# Ordered Restricted and ModelBased Designs for Partial Orders 

## Examples: Patnaik et al. (JCO, Nov. 2000)

- Tox probabilities follow a

| "Combinati <br> on" | Pacitaxel | Carboplatin |
| :--- | :--- | :--- |
| 1 | 54 | 6 |
| 2 | 67.5 | 6 |
| 3 | 81 | 6 |
| 4 | 94.5 | 6 |
| 5 | 67.5 | 7.5 |
| 6 | 67.5 | 9 |

"partial order" : there exist pairs of combinations for which the ordering of toxicity probabilities is not known


## Method of CDP (2004, Biometrics)

- Based on a method of estimating parameters subject to a partial order given by Hwang and Peddada (1994, Annals of Statistics)
- Idea:
- Estimation subject to a simple (complete) order is well worked out
- There are easy-to-do algorithms to get the estimates


## Hwang and Peddada (1994)

- We know all about combinations 1 and 2
- Tox prob with 1 is less than all the others
- Tox prob with 2 is more than 1 , less than the rest
- Let's guess at the orderings we don't know:
- Guess order is 1-2-3-4-5-6
- Estimate probabilities in this simple order
- Keep these for 1 and 2, throw away the rest


## Hwang and Peddada (1994)

- If we didn't have combinations 5 and 6 , we'd have a simple order: 1-2-3-4
- Hold 1 and 2 at previous estimates, use algorithm for simple order to estimate probabilities for 3 and 4
- If we didn't have combinations 3 and 4 , we'd have a simple order: 1-2-5-6
- Hold 1 and 2 at previous estimates, use algorithm for simple order to estimate probabilities for 5 and 6


## Conway, Dunbar, Peddada (2004)

- Compute estimates by taking all possible guesses at the ordering and averaging the results

| Possible guesses consistent with the partial order |
| :---: |
| $1-2-3-4-5-6$ |
| $1-2-3-5-4-6$ |
| $1-2-3-5-6-4$ |
| $1-2-5-3-4-6$ |
| $1-2-5-3-6-4$ |
| $1-2-5-6-3-4$ |

## Dose allocation

- Once the toxicity probabilities have been estimated, allocate next patient to dose level closest to the target
- Had built in stopping rules, etc.


## CRM for Partial Orders Wages, Conaway and O’Quigley (2011)

- Nolan Wages talk will give more detail
- Think of each possible ordering as a "model" within which we can apply usual CRM

| "model" | Ordering |
| :--- | :--- |
| M1 | $1-2-3-4-5-6$ |
| M2 | $1-2-3-5-4-6$ |
| M3 | $1-2-3-5-6-4$ |
| M4 | $1-2-5-3-4-6$ |
| M5 | $1-2-5-3-6-4$ |
| M6 | $1-2-5-6-3-4$ |

## Order restricted method is harder to program

- Estimation algorithm should only be applied to subset of dose combinations that have at least one patient on them
- Depending where you are in the phase I trial, might have a partial order or simple (complete) order
- If have a partial order, need to be able to generate all simple (complete) orders consistent with the partial order.


## CRM for partial orders seems to have better properties

- In terms of identifying the MTD
- Built on CRM for complete orders
- Intuitive argument: Eventually the data weights the correct ordering and the properties of the usual CRM take over.
- Not quite as simple as this....
- Is \% correctly identifying the MTD the right way to compare?


## Different assumptions and diffent

## goals

- Order restricted: only assumes the partial order is correct
- Gives valid estimates of toxicity probabilities for all combinations?
- In theory, yes
- With number of patients in typical phase I?


## Multiple MTDs?

- CRM: Goal is to a find a combination that has toxicity probability close to target
- Not meant to provide estimate of dose-toxicity profile across the combinations
- Should we be looking for one MTD or set of possible MTDs?


## Both rely on specification of simple (complete) orders: can be many of these

|  |  | Agent 2 |  |  |
| :--- | :--- | :--- | :--- | :--- |
|  | Level | 1 | 2 | 3 |
| Agent | 1 | 1 | 2 | 4 |
|  | 2 | 3 | 5 | 7 |
|  | 3 | 6 | 8 | 10 |
|  | 4 | 9 | 11 | 12 |

M1) 1-2-3-4-5-6-7-8-9-10-11-12
M2) 1-3-2-4-5-6-7-8-9-10-11-12
M3) 1-2-3-5-4-6-7-8-9-10-11-12
M4) 1-2-3-6-4-5-7-8-9-10-11-12

## Limiting the number of orders?

- CDP (2004) paper showed that partial order method works well using a subset of all the order
- As long as the correct order is in the subset
- Properties decline if correct order is not among the subset chosen.


## CRM for partial orders

- Anecdotal evidence suggests that properties may decline as well if correct order is omitted
- From my experience in simulating properties for a couple of combination agent studies
- Unknown if this is true in general
- Unknown how the choice of skeleton affects this


## Can the order restricted method be used with the CRM method?

- To help reduce the number of complete orders considered
- To help choose one or more skeletons


## Choose a subset of complete orders?

- At the end of stage 1 :
- Maximize the In-likelihood under the partial order:
- This may not be computationally easy, but it can be done
- Call this L(PO)


## Could the order restricted method be used to help choose a subset of complete orders?

- Start with a number (M) of complete orders
- At the end of stage 1, for each complete order, $1 \leq m \leq M$,
- Compute estimates under the complete order
- Compute the associated likelihood
- Call this L(m)


## Diagnostics for correct orders

- Prior to beginning partially ordered CRM, drop orderings with small log-likelihoods.
- "Small" relative to either

$$
\begin{aligned}
& L(P O) \text { or } \\
& \sum_{i=1}^{M} L(\mathrm{~m})
\end{aligned}
$$

- Could also look at $\sum_{i=1}^{M} L(m)$ relative to $L(P O)$


## Choosing a complete order subset

- Method does not depend on chosen skeleton values
- Could be viewed in Bayesian framework with priors on the orders or the underlying toxicity probabilities


## Choosing a skeleton

- At the end of stage 1, estimate the toxicity probabilities such to the partial order
- Use smoothed versions of this as the skeleton.

