## BRCA 2 Genomic Data Set

| DRCA.sav [DataSet2] - SPSS Data Editor |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
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| 5 : |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
|  | base | count | var | vat | var | var | var | var | var | var | var | var | var | var | var | var | var | var | var | var | var | var |
| 1 | A | 38514 |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| 2 | c | 24631 |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| 3 | G | 25685 |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| 4 | T | 38249 |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| 5 |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
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| 16 <br> 17 |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| 178 |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
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| 28 |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| 29 |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| $\begin{array}{\|c\|} \hline 30 \\ 31 \end{array}$ |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| $\frac{31}{37}$ |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| 32 |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| $\begin{array}{\|c\|} \hline 33 \\ 34 \end{array}$ |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| $34$ |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| $\frac{35}{36}$ |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
|  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| $\mid 37$ |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| $\begin{array}{\|c\|} \hline 38 \\ \hline 39 \end{array}$ |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| $\frac{39}{40}$ |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| 40 |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| 41 |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| 42 |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| 43 |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
|  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |

Use the Data, Weight Cases pulldown selections to define the table


## The dialog gives you the option to define the counts

| In BRCA.sav [DataSet2] - SPSS Data Editor |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
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| 33 |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
|  | base | count | var | var | var | var | var | var | var | var | var | var | yar | var | var | var | var | var | var | var | var | var |
| 1 | A | 38514 |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| 2 | c | 24631 |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| 3 | G | 25685 |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| 4 | T | 38249 |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
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| 6 |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| 7 |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| 8 |  |  | $\square$ Weight Cases $x$ |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| 9 |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| 10 |  |  | $\begin{aligned} & \text { B Nucleotide [base] } \\ & \text { \& Count [count] } \end{aligned}$ |  | © Do not weight cases |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| 11 |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| 12 |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| 13 |  |  |  |  | 1 | Frequency Varables |  | Reset <br> Cancel |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| 14 |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| 15 |  |  |  |  |  |  |  |  | Help |  |  |  |  |  |  |  |  |  |  |  |  |  |
| 16 |  |  | $\qquad$ |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| 17 |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| 18 |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
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| 22 |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
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| 24 |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| 25 |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| 26 |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| 27. |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
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| 29 |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| 30 |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| 31 |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
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| 33 |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
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| 36 |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| 37 |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| 38 |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
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| 42 |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
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| 44 |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |

Select the Weight cases by toggle, and pass the count variable into the Frequency Variable box

| (1) BRCA.sav [DataSet2] - SPSS Data Editor |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
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| 33 |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
|  | base | count | var | var | var | var | var | var | var | var | var | var | var | var | var | var | var | var | var | var | var | var |
| 1 | A | 38514 |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| 2 | c | 24631 |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| 3 | G | 25685 |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| 4 | T | 38249 |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| 5 |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| 6 |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| 7 |  |  | Weight Cases |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| 8 |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| 9 |  |  | \& Nucleotide [base] |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| 10 |  |  |  |  | ODo not weight cases 0 K |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| 11 |  |  |  |  | () Weight cases by |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| 12 |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
| 13 |  |  |  |  | Frequency Variable: O Count [count] |  |  |  | Reset |  |  |  |  |  |  |  |  |  |  |  |  |  |
| 14 |  |  |  |  |  | Cancel |  |  |  |  |  |  |  |  |  |  |  |  |  |
| 15 |  |  |  |  | Curent Status: Weight cases by base |  |  |  | Help |  |  |  |  |  |  |  |  |  |  |  |  |  |
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| $38$ |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |
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Use Analyze, Nonparametric Tests, Chi-Square pulldown selections


Pass the base variable into the Test Variable List, and click OK. here we test against the probability distribution with all probabilities equal.


The results are computed.

Nucleotide

|  | Observed N | Expected N | Residual |
| :--- | ---: | ---: | ---: |
| A | 38514 | 31769.8 | 6744.3 |
| C | 24631 | 31769.8 | -7138.8 |
| G | 25685 | 31769.8 | -6084.8 |
| T | 38249 | 31769.8 | 6479.3 |
| Total | 127079 |  |  |

Test Statistics

|  | Nucleotide |
| :--- | ---: |
| Chi-Square $^{\mathrm{a}}$ | 5522.597 |
| df | 3 |
| Asymp. Sig. | .000 |

a. 0 cells (.0\%) have expected frequencies less than 5. The minimum expected cell frequency is 31769.8.

To test against a different set of probabilities, use the Expected Values toggle


We test against the probability distribution we probabilities ( $0.3,0.2,0.2,0.3$ ). We type these values successively into the Values box, and click Add


The new results are produced.

## Nucleotide

|  | Observed N | Expected N | Residual |
| :--- | ---: | ---: | ---: |
| A | 38514 | 38123.7 | 390.3 |
| C | 24631 | 25415.8 | -784.8 |
| G | 25685 | 25415.8 | 269.2 |
| T | 38249 | 38123.7 | 125.3 |
| Total | 127079 |  |  |

Test Statistics

|  | Nucleotide |
| :--- | ---: |
| Chi-Square $^{\mathrm{a}}$ | 31.492 |
| df | 3 |
| Asymp. Sig. | .000 |

a. 0 cells (.0\%) have expected frequencies less than 5. The minimum expected cell frequency is 25415.8.

## The Eye and Hair colour data set.



Use the Data, Weight Cases pulldown selections to define the table


Use the Analyze, Descriptive Statistics, Crosstabs pulldown selections


Pass eye into the Row(s) list, and hair into the Column(s) list


Click the Statistics button on the Crosstabs dialog, and select Chi-square. Click Continue, then OK.


The results are produced.
Eye Colour * Hair Colour Crosstabulation
Count

|  |  | Hair Colour |  |  |  |  |
| :--- | :--- | ---: | ---: | ---: | ---: | ---: |
| Total |  |  |  |  |  |  |
|  |  | Black | Brown | Red | Blond | Tot |
| Eye | Brown | 68 | 119 | 26 | 7 | 220 |
| Colour | Blue | 20 | 84 | 17 | 94 | 215 |
|  | Hazel | 15 | 54 | 14 | 10 | 93 |
|  | Green | 5 | 29 | 14 | 16 | 64 |
| Total |  | 108 | 286 | 71 | 127 | 592 |

Chi-Square Tests

|  | Value | df | Asymp. Sig. <br> (2-sided) |
| :--- | ---: | ---: | ---: |
| Pearson Chi-Square | $138.290^{\mathrm{a}}$ | 9 | .000 |
| Likelihood Ratio | 146.444 | 9 | .000 |
| Linear-by-Linear | 28.292 | 1 | .000 |
| Association | 592 |  |  |
| N of Valid Cases | 59 |  |  |

a. 0 cells $(.0 \%)$ have expected count less than 5 . The minimum expected count is 7.68 .

