
		0.216	0.033	0.183	61.99	2	0.210	0.032	0.178	60.01	2	0.250	0.032	0.218	59.56	3	0.246	0.032	0.214
		0.214	0.033	0.181	61.99	3	0.241	0.032	0.209	60.01	3	0.246	0.032	0.214	59.56	1	0.251	0.069	0.182
7	200.00	0.212	0.070	0.142	71.59	1	0.204	0.070	0.134	70.78	1	0.233	0.069	0.164	68.93	2	0.233	0.069	0.164
		0.209	0.070	0.139	71.59	2	0.211	0.070	0.141	70.78	2	0.233	0.069	0.164	68.93	3	0.227	0.069	0.158
		0.191	0.070	0.121	71.59	3	0.218	0.070	0.148	70.78	3	0.227	0.069	0.158	68.93	1	0.233	0.069	0.164



Table C7. The average percentage of hydroxyl radical inhibition of Puag-Haad  
(Mean  $\pm$  SD)

No.	final conc ( $\mu\text{g/ml}$ )	% inhibition			Puag-Haad	
		N1	N2	N3	$\bar{x}$	SD
1	0.00	0.00	0.00	0.00	0.00	0.00
2	5.00	29.97	27.42	32.00	29.80	2.29
3	10.00	51.67	50.70	54.35	52.24	1.89
4	20.00	70.44	68.72	73.12	70.76	2.22
5	50.00	84.20	84.64	85.82	84.89	0.84
6	100.00	88.20	91.58	90.41	90.06	1.72

Table C8. The average percentage of hydroxyl radical inhibition of oxyresveratrol  
(Mean  $\pm$  SD)

No.	final conc ( $\mu\text{g/ml}$ )	% inhibition			oxyresveratrol	
		N1	N2	N3	$\bar{x}$	SD
1	0.00	0.00	0.00	0.00	0.00	0.00
2	5.00	40.04	36.00	35.29	37.11	2.56
3	10.00	63.88	61.56	58.21	61.22	2.85
4	20.00	78.90	76.60	76.36	77.29	1.40
5	50.00	89.56	87.54	88.35	88.48	1.02
6	100.00	91.78	90.06	90.18	90.67	0.96

Table C9. The average percentage of hydroxyl radical inhibition of Trolox<sup>®</sup> (Mean  $\pm$  SD)

No.	final conc ( $\mu\text{g/ml}$ )	% inhibition			Trolox	
		N1	N2	N3	$\bar{x}$	SD
1	0.00	0.00	0.00	0.00	0.00	0.00
2	0.50	15.57	13.90	16.01	15.16	1.11
3	1.00	27.24	22.12	24.92	24.76	2.56
4	2.00	40.60	41.45	43.66	41.90	1.58
5	5.00	79.22	86.92	88.75	84.96	5.06
6	10.00	95.59	95.48	97.28	96.12	1.01
7	20.00	97.36	96.46	98.26	97.36	0.90

Table C10. The average percentage of hydroxyl radical inhibition of l-ascorbic acid (Mean  $\pm$  SD)

No.	final conc ( $\mu\text{g/ml}$ )	% inhibition			l-ascorbic acid	
		N1	N2	N3	$\bar{x}$	SD
1	0.00	0.00	0.00	0.00	0.00	0.00
2	0.50	14.37	26.75	22.34	21.15	6.27
3	1.00	39.35	34.65	34.36	36.12	2.80
4	2.00	59.50	57.44	58.16	58.37	1.05
5	5.00	96.06	98.02	94.64	96.24	1.70
6	10.00	97.84	98.62	95.86	97.44	1.42

Table C11. The average percentage of hydroxyl radical inhibition of EGCG (Mean  $\pm$  SD)

No.	final conc ( $\mu\text{g/ml}$ )	% inhibition			EGCG	
		N1	N2	N3	$\bar{x}$	SD
1	0.00	0.00	0.00	0.00	0.00	0.00
2	5.00	41.62	46.20	49.17	45.66	3.80
3	10.00	66.69	63.43	67.20	65.77	2.05
4	20.00	73.35	72.27	76.25	73.96	2.06
5	50.00	81.02	79.87	83.41	81.43	1.81
6	100.00	84.38	82.69	85.29	84.12	1.32

Table C12. The average percentage of hydroxyl radical inhibition of pine bark extract (Mean  $\pm$  SD)

No.	final conc ( $\mu\text{g/ml}$ )	% inhibition			pine bark extract	
		N1	N2	N3	$\bar{x}$	SD
1	0.00	0.00	0.00	0.00	0.00	0.00
2	5.00	5.16	7.21	9.66	7.34	2.25
3	10.00	20.48	21.19	17.49	19.72	1.96
4	20.00	32.43	29.84	30.54	30.94	1.34
5	50.00	49.15	45.89	46.88	47.31	1.67
6	100.00	61.99	60.01	59.56	60.52	1.29
7	200.00	71.59	70.78	68.93	70.43	1.36

Table C13. One-way analysis of variance on the IC<sub>50</sub> values of hydroxyl radical inhibition

ANOVA					
	Sum of squares	df	Mean square	F	Sig.
Between groups	4789.145	5	957.829	622.922	0.000
Within groups	18.452	12	1.538		
Total	4807.597	17			

Table C14. Multiple comparisons on the IC<sub>50</sub> values of hydroxyl radical inhibition

Tukey HSD <sup>a</sup>				
Sample	N	Subset for alpha = 0.05		
		1	2	3
4	3	1.570		
3	3	2.390		
5	3		6.203	
2	3		7.413	
1	3		9.473	
6	3			48.556
Sig.		0.96	0.62	1.000

Means for groups in homogenous subsets are displayed.

a. Uses Harmonic Mean Sample Size = 3.000.

Where; 1 = Puag-Haad, 2 = oxyresveratrol, 3 = Trolox<sup>®</sup>, 4 = l-ascorbic acid, 5 = EGCG and 6 = pine bark extract

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Table C15. One-way analysis of variance on the % inhibition at 5.0  $\mu\text{g/ml}$  of hydroxyl radical test

ANOVA					
	Sum of squares	df	Mean square	F	Sig.
Between groups	17319.270	5	3463.853	347.365	0.000
Within groups	119.662	12	9.972		
Total	17438.930	17			

Table C16. Multiple comparisons % inhibition at 5.0  $\mu\text{g/ml}$  of hydroxyl radical test

Tukey HSD <sup>a</sup>						
Sample	N	Subset for alpha = 0.05				
		1	2	3	4	5
6	3	7.343				
1	3		29.797			
2	3		37.110	37.110		
5	3			45.663		
3	3				84.963	
4	3					96.240
Sig.		1.000	0.118	0.054	1.000	1.000

Means for groups in homogenous subsets are displayed.

a. Uses Harmonic Mean Sample Size = 3.000.

Where; 1 = Puag-Haad, 2 = oxyresveratrol, 3 = Trolox<sup>®</sup>, 4 = l-ascorbic acid, 5 = EGCG and 6 = pine bark extract

Table C17. One-way analysis of variance on the % inhibition at 10.0 µg/ml of hydroxyl radical test

ANOVA					
	Sum of squares	df	Mean square	F	Sig.
Between groups	12742.68	5	2548.537	671.212	0.000
Within groups	45.563	12	3.797		
Total	12788.25	17			

Table C18. Multiple comparisons % inhibition at 10.0 µg/ml of hydroxyl radical test

Tukey HSD <sup>a</sup>					
Sample	N	Subset for alpha = 0.05			
		1	2	3	4
6	3	19.720			
1	3		52.240		
2	3			61.217	
5	3			65.773	
3	3				96.117
4	3				97.440
Sig.		1.000	1.000	0.113	0.956

Means for groups in homogenous subsets are displayed.

a. Uses Harmonic Mean Sample Size = 3.000.

Where; 1 = Puag-Haad, 2 = oxyresveratrol, 3 = Trolox<sup>®</sup>, 4 = l-ascorbic acid, 5 = EGCG and 6 = pine bark extract

Table C19. One-way analysis of variance on the % inhibition at the highest concentration of hydroxyl radical test

ANOVA					
	Sum of squares	df	Mean square	F	Sig.
Between groups	1533.124	5	306.625	178.573	0.000
Within groups	20.605	12	1.717		
Total	1553.729	17			

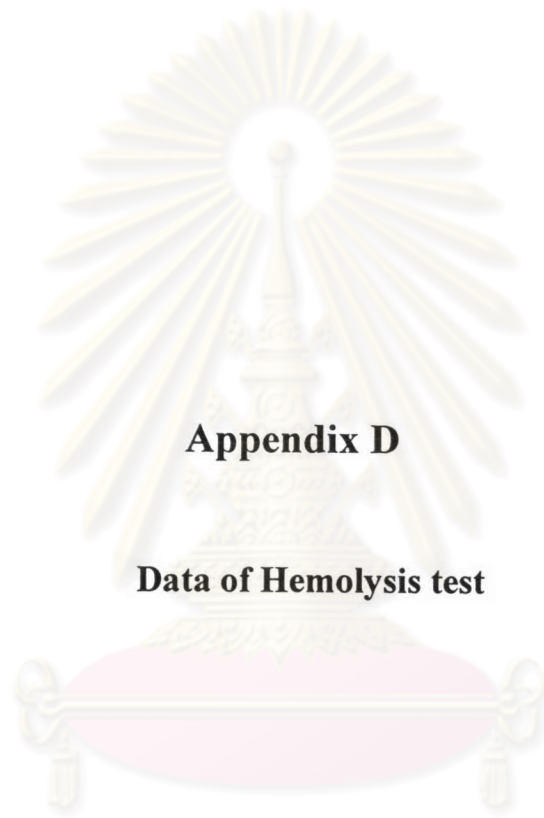
Table C20. Multiple comparisons % inhibition at the highest concentration of hydroxyl radical test

		Tukey HSD <sup>a</sup>			
Sample	N	Subset for alpha = 0.05			
		1	2	3	4
6	3	70.433			
5	3		84.120		
1	3			90.063	
2	3			90.673	
3	3				97.360
4	3				97.440
Sig.		1.000	1.000	0.991	1.000

Means for groups in homogenous subsets are displayed.

a. Uses Harmonic Mean Sample Size = 3.000.

Where; 1 = Puag-Haad, 2 = oxyresveratrol, 3 = Trolox<sup>®</sup>, 4 = l-ascorbic acid, 5 = EGCG and 6 = pine bark extract



## Appendix D

### Data of Hemolysis test

ศูนย์วิทยทรัพยากร  
จุฬาลงกรณ์มหาวิทยาลัย

Table D1. The raw data for the absorbance values and the percentage of hemolysis of Puag-Haad; (a) before UV –irradiation and (b) after UB -irradiation

(a)

Conc ( $\mu\text{g/ml}$ )	n	Sample in water			Sample in PBS			% hemolysis	average	SD
		abs	blank	diff	abs	blank	diff			
0 (control)	1	0.811	0.030	0.781	0.083	0.030	0.053	6.79	5.83	1.03
	2	0.819	0.030	0.789	0.077	0.030	0.047	5.96		
	3	0.831	0.029	0.802	0.067	0.029	0.038	4.74		
200	1	0.802	0.047	0.755	0.060	0.047	0.013	1.72	1.40	0.66
	2	0.821	0.046	0.775	0.051	0.046	0.005	0.65		
	3	0.808	0.048	0.760	0.062	0.048	0.014	1.84		
400	1	0.790	0.060	0.730	0.117	0.060	0.057	7.81	7.94	0.19
	2	0.789	0.063	0.726	0.120	0.063	0.057	7.85		
	3	0.796	0.061	0.735	0.121	0.061	0.060	8.16		
600	1	0.793	0.074	0.719	0.089	0.074	0.015	2.09	0.88	1.05
	2	0.780	0.073	0.707	0.074	0.073	0.001	0.14		
	3	0.794	0.073	0.721	0.076	0.073	0.003	0.42		

(b)

Conc ( $\mu\text{g/ml}$ )	n	Sample in water			Sample in PBS			% hemolysis	average	SD
		abs	blank	diff	abs	blank	diff			
0 (control)	1	0.585	0.029	0.556	0.555	0.029	0.526	94.60	90.06	4.60
	2	0.646	0.029	0.617	0.556	0.029	0.527	85.41		
	3	0.629	0.029	0.600	0.570	0.029	0.541	90.17		
200	1	0.598	0.047	0.551	0.382	0.047	0.335	60.80	61.15	0.67
	2	0.594	0.053	0.541	0.388	0.053	0.335	61.92		
	3	0.599	0.054	0.545	0.385	0.054	0.331	60.73		
400	1	0.581	0.067	0.514	0.230	0.067	0.163	31.71	30.88	0.93
	2	0.607	0.068	0.539	0.229	0.068	0.161	29.87		
	3	0.606	0.062	0.544	0.231	0.062	0.169	31.07		
600	1	0.624	0.077	0.547	0.177	0.077	0.100	18.28	13.48	6.05
	2	0.638	0.085	0.553	0.122	0.085	0.037	6.69		
	3	0.639	0.083	0.556	0.169	0.083	0.086	15.47		



Table D2. The raw data for the absorbance values and the percentage of hemolysis of oxyresveratrol; (a) before UV –irradiation and (b) after UB -irradiation

(a)

Conc ( $\mu\text{g/ml}$ )	n	Sample in water			Sample in PBS			% hemolysis	average	SD
		abs	blank	diff	abs	blank	diff			
0 (control)	1	0.753	0.031	0.722	0.055	0.031	0.024	3.31	3.50	0.49
	2	0.757	0.032	0.725	0.055	0.032	0.023	3.15		
	3	0.761	0.031	0.730	0.061	0.031	0.030	4.06		
200	1	0.751	0.039	0.712	0.056	0.039	0.017	2.43	2.97	0.47
	2	0.767	0.036	0.731	0.059	0.036	0.023	3.19		
	3	0.755	0.046	0.709	0.069	0.046	0.023	3.28		
400	1	0.761	0.050	0.711	0.069	0.050	0.019	2.62	3.23	0.96
	2	0.775	0.055	0.720	0.086	0.055	0.031	4.34		
	3	0.748	0.054	0.694	0.073	0.054	0.019	2.74		
600	1	0.749	0.062	0.687	0.097	0.062	0.035	5.04	5.37	0.30
	2	0.755	0.065	0.690	0.103	0.065	0.038	5.45		
	3	0.746	0.063	0.683	0.101	0.063	0.038	5.62		

(b)

Conc ( $\mu\text{g/ml}$ )	n	Sample in water			Sample in PBS			% hemolysis	average	SD
		abs	blank	diff	abs	blank	diff			
0 (control)	1	0.574	0.029	0.545	0.526	0.029	0.497	91.19	91.54	0.82
	2	0.563	0.032	0.531	0.515	0.032	0.483	90.95		
	3	0.570	0.038	0.532	0.530	0.038	0.492	92.48		
200	1	0.515	0.040	0.475	0.294	0.040	0.254	53.48	51.35	2.27
	2	0.511	0.042	0.469	0.284	0.042	0.242	51.59		
	3	0.519	0.049	0.470	0.279	0.049	0.230	48.97		
400	1	0.518	0.052	0.466	0.228	0.052	0.176	37.75	35.05	2.46
	2	0.517	0.055	0.462	0.214	0.055	0.159	34.46		
	3	0.525	0.060	0.465	0.213	0.060	0.153	32.93		
600	1	0.522	0.072	0.450	0.123	0.072	0.051	11.25	13.34	1.91
	2	0.524	0.069	0.455	0.137	0.069	0.068	15.00		
	3	0.539	0.065	0.474	0.130	0.065	0.065	13.75		

Table D3. The raw data for the absorbance values and the percentage of hemolysis of Trolox<sup>®</sup>; (a) before UV –irradiation and (b) after UB -irradiation

(a)

Conc ( $\mu\text{g/ml}$ )	n	Sample in water			Sample in PBS			% hemolysis	average	SD
		abs	blank	diff	abs	blank	diff			
0 (control)	1	0.808	0.031	0.777	0.066	0.031	0.035	4.50	6.39	1.63
	2	0.820	0.030	0.790	0.088	0.030	0.058	7.34		
	3	0.807	0.028	0.779	0.085	0.028	0.057	7.32		
200	1	0.803	0.029	0.774	0.061	0.029	0.032	4.13	4.80	0.63
	2	0.790	0.028	0.762	0.069	0.028	0.041	5.38		
	3	0.787	0.028	0.759	0.065	0.028	0.037	4.87		
400	1	0.794	0.028	0.766	0.075	0.028	0.047	6.14	6.32	0.54
	2	0.793	0.028	0.765	0.081	0.028	0.053	6.93		
	3	0.791	0.029	0.762	0.074	0.029	0.045	5.91		
600	1	0.793	0.029	0.764	0.077	0.029	0.048	6.28	6.13	1.51
	2	0.776	0.029	0.747	0.063	0.029	0.034	4.55		
	3	0.784	0.029	0.755	0.086	0.029	0.057	7.55		

(b)

Conc ( $\mu\text{g/ml}$ )	n	Sample in water			Sample in PBS			% hemolysis	average	SD
		abs	blank	diff	abs	blank	diff			
0 (control)	1	0.601	0.029	0.572	0.585	0.029	0.556	97.20	96.74	1.10
	2	0.604	0.029	0.575	0.578	0.029	0.549	95.48		
	3	0.596	0.029	0.567	0.582	0.029	0.553	97.53		
200	1	0.456	0.030	0.426	0.448	0.030	0.418	98.12	97.12	1.55
	2	0.459	0.030	0.429	0.450	0.030	0.420	97.90		
	3	0.458	0.030	0.428	0.438	0.030	0.408	95.33		
400	1	0.437	0.034	0.403	0.427	0.034	0.393	97.52	97.02	1.29
	2	0.439	0.034	0.405	0.421	0.034	0.387	95.56		
	3	0.433	0.034	0.399	0.425	0.034	0.391	97.99		
600	1	0.436	0.034	0.402	0.418	0.034	0.384	95.52	94.58	1.67
	2	0.440	0.035	0.405	0.422	0.035	0.387	95.56		
	3	0.441	0.033	0.408	0.411	0.033	0.378	92.65		

Table D4. The raw data for the absorbance values and the percentage of hemolysis of l-ascorbic acid; (a) before UV –irradiation and (b) after UB -irradiation

(a)

Conc ( $\mu\text{g/ml}$ )	n	Sample in water			Sample in PBS			% hemolysis	average	SD
		abs	blank	diff	abs	blank	diff			
0 (control)	1	0.749	0.030	0.719	0.067	0.030	0.037	5.16	4.40	0.75
	2	0.739	0.028	0.711	0.059	0.028	0.031	4.39		
	3	0.779	0.031	0.748	0.058	0.031	0.027	3.66		
200	1	0.778	0.030	0.748	0.051	0.030	0.021	2.83	5.76	2.86
	2	0.771	0.027	0.744	0.071	0.027	0.044	5.90		
	3	0.775	0.028	0.747	0.092	0.028	0.064	8.55		
400	1	0.770	0.029	0.741	0.076	0.029	0.047	6.32	6.57	1.41
	2	0.776	0.030	0.746	0.090	0.030	0.060	8.08		
	3	0.777	0.030	0.747	0.070	0.030	0.040	5.30		
600	1	0.770	0.028	0.742	0.058	0.028	0.030	3.99	4.34	0.57
	2	0.780	0.030	0.750	0.060	0.030	0.030	4.04		
	3	0.776	0.028	0.748	0.065	0.028	0.037	5.00		

(b)

Conc ( $\mu\text{g/ml}$ )	n	Sample in water			Sample in PBS			% hemolysis	average	SD
		abs	blank	diff	abs	blank	diff			
0 (control)	1	0.555	0.030	0.525	0.554	0.030	0.524	99.81	96.60	3.06
	2	0.574	0.031	0.543	0.540	0.031	0.509	93.72		
	3	0.568	0.030	0.538	0.548	0.030	0.518	96.28		
200	1	0.626	0.031	0.595	0.570	0.031	0.539	90.59	92.73	1.88
	2	0.613	0.030	0.583	0.575	0.030	0.545	93.48		
	3	0.608	0.030	0.578	0.574	0.030	0.544	94.12		
400	1	0.641	0.031	0.610	0.580	0.031	0.549	90.00	92.37	2.11
	2	0.622	0.034	0.588	0.587	0.034	0.553	94.05		
	3	0.626	0.034	0.592	0.585	0.034	0.551	93.07		
600	1	0.661	0.036	0.625	0.610	0.036	0.574	91.84	92.46	2.85
	2	0.646	0.036	0.610	0.619	0.036	0.583	95.57		
	3	0.660	0.032	0.628	0.597	0.032	0.565	89.97		

Table D5. The raw data for the absorbance values and the percentage of hemolysis of EGCG; (a) before UV –irradiation and (b) after UB -irradiation

(a)

Conc ( $\mu\text{g/ml}$ )	n	Sample in water			Sample in PBS			% hemolysis	average	SD
		abs	blank	diff	abs	blank	diff			
0 (control)	1	0.788	0.031	0.757	0.069	0.031	0.038	5.02	5.02	0.30
	2	0.781	0.028	0.753	0.068	0.028	0.040	5.31		
	3	0.792	0.029	0.763	0.065	0.029	0.036	4.72		
200	1	0.805	0.040	0.765	0.060	0.040	0.020	2.61	3.41	0.75
	2	0.796	0.039	0.757	0.070	0.039	0.031	4.10		
	3	0.808	0.039	0.769	0.066	0.039	0.027	3.51		
400	1	0.791	0.042	0.749	0.065	0.042	0.023	3.07	3.37	0.53
	2	0.796	0.045	0.751	0.068	0.045	0.023	3.06		
	3	0.800	0.046	0.754	0.076	0.046	0.030	3.98		
600	1	0.776	0.045	0.731	0.066	0.045	0.021	2.87	3.76	1.79
	2	0.789	0.052	0.737	0.071	0.052	0.019	2.58		
	3	0.783	0.044	0.739	0.087	0.044	0.043	5.82		

(b)

Conc ( $\mu\text{g/ml}$ )	n	Sample in water			Sample in PBS			% hemolysis	average	SD
		abs	blank	diff	abs	blank	diff			
0 (control)	1	0.593	0.028	0.565	0.569	0.028	0.541	95.75	95.26	1.05
	2	0.600	0.029	0.571	0.577	0.029	0.548	95.97		
	3	0.601	0.029	0.572	0.567	0.029	0.538	94.06		
200	1	0.589	0.049	0.540	0.508	0.049	0.459	85.00	90.98	5.28
	2	0.599	0.040	0.559	0.571	0.040	0.531	94.99		
	3	0.594	0.042	0.552	0.555	0.042	0.513	92.93		
400	1	0.591	0.049	0.542	0.519	0.049	0.470	86.72	87.72	1.25
	2	0.592	0.048	0.544	0.523	0.048	0.475	87.32		
	3	0.591	0.049	0.542	0.532	0.049	0.483	89.11		
600	1	0.677	0.063	0.614	0.478	0.063	0.415	67.59	71.46	6.89
	2	0.579	0.064	0.515	0.473	0.064	0.409	79.42		
	3	0.675	0.065	0.610	0.476	0.065	0.411	67.38		

Table D6. The raw data for the absorbance values and the percentage of hemolysis of pine bark extract; (a) before UV –irradiation and (b) after UB -irradiation

(a)

Conc ( $\mu\text{g/ml}$ )	n	Sample in water			Sample in PBS			% hemolysis	average	SD
		abs	blank	diff	abs	blank	diff			
0 (control)	1	0.746	0.027	0.719	0.077	0.027	0.050	6.95	6.41	0.47
	2	0.783	0.027	0.756	0.073	0.027	0.046	6.08		
	3	0.770	0.028	0.742	0.074	0.028	0.046	6.20		
200	1	0.873	0.085	0.788	0.109	0.085	0.024	3.05	3.69	0.78
	2	0.873	0.082	0.791	0.118	0.082	0.036	4.55		
	3	0.893	0.086	0.807	0.114	0.086	0.028	3.47		
400	1	0.810	0.054	0.756	0.097	0.054	0.043	-5.69	6.51	1.45
	2	0.832	0.054	0.778	0.098	0.054	0.044	5.66		
	3	0.814	0.056	0.758	0.118	0.056	0.062	8.18		
600	1	1.111	0.101	1.010	0.114	0.101	0.013	1.29	1.33	0.31
	2	1.123	0.101	1.022	0.118	0.101	0.017	1.66		
	3	1.155	0.106	1.049	0.117	0.106	0.011	1.05		

(b)

Conc ( $\mu\text{g/ml}$ )	n	Sample in water			Sample in PBS			% hemolysis	average	SD
		abs	blank	diff	abs	blank	diff			
0 (control)	1	0.546	0.033	0.513	0.542	0.033	0.509	99.22	97.88	2.67
	2	0.541	0.029	0.512	0.539	0.029	0.510	99.61		
	3	0.567	0.029	0.538	0.539	0.029	0.510	94.80		
200	1	0.537	0.091	0.446	0.282	0.091	0.191	42.83	42.97	1.13
	2	0.531	0.094	0.437	0.287	0.094	0.193	44.16		
	3	0.530	0.091	0.439	0.275	0.091	0.184	41.91		
400	1	0.519	0.065	0.454	0.212	0.065	0.147	32.38	32.02	0.33
	2	0.524	0.067	0.457	0.212	0.067	0.145	31.73		
	3	0.522	0.068	0.454	0.213	0.068	0.145	31.94		
600	1	0.553	0.132	0.421	0.163	0.132	0.031	7.36	7.00	1.12
	2	0.556	0.138	0.418	0.162	0.138	0.024	5.74		
	3	0.560	0.142	0.418	0.175	0.142	0.033	7.89		

Table D7. The average value of percent hemolysis for Puag-Haad after correction for the non-UV induced hemolysis.

Conc ( $\mu\text{g/ml}$ )	Average % hemolysis	% Hemolysis	% Hemolysis after 1.5 hr	Average % hemolysis	SD
0 (control)	5.83	94.60	88.78	84.23	4.60
		85.41	79.59		
		90.17	84.34		
200	1.40	60.80	59.40	59.75	0.67
		61.92	60.52		
		60.73	59.33		
400	7.94	31.71	23.77	22.94	0.93
		29.87	21.93		
		31.07	23.13		
600	0.88	18.28	17.40	12.60	6.05
		6.69	5.81		
		15.47	14.59		

Table D8. The average value of percent hemolysis for oxyresveratrol after correction for the non-UV induced hemolysis.

Conc ( $\mu\text{g/ml}$ )	Average % hemolysis	% Hemolysis	% Hemolysis after 1.5 hr	Average % hemolysis	SD
0 (control)	3.50	91.19	87.69	88.04	0.82
		90.95	87.45		
		92.48	88.98		
200	2.97	53.48	50.52	48.38	2.27
		51.59	48.62		
		48.97	46.00		
400	3.23	37.75	34.52	31.82	2.46
		34.46	31.23		
		32.93	29.70		
600	5.37	11.25	5.88	7.96	1.91
		15.00	9.63		
		13.75	8.38		

Table D9. The average value of percent hemolysis for Trolox® after correction for the non-UV induced hemolysis.

Conc (µg/ml)	Average % hemolysis	% Hemolysis	% Hemolysis after 1.5 hr	Average % hemolysis	SD
0 (control)	6.39	97.20	90.82	90.35	1.10
		95.48	89.09		
		97.53	91.14		
200	4.80	98.12	93.33	92.32	1.55
		97.90	93.11		
		95.33	90.53		
400	6.32	97.52	91.20	90.70	1.29
		95.56	89.23		
		97.99	91.67		
600	6.13	95.52	89.39	88.45	1.67
		95.56	89.43		
		92.65	86.52		

Table D10. The average value of percent hemolysis for l-ascorbic acid after correction for the non-UV induced hemolysis.

Conc (µg/ml)	Average % hemolysis	% Hemolysis	% Hemolysis after 1.5 hr	Average % hemolysis	SD
0 (control)	4.40	99.81	95.41	92.20	3.06
		93.72	89.32		
		96.28	91.88		
200	5.76	90.59	84.83	86.97	1.88
		93.48	87.72		
		94.12	88.35		
400	6.57	90.00	83.43	85.81	2.11
		94.05	87.48		
		93.07	86.51		
600	4.34	91.84	87.50	88.12	2.85
		95.57	91.23		
		89.97	85.63		

Table D11. The average value of percent hemolysis for EGCG after correction for the non-UV induced hemolysis.

Conc ( $\mu\text{g/ml}$ )	Average % hemolysis	% Hemolysis	% Hemolysis after 1.5 hr	Average % hemolysis	SD
0 (control)	5.02	95.75	90.74	90.24	1.05
		95.97	90.96		
		94.06	89.04		
200	3.41	85.00	81.59	87.57	5.28
		94.99	91.58		
		92.93	89.53		
400	3.37	86.72	83.35	84.34	1.25
		87.32	83.95		
		89.11	85.74		
600	3.76	67.59	63.83	67.70	6.89
		79.42	75.66		
		67.38	63.62		

Table D12. The average value of percent hemolysis for pine bark extract after correction for the non-UV induced hemolysis.

Conc ( $\mu\text{g/ml}$ )	Average % hemolysis	% Hemolysis	% Hemolysis after 1.5 hr	Average % hemolysis	SD
0 (control)	6.41	99.22	92.81	91.46	2.67
		99.61	93.20		
		94.80	88.38		
200	3.69	42.83	39.14	39.28	1.13
		44.16	40.48		
		41.91	38.22		
400	6.51	32.38	25.87	25.51	0.33
		31.73	25.22		
		31.94	25.43		
600	1.33	7.36	6.03	5.67	1.12
		5.74	4.41		
		7.89	6.56		



Table D13. The average values of percent relative hemolysis of Puag-Haad

Conc ( $\mu\text{g/ml}$ )	% Hemolysis (corrected)	% Relative hemolysis	Average	SD
0 (control)	88.78	105.39	100.00	5.46
	79.59	94.48		
	84.34	100.12		
200	59.40	70.51	70.93	0.79
	60.52	71.85		
	59.33	70.44		
400	23.77	28.22	27.24	1.11
	21.93	26.03		
	23.13	27.45		
600	17.40	20.66	14.96	7.18
	5.81	6.90		
	14.59	17.32		

Table D14. The average values of percent relative hemolysis of oxyresveratrol

Conc ( $\mu\text{g/ml}$ )	% Hemolysis (corrected)	% Relative hemolysis	Average	SD
0 (control)	87.69	99.60	100.00	0.93
	87.45	99.33		
	88.98	101.07		
200	50.52	57.38	54.95	2.58
	48.62	55.23		
	46.00	52.25		
400	34.52	39.22	36.14	2.80
	31.23	35.47		
	29.70	33.74		
600	5.88	6.68	9.05	2.17
	9.63	10.94		
	8.38	9.52		

Table D15. The average values of percent relative hemolysis of Trolox®

Conc (µg/ml)	% Hemolysis (corrected)	% Relative hemolysis	Average	SD
0 (control)	90.82	100.52	100.00	1.22
	89.09	98.61		
	91.14	100.88		
200	93.33	103.29	102.18	1.72
	93.11	103.05		
	90.53	100.20		
400	91.20	100.94	100.39	1.43
	89.23	98.76		
	91.67	101.46		
600	89.39	98.94	97.89	1.85
	89.43	98.98		
	86.52	95.76		

Table D16. The average values of percent relative hemolysis of l-ascorbic acid

Conc (µg/ml)	% Hemolysis (corrected)	% Relative hemolysis	Average	SD
0 (control)	95.41	103.48	100.00	3.32
	89.32	96.87		
	91.88	99.65		
200	84.83	92.00	94.32	2.04
	87.72	95.14		
	88.35	95.83		
400	83.43	90.49	93.07	2.29
	87.48	94.88		
	86.51	93.83		
600	87.50	94.90	95.57	3.10
	91.23	98.95		
	85.63	92.87		

Table D17. The average values of percent relative hemolysis of EGCG

Conc ( $\mu\text{g/ml}$ )	% Hemolysis (corrected)	% Relative hemolysis	Average	SD
0 (control)	90.74	100.55	100.00	1.16
	90.96	100.79		
	89.04	98.67		
200	81.59	90.41	97.04	5.85
	91.58	101.49		
	89.53	99.21		
400	83.35	92.36	93.46	1.38
	83.95	93.02		
	85.74	95.01		
600	63.83	70.73	75.02	7.64
	75.66	83.84		
	63.62	70.50		

Table D18. The average values of percent relative hemolysis of pine bark extract

Conc ( $\mu\text{g/ml}$ )	% Hemolysis (corrected)	% Relative hemolysis	Average	SD
0 (control)	92.81	101.47	100.00	2.92
	93.20	101.90		
	88.38	96.63		
200	39.14	42.79	42.95	1.24
	40.48	44.25		
	38.22	41.79		
400	25.87	28.29	27.89	0.36
	25.22	27.58		
	25.43	27.80		
600	6.03	6.59	6.20	1.23
	4.41	4.82		
	6.56	7.17		

Table D19. One-way analysis of variance on the percent hemolysis at various concentration of Puag-Haad

ANOVA					
	Sum of squares	df	Mean square	F	Sig.
Between groups	9879.839	3	3293.280	223.301	0.000
Within groups	117.985	8	14.748		
Total	9997.825	11			

Table D20. Multiple comparisons on the percent hemolysis at various concentration of Puag-Haad

Tukey HSD <sup>a</sup>					
Sample	N	Subset for alpha = 0.05			
		1	2	3	4
4	3	12.600			
3	3		22.943		
2	3			59.750	
1	3				84.237
Sig.		1.000	1.000	1.000	1.000

Means for groups in homogenous subsets are displayed.

a. Uses Harmonic Mean Sample Size = 3.000.

Where; 1 = 0  $\mu\text{g/ml}$ , 2 = 200  $\mu\text{g/ml}$ , 3 = 400  $\mu\text{g/ml}$ , 4 = 600  $\mu\text{g/ml}$

Table D21. One-way analysis of variance on the percent hemolysis at various concentration of oxyresveratrol

ANOVA					
	Sum of squares	df	Mean square	F	Sig.
Between groups	10217.31	3	3405.771	876.648	0.000
Within groups	31.080	8	3.885		
Total	10248.39	11			

Table D22. Multiple comparisons on the percent hemolysis at various concentration of oxyresveratrol

Tukey HSD <sup>a</sup>					
Sample	N	Subset for alpha = 0.05			
		1	2	3	4
4	3	7.963			
3	3		31.817		
2	3			48.380	
1	3				88.040
Sig.		1.000	1.000	1.000	1.000

Means for groups in homogenous subsets are displayed.

a. Uses Harmonic Mean Sample Size = 3.000.

Where; 1 = 0 µg/ml, 2 = 200 µg/ml, 3 = 400 µg/ml, 4 = 600 µg/ml

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Table D23. One-way analysis of variance on the percent hemolysis at various concentration of Trolox®

ANOVA					
	Sum of squares	df	Mean square	F	Sig.
Between groups	22.785	3	7.595	3.750	0.060
Within groups	16.202	8	2.025		
Total	38.987	11			

Table D24. One-way analysis of variance on the percent hemolysis at various concentration of l-ascorbic acid

ANOVA					
	Sum of squares	df	Mean square	F	Sig.
Between groups	69.781	3	23.260	3.652	0.063
Within groups	50.947	8	6.368		
Total	120.728	11			

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Table D25. One-way analysis of variance on the percent hemolysis at various concentration of EGCG

ANOVA					
	Sum of squares	df	Mean square	F	Sig.
Between groups	924.086	3	308.029	15.800	0.001
Within groups	155.967	8	19.496		
Total	1080.054	11			

Table D26. Multiple comparisons on the percent hemolysis at various concentration of EGCG

Tukey HSD <sup>a</sup>			
Sample	N	Subset for alpha = 0.05	
		1	2
4	3	67.703	
3	3		84.347
2	3		87.567
1	3		90.247
Sigs.		1.000	0.412

Means for groups in homogenous subsets are displayed.

a. Uses Harmonic Mean Sample Size = 3.000.

Where; 1 = 0  $\mu\text{g/ml}$ , 2 = 200  $\mu\text{g/ml}$ , 3 = 400  $\mu\text{g/ml}$ , 4 = 600  $\mu\text{g/ml}$

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Table D27. One-way analysis of variance on the percent hemolysis at various concentration of pine bark extract

ANOVA					
	Sum of squares	df	Mean square	F	Sig.
Between groups	12110.73	3	4036.909	1643.609	0.000
Within groups	19.649	8	2.456		
Total	12130.38	11			

Table D28. Multiple comparisons on the percent hemolysis at various concentration of pine bark extract

Tukey HSD <sup>a</sup>					
Sample	N	Subset for alpha = 0.05			
		1	2	3	4
4	3	5.667			
3	3		25.507		
2	3			39.280	
1	3				91.463
Sig.		1.000	1.000	1.000	1.000

Means for groups in homogenous subsets are displayed.

a. Uses Harmonic Mean Sample Size = 3.000.

Where; 1 = 0 µg/ml, 2 = 200 µg/ml, 3 = 400 µg/ml, 4 = 600 µg/ml

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Table D29. One-way analysis of variance on the percent relative hemolysis at 200 g/ml of Puag-Haad, oxyresveratrol, EGCG and pine bark extract

ANOVA					
	Sum of squares	df	Mean square	F	Sig.
Between groups	4921.140	3	1640.380	152.477	0.000
Within groups	86.065	8	10.758		
Total	5007.206	11			

Table D30. Multiple comparisons on the percent relative hemolysis at 200 g/ml of Puag-Haad, oxyresveratrol, EGCG and pine bark extract

Tukey HSD <sup>a</sup>					
Sample	N	Subset for alpha = 0.05			
		1	2	3	4
4	3	42.943			
2	3		54.953		
1	3			70.933	
3	3				97.037
Sig.		1.000	1.000	1.000	1.000

Means for groups in homogenous subsets are displayed.

a. Uses Harmonic Mean Sample Size = 3.000.

Where; 1= Puag-Haad, 2 = oxyresveratrol, 3 = EGCG, 4 = pine bark extract

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Table D31. One-way analysis of variance on the percent relative hemolysis at 400 g/ml of Puag-Haad, oxyresveratrol, EGCG and pine bark extract

ANOVA					
	Sum of squares	df	Mean square	F	Sig.
Between groups	9089.846	3	3029.949	1090.204	0.000
Within groups	22.234	8	2.779		
Total	9112.080	11			

Table D32. Multiple comparisons on the percent relative hemolysis at 400 g/ml of Puag-Haad, oxyresveratrol, EGCG and pine bark extract

Tukey HSD <sup>a</sup>				
Sample	N	Subset for alpha = 0.05		
		1	2	3
1	3	27.233		
4	3	27.890		
2	3		36.1433	
3	3			93.463
Sig.		0.961	1.000	1.000

Means for groups in homogenous subsets are displayed.

a. Uses Harmonic Mean Sample Size = 3.000.

Where; 1= Puag-Haad, 2 = oxyresveratrol, 3 = EGCG, 4 = pine bark extract

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Table D33. One-way analysis of variance on the percent relative hemolysis at 600 g/ml of Puag-Haad, oxyresveratrol, EGCG and pine bark extract

ANOVA					
	Sum of squares	df	Mean square	F	Sig.
Between groups	9613.543	3	3204.514	110.473	0.000
Within groups	232.057	8	29.007		
Total	9845.600	11			

Table D34. Multiple comparisons on the percent relative hemolysis at 600 g/ml of Puag-Haad, oxyresveratrol, EGCG and pine bark extract

Tukey HSD <sup>a</sup>			
Sample	N	Subset for alpha = 0.05	
		1	2
4	3	6.193	
2	3	9.047	
1	3	14.960	
3	3		75.0233
Sigs.		0.266	1.000

Means for groups in homogenous subsets are displayed.

a. Uses Harmonic Mean Sample Size = 3.000.

Where; 1= Puag-Haad, 2 = oxyresveratrol, 3 = EGCG, 4 = pine bark extract

## VITA

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