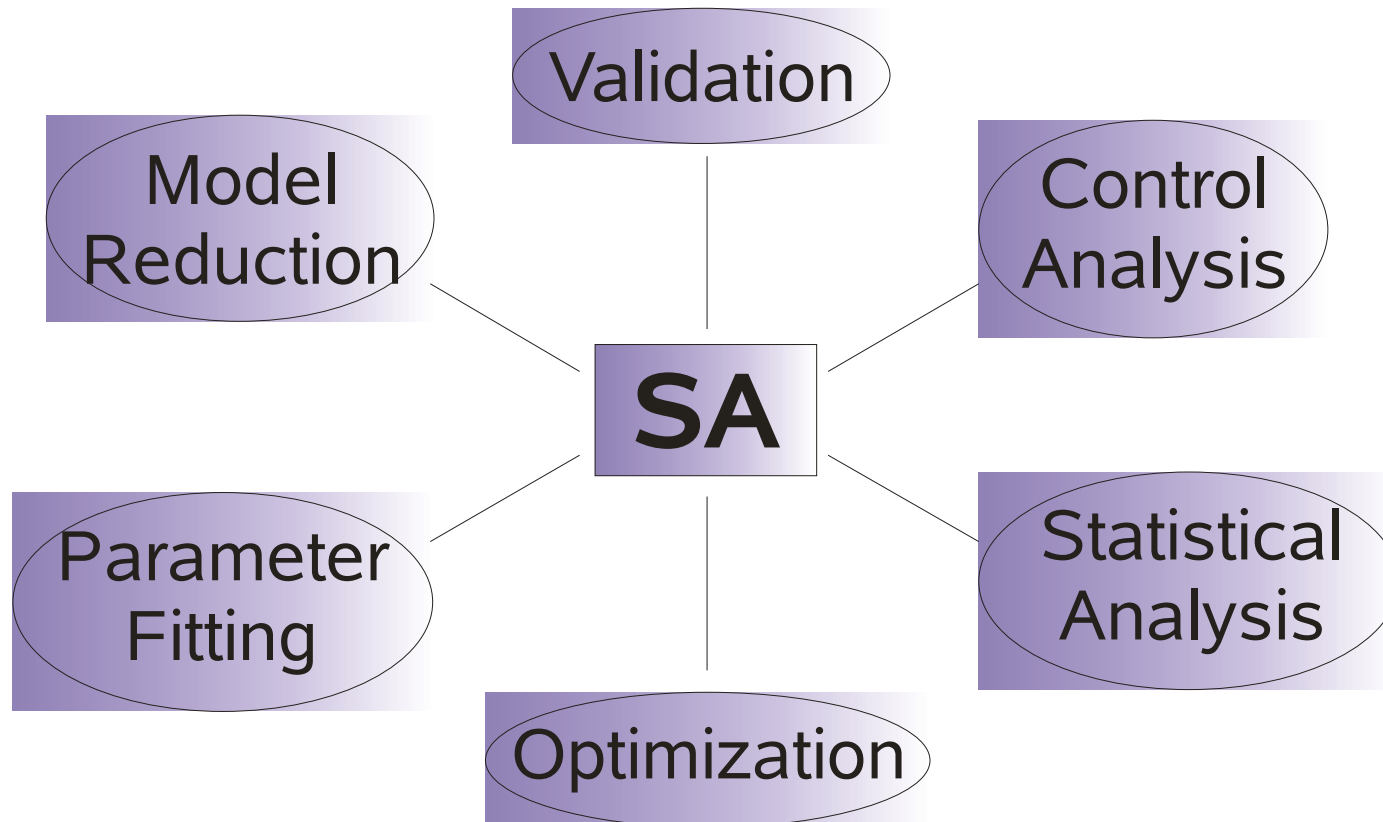


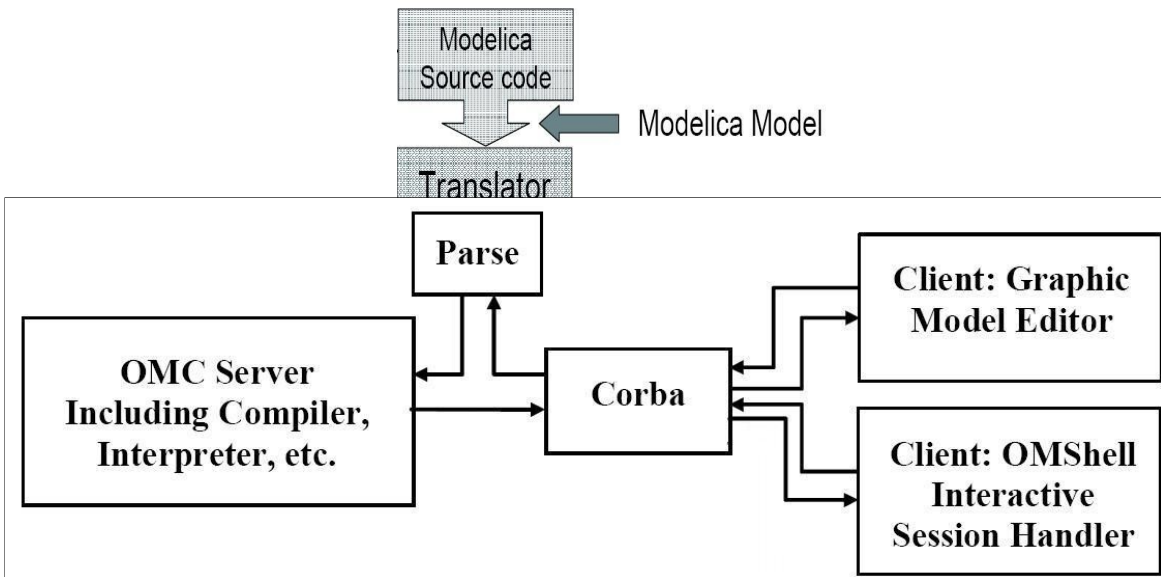
- **Roland Reichardt**
 - Simulation of Mechanical Alloying
- **Thomas Haschke**
 - Numerical Simulation of the Evaporation of Solute Droplets on Polymer Polymer Substrates
- **Marc Kalkuhl**
 - Generation of synthetic SAR-rawdata - Problems and Methods
- **Frieder Hadlich**
 - Simplification of Biochemical Network Models
- **Attiya Elsheikh**
 - Automatic Differentiation of Modelica Models
- **Michael Weitzel**
 - The Topology of Metabolic Isotope Labeling Networks



$S_1 + S_2 \rightarrow P \equiv$

```

connector Metabolite
  Concentration C;
  Parameter Integer M, N;
  Reaction Rin[N] "in reactions";
  Reaction Rout[M] "out reactions";
equation
  Metabolite S1 (N=0, M=1, C=1);
  der(C) = sum(Rin.v) - sum(Rout.v);
end Metabolite;
connector Reaction
  ReactionRate v;
  Parameter Integer M, N;
  Metabolite S[N] "substrates";
  Metabolite P[M] "products";
  Parameter Real alpha;
  ... // other parameters
equation
  v = ... ; // a kinetic formula
end Reaction;
end BiUniReaction
  
```

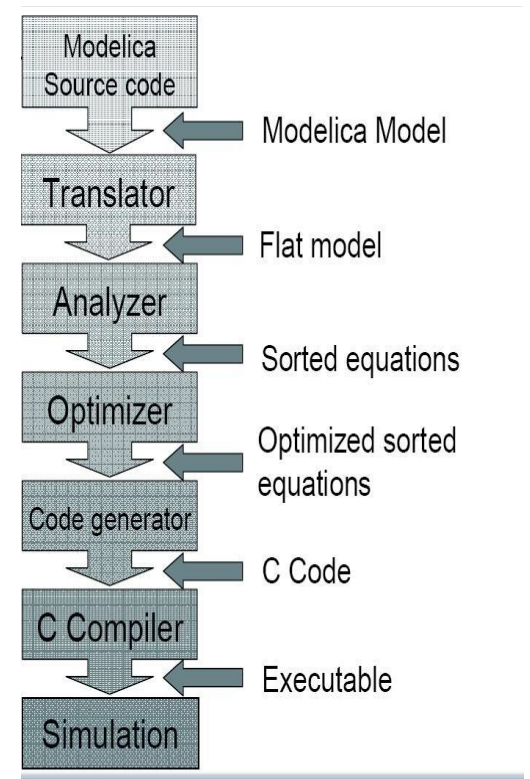
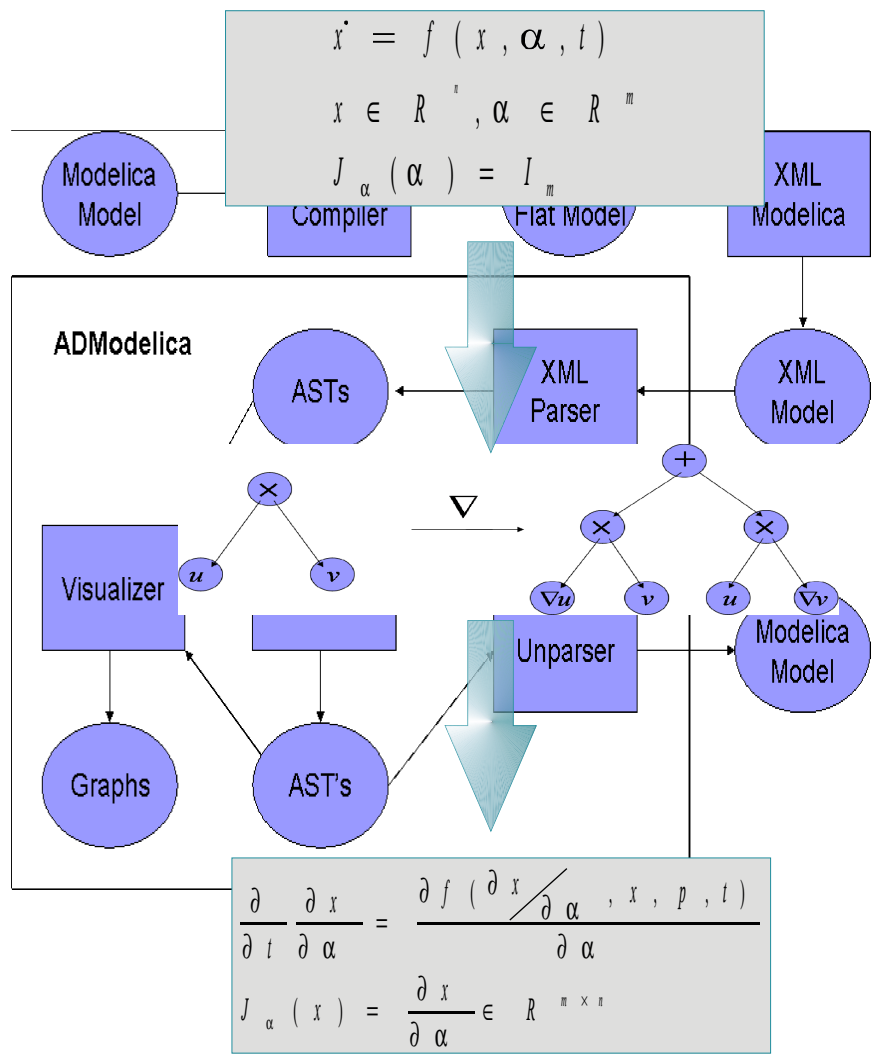


Peter Fritzson



Simulation

Peter Fritzson   



Modelica Model

Flat model

Sorted equations

Optimized sorted equations

C Code

Executable