

AcF3'5'H Recombonant Protein

	Contents				
	1	2	3	Mean	SD
WT	0,0004	0,0006	0,0007	0,0006	0,0002
T01	0,212	0,191	0,207	0,2033	0,0110
T03	0,315	0,289	0,326	0,3100	0,0190
T05	0,203	0,218	0,196	0,2057	0,0112
T014	0,329	0,314	0,332	0,3250	0,0096

WT: Cây thuốc lá không chuyển gen

T01, T03, T05, T014: Cây thuốc lá chuyển gen AcF3'5'H

		Descriptives		
rProtein			Statistic	Std. Error
Hamluong	1	Mean	,20333	,006333
		95% Confidence Interval for Mean		
		Lower Bound	,17608	
		Upper Bound	,23058	
		Median	,20700	
		Variance	,000	
		Std. Deviation	,010970	
		Minimum	,191	
		Maximum	,212	
		Range	,021	
		Interquartile Range		
		Skewness	-1,336	1,225
		Kurtosis		
			2	Mean
95% Confidence Interval for Mean				
Lower Bound	,26280			
Upper Bound	,35720			
Median	,31500			
Variance	,000			
Std. Deviation	,019000			
Minimum	,289			
Maximum	,326			
Range	,037			
Interquartile Range				
Skewness	-1,102			1,225
Kurtosis				
	3			Mean
		95% Confidence Interval for Mean		
		Lower Bound	,17775	
		Upper Bound	,23359	

	Median		,20300	
	Variance		,000	
	Std. Deviation		,011240	
	Minimum		,196	
	Maximum		,218	
	Range		,022	
	Interquartile Range			
	Skewness		1,008	1,225
	Kurtosis			
4	Mean		,32500	,005568
	95% Lower Bound		,30104	
	Confidence Upper Bound		,34896	
	Interval for 5% Trimmed Mean			
	Median		,32900	
	Variance		,000	
	Std. Deviation		,009644	
	Minimum		,314	
	Maximum		,332	
	Range		,018	
	Interquartile Range			
	Skewness		-1,545	1,225
	Kurtosis			

Ham Luong

Subset for alpha = 0.001

rProtein		N	1	2
Duncan ^a	1	3	,20333	
	3	3	,20567	
	2	3		,31000
	4	3		,32500
	Sig.		,834	,203

Means for groups in homogeneous subsets are displayed.

a. Uses Harmonic Mean Sample Size = 3,000.

ug/uL

	Trung bình	Sx	
WT	0,0000	0,0000	1
T01	0,2033	0,0063	20.33
T03	0,3100	0,0109	31.00
T05	0,2057	0,0065	20.57
T014	0,3250	0,0056	32.50

