

My name . . .

Partners name . . . (if applicable)

average atomic mass simulation lab

honors chem

period x

introduction

1. A simulation mimics the behavior of a system. [“citation not needed since didn’t need to look it up”]

purpose

2. Phet: blah, since given free points

AACT: blah

3. Based on the output of the Excel simulation

atomic mass simulation lab using beans to represent isotopes	
to simulate running multiple trails, press F9 key for each trail	
output in green high-lighted cells	
number of beans	
lima =	15
pinto =	42
navy =	59
total mass of sample =	42.83 grams

use below average mass of a single bean in your calculations

	Bean 1 (Lentils)	Bean 2 (Navy)	Bean 3 (Pinto)	Bean 4 (Lima)
Mass of 100 beans plus cup (g)	18.16	28.65	52.55	113.15
Mass of 100 beans (g)	6.01	16.50	40.40	101.00
Average mass of one bean (g)	0.0601	0.165	0.404	1.01

to find the average bean mass, using equation 1,

$$\text{average bean mass} = \frac{\# \text{ g beans in sample}}{\# \text{ beans in sample}} = \frac{42.83 \text{ g}}{(15+42+59)\text{beans}} = 0.369 \frac{\text{g}}{\text{bean}}$$

and using equation 2,

$$\begin{aligned} \text{average bean mass} &= \sum m_i p_i \\ &= m_{\text{lima}} p_{\text{lima}} + m_{\text{pinto}} p_{\text{pinto}} + m_{\text{navy}} p_{\text{navy}} \\ &= 1.01 \left(\frac{15}{15+42+59} \right) + 0.404 \left(\frac{42}{15+42+59} \right) + 0.165 \left(\frac{59}{15+42+59} \right) \\ &= 0.229 \text{ g / bean} \end{aligned}$$

seems to have different values.

results

4. Running the Excel simulation, 8 more times, to find the average bean mass (in g / bean) produced the below results.

Use eqn 1	Use eqn 2
0.341	0.337
0.341	0.359
0.325	0.346
0.308	0.325
0.318	0.329
0.340	0.355
0.349	0.360

The results of using the independent sample t-test (<http://vassarstats.net/>), where column A was from eqn 1, while column B is from eqn 2, is shown below.

Data Summary			
	A	B	Total
n	8	8	16
ΣX	2.649	2.76299999	5.41199999
ΣX^2	0.87920700	0.95556099	1.834768
SS	0.0021	0.0013	0.0042
mean	0.3311	0.3454	0.3382

Results

Mean _a —Mean _b	t	df	p	one-tailed	two-tailed
-0.0142	-1.84	14		0.043532	0.087064

For independent samples, these results pertain to the "usual" t-test, which assumes that the two samples have equal variances.

F-Test for the Significance of the Difference between the Variances of the Two Samples

df ₁	df ₂	F	P
7	7	1.59	0.277767

[Applicable only to independent samples.]
P>.05 indicates no significant difference detected between the variances of the two samples.

as the 2-tail p-value > 0.05, the data is the same; i.e. eqn 1 and eqn 2 produce the same results.

discussion

4. Show that

$$\text{eqn1: average bean mass} = \frac{\# g \text{ beans in sample}}{\# \text{ beans in sample}}$$

$$\text{eqn 2: average bean mass} = m_{\text{lima}} p_{\text{lima}} + m_{\text{pinto}} p_{\text{pinto}} + m_{\text{navy}} p_{\text{navy}}$$

are the same. Equation 1 is

average bean mass

$$\begin{aligned} &= \frac{\# g \text{ beans in sample}}{\# \text{ beans in sample}} = \frac{m_{\text{lima}} + m_{\text{pinto}} + m_{\text{navy}}}{n_{\text{total}}} \\ &= \frac{m_{\text{single lima}} n_{\text{lima}} + m_{\text{single pinto}} n_{\text{pinto}} + m_{\text{single navy}} n_{\text{navy}}}{n_{\text{total}}} \\ &= \frac{m_{\text{single,lima}} n_{\text{lima}}}{n_{\text{total}}} + \frac{m_{\text{single,pinto}} n_{\text{pinto}}}{n_{\text{total}}} + \frac{m_{\text{single,navy}} n_{\text{navy}}}{n_{\text{total}}} \\ &= m_{\text{lima}} p_{\text{lima}} + m_{\text{pinto}} p_{\text{pinto}} + m_{\text{navy}} p_{\text{navy}} \end{aligned}$$

which is eqn 2; therefore, eqn 1 and eqn 2 are the same.