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# Uncertainty Analysis of RELAP5-3D© 

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#### Abstract

As world-wide energy consumption continues to increase, so does the demand for the use of alternative energy sources, such as Nuclear Energy. Nuclear Power Plants currently supply over 370 gigawatts of electricity, and more than 60 new nuclear reactors have been commissioned by 15 different countries. The primary concern for Nuclear Power Plant operation and lisencing has been safety. The safety of the operation of Nuclear Power Plants is no simple matter- it involves the training of operators, design of the reactor, as well as equipment and design upgrades throughout the lifetime of the reactor, etc. To safely design, operate, and understand nuclear power plants, industry and government alike have relied upon the use of best-estimate simulation codes, which allow for an accurate model of any given plant to be created with welldefined margins of safety. The most widely used of these best-estimate simulation codes in the Nuclear Power industry is RELAP5-3D. Our project focused on improving the modeling capabilities of RELAP5-3D by developing uncertainty estimates for its calculations. This work involved analyzing high, medium, and low ranked phenomena from an INL PIRT on a small break Loss-Of-Coolant Accident as well as an analysis of a large break Loss-Of-Coolant Accident. Statistical analyses were performed using correlation coefficients. To perform the studies, computer programs were written that modify a template RELAP5-3D input deck to produce one deck for each combination of key input parameters. Python scripting enabled the running of the generated input files with RELAP5-3D on INL's massively parallel cluster system. Data from the studies was collected and analyzed with SAS. Summaries of the results of our studies are presented.


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## 1 Nomenclature [1]

| Accronym | Definition |
| :---: | :--- |
| ADS | Automatic Depressurization System |
| ALWR | Advanced Light-Water Reactor |
| AP600 | Advanced Passive 600 MWe Reactor |
| CMT | Core Makeup Tank |
| ECCS | Emergency Core Coolant System |
| INL | Idaho National Laboratory |
| IRWST | In-containment Refueling Water Storage Tank |
| LBLOCA | Large Break Loss-of-Coolant Accident |
| LOCA | Loss-of-Coolant Accident |
| MSLB | Main Steam Line Break |
| NPP | Nuclear Power Plant |
| NRC | United States Nuclear Regulatory Commission ${ }^{1}$ |
| PIRT | Phenomena Identification and Ranking Table |
| PPMCC | Pearson Product Moment Correlation Coefficient |
| PRHR | Passive Residual Heat Removal |
| PWR | Pressurized Water Reactor |
| SBLOCA | Small Break Loss-of-Coolant Accident |
| SG | Steam Generator |
| $\hat{\rho}$ | The estimated value of PPMCC |
| $\hat{\tau}$ | The estimated value of Spearman's rho |
| The estimated value of Kendall's tau |  |

[^1]
## 2 Motiviation

As scientists, we wish to confidently provide the answer to any question quantitatively. However, every measured value has uncertainty attached to it and thus any answer we provide must include uncertainty. Uncertainty in measurement is not the only uncertainty we must concern ourselves with. For any system we construct, outside factors will play some role within our calculations and findings, and therefore additional uncertainties must be accounted for in any solutions we provide.

In practice it is not reasonable to assume that we could quantify all uncertainties for a particular solution. Even if it were possible to quantify all possible solutions, combining all the uncertainties together could cause our data to lose significance, i.e. to become "washed out" in terms of significance. Therefore we must utilize practices that limit the number of uncertainties we consider for any particular problem. This is where the studies of Statistical Uncertainty Analysis and Sensitivity Analysis have resulted from. [8]

Uncertainty (and Sensitivity) Analysis continue to be vitally important to Nuclear Power Plants (NPPs), as well as many other fields. For NPPs in particular, the fields of uncertainty and sensitivity analysis are vitally important, as margins of safety for plant operation are a critical aspect of licensing and plant operation.

In 1988, the USNRC (United States Nuclear Regulatory Commission) issued a revision to the emergency core coolant system (ECCS) which allows for the use of best estimate plus uncertainty methods in safety analysis of LWRs (Light Water Reactors). In support of this licensing revision, the code scaling, applicability and uncertainty (CSAU) methodology was developed. As a part of that methodology, the Phenomena Identification and Ranking Table (PIRT) was developed. [14]

The PIRT process is a structured and facilitated elicitation process in which experts are
asked to rank various phenomena pertaining to a particular scenario. The phenomena are typically classified as "high", "medium", or "low". [6] The PIRT process of today also typically includes the utilization of best-estimate codes to assist in the ranking process of phenomena.

We wish to find a way to mathematically evaluate the accuracy of PIRTs, and to provide insight to validate and/or to make suggested changes to a given PIRT.

To do this, we utilized a RELAP5-3D input deck and a corresponding PIRT to determine variables of interest. Using those, we were able to create a variable specification file (spec file), as well as a template input file. We designed a program which was able to take the template input file and a specification file for the study and generate more specification files which would allow the study to be broken into sets. Then each set was run individually on the INL's supercomputer cluster (using a Python script), where up to 1,737 input files were created using our program, and run with a RELAP5-3D executable. The data was collected using the python script, and then a statistical analysis was conducted. A depiction of this process can be seen in Figure 1.


Figure 1: Algorithmic Design

We will begin by listing general statistical definitions in Section 3.1 on Page 8, defining project specific definitions in Section 3.2 on Page 10, and then discussing Coefficients of Correlation in Section 4 on Page 11. We will discuss general properties of correlation coefficients in Section 4.1 on Page 11, and discuss three types of correlation coefficients in Sections 4.2, 4.3, and 4.4 on Pages 14, 15, and 16. The relationship between the three is discussed in Section 4.5 on Page 19. We then discuss our experiments in Section 5 on Page 20. We will conclude by discussing our results in Section 6 on Page 35, summarizing our findings and discussing future work following from this project in Section 7 on Page 37 ,

## 3 Standard Formulas

### 3.1 General Definitions

For a general closed non-degenerate interval, $[c, d]$, with $n$ points equally spaced within the interval (i.e. uniformly distributed), the increment size of the interval, $\xi$, is $\xi=\frac{d-c}{n-1}$.

The probability that a sample point $x$ will occur is the proportion of ocurrences of the sample point in a long series of experiments, and is denoted by $P(x) . P(x) \in[0,1]$.

The probability density function (PDF) of a continuous random variable $X$, denoted by $f(x)$, is defined such that $\int_{-\infty}^{\infty} f(x) d x=1$. The probability that an observation lies between $x_{1}$ and $x_{2}$ is defined as: $\mathrm{P}\left(x_{1}<X<x_{2}\right)=\int_{x_{1}}^{x_{2}} f(x) d x$. The mean $\left(\mu_{x}\right)$ or expected value $E[X]$ of the continuous random variable $X$, is $E[X]=\int_{-\infty}^{\infty} x f(x) d x$.

Other expectations are mathematically useful and important. We may define expecation more generally for $g(X)$, a function of the continuous random variable $X$, by:

$$
E[g(X)]=\int_{-\infty}^{\infty} g(x) f(x) d x
$$

The variance of the continuous random variable $X$, where $\mu_{x}=E[X]$ is constant, is defined as:

$$
\operatorname{variance}(X)=\sigma_{x}=E\left[\left(X-\mu_{x}\right)^{2}\right]=\int_{-\infty}^{\infty}\left(x-\mu_{x}\right)^{2} f(x) d x=E\left[X^{2}\right]-\left\{\mu_{x}\right\}^{2}
$$

The covariance of two random variables, $X$ and $Y$ with $\operatorname{PDF} f(x, y)$, is defined as:

$$
\begin{gathered}
\operatorname{Cov}(X, Y)=E\left[\left(X-\mu_{x}\right)\left(Y-\mu_{y}\right)\right]=\int_{-\infty}^{\infty} \int_{-\infty}^{\infty}\left(X-\mu_{x}\right)\left(Y-\mu_{y}\right) f(x, y) d y d x= \\
\int_{-\infty}^{\infty} \int_{-\infty}^{\infty} x y f(x, y) d x d y-\mu_{x} E[X]-\mu_{y} E[Y]+\mu_{x} \mu_{y}=E[X Y]-\mu_{X} \mu_{Y}
\end{gathered}
$$

Let $X$ be a random variable with $n$ data points. Let each point be denoted by $x_{i}$ for $i=1,2, \ldots, n$. (i.e. the data points are $x_{1}, x_{2}, \ldots, x_{n}$.)

The sample mean of $X, \hat{\mu_{x}}$, is defined as ${ }^{2}$,

$$
\hat{\mu_{x}}=\frac{1}{n} \sum_{i=1}^{n} x_{i} \cdot[2, \mathrm{p} 39,81-82,87,106-113]
$$

The sample variance of $X, \hat{\sigma}^{2}$, where $\tilde{s}^{2}$ is an unbiased estimate of $\hat{\sigma}^{2}$, is defined as:

$$
\hat{\sigma_{x}^{2}}=E\left[\tilde{s}^{2}\right]=\frac{\sum_{i=1}^{n}\left(x_{i}-\mu_{x}\right)^{2}}{n-1}=\frac{\sum_{i=1}^{n}\left(x_{i}\right)^{2}-\frac{1}{n}\left(\sum_{i=1}^{n} x_{i}\right)^{2}}{n-1} \text {. [11, p. 11] }
$$

The sample covariance of two random variables, $X$, and $Y$, each with $n$ data points, where each point of $X$ is denoted by $x_{i}$ and each point of $Y$ is denoted by $y_{i}$ for $i=1,2, \ldots, n$, is

$$
\widehat{\operatorname{Cov}}(X, Y)=\frac{1}{n-1} \sum_{i=1} n\left[\left(x_{i}-\hat{\mu_{x}}\right)\left(y_{i}-\hat{\mu_{y}}\right)\right] .
$$

[^2]
### 3.2 Project Specific Definitions

In our input modification program, we are given a base file name, the number of variables, the number of sets the study will be divided into, the number of groups of variables $3^{3}$ as well as a list of variables and some basic variable information. The variable information includes the minimum and maximum values of the interval over which the variable ranges, the number of points within the interval that the variable will have (i.e. sample size), the standard deviation of the variable, the type of probability distribution the points will have within the interval (currently only the uniform is available in our program), and the group number of the variable.

For a specific variable, v , with $n$ points, a minimum of $a$, and a maximum of $b$ (where $a<b$ ), for which the points are distributed uniformly throughout the interval, the following formulas follow from the general definitions.

$$
\begin{aligned}
\xi_{v} & =\frac{b-a}{n-1} \\
\hat{\mu_{v}} & =\frac{b-a}{2}
\end{aligned}
$$

$$
\begin{align*}
\hat{\sigma_{v}} & =\frac{1}{n} \sqrt{\frac{1}{n-1} \sum_{i=1}^{n}\left[a+\xi_{\mathrm{v}}(i-1)-\hat{\mu_{v}}\right]^{2}}=\frac{1}{n} \sqrt{\frac{1}{n-1} \sum_{i=1}^{n}\left[a+\frac{b-a}{n-1}(i-1)-\frac{b-a}{2}\right]^{2}} \\
& =\frac{1}{2 n(n-1)} \sqrt{\frac{1}{n-1} \sum_{i=1}^{n}[3 a n-a-b n-b-2 a i+2 b i]^{2}} \tag{1}
\end{align*}
$$

[^3]
## 4 Coefficients of Correlation

Let $X$ and $Y$ be two random variables with a bivariate probability distribution.

The covariance of $X$ and $Y$, denoted by $\operatorname{Cov}(X, Y)$, is a measure of the amount of association (or, equivalently, correspondence) and the direction of association between $X$ and $Y$.

The most important measure of the degree of correlation between two variables is the correlation coefficient. It standardizes the covariance in such a way as to eliminate the dependency that covariance has on the scale of measurement for the data set. [2, 187]

### 4.1 General Properties of Correlation Coefficients

A correlation coefficient is said to measure the strength of the relationship between variables. It is not an inferential statistical test. Rather, it is a descriptive statistical measure that depicts the strength of the relationship between two or more variables.

Let $\zeta$ denote a correlation coefficient of $X$ and $Y$.
$\zeta$ should satisfy four key criteria. [3]

1. $\zeta \in[-1,1]$
2. If the larger (smaller) values of $X$ tend to be paired with the larger (smaller) values of $Y \Rightarrow \zeta>0, \zeta \rightarrow+1$ if the correlation is strong. $\Rightarrow \exists$ a positive correlation between $X$ and $Y$.
3. If the larger (smaller) values of $X$ tend to be paired with the smaller (larger) values of $Y, \Rightarrow \zeta<0$ and $\zeta \rightarrow-1$ if the correlation is strong. $\Rightarrow \exists$ a negative correlation between $X$ and $Y$.
4. If the values of $X$ tend to be randomly paired with the values of $Y \Rightarrow \zeta \rightarrow 0 .{ }^{4}$ When $\zeta \approx 0, \Rightarrow X$ and $Y$ are uncorrelated (or, equivalently have no correlation, or zero correlation.)

Correlational information does not provide any conclusions regarding cause and effect, rather it indicates the degree of statistical relationship between two variables. [11, p 72]

The absolute value of $\zeta,|\zeta|$, indicates the strength of the relationship between $X$ and $Y$. As $|\zeta| \rightarrow 1$, the stronger the relationship between $X$ and $Y$, and the more accurately a researcher can predict the value of $y_{i}$ given $x_{i}$. As $|\zeta| \rightarrow 0$, the weaker the relationship between $X$ and $Y$, and the less accurately a researcher can predict a particular $y_{i}$ given $x_{i}$. When $\zeta=0$, the relationship between $X$ and $Y$ cannot be predicted as $\zeta$ is no more accurate than a prediction based purely upon chance.[11, p 945-946]

The sign of $\zeta$ indicates the direction of the relationship between $X$ and $Y . \zeta>0$ indicates a direct relationship and $\zeta<0$ indicates an inverse relationship.

Some general guidelines for the interpretation of $\zeta$ are:

- $\zeta \in[0.7,1] \Rightarrow$ strong direct relationship
- $\zeta \in[0.3,0.69] \Rightarrow$ moderate direct relationship
- $\zeta \in[0.01,0.29] \Rightarrow$ weak direct relationship
- $\zeta \approx 0 \Rightarrow$ no consistent pattern which allows for prediction of one variable's values based upon knowledge of the other variable's values
- $\zeta \in[-0.29,-0.01] \Rightarrow$ weak indirect relationship
- $\zeta \in[-0.69,-0.3] \Rightarrow$ moderate indirect relationship

[^4]- $\zeta \in[-1,-0.7] \Rightarrow$ strong indirect relationship. [11, p 72 ]

This allows us to make a fairly intuitive comparison of the strength of a correlation coefficient to a PIRT ranking: $|\zeta| \in[0.7,1] \Rightarrow$ 'high', $|\zeta| \in[0.3,0.7) \Rightarrow$ 'medium', and $|\zeta| \in[0,0.3)$ 'low'. We use this in our AP600 and LOFT studies, the results of which are discussed in Section 6 on Page 35

As with any statistical computation, it is important to determine the significance of the calculation, or in other words, it is crucial that we determine whether the correlation coefficient that we've computed is statistically significant. To determine whether the correlation coefficient is significant, it is common practice to perform inferential statistical tests to evaluate one or more hypothesis concerning the correlation coefficient. [11, p 946] In our studies, we utilized the p-value, sometimes referred to as a prob value or the associated probability or the significance probability [4, p. 18-19], as the inferential statistical test to evaluate the statistical significance of the correlation coefficient. We considered a p-value $\leq 0.05$ to indicate statistical significance of the correlation coefficient. The null hypothesis, $\mathrm{H}_{o}$, is: $\mathrm{H}_{o}: \zeta=0$, p -value $\leq 0.05 \Rightarrow$ significant evidence that there is correlation between our two variables. On the other hand, for the same $H_{o}$ if the p-value $>0.05 \Rightarrow$ we fail to reject the null hypothesis, i.e. we do not have sufficient evidence that the two variables are correlated. In our studies, if the p-value was too large, we were unable to make any conclusions about that phenomena based on its correlation coefficient.

There are many different methods of calculating correlation coefficients. The three we explore (Pearson, Spearman, and Kendall) are the most frequently used.

### 4.2 Pearson's Product Moment Correlation Coefficient

The Pearson Product Moment Correlation, denoted by $\rho$, is a measure of the linear relationship between $X$ and $Y$. It is defined as: $\rho(X, Y)=\frac{\operatorname{Cov}(X, Y)}{\sigma_{X} \sigma_{Y}}$. It is a correlation coefficient (and thus meets all four criteria specified in the section above.) In the bivariate normal case, $\rho(X, Y)=0 \Rightarrow$ independence of $X$ and $Y$.

The Pearson Product Moment Correlation Coefficient is the most commonly used measure of correlation.[11, p 71]
$\rho$ is invariant under positive linear transformations of the random variables but it is not invariant under all order-preserving transformations, and requires that the two variables have a bivariate normal distribution. [11, p. 947] If $X$ and $Y$ are not approximately normally distributed, then another correlation coefficient should be used.

The most commonly evaluated hypothesis for the PPMCC is: in the population represented by the sample, the two variables have no correlation. [11, p 945]

The statistic computed for the PPMCC will be denoted by $\hat{\rho}$.

The coefficient of determination is $\hat{\rho}^{2}$, and it represents the proportion of variance on one variable which can be accounted for by variance on the other variable.[11, p 953]

For $X$ and $Y$ of sample size $n$, using the notation from section 3.1:
$\hat{\rho}=\frac{\sum_{i=1}^{n} x_{i} y_{i}-\frac{1}{n} \sum_{i=1}^{n} x_{i} \sum_{i=1}^{n} y_{i}}{\sqrt{\left\{\left[\sum_{i=1}^{n} x_{i}^{2}-\frac{1}{n}\left(\sum_{i=1}^{n} x_{i}\right)^{2}\right]\left[\sum_{i=1}^{n} y_{i}-\frac{1}{n}\left(\sum_{i=1}^{n} y_{i}\right)^{2}\right]\right\}}}$. [11, p 950$]$.
We compute the p-values for the Pearson Product Moment Correlation Coefficient by treating $t=\sqrt{\frac{\hat{\rho}^{2}(n-2)}{1-\hat{\rho}^{2}}}$ as coming from a t-distribution with $n-2$ degrees of freedom.[10, p

18-19]

### 4.3 Spearman's Rho

Spearman was developed in 1904 and is a bivariate measure of correlation which is used with rank-order data.[11, p 1061] Spearman's is the application of PPMCC to ranked data.

Spearman's determines the degree to which a monotonic relationship exists between two variables. [11, p 1062]

The underlying null hypothesis for Spearman's Rank-order Correlation Coefficient is: in the population represented by the sample, the correlation between the ranks of the two is 0. [11, p 1063]

The sample statistic computed to estimate the value of Spearman's rho (or Spearman's r) will be dentoed by $\hat{\mathfrak{r}_{s}}$.

For $X$ and $Y$ of sample size $n$, using the notation from section 3.1 and letting $R_{x i}$ denote the rank of variable $x_{i}$, and $R_{y i}$ denote the rank of variable $y_{i}$ :
$\hat{\mathfrak{r}_{s}}=1-\frac{6 \sum_{i=1}^{n}\left(R_{x i}-R_{y i}\right)^{2}}{n\left(n^{2}-1\right)}$.
SAS computes $\hat{\mathfrak{r}_{s}}$ by ranking the data and using the ranks in the Pearson formula. When ties occur, the averaged ranks are used. The p-value for Spearman is computed by treating $t=\sqrt{\frac{{\hat{\mathfrak{r}_{s}}}^{2}(n-2)}{1-\hat{\mathfrak{r}}_{s}{ }^{2}}}$ as coming from a $t$-distribution with $n-2$ degrees of freedom. [10, p 19 ] This is the same way the p-value was computed for Pearson, except that $\hat{\mathfrak{r}_{s}}$ is used in place of $\hat{\rho}$.

### 4.3.1 Generic Example of Computing Spearman's Rho and the Corresponding P -value

Suppose we have $n=4$, and the following points, $\left(x_{i}, y_{i}\right)$, within our data:
$(0.3,1.5),(0.5,2.5),(0.6,2.3)$, and (0.9, 4.6).

Ranking these points, $\left(R_{x i}, R_{y i}\right):(1,1),(2,3),(3,2)$, and $(4,4)$.
$\Rightarrow \hat{\mathfrak{r}_{s}}=1-\frac{6\left((1-1)^{2}+(2-3)^{2}+(3-2)^{2}+(0-0)^{2}\right)}{4\left(4^{2}-1\right)}=0.8$, and $t=\sqrt{\frac{0.8^{2}(4-2)}{1-0.8^{2}}} \approx$
1.88562 , and using a table of selected values $\Rightarrow \mathrm{p}$-value $\approx 0.9$.

### 4.4 Kendall's Tau

Kendall's tau was developed in 1938 and is a bivariate measure of correlation used with rank-order data. The population parameter estimate is denoted by $\tau$. The sample statistic computed to estimate the value of $\tau$ will be represented by $\hat{\tau}$.

Kendall's tau measures the degree of agreement between two sets of ranks with respect to the relative ordering of all possible pairs of subjects/objects.[11, p 1079]

For $X$ and $Y$ of sample size $n$, using the notation from section 3.1 and letting $R_{x i}$ denote the rank of variable $x_{i}$ and $R_{y i}$ denote the rank of variable $y_{i}$.

A pair, $\left(R_{x i}, R_{y i}\right)$ and $\left(R_{x j}, R_{y j}\right)$ is said to be concordant if $\left(R_{x i}-R_{x j}\right)\left(R_{y i}-R_{y j}\right)>0$, or discordant if $\left(R_{x i}-R_{x j}\right)\left(R_{y i}-R_{y j}\right)<0$. Let $n_{c}$ denote the number of concordant pairs of ranks and let $n_{d}$ denote the number of discordant pairs in the ranks. Then $\hat{\tau}$ is defined as: $\hat{\tau}=\frac{n_{c}-n_{d}}{\left\{\frac{n(n-1)}{2}\right\}}$.

Let $t_{k}$ denote the number of tied $x$ values in the $k$ th group of tied $x$ values, $u_{l}$ denote the number of tied $y$ values in the $l$ th group of tied $y$ values, $n$ denote the sample size, and define $\operatorname{sgn}(z)$ as:
$\operatorname{sgn}(z)= \begin{cases}1, & \text { if } z>0 \\ 0, & \text { if } z=0 . \\ -1, & \text { if } z<0\end{cases}$

Let $T_{0}=\frac{1}{2} n(n-1), T_{1}=\sum_{k} \frac{1}{2} t_{k}\left(t_{k}-1\right)$, and $T_{2}=\sum_{l} \frac{1}{2} u_{l}\left(u_{l}-1\right)$.

Then we can define $\hat{\tau}$ as: $\hat{\tau}=\frac{\sum_{i<j}\left(\operatorname{sgn}\left(x_{i}-x_{j}\right) \operatorname{sgn}\left(y_{i}-y_{j}\right)\right)}{\sqrt{\left(T_{0}-T_{1}\right)\left(T_{0}-T_{2}\right)}}$.

Let $s=\sum_{i<j}\left(\operatorname{sgn}\left(x_{i}-x_{j}\right) \operatorname{sgn}\left(y_{i}-y_{j}\right)\right)$.

Define the variance of $s, V(s)$, as: $V(s)=\frac{v_{0}-v_{t}-v_{u}}{18}+\frac{v_{1}}{2 n(n-1)}+\frac{v_{2}}{9 n(n-1)(n-2)}$, where $v_{0}=n(n-1)(2 n+5), v_{t}=\sum_{k} t_{k}\left(t_{k}-1\right)\left(2 t_{k}+5\right), v_{u}=\sum_{l} u_{l}\left(u_{l}-1\right)\left(2 u_{l}+5\right)$,
$v_{1}=\left(\sum_{k} t_{k}\left(t_{k}-1\right)\right)\left(\sum_{l} u_{l}\left(u_{l}-1\right)\right)$, and $v_{2}=\left(\sum_{k} t_{k}\left(t_{k}-1\right)\left(t_{k}-2\right)\right)\left(\sum_{l} u_{l}\left(u_{l}-1\right)\left(u_{l}-2\right)\right)$.
We compute the p -values by treating $\frac{s}{\sqrt{V(s)}}$ as coming from a standard normal distribution. [10, p. 20-21]

### 4.4.1 Generic Example of Computing Kendall's Tau and the Corresponding P -value

To illustrate the computation of Kendall's tau, as well as the corresponding p-value, we will utilize the same example as was used in the generic example of computing Spearman's rho and its corresponding p-value in Section 4.3.1 on Page 16.

Suppose we have $n=4$, and the following points, $\left(x_{i}, y_{i}\right)$, within our data:
$(0.3,1.5),(0.5,2.5),(0.6,2.3)$, and (0.9, 4.6).

Ranking these points, $\left(R_{x i}, R_{y i}\right):(1,1),(2,3),(3,2)$, and $(4,4)$.

The concordant pairs are: $(1,1),(2,2) \&(1,1),(3,2) \&(1,1),(4,4) \&(2,3),(4,4)$ and $(3,2),(4,4)$.

The discordant pair is: $(2,3),(3,2)$.
$\Rightarrow n_{c}=5$ and $n_{d}=1$.
$\Rightarrow \hat{\tau}=\frac{5-1}{\left\{\frac{4(4-1)}{2}\right\}}=\frac{2}{3}$

If we use the second formula for $\hat{\tau}$, we get the same result.
$s=5-1=4, t_{k}=0$, and $u_{l}=0$.
$T_{0}=\frac{1}{2} \times 4 \times(4-1)=6, T_{1}=0$, and $T_{2}=0$.
$\Rightarrow \hat{\tau}=\frac{5-1}{\sqrt{(6-0)(6-0)}}=\frac{2}{3}$.

We can now compute $V(s)$.
$v_{o}=4 \times(4-1) \times(2 \times 4+5)=156, v_{1}=0$, and $v_{2}=0$.
$\Rightarrow V(s)=\frac{156-0-0}{18}=\frac{26}{3}$
$\Rightarrow \frac{s}{\sqrt{V(s)}}=\frac{4}{\left\{\sqrt{\frac{26}{3}}\right\}} \approx 1.35873$

Using a table of values for the standard normal distribution, $\Rightarrow \mathrm{p}$-value $\approx 1-0.3708=$ 0.6292 .

### 4.5 Discussion of Pearson, Spearman, and Kendall

It is recommended that for interval or ratio data in the bivariate case that the Pearson product-moment correlation coefficient be used, for ordinal or rank order data of bivariate case that Spearman's rank-order correlation coefficient or Kendall's tau be used.[11, p 117]

Pearson is a stronger correlation coefficien ${ }^{5}$ than Spearman or Kendall, but has additional restrictions, such as the requirement of parametric data ${ }^{6}$ Spearman's rho is more commonly used than Kendall's tau for two primary reasons. The computations for tau tend to be more tedious than those completed when calculating rho.711, p. 1080] The second reason is that when the sample is derived from a bivariate normal distribution, Spearman's rho tends to provide a reasonably good approximation of Pearson, while Kendall's tau will not. An advantage of Kendall's tau is that it has a sampling distribution that approaches normality very quickly. Spearman's rho, on the other hand, needs a fairly large sample size to employ the normal distribution to approximate the sampling distribution of rho.[11, p. 1080] As a general rule, $\left|\tilde{\tau}_{c}\right|<\left|\tilde{\mathfrak{r}}_{s}\right|$ for a set of data, and $\left.\frac{\hat{\tau}}{\hat{\mathfrak{r}_{s}}} \rightarrow(0.67)^{2} \cdot[11, \text { p. } 1080]^{8}\right]$

In Sections 4.3.1 and 4.4.1 we had $\hat{\tau}=\frac{2}{3}$ and $\hat{\mathfrak{r}_{s}}=0.8, \Rightarrow|\hat{\tau}|<\left|\hat{\hat{r}_{s}}\right|$, which follows the general rule. However, $\frac{\hat{\tau}}{\hat{\mathfrak{r}_{s}}} \approx 0.83333 \neq(0.67)^{2}$, yet, it does not disprove the general rule as we just used a generic example which had a very small sample space.

[^5]
## 5 Experimental Information

### 5.1 RELAP5-3D

### 5.1.1 Terminology and Basic Deck Requirements[7, p. A1-1]

| Term/Accronym | Definition/Description |
| :---: | :--- |
| Card | Best-Estimate |
| an 80-character record in the input deck |  |
| Comment Card | the first field on the data card. It is an unsigned <br> integer, and is used to specify the component number <br> the card supplies information about. |
| identified by a '*' or a ' $\$$ ' as the first nonblank |  |
| character. Blank cards are treated as comment cards. |  |
| Continuation Card | With the exception of printing out their contents, <br> there is no processing of comment cards. |
| identified by a '+' as the first nonblank character on |  |
| the card, may follow a data card or another |  |
| continuation card. Fields on each card must be |  |
| completed on that particular card (i.e. a field may not |  |
| continue from one card onto the next.) |  |

A RELAP5-3D input deck typically consists of at least one title card, optional comment cards, data cards and a terminator card.

### 5.1.2 Safety Analysis of NPPs \& BE Codes

To ensure safety in the reactor design, operator training, and upgrades to the Nuclear Power Plant (NPP), Best Estimate (BE) codes are frequently used to model NPPs and to analyze their safety.

A Best Estimate code is a system code that is capable of predicting/ modeling physical phenomenon, free of any conservatism regarding selected acceptance criteria, and able to provide a sufficiently detailed model to describe the relevant processes. 9, p. 193] RELAP53 D is the BE code used in this study.

### 5.1.3 About

RELAP5-3D is a fully integrated, multi-dimensional thermal-hydraulic and kinetic modeling program. It is a BE code developed at the INL, which serves as a modeling and simulation tool to support engineering design and safety analysis of nuclear reactors. It is also used for non-nuclear fields such as fossil power plants, oil and gas pipelines, municipal steam supply systems.

### 5.1.4 Running RELAP5-3D

Upon performing the calculations of the input model, RELAP5-3D produces three output files: a '.p' file, a '.plt' file, and a '.r' file. The '.p' file is a printed output file, the '.plt' file is a plot file, and the '. r ' file is the restart file. In our studies, we were primarily concerned with examining the printed output file. We wrote a Python script, which is described in Section 5.5 on Page 33. This script locates our desired ouput parameter value from the '.p' file, and then places that value into a parameter value file which is created with the input modification program described in Section 5.4 on Page 29

### 5.2 AP600 Deck



### 5.2.1 Background on the AP600 Nuclear Power Plant

The AP600 Nuclear Power Plant (NPP) was designed by Westinghouse as a part of the cooperative U.S. Department of Energy (DOE) and the Electric Power Research Institute (EPRI) Advance Light Water Reactor Program (ALWR). 12]

For safety, the AP600 relies on operation of passive systems. The Phenomena Identification and Ranking Table (PIRT) compiled by Burtt, et al (used in this paper, see the references section, number [1]) addresses AP600 behavior expected during small break loss-of-coolant, main steam line break, and steam generator tube rupture accidents. [1]

### 5.2.2 Variable Names, Representation, \& Corresponding PIRT[1] Phenomena

| Variable | Representing | Corresponding to (in PIRT) | PIRT <br> Ranking |
| :--- | :--- | :--- | :--- |
| VAR1 | Fission- <br> Product Yield <br> (i.e. <br> decay-heat) | Core Power | High |
| VAR2 | Roughness | Flow Resistance in the IRWST | High |
| VAR3 | Form Loss | Flow in Accumulator (pressurized <br> volume) | High |
| VAR4 | Form Loss | Flow in Accumulator (pressurized <br> volume) | High |
| VAR5 | Area (ADS-4) | ADS Energy Release | High |
| VAR6 | Area (Break <br> Valve) | Mass Flow in Break 9 | High |
| VAR7 | Area (Orifices) | Injection Line - CMT | High |
| VAR8 | Initial Level of <br> Pressure | Level in CMT | High |
| VAR9 | Roughness | ADS Flow Resistance | Medium |
| VAR10 | Roughness | ADS Flow Resistance | Medium |
| VAR11 | Fouling Factor | PRHR (Heat Transfer between <br> PRHR \& IRWST) | Medium |
| VAR12 | Fouling Factor | Steam Generators (Primary to <br> Secondary Heat Transfer) | Medium |
| VAR13 | Roughness | PRHR-Flow Resistance \# 1 | Low |
| VAR14 | Roughness | PRHR-Flow Resistance \#2 | Low |
| COREMIN | Control <br> Variable 116 | Minimum Value of the Core <br> Collapsed Liquid Level 10 | N/A |

### 5.2.3 Studies with the AP600 Deck

Using the AP600 deck, we conducted four studies. These studies were based upon the size of break in the cold leg. Each of the 13 variables ${ }^{117}$ had three values which were evenly distributed throughout the interval (i.e. a minimum, a mean, and a max). Variable 6 denoted the size of the break and was used to create the four different studies. To have

[^6]every possible combination of variable values, this would mean creating $3^{13}=1,594,323$ input files.

Performing even a single study with $3^{13}$ runs of 400 seconds would require 7,381 days on a single processor. Even utilizing the INL cluster, described in Section 5.5 on Page 33, which has 32 cores per node and 12 nodes, it would require 19 days to run the study. There are many ways to reduce the number of runs in a study. We decided to investigate the use of grouping our studies into smaller sets to reduce the number of required runs for each study. We first separated each study into two or three sets. The variables that had correlation coefficients that were ranked high or medium among the sets would then be grouped together and run again. Through experimentation, we found that separating the studies into smaller sets did not significantly affect the calculations of the relative correlation coefficients and substantially reduced the amount of time needed to run our studies.

The size of the break dictates how long the transient runs. In order to further decrease the run-time of our studies, we plotted each of the four transients ( 2 inch, 4 inch, 6 inch, 8 inch) to determine what the duration of the transient should be for each respective study. The duration of the transient that was selected for each study encompassed the time when the collapsed core level reached its lowest height. These plots can be seen in Figures 2, 3, 4. and 5 , respectively ${ }^{12}$ We determined that it was effective to run the 2 inch break transient for 400 seconds, the 4 inch break transient for 200 seconds, the 6 inch break transient for 200 seconds, and the 8 inch break transient for 75 seconds.

The results from the 'top study' of each of the cold-leg break studies are presented in Section 6.1.1 on Page 35.

[^7]

Figure 2: 2 inch Break Transient Plot


Figure 3: 4 inch Break Transient Plot

Our first study that we conducted was on a four inch break. To decrease run-time, we broke the study into two sets: the first set had only variables $1-5$ and $7-8$ (i.e. the PIRT-ranked 'high' variables, note that as stated previously, variable 6 is the break size and thus will not be adjusted within the 4 inch study) varying, while the other 6 variables were held at the nominal (or mean) value. The first set was $3^{7}=2,187$ input files. The second set had only variables $9-14$ (i.e. the PIRT-ranked mediums and lows) varying, while the other 7 variables were held at their respective nominal values. The second set consisted of $3^{6}=729$ input files. After both sets were complete, and analyzed using SAS, we created a new study, which was based on the top 5 most correlated variables from both 4 inch studies: VAR1, VAR8, VAR11, VAR12, and VAR13. Just as before, all five variables varied between the three values while the other 8 variables were held at their respective nominal values. This created $3^{5}=243$ input files.

Our second study was on a 2 inch cold leg break. We put this study into three sections (the other variables are held at their respective nominal values): variables 1-5 varying $\left(2^{5}=243\right.$ input files), variables $7-10$ varying ( $2^{4}=81$ input files ), and variables 11-14 varying ( $2^{4}=81$ input files). After all three sets were analyzed using SAS, we created a new study, based on


Figure 4: 6 inch Break Transient Plot


Figure 5: 8 inch Break Transient Plot
the top 4 most correlated variables from the three sets: VAR1, VAR8, VAR12, and VAR13. As with the four inch study, we varied these variables between the three variables, while the other 9 variables were held at their respective nominal values. This created $3^{4}=81$ input files.

Our third study was on a 6 inch cold leg break. We split the study into 3 sections, which were identical to the sections for the 2 inch break study. After the three sets were analyzed with SAS, a new study was created based on the top 5 most correlated variables from the three sets: VAR1, VAR7, VAR8, VAR12, and VAR13. These were varied as with the 2 inch and 4 inch break studies: the 5 variables were varied by 3 values, and the other 8 were held constant at their respective nominal values. This created $3^{5}=243$ input files.

Our final study of the AP600 deck was on an 8 inch cold leg break. As with the 2 inch and 6 inch break studies, we split the 8 inch study into the same sets. After the three sets were analyzed using SAS, a new study was created based on the top 4 most correlated variables from the three sets: VAR1, VAR8, VAR12, and VAR13, creating $3^{4}=81$ input files for this section of the study.

In all of the studies, there were variables which had their respective p -value $>0.05$. This does not indicate that these variables were statistically insignificant, or that they were unimportant to the minimum core level. Rather, it indicates that we cannot make a conclusion regarding their correlation to the minimum core level, and that further investigation is needed for these variables.

### 5.3 LOFT Deck

### 5.3.1 Background Information on the LOFT Nuclear Power Plant

Following the 1988 revised emergency core cooling system rule for LWRs which allows the use of best estimate plus uncertainty methods in safety analysis, a study was conducted at the Idaho National Laboratory (INL) based on a cold leg large break loss of coolant accident test in the Loss of Fluid Test (LOFT) experimental facility. The LOFT facility was a 50 MW PWR that was designed to model a large break loss of coolant (LBLOCA) in a commercial pressurized water reactor (PWR). The facility was operational from the late 1970's to the mid 1980's. 15]

The input deck used in this study was the same input deck used by G.E. Wilson and C.B. Davis in their "Best Estimate Versus Conservative Model Calculations of Peak Clad Temperature: An Example Based on the LOFT Facility" paper (see reference [15].)

### 5.3.2 Variable Names and Corresponding Phenomena [15]

| Group Number | Variable(s) | Representing |
| :---: | :--- | :--- |
| 1 | VAR1-VAR24 | Peaking Factor (i.e. core power fractions) |
| 2 | VAR 25 | Fuel Clad Gap Width |
| 3 | VAR26-VAR43 | Fuel Thermal Conductivity |
| 4 | VAR 44 | Clad to Coolant Heat Transfer (i.e. fouling <br> factor) |
| 5 | VAR45-VAR46 | Break Discharge Coefficient |
| 6 | VAR47-VAR51 | Pump Degradation |
| $\mathrm{n} / \mathrm{a}$ | Control Variable <br>  234 | Peak Clad Temperature |

### 5.3.3 LOFT Studies

We conducted two studies using the LOFT deck. Both studies values were selected based upon G.E. Wilson and C.B. Davis' paper. For both studies, the key output parameter in the study was Peak Clad Temperature (PCT).

In the first study, each of the six groups of variables had 3 points uniformly distributed (i.e., minimum, mean, and maximum). All groups used minimum, mean, and maximum for their three values. However, the variables within each group often had differing maxima, minima, and means. For example, in group one there were 24 different sets of minimum, mean, and maximum, yet all variables in the group needed to vary in the same way. For instance, each of the 24 variables in group one all needed to experience their respective minimum simultaneously. Likewise, each of the 24 variables in group one had to achieve their mean simultaneously. Similarly, the 24 variables all experienced their respective maxima simultaneously. There were $3^{6}=729$ input files in the first study.

In the second study, each of the six groups had 5 different values (i.e., minimum, lowmean, mean, high-mean, and maximum.) The groups of variables functioned the same way for the second study as the first study, just with additional values. In other words, for

[^8]group one, the 24 variables were varied simultaneously, with each variable within the group experiencing it's particular low-mean simultaneously. Likewise, each of the 24 variables experienced its respective high-mean simultaneously. There were $5^{6}=15,625$ input files in the second study.

The results from both studies with the LOFT deck are presented in Section 6.2 on Page 36

### 5.4 Input Modification Program

In order to run the AP600 study and the LOFT study, a Fortran $90^{\circledR}$ program was created. This program uses a marked input deck and study specification file(s) to create input files with input values varying as specified in the specification file(s).

### 5.4.1 Marked Input File

After an input deck ${ }^{[14}$ and the variables of interest (sometimes referred to as input parameters) as well as the key output parameter have been selected, the user must determine how these selected input parameters should vary. Once that is determined, the user should create a 'spec file' (or Specification File). Specification Files are discussed in Section 5.4.2. The base input deck should then be 'marked' by the user- this consists of locating the variables of interest (i.e. card number and word) within the deck, and then placing a ' $\$ X_{X X X} \sqrt{15}$ in place of the current value on the card in the deck ${ }^{\left[{ }^{16}\right.}$ This process should be undertaken simultaneously with the creation of the 'spec file' so as to avoid mistakes. ${ }^{17}$ After the base deck is completely marked, we refer to it as the "marked input deck."

[^9]
### 5.4.2 Specification Files

The specification file(s) or 'spec file(s)', contains all of the necessary information for a given study. The first line of a spec file indicates the type of spec file it is. There are two types of spec files: (1) a Generator Specification File and (2) a Node Specification File. These are discussed in the Subsections 5.4.2(a) and 5.4.2(b) respectively. Both types of spec files contain: the base name of the files that are generated (this is also the name (without the extension) of the marked input file), the number of variables in the study, the number of nodes in the study ${ }^{18}$, which node (or set) of this spec file corresponds to, the number of groups in the study ${ }^{19}$, as well as variable nam ${ }^{20}$, the minimum value of the variable, maximum of the variable, standard deviation of the variable, number of variations of the variable (i.e. number of points for the variable), the distribution type of the variable, and the group number the variable belongs $t{ }^{211}$. A template of a spec file can be seen in Appendix B on Page 51.

### 4.5.2(a) Generator Specification File A "Generator Specification File" or "generator

 spec file" is a spec file with generate on the first line of the spec file. When combined with the executable produced by the input modification program, described in Section 5.4.3, it produces a specified number of node spec files, each of which has a different node number ${ }^{[22}$,The node number corresponds to the INL cluster node, where each node has multiple cores.
The INL cluster is discussed more in Section 5.5 on Page 33. To see an example of a

[^10]generator spec file, see Appendix Con Page 52 .

The motivation for generating separate node spec files was cluster supercomputer limitations. Generating thousands of input files on a single processor, then moving them to the nodes where they would run ties up communication resources need for other data flow and can seriously impair performance. It is far more efficient to generate the RELAP5-3D input files on the nodes where they will run, thereby eliminating all such input file movement.
4.5.2(b) Node Specification File A "Node Specification File" or "node spec file" is a spec file with run on the first line of the spec file. When combined with the executable produced by the input modification program, described in Section 5.4.3, it produces the input files corresponding the particular node number of the node spec file. To see an example of a node spec file, see Appendix D.

### 5.4.3 Main Program

The main program, input_mod_gen.f90, is included as Appendix $A$ on Page 40. Its purpose is to create an entire statistical study by generating input relating to a PIRT analysis. It has three purposes:

1. Interpret the "generator specification file" and create a number of "node specification files."
2. Produce all RELAP5-3D input files for a given node of the cluster supercomputer from the "node specification file" and a single RELAP5-3D input template file.
3. Handle the situation where a group of input parameters vary together (are 100\% correlated).

The main program is run first to create a node specification file for each cluster node; a node may have up to 32 cores that can each run RELAP5-3D independently and simultaneously.

The creation of the node spec files from a generator spec file is done through the command line prompt:
input_mod_gen.exe -i generator_spec_file_of_my_study

This creates the corresponding node spec files and these node spec files are exported to each node along with a copy of the template input file. The main program is run on each node to produce the specific input files of the statistical study, and is run on the cluster via a Python script which is described in Section 5.5 on Page 33 . The complete set of possible combinations of input values can be ordered as a set of $n$-tuples. The set of n tubles is generated by a recursive algorithm and therefore works for any number of input parameters.

The ordinal number of the n-tuple that is used to generate an input file is assigned as the input file's sequence number and becomes part of its name. Concatenating its sequence number to the base name of the statistical study forms its name. All input files have a unique name, regardless of the node on which it runs. The input files are created by copying the template input file then substituting the variable values corresponding to its n-tuple into the input file for that variable's marker. Markers had the form \$VAR1, \$VAR2, ... $\$$ VAR $K$. When some variables are $100 \%$ correlated, $K$ is larger than $n$ (the size of the n-tuples).

The values of the correlated variables are varied together. For example, if group one has 20 variables and a uniform distribution with 3 "levels" in use, namely minimum, mean, and maximum value, the group counts as one variable in the first position of the n-tuple previously discussed. The group has three levels: minimum, mean, and maximum. However, when an input file is generated, the particular "maximum value" of each variable is substituted for its marker (\$VAR1 through \$VAR20). Input files are created on node 3 with the following command:

> input_mod_gen.exe -i node3_spec_file_of_my_study
. The number of input files generated is a function of the number of groups and the number of value levels within the group. For example, if group 1 has 6 variables and 3 levels, group 2 has only one variable with 8 levels, and group 3 has 9 levels, there would be $3 * 8 *=216$ combinations in the entire statistical study. The program determines the number of input files that are created by each node spec file based upon the \#nodesInStudy (combSet in program) by taking the celing function of $\left\{\frac{\# \text { of Input Files in Study }}{\# \text { nodesInStudy }}\right\}$. The second LOFT study had 6 groups of variables, each with 5 values for a total of 15,625 runs. Splitting these runs among 9 node spec files placed 1736 runs each on nodes 1-8 and 1737 runs on node 9 .

### 5.5 Running on the Cluster \& Python Scripting

Studies were run on one of the INL's supercomputers, Quark, which has 12 nodes. Users submit jobs, such as our statistical studies, via a batch queuing system.

Due to limited resources particularly on moving large amounts of data from the head node to computational nodes, it is much more efficient to generate the input files on the node as explained in Section 5.4.3 on Page 31.

The Python scripts are included in Appendix E on Page 53. It runs the main program on the head node of the cluster, moves the node specification file onto the cluster's computation nodes, runs RELAP5-3D in parallel on the input files, collects the output from the RELAP53D output files, and plaes it in a "study output file" on the line corresponding the input's n-tuple. These files are ported back to the head node and are then combined into a single "study output file."

Thereafter, SAS can be used to perform the statistical analysis.

### 5.6 Using SAS ${ }^{\circledR}$

Once all of the res files have been created, as described in Section 5.5, we copy them into the local desktop, relabeling the files' extensions as '.csv' (as opposed to '.res'), and adding an additional line to the beginning of the file which typically looks like 'runNumber,VAR1,VAR2,VAR3,....,KEYOUTPUT'. We then import each file into SAS, a statistical software package. We used the SAS package, SAS EnterpriseGuide 4.3, which offers a graphical user interface of SAS 9.2. We then combined the res files from all of the nodes into one data set, through utilization of the 'append table' feature, "tasks $\backslash$ data $\backslash$ append table". We then calculated the correlation coefficients using the correlation coefficient function, "tasks $\backslash$ Multivariate $\backslash$ Correlation Coefficient". We selected the variables as analysis variables, and correlated them with the key output parameter. The options we selected were Pearson, Hoeffding, Kendall, and Spearman, and we included the Pearson correlation options of covariances, sums of squares and crossproducts. These produced SAS reports, which are found in Appendix F. We were then able to rank the variables accordingly.

## 6 Results

### 6.1 AP600 Studies

### 6.1.1 AP600 Correlation Coefficient Classification

| Break | High $\zeta$ | Medium $\zeta$ | Low $\zeta$ |  |  |  |
| :---: | :--- | :--- | :--- | :--- | :--- | :--- |
| 2 inch | Core Power <br> (PIRT High) |  | PRHR-Flow Resistance \# 1 (PIRT Low) |  |  |  |
| 4 inch | Core Power <br> (PIRT High) |  |  |  |  |  |
| 6 inch |  | Core Power <br> (PIRT High) <br> Level in CMT <br> (PIRT High) | SG-Heat | Transfer | (PIRT | Medium) |
|  |  | PRHR-Flow <br> Resistance \#1 <br> (PIRT Low) |  |  |  |  |
| 8 inch $^{23}$ | Core Power <br> (PIRT High) <br> Level in CMT <br> (PIRT High) | SG-Heat | Transfer | (PIRT | Medium) |  |

### 6.1.2 Discussion of the Rankings of the Phenomena

All variables which were not listed in Section 6.1.1 had their respective p-value $>0.05$.
In the 2 inch and 4 inch studies, the findings were as we expected based upon the PIRT recommendations. In the 6 inch and 8 inch studies, there were some surprising results. In the 6 inch study, two PIRT-ranked 'highs' had a correlation coefficient classifications of 'medium', and a PIRT-ranked 'medium' had a low correlation coefficient classification, of particular interest was that one PIRT-ranked 'low' had a medium correlation coefficient classification. In the 8 inch study, we saw a similar ranking pattern to that of the 6 inch study, with the exception being that all rankings in the 8 inch study were less than those of the PIRT, specifically, none of the rankings of the correlation coefficient classification

[^11]were greater than that of the PIRT. The rankings we saw in the 6 inch study (and in the 8 inch study) which were less than or equal to the PIRT rankings do not cause too much concern as the PIRT was conservative, i.e. if the panel was unsure as to whether something should be ranked as 'high' or 'medium', they ranked it as 'high'. The result that surprised us was the phenomena in the 6 inch study which was ranked higher than the PIRT ranking in terms of correlation coefficients.

### 6.2 LOFT

### 6.2.1 LOFT Correlation Classification

Classification Group Name(s)

| High $\zeta$ | Fuel Clad Gap Width |
| :---: | :---: |
| Medium $\zeta$ | Clad to Coolant Heat Transfer \& Peaking Factor |
| Low $\zeta$ | Break Discharge Coefficient \& Fuel Thermal Conductivity |
| p-value $>0.05$ | Pump Degradation |

### 6.2.2 Ranking the Phenomena Following the Correlation Computation

After completing the correlation analyses for the LOFT study, we were able to numerically rank the phenomena from 1-5, with 1 representing the variable which was most strongly correlated with PCT and 5 being the variable which was least correlated with PCT. From an engineering perspective, we would rank the phenomena by absolute value of the change in PCT from the minimum value to the maximum value of the phenomena. To approximate that computation, we held the other phenomena at their respective nominal values. $\approx \Delta \mathrm{t}$ corresponds to the approximate change in temperature when we compare the PCT of the minimum value of the variable to the PCT of the maximum value of the variable, where all other variables are held constant at their respective nominal values. The following table presents the rankings:

| Rank | Phenomena | $\approx \Delta \mathbf{t}($ Rank by $\approx \Delta \mathbf{t})$ |
| :---: | :--- | :--- |
| 1 | Fuel Clad Gap Width <br> (G2) | $200(1)$ |
| 2 | Clad to Coolant Heat <br> Transfer (G4) | $80(3)$ |
| 3 | Peaking Factor (G1) | $50(4)$ |
| 4 | Break Discharge <br> Coefficient (G5) | $90(2)$ |
| 5 | Fuel Thermal <br> Conductivity (G3) | $45(5)$ |

The ranking by correlation coefficients almost matches the ranking we get by considering $\Delta$ t- with the exception being that group 5 should be ranked number 2 , and then the others would adjust accordingly. This seems to demonstrate that ranking using the correlation coefficients is fairly reasonable. Additionally, it may be worth noting that by the Apendix K ruling of the NRC, all of these correlation coefficients change in temperature would require that these phenomena be ranked 'high.'

## 7 Final Conclusions and Potential Future Work

Overall, in the AP600 studies, we found that Core Power was the dominating input parameter which most strongly affected our key output, but found that the significance of the variables after that were largely dictated by the break size. For the most part, we found that our statistical rankings were generally the same or less than the PIRT ranking, with the exception of VAR13 in one transient (the 6 inch break.) This makes sense as the PIRT was conservative, i.e. if the experts doing the ranking were unsure as to whether or not a particular phenomenon was medium or high, they tended to rank it high. With regards to the one odd case of the variable 13 ranking medium in the 6 inch break, this indicates that we should further examine this variable and phenomenon. It may be that changing this variable actually relates to another phenomena not covered in the PIRT, or that this particular variable is more complex than we originally thought. One of the limitations of our study is that we cannot be sure how accurately our variables represent the phenomena. We
were limited by what quantities it made sense to adjust, as well as the fact that there were cases where we were unable to easily select variables to vary to correspond to phenomenon. In any case, the study does demonstrate great potential for further studies, in particular in applications to other PIRT analyses.

In the LOFT studies, we found that fuel clad gap width was most strongly correlated with the PCT, and interestingly enough, found that the pump degradation seemed to have no relation to Peak Clad Temperature.

Through utilization of PIRTs and statistical methods, we believe that a more accurate uncertainty analysis can be performed.

Further studies may demonstrate that the utilization of statistical methods may improve upon the accuracy of PIRTs produced by a panel of experts. Other future work may utilize the concepts of sampling reduction techniques to vary a greater number of parameters which maintaining a relatively small run-time for studies. Additional improvements to the input file generator may include the addition of the ability for the user to specify a larger number of distributions for the values of various variables.

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## Appendices

## A Input Modification Program

```
program input_mod
!COGNIZANT: Alexandra E. Gertman
!CREATED: 2/22/2012
!UPDATED: 5/02/2012
!PROGRAM DESCRIPTION
    This program was created to generate many input files with varying
    parameters (i.e. phenomena or variables).
    The program generates an executable (a.out) upon compilation.
    The executable (a.out) is combined with a marked input file as
```

```
well as a study specification file (spec file) to create the
    new input files.
    A spec file includes:
    -if the spec_file should generate new spec_files OR
        if it should generate input files
    -the base name of each generated input file
        (base name also serves to indicate the name of
        the marked input file)
    -the number of variables
    -the number of jobs the combinations will be split
    into
    -which set of combinations this particular spec_file
    will generate
    -the number of groups of variables
        (there are some cases when variables MUST be changed
        simultaneously- in those cases, those variables
        will all be grouped together)
    -the name of each variable
    -variable information such as distribution which
        allow the program to make necessary calculations
        to calculate the values each variable in the
        spec file should experience, as well as the group
        number each variable belongs to.
    The executable will produce a file containing the run information
    (run_layout) which specifies the number of input files that will
    be built in each of the 'input file generator, spec files.
    If the spec file is a 'spec file generator, spec file, when
    compiled with the executable, it will create 'input file
    generator, spec files (the number of which is specified in the
    'spec file generator, spec file
    If the spec file is an 'input file generator, spec file, when
    compiled with the executable, it will create new input files
    based upon the marked input file and the specifications in
    the spec file itself. Once compiled using the executable, it
    will create a comb file (which it writes to as it creates each
    of the input files) that contains all of the 'run information
    for each of the input files being generated.
!PROGRAM OUTLINE:
!1. Initialize variables, obtain and bypass the name of the executable file,
    and read the command line option.
!2. Open the files
    2.1 Open spec_file - a file that has info about the study
        2.1.1 Read spec_file and the following information
            2.1.1.a. Whether the spec file is an 'input file generator, OR
                a 'spec file generator.'
            2.1.1.b. BASE Name for the input files
            2.1.1.c. Number of variables, number of 'input file generator'
                spec files that will be used in the study (i.e. njobs), which
                particular 'input file generator, the spec file is (i.e
                combSet), and the number of groups of variables in the study
            2.1.1.d. Name of each variable (as it is marked in the file- except
                for the '$'- that is added immediately after the data is
                read-in. i.e. 'VAR1', becomes '$VAR1'.)
            2.1.1.e. Minimum and Maximum values of each variable
            2.1.1.f. Standard Deviation of each variable
            2.1.1.g. Number of data points within the range
            2.1.1.h. Probability function name (uniform, nromal, lognormal, ....).
                NOTE: Currently only uniform is available.
    2.1.1.i. Group number each variable belongs to
    2.1.2 Calculate the number of possible combinations, and create an array based on njobs
        and nposscombs which specifies which runs will take place on each combSet.
    2.1.3 Create the file run_layout. Write nposscombs on line 1, and then write the
        number of combinations in each combSet onto line 2. The file is then closed
    2.1.4 Based on the first line of the spec_file, one of 3 things will occur
        (1) If the spec_file's first line says "generate" send to the subroutine
        spec_gen. spec_gen will generate njobs of spec_files which have "run"
        instead of "generate" and have different combSet numbers. i.e. only
        the first two lines of the newly generated spec_files will differ
        the first two lines of the newly generated spec_files will differ rill
        from the o
        (2) If first line doesn't say 'run' or 'generate,', program will terminate
        (3) If the first line is "run", the program will execute normally. i.e.,
                Follow the remainder of the outline.
    2.2 Name of marked_input_file is created.
    2.3 Open marked_input_file
3. Generate input files
3.1 Calculate the variable values for each interval with 'uniform' distribution.
    3.1.1 Calculate the interval length and increment size
```

```
    3.1.2 Use the probability function, minimum, interval length,
        and increment size to calculate all the values the variable
        will have and save these in an array
3.2 Input file generation through loops and subroutines
    3.2.1 Make columns an array of ones (of size ngroups.)
    3.2.2 Open var_val_comb (i.e. comb file)
3.3 Input file name generation and input file creation through subroutines
    3.3.1 Generate a file names based on combination number or var value
        For example: 'edhtrk99' vs 'edhtrk_11_3_3
    3.3.2 Call the subroutine index_gen to generate all the files
            -As each file is generated, the run number (i.e. comb_number) and
                the values for each variable in that particular run/input file will
                be written to the comb file
4. Close all files (with the exception of file 8, fileName, which is closed in the
subroutine copy_file or the subroutine spec_gen).
Data Dictionary
baseFileName = a character (of size 10) indicating the name of RELAP5-3D
                        input file, will serve as beginning letters of all generated
                        input files. Additionally, it will serve as the beginning of
                        the name of the file var_val_comb
charCombNum = a character (of size 8) which is a character conversion of
    the combination number of the run so that each line in the
    corcombination number of the run so that each line 
    specific combination of variables. It is a local variable
    in the recursive subroutine index_gen. It is padded with 0's
    to ensure that each run is EXACTLY 8 digits
charCombSet = a character of size 3 which is the character conversion of
    combSet. It is used so that the spec_file can be appended with
    the number corresponding to the set of combinations the spec
    file needs to generate. It,s a local variable used in the
    subroutine spec_gen. It's also used in the main program to
    make the comb file specific to the combSet
char_i = a character (of size 8) which is a character conversion of
    columns(1:nvar) or (a character conversion of the combination
    number of the run) so that newFile can be appended with each
    particular variable's corresponding index for that
    combination. It is a local variable used in the subroutine
    name_gen, and is padded with 0's to ensure it each name is
    baseFileName_XXXXXXXX, where XXXXXXXX is EXACTLY 8 digits.
charlen = a local integer in subroutine var_finder which is used to store
    the character conversion of the length of a particular var(i).
    It is used to re-write line with the variable value substituted
    in place of '$VARX'
charvalu = a local variable (of size 12) in subroutine var_finder which is
    the character conversion of valu. It allows valu to be appended
    to the file name
columns = an array of integers (of size nvar- originally set to 55) which
    stores the indices of variables in any given combination.
    stores the indices of variables in any given combin cor variable j
comb_number = an integer indicating the combination number a particular
    generated input file corresponds to
combSet = an integer of size 3 indicating which particular set of
    combinations the given spec_file will need to generate.
ex = logical flag indicating existance of spec_file and
exfile = a character of size 10 indicating the name of the executable
    marked_input_file
    file (usually a.out unless otherwise specified at run time for
    input_mod.f90. NOTE: if the executable is changed, the user is
    advised to make corresponding changes to the python script
    (run_xxxx.py) if tests will be made on the INL cluster
flag = a character of size 2. It is entered by user after executable file
If flag is not -i program will send error message to user and quit
found = a logical variable, used in the subroutine copy_file to test whether
o or not '$VAR' occured in the line (card) being copied in copy-file.
genRun = character of size '10' which should be either 'run' or 'generate',
    = character of size it is found on the first line of the spec_file and if genRun is
                        It is found on the first line of the spec_file and if genRun is , in , generate, the program will call the subroutine spec-gen. If it,s
                        ,run, then the program will run normally
groupNumb = integer array of size 55 which stores the group number of each var.
i
    = integer used in various arguments (i.e. getarg, do loops, etc).
ierr = integer used to detect an input error in reading the spec file,
    marked input file fileName (subroutine copy file and subroutin
    marked_input_file, fileName (subroutine copy_file and subroutine
    spec_gen), and var_val_comb files
increment_size_var = array of real numbers indicating the size of increment each
    array of real numbers indicating the size of 
colindex = integer indicating the index of a particular column. It is used in
    if statements in the subroutine indexgen
int_length_var = array (of size 55) of real numbers indicating the size of the
    range of the values for each particular variable.
is = integer used to denote the number of command line arguments when
    the executable file is run (usually a.out is the executable)
j = integer used in various arguments (i.e. do loops, etc.)
job_inc = real number which is used to determine the number of jobs that
    should be in each combSet. (i.e. job_inc = nposscombs/njobs).
```



```
where
= an integer corresponding to a local variable in subroutin
    var_finder which holds the value returned by the index function
    (either a '0' or a',') which indicates if the variable flag
    (ex: $VAR1) is within a particular line
```

files
file unit number $=$ file name
$\begin{aligned} 1= & \text { spec_file (opened and closed in input_mod, utilized in the } \\ & \text { subroutine spec_gen if genRun }=\text {, generation } \text {.) }\end{aligned}$
$\begin{aligned} 1= & \text { spec_file (opened and closed in input_mod, utilized in the } \\ & \text { subroutine spec_gen if genRun }=\text {, generation } \text {.) }\end{aligned}$
$2=\underset{\text { the subroutine copy file.) }}{2}$ (opened and closed in input_mod, read in
the subroutine copy_file.)
var_val_comb file (opened and closed in input_mod, written
$3 \quad=\begin{aligned} & \text { var_val_comb file (opened and closed in } \\ & \text { in the recursive subroutine index_gen.) }\end{aligned}$
$8=$ fileName. This is the file which is a copy all of information
$=\begin{aligned} & \text { filen mer } \\ & \text { from the marked in put_file or the information for spec_file if }\end{aligned}$
from the marked_input_file or the information for spec_file it
genRun $=$ generate ${ }^{\prime}$. (It is opened, written, and closed
genRun $=$ 'generate, (It is opened, written, and closed
in the subroutine copy_file, or in the subroutine spec-gen.)
$9 \quad=r u n \_l a y o u t$. This is the file which contains the information about
the runs it allows the user to double check that the comb files
generated contain the correct number of runs. The first line is
nposscombs. The second line is the array runLayout, with each
integer in the array being size 4. It was originally created to
integer in the array being size 4. It was originally created to
be used in the python script, but we ended up not using it in
be used in th
program/subroutine names and descriptions
name $\quad=$ description
copy_file $\quad=$ subroutine which is called by index_gen. It is passed (and
returns) fileName, var, nvar, and valu. It is used to generate
a copy of a marked input deck. In order to substitute in correct
variable values, it calls the subroutine var_finder
index_gen $\quad=\quad$ recursive subroutine called by input_mod. It is passed (and
returns) baseFileName, index, columns, maxcolumn, nvar,
nposscombs, val, var, comb_number, and nameGenType. It
recursively generates all of the possible combinations of
variables and values. It calls the subroutines copy_file (which
calls var_finder) and name_gen to generate all of the
calls var_finder) and name_gen to generate all of the
input_mod $\quad=\quad$ main program. It opens (and saves information from) spec_file
It generates the points for a uniform distribution, and it
calls the recursive subroutine index_gen to generate the
files corresponding to the variable/value combinations unless
the first line of the spec file is 'generate, in which case
the program will call spec_gen to generate all of the , input
the program will call spec_gen to generate all of the 'input
name_gen $\quad=$ subroutine called by index_gen. It is passed (and returns)
originalFile, columns, nvar, newFile, comb_number, and
nameGenType. It is used to generate a newFile name which
indicates the particular variable value combination of the
file being created/copied if nameGenType $=$ 'var', otherwise
nameGenType $=$ 'comb' and the generated name will correspond to
spec_gen $=$ subroutine which is passed (and returns) spec_file, genRun
$=$ subroutine which is passed (and returns) spec -file, genRun,
baseFileName, nvar, and combSet. It is called by the main program
if genRun $=$ generate". It will, create njobs specfiles which have
the first line changed to 'run' as opposed to 'generate, so tha
when these new spec files are run, they will call index-gen and
are able to generate the new input files. Additionally, it will
generate njobs of them, each of which will have a different
combSet value. (e.g. if njobs $=9$, then one file will have combSet $=1$,
$=2, \ldots$, one file $\quad$,
$=$ subroutine which is passed (and returns) line, var, nvar and valu
It is called by copy_file, and uses the above information to find
It is called by copy file, and uses the above information to find
the VARIABLE MARKERS in the file and substitutes the desired values
into the line and then passes the edited line back to copy_file
INFORMATION FOR THE USER

1. THE MARKED INPUT FILE AND SPEC FILE (VARIABLE NAMING)
Add a ' $\$$, to the beginning of EVERY variable name in your marked input file
Add a '\$' to the beginning of EVERY variable name in your marked input file
$($ i. e. template input file). In the SPEC FILE, place the variable WITHOUT the
' (i.e. in MARKED INPUT '\$VAR1' in SPEC FILE 'VAR1') as a '\$' is added in
in the main program immediately following the read statement (see sec 2.2 .1 ).
2. VARIABLE NAMES:
In the program, variable names are set to be of size 8. HOWEVER, due to the
addition of ' $\$$ ' to the beginning of the variable name (see \#1 directly above)
and due to the SPACE added to the end of the variable name (see the subroutine
var_finder for a more detailed explanation) variable names MUST be no more
var_finder for a more
than 6 CHARACTERS LONG.
3. NUMBER OF VARIABLES OCCURING WITHIN A CARD:
Currently ONLY 7 instances of variables may appear in a given line (regardless
```
    of which variables - it could be the same variable multiple times plus a few
    others, or all one variable, or all different variables a single line, i.e.
    card, may only have 7 substitutions). For more details, see the subroutine
    var_finder
    4. CASE OF A VARIABLE WITH A SINGLE POINT (i.e. stays constant):
    In the SPEC FILE, simply write the value the user desires to remain constant
    for that variable in the place where the MIN (i.e. mn) should go.
    5. MAX NUMBER OF RUNS:
        Currently ONLY runs of up to 8 digits long are permitted. To change this, and
        for additional information see NOTE in recursive subroutine index_gen.
    6. MAXIMUM NUMBER OF VARIABLES SPECIFIED in SPEC FILE
    Currently a spec file can ONLY contain information for 55 variables. To
    change this, the user will need to change the size of several arrays
    prob(XX), var(XX), columns(XX), groupNumb(XX), npts(XX), nptsGroup(XX),
    nruns(XX), runLayout(XX), increment_size_var(XX), int_length_var(XX)
    mn(XX), mx(XX), stdev(XX), val(20,XX), as well as in a write statement
    mn(XX), mx(XX),stdev(XX), val(20,XX), as well as in a write statemen
    [write (9, '(XXi4)')]. Additionally, changes should be made in the 
    ALSO should more than 99 variables be in a given file, additional changes will
    need to be made in var_finder. See the subroutine for further details
```


## Declarations

```
implicit none
    character *10 baseFileName, exfile, genRun, run_layout
    character *3 charCombSet
    character *2 flag
    character *25 marked_input_file
    character *4 nameGenType
    character *8 prob(55), var(55)
    character *20 spec_file, var_val_comb
    integer combSet, comb_number, i, ierr, is, j, k, maxpts, nposscombs, ngroups, njobs, nvar
    integer columns(55), groupNumb(55), npts(55), nptsGroup(55), nruns(55), runLayout(55)
    integer columns(55), groupNumb(55), npts(55), nptsGroup(55), nruns(55), runLayout(55)
    real increment_size_var(55), int_length_var(55), mn(55),mx(55), stdev(55), val(20,55)
    real job_inc
    Executable code
    1.0 Initialize
    comb_number = 0
    genRun =, hello,
    i}=
    j}=
    maxpts =1
    nameGenType = 'type
    nposscombs = 1
    var_val_comb}=,s
    1.1 Obtain and bypass the name of the executable file
    call getarg (i, exfile)
    is = iargc()
    1.2 Read the command option
    i = 1
    call getarg (i, flag)
    if (flag(1:2)== "-i") then
    i = 2 (1:2) == -1") then
    call getarg (i, spec_file)
    else
        write (*,*) "Error:_no--i_was_specified"
        stop "-1"
    endif
    2.0 Open the files
    2.1 Open spec_file
    inquire (file= spec_file, exist = ex)
    if (.not. ex) then
        write (*,*) "Error:-file_not_found, -", spec_file
        go to 100
    endif
    open (unit = 1, file = spec_file, action = "read", status = "old", iostat = ierr)
1 2.1.1 Read the spec_file and save the relevant variable information. Then rename
        each variable to include marker (i.e.'$').
    read (1, "(a10)") genRun
    read (1, "(a10,i3,i3,i3,i3)") baseFileName, nvar, njobs, combSet, ngroups
    do i = 1, nvar
        read (1, "(a8,_3e12.4,_i3,_a8,_i3)", end = 10) var(i), mn(i), mx(i), stdev(i), npts(i),&
prob(i), groupNumb(i)
        var(i) ="$"// trim(var(i))
    end do
10 continue
    rewind 1
```

!
$!$

```
    2.1.2 Calculate the number of possible combinations, and create an array based on njobs
    and nposscombs which specifies which runs will take place on each combSet.
    For example: if there are 27 possible combinations, and njobs = 3, then
    nruns( ) = [\begin{array}{lll}{3}&{6}&{9}\end{array}], so we know combinations 1-3 will take place if combSet = 1,
    combinations 4-6 will take place if combset = 2, and combinations 7-9 occur when
    combSet = 3. Since the ceiling function is used, any number of njobs may be used
    without concern for the case where njobs does not evenly divide nposscombs. In 
    that case, we will just have a different number, i.e. +/- 1, of runs generated
    in one of the, input deck generator, spec files.)
    k}=
    nptsGroup(1) = npts(1)
    do i}=1, nva
        if (groupNumb(i) /= groupNumb(k)) then
        k}=\textrm{k}+
        nptsGroup(k) = npts(i)
        M\mp@code{npts}
    end do
    uncomment the line below to write out the group number of each variable
    write (*,"(20i4)") groupNumb
    nposscombs = 1
    do i = 1, ngroups
    nposscombs = nptsGroup(i)*nposscombs
    end do
    job_inc=(nposscombs)/(njobs)
    do i}=1, njobs-
    nruns(i) = i*ceiling(job_inc)
    end do
    nruns(njobs) = nposscombs
    2.1.3 Open the file "run_layout". It will be used by the python script.
    For the file, we'll print out the number of possible combinations
    (i.e. the total number of tests). Then we'll calculate exactly how
    many runs are expected for each of the nodes (i.e. for each combSet),
    and then write that onto the second line. (each will be of size 4).
    Note: npossCombs will be of size 12. The file will then be closed
    run_layout = "run_layout"
    open(unit = 9, file= run_layout, action = "write", status = "replace", iostat = ierr)
    write(9, "(i12)") nposscombs
    runLayout(1)= nruns (1)
    do i = 2, njobs
        runLayout(i) = nruns(i) - nruns(i - 1)
    end do
    write(9, '(55i4)') runLayout(1: njobs)
    close(unit = 9)
    2.1.4 Based on genRun, program will follow one of three options.
        (1) If spec_file is a spec_file with a genRun= "generation", send the spec_file
        to the subroutine spec-gen which will generate njobs of spec_files. (first if)
        (2) If spec_file's genRun is not equal to "run" program will terminate
        (3) Otherwise, execute normally, utilizing the subroutine index_gen
    if (trim(genRun) == 'generate') then
    call spec_gen(spec_file, genRun, baseFileName, nvar, njobs, combSet, ngroups)
    calospoc
    elseif (trim(genRun) /= 'run') then
    go to 100
    endif
    2.2 Use the baseFilename as the marked_input_file
    marked_input_file = trim(baseFileName) / , .i,
    2.3 Open the marked_input_file
    inquire (file = marked_input_file, exist= ex)
    if (.not. ex) then
    write (*,*) "Error:^file_not_found, "", marked_input_file
    go to }10
    endif
    open(unit =2, file = marked_input_file, action = "read", status = "old", iostat = ierr)
    3. Generate input files
    3.1 Calculate the variable values for each interval with 'uniform' dist
    3.1.1 Calculate the interval length and increment size
do i = 1, nvar
    int_length_var(i) = mx(i) - mn(i)
    if (npts(i) <= 1) then
        increment_size_var(i) = 0.0
    else
        increment_size_var(i) = int_length_var(i) / (npts(i) - 1)
```



```
!\mp@code{1. size of char_i in declarations to '*X',}
    Declarations
    Arguments
    implicit none
    character *10 baseFileName
    character *4 nameGenType
    character *80 newFile
    character *8 var(*)
    integer columns(*), groupNumb(nvar), maxcolumn(*), nruns(*)
    integer combSet, comb_number, colindex, nvar, nposscombs
    real val(20, 55)
!
! Local Variables
    integer i, j
    real valu(nvar)
    character *8 charCombNum
    newFile = baseFileName
!
! Executable Code
    if (colindex == 1) then
        do i = 1, maxcolumn(1)
            columns(1) = i
            do j = 1, ngroups
                where (groupNumb == j) valu = val(columns(j),:)
            end do
            comb_number = comb_number + 1
    ! if (comb_number > 6) stop
    Remove line above to generate more than 2 input files
            if (comb_number > nruns(combSet)) then
                stop
            elseif (combSet == 1) then
                call name_gen(baseFileName, columns(1: nvar), nvar, newFile, comb_number, nameGenType)
                call copy_file(newFile, var, nvar, valu)
                write (charCombNum, '(i8.8)') comb_number
                write (3,62) charCombNum, valu(1: nvar)
                62 format (a8, 55 es12.4)
                ! write (3,621) charCombNum, (columns(j), j=1, ngroups)
                ! uncomment line above (and below) to write the index of each group for each run
                ! 621 format (a8, 55i3)
            elseif (comb_number > nruns(combSet-1)) then
                call name_gen(baseFileName, columns(1: nvar), nvar, newFile, comb_number, nameGenType)
                call copy_file(newFile, var, nvar, valu)
                write (charCombNum, '(i8.8),) comb_number
                write (3,62) charCombNum, valu(1: nvar)
                ! write (3,621) charCombNum, (columns(j), j=1, ngroups)
                uncomment line above (as well as the format line) to write the index of each group
                for each run
            endif
        end do
    else
        do i = 1, maxcolumn(colindex)
            call index_gen (baseFileName, colindex - 1, columns, maxcolumn, nvar, nposscombs, val,&
var, comb_number, nameGenType, nruns, combSet, groupNumb)
                columns(colindex) = columns(colindex) + 1
            end do
        if (columns(colindex) > maxcolumn(colindex)) then
            columns(colindex)=1
        endif
    endif
    if (comb_number > nposscombs) then
        stop
    end i
return
ret
end subroutine index_gen
!-
subroutine copy_file(fileName, var, nvar, valu)
! CREATED: Mar 6, 2012
    UPDATED: Apr 11, 2012
    SUBROUTINE DESCRIPTION:
    Make (i.e. open and write to) a copy of the marked-up input file
    with the generated name. Then call var_finder to modify fileName
    so it's variable values correspond to the file name. Also appends
    the file name to include ,.i,
    Declarations
```

```
シゴ
Arguments
Arguments
    implicit none
    character *80 fileName
    character *8 var(*)
    integer nvar
    integer nvar
!
! Local Variables
    character *132 line
    integer ierr
    logical found
!
! Executable Code
    open (unit = 8, file = fileName, action = "readwrite-", position = "rewind", &
status = "replace", iostat = ierr)
    do
        read (2, "(a132)", end = 999) line
        found = index (line, "$VAR") > 0
        call var_finder(line, var, nvar, valu)
        write (8, "(a)") trim(line)
            UNCOMMENT the text below to print out the line that was altered by
            the subroutine var_finder and written into the NEW input file.
            if (found) then
                    write (*,*) "Copy File: line = ", line
        end if
    end do
    999 continue
    close (unit = 8)
    rewind 2
    return
end subroutine copy_file
!-
subroutine name_gen(originalFile, columns, nvar, newFile, comb_number, nameGenType)
!
    CREATED: Mar 7, 2012
    UPDATED: Mar 19, 2012
    SUBROUTINE DESCRIPTION:
    Creates name for input files based on base name AND variables if
    Creates name for input files based on base name AND variables if ' ' ' var', otherwise nameGenType = 'comb' and the name generated will
    be based on the base name and combination number the var/val corresponds to.
    Declarations
    Arguments
    implicit none
    character *4 nameGenType
    character *4 nameGenTyp
    character *80 newFile
    integer columns(1: nvar)
    integer comb_number, nvar
!
    Local Variables
    character *8 char_i
    character
!
    Executable Code
    newFile = originalFile
    if (nameGenType == 'comb') then
        write(char_i, '(i8.8)') comb_number
        newFile = trim(newFile) // " _" //trim(adjustl(char_i))
    else
        do j = 1, nvar
            write(char_i, '(i2)') columns(j)
            newFile = trim(newFile) // " -" // trim(adjustl(char_i))
        end do
    endif
    newFile = trim(newFile) // '.i,
    return
end subroutine name_gen
```

```
--
subroutine var_finder(line, var, nvar, valu)
! CREATED: March 07, 2012
    UPDATED: April 12, 2012
    SUBROUTINE DESCRIPTION.
    Find the lines in the input file which contain a variable MARKER
    (i.e. '$VAR1') and then replace the MARKER with the desired value.
    NOTE:
    The variable name can have a double digit number, so special care
    must be taken when forming the new line to prevent part of the
    name from appearing in the new line. To handle this case, an
    additional SPACE has been added to the END of the value replacing
    the marked variable in the line. THIS could potentially cause errors
    in input processing as it may cause there to be too many characters
    on a card in the new input file.
    LET THE USER BEWARE:
    Currently, this subroutine is set to account for up to 7 occurences
    of a variable within a single card (line) in the input deck
    IF more occurences exist in the user's modified deck, CHANGE the
    outermost do-loop from 1 to 7 TO 1 to X (greatest number of
    instances of variables occuring on a single card (line) in the marked
    input deck.
    ADDITIONAL WARNING:
    ALL VARIABLES ($XXXXXX, where XXXXX is input by the user) are PADDED
    with a SPACE (i.e. from '$XXXXX' to '$XXXXX ', within this subroutine)
    so as to ENSURE that the program will not read in variables such as
    '$VAR1' and '$VAR11', (assuming the user puts them in sequentially in
    '$VAR1' and '$VAR11', (assuming the user puts them in sequentially in l
    values. The user should also be aware that we insert an extra space
    in the line immediately following the value insertion. If the user
    should have more than 99 variables, it is recommended that an extra
    space be inserted for each additional digit of nvar (e.g. for 2000,
    add two extra spaces, for 999 add one extra space.)
    Declarations
    Arguments
    implicit none
    character *132 line
    character *8 var(*)
    integer nvar
    real valu(*)
!
    Local Variables
    integer charlen, i, j, loc
    character *12 charvalu
    character *1 sp
!
    Executable Code
    sp = " ""
    sp = "-",
        do i}=1, nva
            loc = index (line, trim(var(i)) // sp)
            if (loc .ne.0) then
            charlen = len_trim(var(i))
            write (charvalu,"(es12.4)") valu(i)
    UNCOMMENT the text below to check which variable and value is being
    substituted into a particular line. NOTE that it will print out EACH
    occurence of the variable being replaced.
        write (*,*) "Var Finder: var(i) = ", var(i), i
            write (*,*) "Var Finder : line before = ", line
            line = line(1: loc - 1) // charvalu // sp // line(charlen+loc +1:)
            write (*,*) "Var Finder : line after = ", line
        endif
        end do
    end do
    return
end subroutine var_finder
subroutine spec_gen(spec_file, genRun, baseFileName, nvar, njobs, combSet, ngroups)
    CREATED: Mar 15, 2012
    CREATED: Mar 15, 2012
```

```
    SUBROUTINE DESCRIPTION
    Create njobs specfiles which have the first line changed to 'run'
    as opposed to 'generate,, so that when these new spec files are run,
    they will call index_gen and actually generate the new input files.
    Declarations
    Arguments
    implicit none
    character *10 baseFileName, genRun
    character *20 spec_file
    integer combSet, ngroups, njobs, nvar
!
    Local Variables
    character *3 charCombSet
    character *20 fileName
    character *20 fieN
    character *132 line
!
    Executable Code
    do i}=1, njob
        combSet = i
        write(charCombSet,'(i3)') combSet
        fileName = trim(spec_file) // , -' // adjustl(charCombSet)
        fileName = trim(fileName)
        open (unit = 8, file= fileName, action = "readwrites", position = "rewind",&
status = "replace", iostat = ierr)
    read in the first two lines of spec_file. At this point, we won't be
    doing anything with them, but after we read them in, we' ll be all set
    to read in the rest of the lines (and print them out) in a loop.
        read (1, "(a132)", end = 999) line
        read (1, "(a132)", end = 999) line
    write the first two lines of the new spec_file- replacing "generate" with
    "run" and changing the combSet number appropriately
        write (8, "(a3)") "run"
        write(8, 23) baseFileName, nvar, njobs, combSet, ngroups
        23 format(a10, i3, i3, i3, i3, i3)
        999 continue
!
    now we read in the rest of the lines from file 1 and write with no
    modifications onto the spec_file we're generating.
        do j = 1, nvar
            read (i, "(a132)", end = 99) line
            write (8, "(a)") trim(line)
        end do
        99 continue
        rewind 8
        close (unit = 8)
        rewind 1
    end do
    return
end subroutine spec_gen
end program input_mod
```


## B Template Spec File

```
typeOfSpecFile
baseFileName #vars #nodesInStudy currentNode #groups
varX min max stdDev #pts distribution group#
```



## C Example of a Generator Spec File



## D Example of a NodeSpec File

| 1 | run |  |  |  |  |  |  |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 2 | apsbs | $8 \quad 9 \quad 9$ | 8 |  |  |  |  |
| 3 | VAR1 | 0.9 | 1.1 | 0.0471404520 | 3 | uniform | 1 |
| 4 | VAR2 | 0.00135 | 0.00165 | 0.0000707106 | 3 | uniform | 2 |
| 5 | VAR3 | 8.95959 | 10.95061 | 0.4692879144 | 3 | uniform | 3 |
| 6 | VAR4 | 0.043749 | 0.053471 | 0.0022914973 | 3 | uniform | 4 |
| 7 | VAR5 | 0.193752 | 0.236808 | 0.0101483965 | 3 | uniform | 5 |
| 8 | VAR6 | 0.04419 | 0.05401 | 0.0023145962 | 3 | uniform | 6 |
| 9 | VAR7 | 0.000135 | 0.000165 | 0.0000070710 | 3 | uniform | 7 |
| 10 | VAR8 | . 19493874 | 0.23825846 | 0.0102105559 | 3 | uniform | 8 |

## E Python Script for Running Studies on INL's Supercomputer, Quark

```
#!/apps/local/python/activestate/2.7.1.4/bin/python
# # Purpose:
Note: 
# Note: Pure Python script, no use of Linux scripts or commands.
    To adjust this run_file for your particular test, go to
    main program, section 1.0 and adjust the file names (and
    keyOutputString) accordingly. The user is also encouraged
    to adjust the walltime request to better fit the user's
    study
    Authors: Dr. George L Mesina, Alexandra E. Gertman
    J. Shelley
    Created: Mar 16, 2012
    Updated: May 02, 2012
Pro Batch Scheduler (PBS) commands
#PBS -N relap5
#PBS -1 select=1:ncpus=12:mpiprocs=12:mem=23gb
#PBS -1 place=excl
#PBS -1 walltime=10:00:00
#PBS -V
#PBS - j oe
#PBS -q general
#
###################################
# 1.0 Imports
import glob, os
from tempfile import mkdtemp
import shutil, subprocess, sys
from string import join
import time
try:
    import forkmap as fm
    FORKMAPavail = "yes"
except Exception, e:
    FORKMAPavail = "no"
```




```
def CreateDirs(curPath,outPath, filePre, tmpPath):
# Make the directories where each node's (or combSet's)
# cases will be run
# tmp_dir = mkdtemp(prefix=filePre, dir=tmpPath)
    return tmp_dir
return tmp_dir
###################################
def GetInputFiles(inputPre):
# Grab all of files (each of which is named inputPre) from our current path
    Then return all of those files
    Note: inputPre = specFilePre + ,*, for specFiles AND
            inputPre = inputFilePre%"*" for inputFiles
    inputs = glob.glob(inputPre)
    return inputs
```




```
def PrepRunGetClean(index, currentPath, relap_exe, tmpPath):
#
Runs each input file passed to it (using relap executable rund)
    Currently, only 12 input files are run at a time (since we're running
    our tests on quark.) If a different number should be run through each
    # time, appropriate changes should be made in the parallel section, as
# as in the beginning lines of this program (the PBS - l select line)
#
    cmd = "%s_%s"%(relap_exe,index)
    output = subprocess.Popen(cm, shell=True, stdout=subprocess.PIPE).communicate() [0]
    print >> sys.stderr, 'cmd:_', cmd
    return
#######################################
##################################
if _-name_- ==,_-main__':
# DESCRIPTION: Create working directory in temporary-space with subdirs
                    one per cluster node. Run input_mod_gen.f90 to create 1
    spec-file for each subdirectory by dividing the total no.
    of combinations equally among the cluster-nodes. Copy the
```

```
############
COGNIZANT: Alexandra E. Gertman, Dr George Mesina, Jon Shelley
CREATED: Mar 16, 2012
UPDATED: Mar 27, 2012
PROGRAM OUTLINE:
1.0 Initialize and set-up directories
1.1 Construct & Populate subdirectories
    Construct subdirectory structure explained above
    1.1.1 Change the current working directory to currentPath
    1.1.2 Collect all of the spec files we'll use in this study
    1.1.3 If the directory (tmpPath) does not yet exist, create it .
            Then create the directory which will hold all of the runs
            for each combset (i.e. for each node)
1.2 Obtain list of computer nodes
1.3 Input file generation
    1.3.1 Switch the current working directory to tmp_run_dir
    (i.e. the temporary directory)
    1.3.2 In each temporary directory, create a link to the needed files:
        the RELAP5-3D executable (rund), relap5.x, its license and property
        files, and the appropriate specfile
    1.3.4 Collect all of the input files
2.0 Output file generation through parallelization
2.1 Run Parallel: All of the input files that were
    generated will be run with the relap executable
    2.1.1 Grab 12 of the input files at a time, passing them
        to the function PrepRunGetClean where the output
        for those 12 input files will be created.
3.0 Results
3.1 Collect the newly generated output files
    3.1.1 Create a results dictionary where we'll store
        the run number and the corresponding desired
        output parameter
    3.1.2 Loop through output files to find the key output
                parameter, and then put that value into the results
            dictionary along with the corresponding run number
3.2 Find the file with the variable information for each
    run, copy that information and put into a new file
    appending the run information with the desired output
    from the results dictionary. Then save the results file
    o the current directory
    3.2.1 Open and read the comb file for this particular combSet/node
    3.2.2 Create the list outline
        3.2.2.1 Loop through each line of combline
            3.2.2.1.1 Split each line in each of the combFiles
                            3.2.2.1.2 Append each line with it's corresponding result
                            3.2.2.1.3 Place modified cline into the list outline
    3.2.3 Save all of outline to res_text
    3.2.4 Open combFile.res, write all of res_text to it, then close the file
    3.2.5 Copy the file to the folder 'results, in the home directory
4.0 Remove temporary directory and exit program
DATA DICTIONARY
array_idx = array of cluster nodes. Should correspond to the number of
                        jobs specified in spec file. (3rd word on line 2 of the
        spec file.)
cline = variable denoting each specific line in combLine. It is
    stripped and split, appended, and then put back together
cmd = variable which denotes arguments we'll be passing to the
                                    command line.
                                    name of the comb file for the particular combSet/node
                                It is combFilePre with arry_idx.
                            = base name for each comb file
                                    For ex: 'apsbs2_comb'. No additional string w/in a string
                is specified
combLine = where each line of the combFile is stored (each line is a
    string.)
    the run number of cline
    = the run number of clin
    = PBS working directory 
    = where the information for an output file is stored
    = base name used in all of the input files
    For example: 'apsbs_%s.i', where the %s indicates that we
    will specify a what string will get placed in between the
    underscore and the ".i"
index = variable used to denote 12 different strings (each of which
    fil strings
    corresponds to a specific input file.) It is used in the
    parallel section, and passed to PrepRunGetClean so that each
    of the 12 input files will be run with rund.
inputFiles = all of the generated input files (i.e. spec files which contain
```



```
        os.mkdir(tmpPath)
    tmp_run_dir = CreateDirs(currentPath, outputPath, specFilePre+str(arry_idx)+'_', tmpPath)
##
1.2 Obtain list of computer nodes
    Each node has a number of cores
    nodes = open(os.environ['PBS_NODEFILE'],'r').readlines()
    numNodes = len(nodes)
    print "Number_of_threads:_",numNodes
    1.3 Input file generation
        1.3.1 Switch the current working directory to tmp_run_dir
        (i.e. the temporary directory)
    os.chdir(tmp_run_dir)
    1.3.2 In each temporary directory, create a link to the needed files:
        the RELAP5-3D executable (rund), relap5.x, its license and property
        files, and the appropriate specfile
    for lnfile in ['rund','relap5.x','tpfh2o',',rellic.bin', specFilePre+'%d'%arry_idx,
                markedInputFile]:
    cmd =, ln--s %%s/%s'%(currentPath, lnfile)
    print "cmd:-%s"%cmd
    output = subprocess.Popen(cmd, shell=True, stdout=subprocess.PIPE).communicate()[0]
        1.3.3 In each temporary directory, create the command to
        run the executable from input_mod_gen.f90 (a.out),
        passing it '-i, (necessary for executable to work),
        and specFilePre arry_idx
    cmd = currentPath+os.sep+'a.out_-i _%s%d'%(specFilePre, arry_idx)
    print "cmd:_%s"%cmd
    output = subprocess. Popen(cmd, stdout=subprocess.PIPE, shell=True).communicate()[0]
    1.3.4 Collect all of the input files
    inputFiles=GetInputFiles(inputFilePre%"*")
    #inputFiles = inputFiles [:12] #Uncomment for smaller test
    print inputFiles
    2.0 Output file generation through parallelization
    2.1 Run Parallel: This is where all of the input files that
    were generated will be run with the relap executable rund
    ####################################
    ## Parallel section
    #################################
        2.1.1 Grab 12 of the input files at a time, passing them
        to the function PrepRunGetClean where the output
        for those 12 input files will be created.
    @fm.parallelizable(12)
    def controller(index)
        print >> sys.stderr, 'index=', index
        print >> sys.stderr(,
        PrepRunGetClean(index, currentPath, relap_exe, tmp_run_dir)
        return time.time() - start2
    runtime = fm.map(controller, [x for x in inputFiles])
```



```
    ## End Parallel section
```



```
    ##
#
##
# 3.1 Collect the newly generated output files
    outputFiles = glob.glob(outputFilePre%'*')
    print "Output_Files:_",outputFiles
        3.1.1 Create a results dictionary where we'll store
        the run number and the corresponding desired
        output parameter.
    res_dict = dict()
    3.1.2 Loop through output files to find the key output
                parameter, and then put that value into the results
                dictionary along with the corresponding run number
    for outFile in outputFiles:
        data=open(outFile).read()
        fidx = data.rfind(keyOutputString)
        keyOutputParam = (data[fidx: fidx + 80]).split ()[2]
        #print "%s Coremin: %s"%(outFile, pct)
        res_dict[outFile.split(',',)[1].split(',') [0]]= keyOutputParam
    print "Results_Dictionary:\n", res_dict
# 3.2 Find the file with the variable information for each
    run, copy that information and put into a new file,
    appending the run information with the desired output
    from the results dictionary. Then save the results file
    to the current directory.
```



## F SAS Reports

## F. 1 AP600 2 inch Break, Top 4 Variables

## Correlation Analysis

The CORR Procedure

| 1 With Variables: |  |  |  |  |  | coremin |
| :--- | :--- | :--- | :--- | :---: | :---: | :---: |
| 4 Variables: | VAR1 | VAR8 | VAR12 |  |  |  |
| VAR13 |  |  |  |  |  |  |
| SSCP Matrix |  |  |  |  |  |  |
| coremin | 6607.808270 | VAR1 | VAR8 |  |  |  |


| CSSCP Matrix |  |  |  |  |
| :--- | ---: | ---: | ---: | ---: |
|  | VAR1 | VAR8 | VAR12 | VAR13 |
| coremin | -5.60623000 | -12.48935000 | 4.69284000 | -0.0078711 |


| Covariance Matrix, DF = 80 |  |  |  |  |
| :--- | ---: | ---: | ---: | ---: |
|  | VAR1 | VAR8 | VAR12 | VAR13 |
| coremin | -.0700778750 | -.1561168750 | 0.0586605000 | -.0000983896 |


| Simple Statistics |  |  |  |  |  |  |
| :--- | ---: | ---: | ---: | ---: | ---: | ---: |
| Variable | N | Mean | Std Dev | Median | Minimum | Maximum |
| coremin | 81 | 81.64709 | 1.88027 | 81.67210 | 78.39860 | 84.71310 |
| VAR1 | 81 | 1.00000 | 0.04108 | 1.00000 | 0.95000 | 1.05000 |
| VAR8 | 81 | 0.50000 | 0.41079 | 0.50000 | 0 | 1.00000 |
| VAR12 | 81 | 1.00000 | 0.24648 | 1.00000 | 0.7000 | 1.30000 |
| VAR13 | 81 | 0.0002525 | 0.0002033 | 0.0002525 | $5 \mathrm{E}-6$ | 0.0005000 |


| Pearson Correlation Coefficients, $\mathbf{N}=\mathbf{8 1}$ <br> Prob $>\|\mathbf{r}\|$ under H0: Rho=0 |  |  |  |  |
| :--- | ---: | ---: | ---: | ---: |
|  | VAR1 | VAR8 | VAR12 | VAR13 |
| coremin | -0.90727 | -0.20212 | 0.12658 | -0.25734 |
|  | $<.0001$ | 0.0704 | 0.2602 | 0.0204 |


| Spearman Correlation Coefficients, $\mathbf{N}=81$ <br> Prob $>\|\mathbf{r}\|$ under H0: Rho=0 |  |  |  |  |
| :--- | ---: | ---: | ---: | ---: |
|  | VAR1 | VAR8 | VAR12 | VAR13 |
| coremin | -0.90408 | -0.21729 | 0.13451 | -0.25932 |
|  | $<.0001$ | 0.0513 | 0.2312 | 0.0194 |


| Kendall Tau b Correlation Coefficients, $\mathbf{N}=81$ <br> Prob > \|taul under H0: Tau=0 |  |  |  |  |
| :--- | ---: | ---: | ---: | ---: |


| Hoeffding Dependence Coefficients, $\mathbf{N}=81$ <br> Prob $>$ D under H0: $\mathbf{D}=\mathbf{0}$ |  |  |  |  |
| :--- | ---: | ---: | ---: | ---: |
|  | VAR1 | VAR8 | VAR12 | VAR13 |
| coremin | 0.31322 | 0.00025 | -0.00573 | 0.00603 |
|  | $<.0001$ | 0.3472 | 0.9429 | 0.1216 |

## F. 2 AP600: 4 inch Break, Top 5 Variables



| Spearman Correlation Coefficients, $\mathbf{N}=\mathbf{2 4 3}$ <br> Prob $>\|\mathbf{r}\|$ under H0: Rho=0 |  |  |  |  |  |
| :--- | ---: | ---: | ---: | ---: | ---: |
|  | VAR1 | VAR8 | VAR11 | VAR12 | VAR13 |
|  | -0.76563 | -0.03280 | -0.01617 | -0.00007 | -0.05008 |
|  | $<.0001$ | 0.6109 | 0.8020 | 0.9991 | 0.4371 |


| Kendall Tau b Correlation Coefficients, $\mathbf{N}=\mathbf{2 4 3}$ <br> Prob > $\mid$ tau under H0: Tau=0 |  |  |  |  |  |
| :--- | ---: | ---: | ---: | ---: | ---: |


| Hoeffding Dependence Coefficients, $\mathbf{N}=243$ Prob > D under H0: D=0 |  |  |  |  |  |
| :---: | :---: | :---: | :---: | :---: | :---: |
|  | VAR1 | VAR8 | VAR11 | VAR12 | VAR13 |
|  | 0.18111 | -0.00235 | -0.00234 | -0.00240 | 0.00021 |
| coremin | <. 0001 | 0.9977 | 0.9974 | 0.9987 | 0.3226 |

Generated by the SAS System ('SASApp', X64_ESRV08) on April 12, 2012 at 4:42:07 PM

## F. 3 AP600: 6 inch Break, Top 5 Variables

Thursday, April 12, 2012 5:32 PM 1


| Spearman Correlation Coefficients, $\mathbf{N = 2 4 3}$ <br> Prob > $\mid$ r\| under H0: Rho=0 |  |  |  |  |  |
| :--- | ---: | ---: | ---: | ---: | ---: |
|  | VAR1 | VAR7 | VAR8 | VAR12 | VAR13 |
|  | -0.52077 | -0.00833 | 0.45018 | 0.24533 | -0.30260 |
|  | $<.0001$ | 0.8972 | $<.0001$ | 0.0001 | $<.0001$ |


| Kendall Tau b Correlation Coefficients, N = 243 <br> Prob $\boldsymbol{>} \mid$ taul under H0: Tau=0 |  |  |  |  |  |
| :--- | ---: | ---: | ---: | ---: | ---: |
|  | VAR1 | VAR7 | VAR8 | VAR12 | VAR13 |
|  | -0.41714 | -0.00594 | 0.35799 | 0.19097 | -0.23436 |
|  | $<.0001$ | 0.9046 | $<.0001$ | 0.0001 | $<.0001$ |


| Hoeffding Dependence Coefficients, $\mathbf{N}=243$ <br> Prob > D under H0: D=0 |  |  |  |  |  |
| :---: | :---: | :---: | :---: | :---: | :---: |
|  | VAR1 | VAR7 | VAR8 | VAR12 | VAR13 |
|  | 0.07638 | -0.00338 | 0.05917 | 0.01273 | 0.01938 |
| coremin | <. 0001 | 1.0000 | <. 0001 | 0.0010 | <. 0001 |

Generated by the SAS System ('SASApp', X64_ESRV08) on April 12, 2012 at 5:32:43 PM

## F. 4 AP600: 8 inch Break, Top 4 Variables

Friday, April 13, 2012 4:41 PM 1


| Pearson Correlation Coefficients, $\mathbf{N}=\mathbf{8 1}$ <br> Prob $>\|\mathbf{r}\|$ under H0: Rho=0 |  |  |  |  |
| :--- | ---: | ---: | ---: | ---: |
|  | VAR1 | VAR8 | VAR12 | VAR13 |
|  | 0.38637 | 0.28539 | -0.07139 | -0.04129 |
|  | 0.0004 | 0.0098 | 0.5265 | 0.7144 |


| Spearman Correlation Coefficients, $\mathbf{N = 8 1}$ <br> Prob $>\|\mathbf{r}\|$ under H0: Rho=0 |  |  |  |  |
| :--- | ---: | ---: | ---: | ---: |
|  | VAR1 | VAR8 | VAR12 | VAR13 |
| coremin | 0.32918 | 0.46694 | -0.23755 | -0.18540 |
|  | 0.0027 | $<.0001$ | 0.0327 | 0.0975 |


| Kendall Tau b Correlation Coefficients, $\mathbf{N}=81$     <br> Prob $>\mid$ tau     <br> under H0: Tau=0     |  |  |  |  |
| :--- | ---: | ---: | ---: | ---: |
|  | VAR1 | VAR8 | VAR12 | VAR13 |
| coremin | 0.25619 | 0.31788 | -0.19474 | -0.14414 |
|  | 0.0032 | 0.0002 | 0.0255 | 0.0983 |


| Hoeffding Dependence Coefficients, $\mathbf{N}=81$ <br> Prob $>$ D under H0: $\mathbf{D}=\mathbf{0}$ |  |  |  |  |
| :--- | ---: | ---: | ---: | ---: |
|  | VAR1 | VAR8 | VAR12 | VAR13 |
|  | 0.01622 | 0.04746 | 0.01046 | 0.00160 |
|  | 0.0242 | 0.0003 | 0.0588 | 0.2683 |

## F. 5 LOFT: 3 Values per Variable



## F. 6 LOFT: 6 Values per Variable

Correlation Analysis
The CORR Procedure

1 With Variables: |PCT
6 Variables: VAR1 VAR25 VAR26 VAR44 VAR45 VAR47

| SSCP Matrix |  |  |  |  |  |  |
| ---: | ---: | ---: | ---: | ---: | ---: | ---: |
|  | VAR1 | VAR25 | VAR26 | VAR44 | VAR45 | VAR47 |
| PCT | 741345.7 | 70880.1 | 128445869.5 | 16928250.1 | 14118361.3 | 1523872.0 |


| CSSCP Matrix |  |  |  |  |  |  |
| :--- | ---: | ---: | ---: | ---: | ---: | ---: |
|  | VAR1 | VAR25 | VAR26 | VAR44 | VAR45 | VAR47 |
| PCT | -354.6168 | -66.9270 | -170739.5504 | -215551.4121 | -53847.9969 | -21.4771 |


| Covariance Matrix, DF = 15624 |  |  |  |  |  |  |  |  |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  | VAR1 | 1 VAR25 |  | VAR26 VAR44 |  | \| VAR45 |  | VAR47 |
| $\overline{\text { PCT }}$ | -0.02269692 | 2 -0.00 | 0428360-10 | 0.92803062 | -13.7961733 | -33 -3.44649 | 9237 | -0.00137462 |
|  | Simple Statistics |  |  |  |  |  |  |  |
|  | Variable | N | Mean | Std Dev | Median | Minimum | Maxim | mum |
|  | PCT | 15625 | 975.29182 | 138.06833 | 960.68800 | 723.60100 |  | 1476 |
|  | VAR1 | 15625 | 0.04867 | 0.0005002 | 0.04867 | 0.04796 |  | 04938 |
|  | VAR25 | 15625 | 0.00466 | 0.0000540 | 0.00466 | 0.00458 |  | 0473 |
|  | VAR26 | 15625 | 8.44000 | 0.59682 | 8.44000 | 7.59600 |  | 28400 |
|  | VAR44 | 15625 | 1.12500 | 0.26517 | 1.12500 | 0.75000 |  | 50000 |
|  | VAR45 | 15625 | 0.93000 | 0.13153 | 0.93000 | 0.74400 |  | 1600 |
|  | VAR47 | 15625 | 0.10000 | 0.03536 | 0.10000 | 0.05000 |  | 5000 |


| Pearson Correlation Coefficients, $\mathbf{N}=15625$ <br> Prob $>\mid$ r\| under H0: Rho=0 |  |  |  |  |  |  |
| :--- | ---: | ---: | ---: | ---: | ---: | ---: |
|  | VAR1 | VAR25 | VAR26 | VAR44 | VAR45 | VAR47 |
|  | -0.32863 | -0.57473 | -0.13262 | -0.37682 | -0.18979 | -0.00028 |
| PCT | $<.0001$ | $<.0001$ | $<.0001$ | $<.0001$ | $<.0001$ | 0.9719 |


| Spearman Correlation Coefficients, $\mathbf{N}=15625$ <br> Prob $>\mid \mathbf{r \|}$ under H0: Rho=0 |  |  |  |  |  |  |
| :--- | ---: | ---: | ---: | ---: | ---: | ---: |
|  | VAR1 | VAR25 | VAR26 | VAR44 | VAR45 | VAR47 |
|  | -0.25152 | -0.80646 | -0.17079 | -0.31533 | -0.25236 | -0.00048 |
| PCT | $<.0001$ | $<.0001$ | $<.0001$ | $<.0001$ | $<.0001$ | 0.9517 |


| Kendall Tau b Correlation Coefficients, $\mathbf{N}=15625$ <br> Prob $>\mid$ taul under H0: Tau=0 |  |  |  |  |  |  |
| :--- | ---: | ---: | ---: | ---: | ---: | ---: |
|  | VAR1 | VAR25 | VAR26 | VAR44 | VAR45 | VAR47 |
|  | -0.18691 | -0.66938 | -0.12528 | -0.23445 | -0.18548 | -0.00035 |
| PCT | $<.0001$ | $<.0001$ | $<.0001$ | $<.0001$ | $<.0001$ | 0.9523 |


[^0]:    *Mathematics Department, Boise State University
    ${ }^{\dagger}$ Idaho National Laboratory

[^1]:    ${ }^{1}$ Also referred to as USNRC

[^2]:    ${ }^{2} \hat{\mu_{x}}$ is also denoted $\bar{x}$

[^3]:    ${ }^{3}$ In some cases, groups of variables may need to be changed simultaneously in input file generation, and in those cases, groups are needed. In the case where a set of variables needs to be simultaneously changed all of those variables will be in one group, and in the case where only one variable needs to be changed it will be placed in its own group. For instance, in the LOFT study discussed later in the paper, variables 1-24 represent peaking factor and are in group one, variable 25 represents fuel clad gap width and is in group 2, variables $26-43$ represent fuel thermal conductivityand belong to group 3, etc.

[^4]:    ${ }^{4} \zeta \approx 0$ when $X$ and $Y$ are independent. $\zeta=0 \nRightarrow X$ and $Y$ are independent

[^5]:    ${ }^{5}$ 'Stronger' in this case is referring to a higher degree of statistical significance.
    ${ }^{6}$ Recall, it specifically requires a bivariate distribution which is approximately normal.
    ${ }^{7}$ With the advances in computing, and the introduction of statistical software packages, this has become less of an issue.
    ${ }^{8}$ This is not the case when the value of Spearman's rho is 1 or -1 . In that case, Kendall's tau will also be 1 or -1 respectively. This is the same for the case when Kendall's tau is known to be 1 or -1 .

[^6]:    ${ }^{9}$ This was used to create the four different studies: 2 inch Break, 4 inch Break, 6 inch Break, and 8 inch Break.
    ${ }^{10}$ COREMIN is the key output parameter used in the AP600 studies
    ${ }^{11}$ i.e. VAR1-5, VAR7-14. VAR 6 was used to vary the break size.

[^7]:    ${ }^{12}$ coremin represents the key output parameter, which is a control variable we created that takes the minimum of control variable 114 (collapsed core liquid level). Control Output() is the (Percentage of) Collapsed Core Liquid Level.

[^8]:    ${ }^{13}$ Peak Clad Temperature is the key output parameter used in the LOFT study.

[^9]:    ${ }^{14}$ To avoid confusion, we will refer to this input deck as the base input deck.
    ${ }^{15}$ XXXX denotes the variable name of that particular input parameter
    ${ }^{16}$ In our studies, we selected 'VARX', where $\mathrm{X}=1,2,3, \ldots$ to denote the various variables.
    ${ }^{17}$ There are very specific parameters for the length of character name, so care should be taken. Details of this can be found within the main program- see Appendix A for further details. Additional care should be taken when replacing values as a card may only contain 80 characters, so the user may wish to consider adjusting spaces within a given card or splitting the card (i.e. making a continuation card.) See reference [7.

[^10]:    ${ }^{18}$ The 'number of nodes' refers to the number of studies within the study. i.e. We break each study into multiple sets (or studies) so that it is easier to run the study on the cluster. This is discussed in greater detail in Section 5.5 .
    ${ }^{19}$ The 'number of groups' refers to the number of variables that vary INDEPENDENTLY within the study- i.e., in our LOFT studies, we had variables which had to vary simultaneously (dependently). For instance, variables 1-24 all varied simultaneously, variable 25 varied independently, variables $26-43$ varied simultaneously, variable 44 varied independently, variables $45-46$ varied simultaneously, and variables 4751 varied simultaneously, so we had 6 groups. In the case of the AP600 studies, all 14 variables varied independently and so the number of groups was equivalent to the number of variables, 14.
    ${ }^{20}$ Note that the ' $\$$ ' is not written into the spec file- though it is marked that way in the marked input file. The ' $\$$ ' is appended within the main program, see Appendix A for more information.
    ${ }^{21}$ i.e. VAR1-24 in LOFT study are in group 1, and so a 1 is written in that variable's line.
    ${ }^{22}$ Using the notation from the template spec file in Appendix B the specified number of spec files is \#nodesInStudy, and the node number corresponds to 'currentNode'.

[^11]:    ${ }^{23}$ In the 8 inch break, the average of the correlation coefficients for the various phenomena were used. In the other studies, the three correlation coeffients were in agreement. The exact correlation coefficient values can be found in the $\mathrm{SAS}^{\circledR}$ Reports in Appendix F on Page 58 .

