

CHAPTER 6
TWACKING THROUGH
A PLETHORA OF CONNUCOPIA

OR

Browsing through the field with the intention of locating the meadow muffins so that, when you have to make tracks later, they are not dark brown.

OR

How to drain swamps and avoid alligators.

At this point, we will try to put the popular space-time-computation methods in a family tree. This will help to keep things in perspective when we are whapping around for a method to solve a particular problem. Equally important, it will help us avoid discussing the method that we do finally choose. For further research in this field, we should have, by the end of this chapter, all the basic insight we need to understand the sense of just about any paper on space-time kinetics, find its technical merit, 2) be able to place it on the family tree and 3) assess the good and bad points, assumptions, etc.

The concept of space is not quite so overwhelming if you consider that:
No matter where you go, there you are!

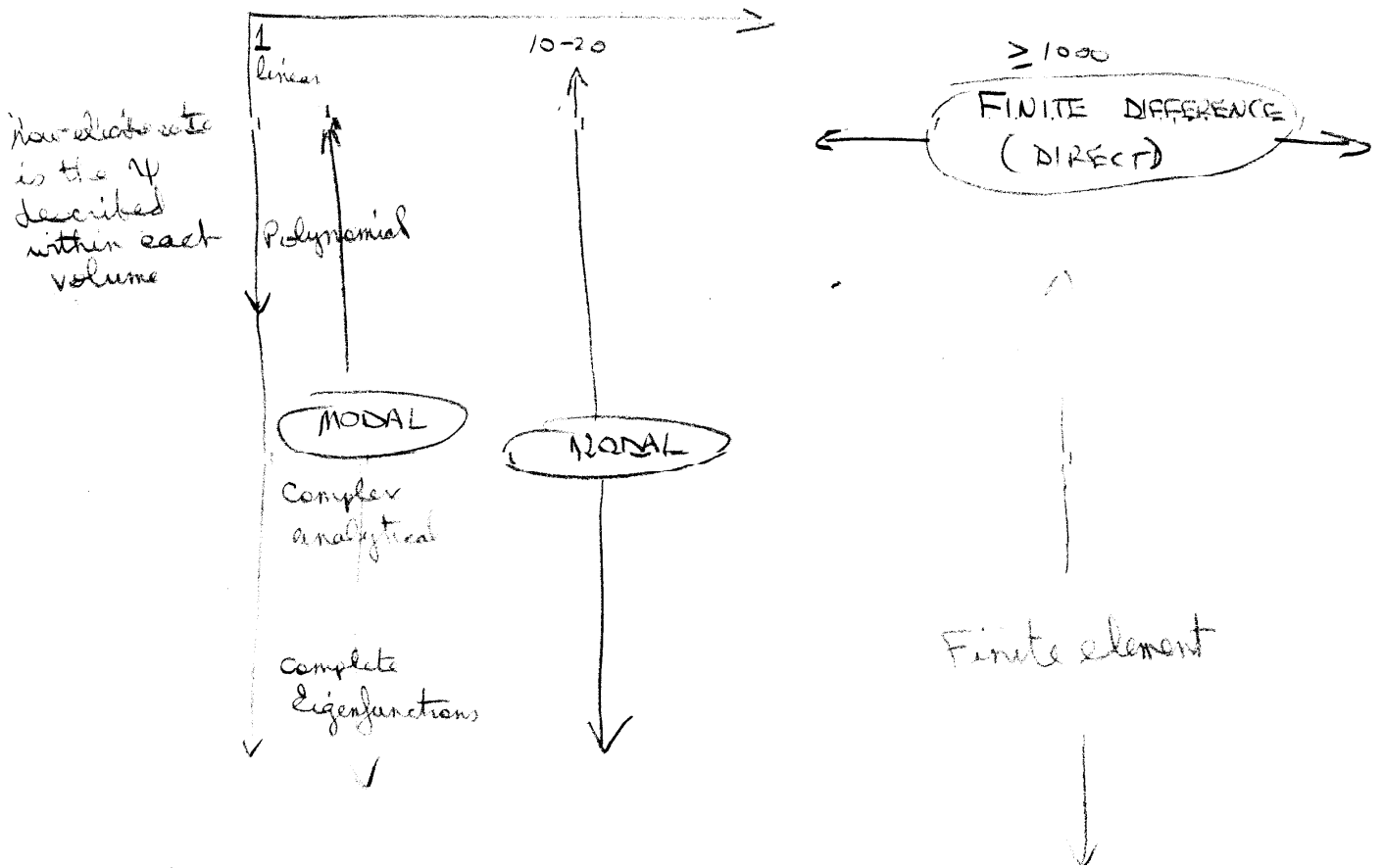
In Hettrick's book (reference 1), Hettrick gives an overview of the field. I've condensed his overview on figure 1. His whole paper, p9-21 is well worth reading.

I guess people like to put things in specific categories; it is convenient. But, at the same time, it can be stifling because it tends to obscure the links and similarities behind all these methods.

In my opinion,

the categories arise basically because of the way the space component is handled. There are really only two aspects of the space issue which describe all the categories:

how many volumes is the core divided up into?



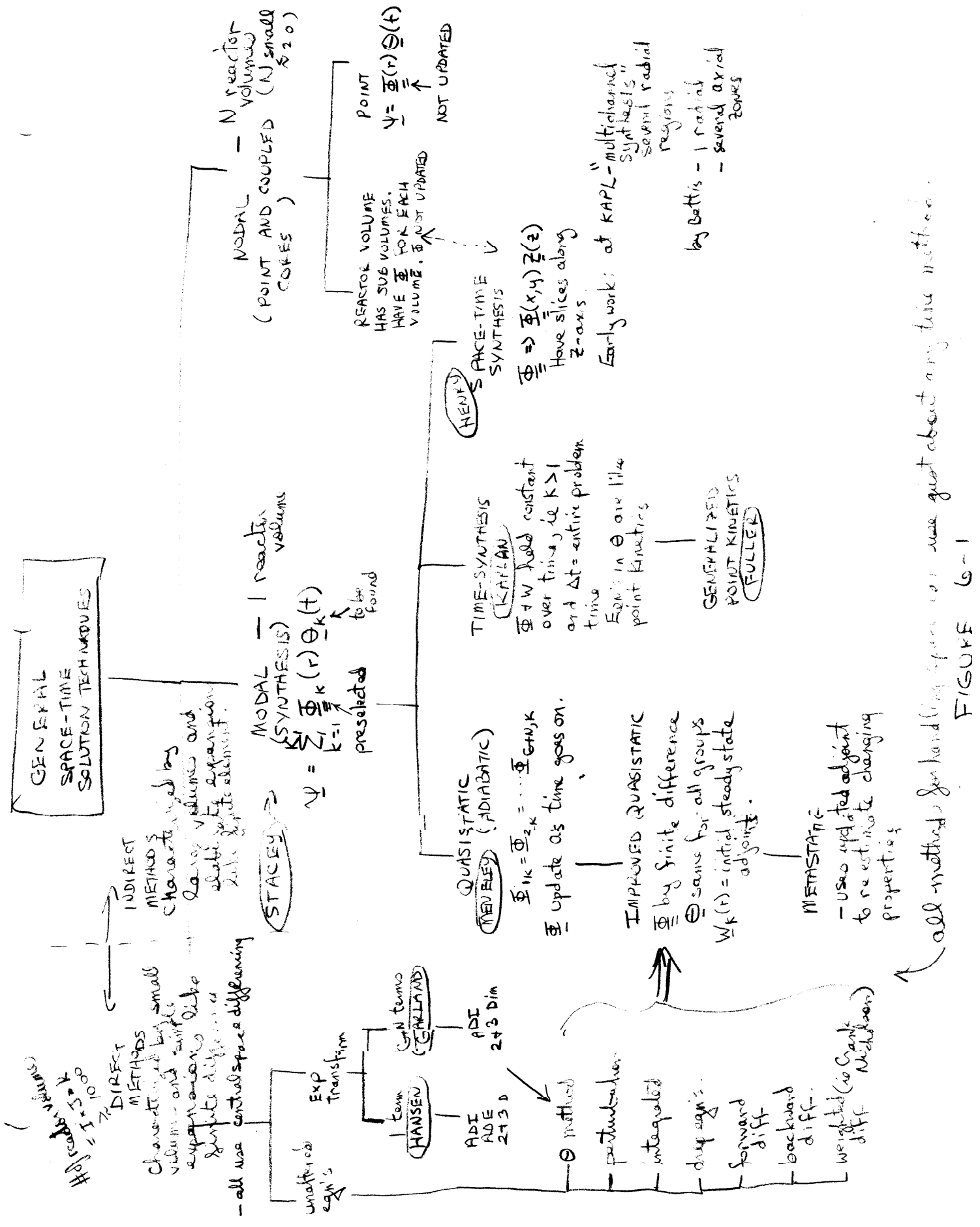


FIGURE 6-1

all methods for handling space are more quiet about any time method.

6-3

see, for instance, in Hetrick's book (ref 1),
Henry p 14 + Hansen p 165-166

There seems to be an ongoing battle as to the best approach. Stacey and Henry are the upholders of the synthesis (nodal) and nodal schools while Hansen believes the best approach is the finite difference approach. Let's summarize the relative merits:

- Finite Difference -
- slow & expensive
 - can estimate error by truncation analysis
 - used for bench marking
 - flexible
 - can easily change Φ eqn's without changing technique much.
 - don't need much intuition.

- Nodal and nodal -
- fast & cheap
 - need lots of intuition to choose Φ
 - can be used to generate insight into harmonics of Φ .
 - method of generating flux and volume averaged parameters are in dispute.
 - no error estimate available.

In the limit, as more variations on old themes hit the marketplace, a continuum of techniques forms. For instance, as higher order approximations are used in finite difference techniques, we approach finite elements. If the finite element volumes were made larger, we move toward nodal techniques. Likewise if you have a nodal technique and simplify the elaborate

- scheme to find $\underline{\Phi}$, $\underline{\Theta}$, etc, and apply it over smaller volumes, you move into nodal and eventually into finite difference techniques.

Now, if you throw in all the possible time solution schemes, you end up with countless variations on a theme. Of course, each investigator of scheme X, or should I say XYZT, claims superior results with his or her method. It is a bit like the old adage - for a limited time...

At this point in time, I would recommend the consumer to buy ^{as a first purchase} a good used cluster which will always crank over even on the coldest morning. Pick a nice stable, central differenced in space, weighted difference in time (or some variant thereof) general E+N, 1, 2 + 3 package that has good I/O including graphics. As an essential extra you'd need a good property library which will do group integration or collapsing of general energy dependent cross sections. You should look for one that will allow interfacing with a thermal/hydraulic package. Nothing fancy, but it will probably beat the marks of the competitors when evaluated on the basis of overall ability to get the total job done from start to finish within a tight, preset job time (including, problem definition, equation setup, programming, debugging, report writing, error analysis, etc).

If you pick one that has an exponential transformation in time using the dominant eigenvalue and one that satisfies the "strongly H-stable" criteria,

- then you may get quite acceptable performance in terms of Δt 's.

As time goes on, you may see the need to pick up one of the nodal or nodal methods. They are faster, but harder to program and are more fickle. But by this time you've gained some experience in the field so that you can pick the $\underline{\Phi}$, etc., appropriate for your problem and you always have your clunker to bench mark against.

Let's take a closer look at some of the more favored methods.

DIRECT (finite difference) as per Hansen, vol. 1 p 65 & fig 1.

→ WIGLE, a 1-D code and TWIGLE, its 2-D counterpart, are implicit based codes which use the following algorithm:

$$\underline{A} = \underline{A}_1 + \underline{A}_2$$

$$\frac{\underline{\Psi}^{j+1} - \underline{\Psi}^j}{\Delta t} = \underline{A}_1 \underline{\Psi}^{j+1} + \underline{A}_2 \underline{\Psi}^j, \quad j \Leftrightarrow \text{time slice}$$

$$\text{or } \underline{\Psi}^{j+1} = [\underline{I} - \Delta t \underline{A}_1]^{-1} [\underline{I} + \Delta t \underline{A}_2] \underline{\Psi}^j \quad 6-1$$

The truncation error is given as:

$$(T.E.) = \frac{\Delta t^2}{2} [\underline{A}^2 - 2\underline{A}_1 \underline{A}_2] + \frac{\Delta t^3}{6} [\underline{A}^3 - 6\underline{A}_1^2 \underline{A}_2] + O(\Delta t^4)$$

∴ $\underline{A}_1 = \underline{A}_2 = \underline{A}/2$ we get Crank-Nicholson.

For 1-D problems this method is attractive since the matrices are easy to invert. For 2 and 3 D, it is better to go with iterative methods, (see below).

→ GARKIN - a 1-D code using an exponential transformation:

$$\underline{\Psi}(H) = e^{-\underline{\Omega}t} \underline{\Theta}(H) \Rightarrow \frac{d\underline{\Theta}}{dH} = e^{-\underline{\Omega}t} (\underline{A} - \underline{\Omega}) e^{\underline{\Omega}t} \underline{\Theta} \quad \leftarrow 6-2$$

But what is $\underline{\Omega}$? How do you choose it?

One popular method is to make $\underline{\Omega}$ a scalar and select its value based on past history. This has good stability characteristics.

Equation 6-2 is rewritten:

$$\frac{d\underline{\Theta}}{dH} + (\underline{Q} - \underline{\Gamma}) \underline{\Theta} = e^{-\underline{\Omega}t} (\underline{P} + \underline{L} + \underline{U}) e^{\underline{\Omega}t} \underline{\Theta} \quad 6-3$$

↑ non-diffusion diagonal
↑ diffusion
↑ upper+lower triangular

Equation 6-3 is then integrated analytically over the time step.

$$\underline{\Theta}_0(T.E.) = -\frac{\Delta t^2}{2} \left[\delta (\underline{W}_0 \underline{I} - \underline{\Gamma} - 2\underline{P}) \right] \underline{U}_0 \quad 6-4$$

where $\delta = \underline{W}_0 \underline{I} - \underline{\Omega} \Leftarrow$ nice vector dimension!

$\underline{U}_0 =$ fundamental mode

$\underline{W}_0 =$ " " " eigenmatrix

LUMAC is a 2-D extension of GARKIN. Replace \underline{P} by

$$P(x) + P(y) \Rightarrow \underline{P} \underline{\Psi} = \underline{P}(x) \underline{\Psi} + B_y^2 \underline{\Psi} \quad 6-5$$

↑ known buckling.

→ MITKIN - 2-D using alternating-direction explicit (ADE) method:

$$\frac{\underline{\psi}^{j+1} - \underline{\psi}^j}{\Delta t} = e^{-\frac{\underline{\Omega}}{2}\Delta t} \left[\underline{A}_1 + \underline{Q}_1 - \underline{\Omega} \right] e^{\frac{\underline{\Omega}}{2}\Delta t} \underline{\psi}^{j+1} + e^{-\frac{\underline{\Omega}}{2}\Delta t} \left[\underline{A}_2 + \underline{Q}_2 \right] e^{\frac{\underline{\Omega}}{2}\Delta t} \underline{\psi}^j$$

and

$$\frac{\underline{\psi}^{j+2} - \underline{\psi}^{j+1}}{\Delta t} = e^{-\frac{\underline{\Omega}}{2}\Delta t} \left[\underline{A}_1 + \underline{Q}_1 \right] e^{\frac{\underline{\Omega}}{2}\Delta t} \underline{\psi}^{j+1} + e^{-\frac{\underline{\Omega}}{2}\Delta t} \left[\underline{A}_2 + \underline{Q}_2 - \underline{\Omega} \right] e^{\frac{\underline{\Omega}}{2}\Delta t} \underline{\psi}^{j+1}$$

where $\underline{A}_1 + \underline{Q}_1 + \underline{A}_2 + \underline{Q}_2 = \underline{A}$
 $\underline{Q}_1 + \underline{Q}_2 = \underline{P}$

6-6

Choose partitioning to make inversion easy, example:

$\underline{Q} = 5$ stripe in 2-D; set $\underline{Q}_1 =$ lower $+\frac{1}{2}$ diagonal

$\underline{Q}_2 =$ upper $+\frac{1}{2}$ diagonal

Trotter algorithm

$$\underline{\psi}^{j+2} = e^{-\frac{\underline{\Omega}}{2}\Delta t} \left[\underline{I} - \Delta t (\underline{A}_2 - \underline{Q}_2 - \underline{\Omega}) \right]^{-1} \left[\underline{I} + \Delta t (\underline{A}_1 + \underline{Q}_1) \right] \times$$

$$\left[\underline{I} - \Delta t (\underline{A}_1 + \underline{Q}_1 - \underline{\Omega}) \right]^{-1} \left[\underline{I} + \Delta t (\underline{A}_2 + \underline{Q}_2) \right] e^{-\frac{\underline{\Omega}}{2}\Delta t} \underline{\psi}^j = \underline{G}_1 \underline{\psi}^j$$

6-7

$$(T.E) = \Delta t^2 \left[e^{\frac{\underline{\omega}_0}{2}\Delta t} (\delta \underline{A} + \delta^2) \right] \underline{U}_0$$

Hansen says this is explicit but it sure looks semi-implicit to me (!?!).

On digging into some papers (ref 2,3 + 4) I found that indeed, MITKIN is semi-implicit but that this particular formalism is called ADE (it is explicit part of the time).

By comparison, the ADI method splits the diffusion term into $X + Y$ parts rather than upper and lower part.

-- STKADI - similar to MITKIN except its implicit (!?) and has a different split for Ψ .

Hansen disagrees if we seem hand waving on why the (T.E.) is not the be-all and end-all for choosing a method. He argues that one should stay away from methods which partition off just the P part or some other part of terms. He says we should keep the terms intact since we know that the norm:

$$\| \underline{A} \underline{\Psi}^i \| = \left\| \frac{\partial \Psi^i}{\partial t} \right\| = \epsilon \text{ where } \epsilon \text{ is small.}$$

That is, the order of magnitude of $\underline{A} \underline{\Psi}^i$ is small compared to $\underline{\Psi}^i$ and we can't say the same thing about such combinations as $\underline{P} \underline{\Psi}^i$ etc.

He says that the exponential transformation operates on the full matrix and makes it smaller since we effectively replace $\underline{\Psi}$ with a more slowly varying function $\underline{\Theta}$. He may be right.

But I'm more inclined to think along the lines of the stability analysis that we've done because I did carry over Hansen's line of thought and tried a full exponential transformation and got really low errors. This turned out to be okay if small time steps were required because of rapidly changing fluxes. But stability problems arise just as Devought predicted because of carrying all the insignificant (large negative) eigenvalues.

MODAL and NODAL

These methods employ:

$$\underline{\Psi} = \sum_{k=1}^K \underline{\Phi}_k(\underline{r}) \underline{\Theta}_k(t) = \sum_{k=1}^K \begin{bmatrix} \underline{\Phi}_{1k} & 0 & 0 & 0 \\ 0 & \underline{\Phi}_{2k} & & \\ & & & \\ & & & \end{bmatrix} \begin{bmatrix} \underline{\Theta}_{1k} \\ \underline{\Theta}_{2k} \\ \vdots \end{bmatrix} \quad 6-8$$

Substitution into $\frac{\partial \underline{\Psi}}{\partial t} = \underline{A} \underline{\Psi}$ gives:

$$\sum_{k=1}^K \underline{\Phi}_k(\underline{r}) \frac{\partial \underline{\Theta}_k(t)}{\partial t} = \sum_{k=1}^K \underline{A} \underline{\Phi}_k(\underline{r}) \underline{\Theta}_k(t) \quad 6-9$$

$$\text{or } \underline{\Phi}_k(\underline{r}) \frac{\partial \underline{\Theta}_k(t)}{\partial t} = \underline{A} \underline{\Phi}_k(\underline{r}) \underline{\Theta}_k(t)$$

$$\text{or } \frac{\partial \underline{\Theta}_k(t)}{\partial t} = [\underline{\Phi}_k(\underline{r})]^{-1} \underline{A} \underline{\Phi}_k(\underline{r}) \underline{\Theta}_k(t) \quad 6-10$$

For each k , we have a weighting function, $\underline{W}_k(\underline{r})$.

Equation 6-10 is premultiplied by $\underline{W}_k(\underline{r}) = \begin{bmatrix} W_{1k}(\underline{r}) & 0 \\ 0 & W_{2k}(\underline{r}) \\ & & \ddots \end{bmatrix}$

and the equation is integrated over the entire reactor volume, leaving K vector equations (of dimension $G+N$) in K vector unknowns (of dimension $G+N$).

The variations on the theme give the family tree as shown in Figure 6-1.

A conclusion is the place where you get tired of thinking.

6-10

CONCLUSION

Enough written, I think. You should know a meadow muffin when you step in one by now, which was the small intent of this segment of the course.

Points to contemplate:

1) In light of this course, finish the statement:

" And it goes without saying that ... "

2) What do you think I should have done for this course? Why?

3) What do you think I should do next? Why?

4) How has your thinking on space-time kinetics changed as a result of this course?

5) Could you apply this knowledge elsewhere?

6) Do you think that your propagator is in better shape now? You do?! ... What if I told you that all of the above was crap? It isn't? $\leftarrow ? \rightarrow$ It is?

How do you know? - maybe we don't you recall that I said Stacey + Hansen made errors, expressed opinions which didn't stand the test of time?

Well, then, let's hear your version.

References

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- 3) Wight, Hansen & Ferguson, Nucl. Sci. & Eng., 44, p239,
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- 4) Ferguson & Hansen, Nucl. Sci. & Eng., 51, p183, 1972