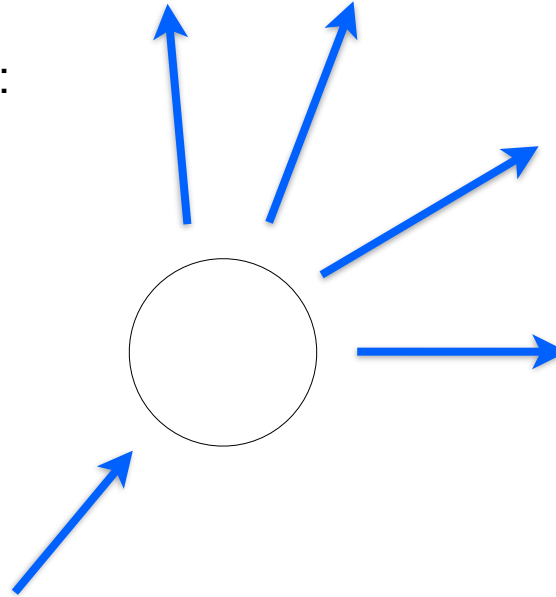


## Advanced language features 2: **SPLIT, EXTEND, WHEN & GROUP**



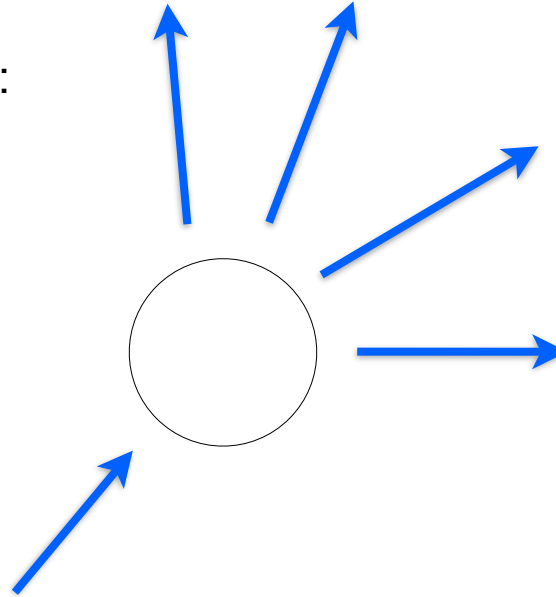
# SPLIT

- Increase statistics beyond this point in the instrumentfile
- $\text{SPLIT } n \text{ MyArm} = \text{Arm}()$
- AT somewhere
- will “formulate an if-statement”:
- for  $j=1:n$
- comp1
- comp2
- comp3
- ...
- end (of instrument)
- ONLY meaningful in case of Monte Carlo choices after SPLIT point...



# SPLIT

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*slight sidetrack....*

# Problem: McStas Single\_crystal.comp “slow” for large unit cell diffraction studies

- Example: Rubredoxin

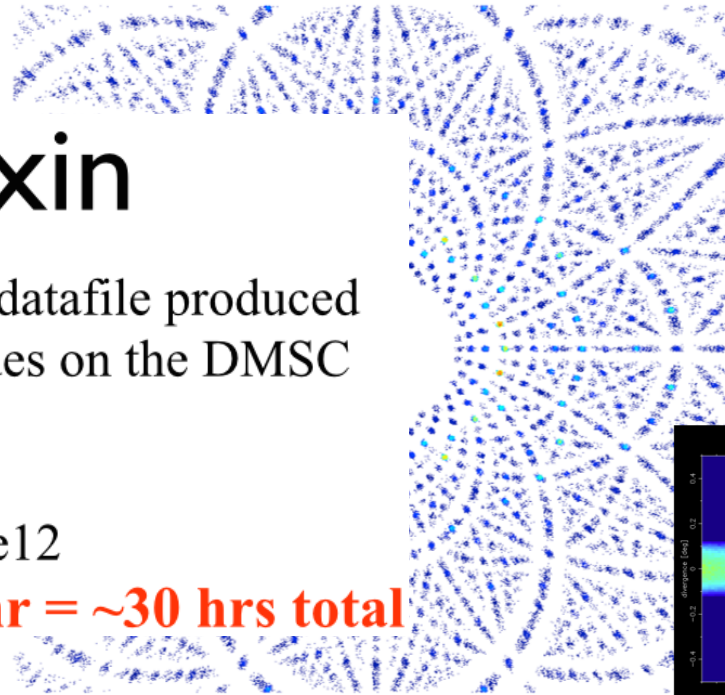
1 timebin, 1000 x,y-bins

## Rubredoxin

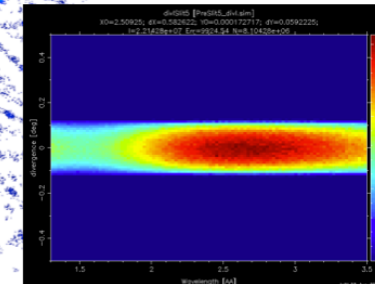
Images created from simulated datafile produced  
August 20th 2012 using 25 nodes on the DMSC  
cluster.

Neutron count: 1e12

**Simulation time: ~10 + ~20 hr = ~30 hrs total**

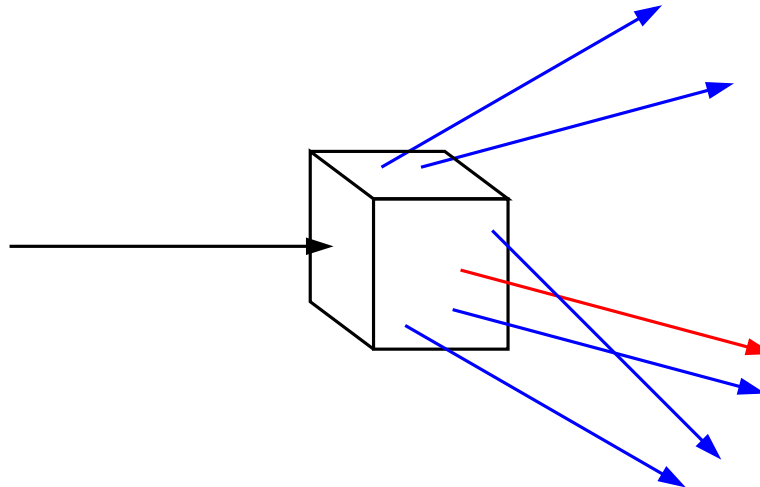


Neutroncount: 1e12  
No gravitation  
Xtal size: 0.5 mm  
Xtal mosaicity: 12'  
Detector: 50 x 50 cm flat  
Detector-to-sample  
distance: 20 cm  
Guide length: 131 m  
Guide dimensions: 9.5 cm  
 $\lambda_{\min} = 1.3 \text{ \AA}$   
 $\lambda_{\max} = 3.5 \text{ \AA}$   
Timespan: 51.39 to 143.4 ms  
Divergence = 0.2 degs



# Algorithm improvement: Use incoming neutrons more efficiently - scatter each one on all possible reflections

- **Red**: Original algorithm, one incoming neutron used only once
- **Blue**: Improved algorithm, each incoming neutron scattered (via SPLIT keyword) all possible times
- Component makes **estimate on average number of “active” diffraction spots** - in the case Rubredoxin this is around **50!**



# GROUP - components working in parallel



*AT (0,0,-LMM) RELATIVE Cradle ROTATED (0,A1/2,0) RELATIVE Cradle  
GROUP IN6Monoks*

*AT (0,0,0) RELATIVE Cradle ROTATED (0,A2/2,0) RELATIVE Cradle  
GROUP IN6Monoks*

*- One comp after the other is “tried” in sequential order until the neutron was  
SCATTERED.*

# EXTEND

- Enrich component behaviour using EXTEND:

```
COMPONENT Mono1 = Monochromator_curved(...)
AT (0,0, -LMM) RELATIVE Cradle ROTATED (0,A1/2,0) RELATIVE Cradle
GROUP IN6Monoks
```

```
EXTEND
```

```
%{
  if (SCATTERED) { myvar = 1; }
}%
```

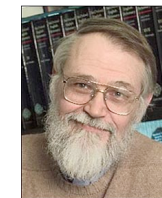
```
...
```

```
COMPONENT Mono2 = Monochromator_curved(...)
AT (0,0, 0) RELATIVE Cradle ROTATED (0,A2/2,0) RELATIVE Cradle
GROUP IN6Monoks
```

```
%{
  if (SCATTERED) { myvar = 2 ;}
}%
```



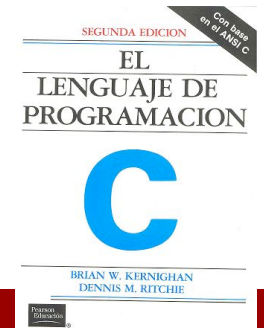
K & R. / GNU



Brian Kernighan



Dennis Ritchie





# WHEN

- Syntax:

COMPONENT Mine = Yours(blah, blah)

WHEN (c-expression) AT (....)

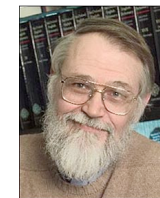
- Is very powerful when combined with EXTEND and user variables, or as a method to let input parameters select if certain components are active.
- Example: Use EXTEND to flag if neutron was scattered on one monochromator blade or another. Then later use WHEN to only show contribution from blade N at sample position?

COMPONENT Mon = PSD\_monitor(...)

WHEN (myvar==1) AT (0,0,0) RELATIVE Sample



K & R. / GNU



Brian Kernighan



Dennis Ritchie

