

Experiences from the application of distributed simulation and their possible relevance for computational systems biology

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- **State-of-the-Art in Distributed Simulation**
- Experiences
 - Case Study “Supply Chain Simulation”
 - Case Study “Training Simulation”
 - Case Study “Digital Manufacturing”
 - Case Study “Agent-based Simulation”
- Trends and Potential Impact
- Summary

Distributed Simulation

- State-of-the-Art

- Distributed Simulation (DS) – a definition:
 - technology that enables a simulation program to be executed on distributed computer systems [Fujimoto 2000]
- Core algorithms have been around for many years
 - Synchronization: conservative/optimistic/hybrid
 - Efficient Data Distribution: Publish/Subscribe Paradigms, Routing Spaces, Dead Reckoning
- Different motivations for DS: From speed-up to interoperability

Distributed Simulation

- State-of-the-Art

- HLA as the leading (but not undisputed) IEEE standard for distributed simulation
 - HLA: High Level Architecture for Modeling and Simulation
 - IEEE 1516-2010 (“HLA Evolved”) as latest version
 - Advantages:
 - Robust industry standard
 - Out of the box solutions for synchronization and efficient data exchange
 - Commercial as well as open source software available
 - Disadvantages:
 - Complex & heavy
 - Limited extensibility

Contents

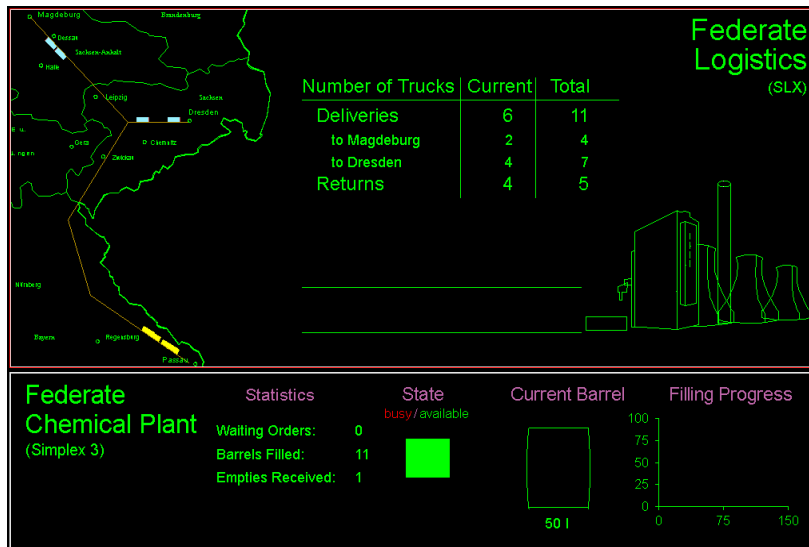
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Case Studies - Background

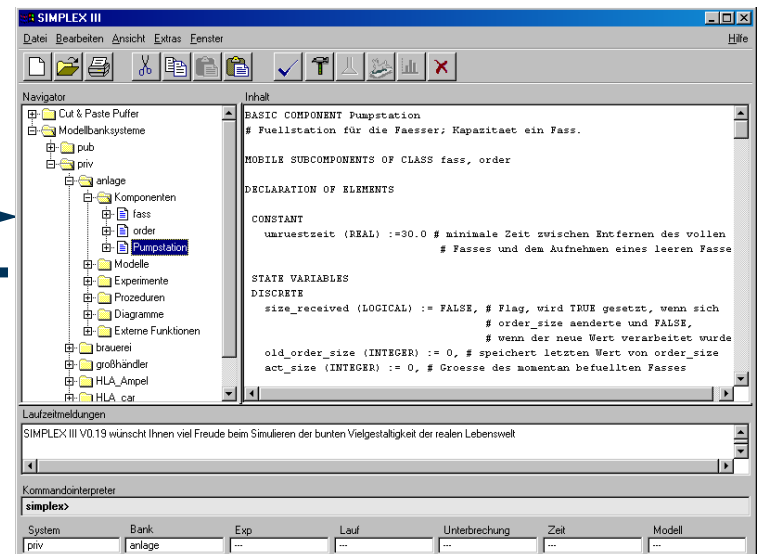
- Involvement with HLA since 1997
- First prototypes for “federating” discrete-event simulation systems (“COTS simulators”) including SLX, Simplex III, Pro Model, Automod, QUEST and IGRIP
- Applications in production and logistics
- From prototypes to industrial applications: Deere & Co. as latest industry partner

Case Study I – Supply Chain Simulation

- Combination of (discrete) logistic model with (continuous) model of chemical production facility

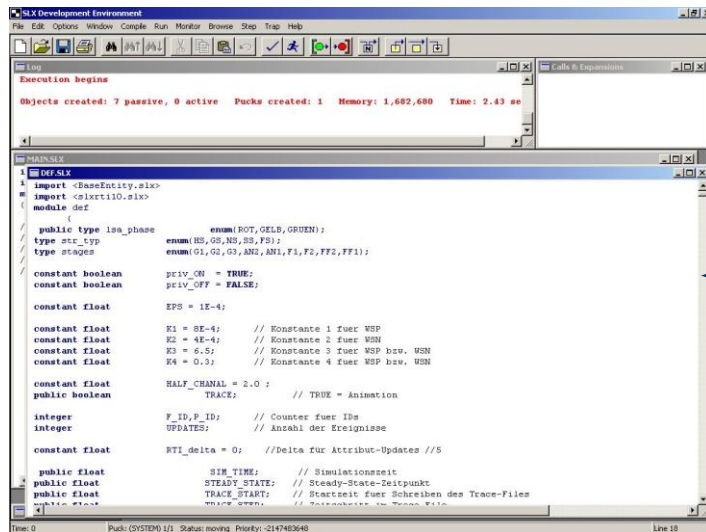


HLA



Case Study II – Training Simulation

- Combination of a real-time driving simulator with an event-based traffic simulation

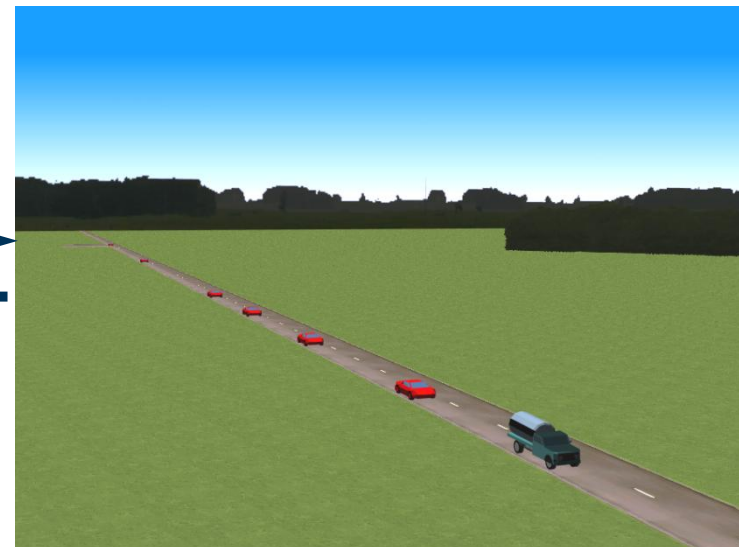


```
SLX Development Environment
File Edit Options Window Compile Run Monitor Browse Step Trap Help

Log
Execution begins
Objects created: 7 passive, 0 active Pucks created: 1 Memory: 1,692,680 Time: 2.43 sec

MAINSEY
1 in 04.slx
import <classFactory.slx>
import <slxrt110.slx>
module def
{
public type lsa_phase enum(R0T,GELB,GRUEN);
type str_type enum(ES,GS,NS,SS,FS);
type stages enum(G1,G2,G3,AN2,AN1,F1,F2,FF2,FF1);
constant boolean PRIV_ON = TRUE;
constant boolean PRIV_OFF = FALSE;
constant float EPS = 1E-4;
constant float K1 = 8E-4; // Konstante 1 fuer WSP
constant float K2 = 4E-4; // Konstante 2 fuer WSN
constant float K3 = 6.5; // Konstante 3 fuer WSP bzw. WSN
constant float K4 = 0.3; // Konstante 4 fuer WSP bzw. WSN
constant float HALT_CHANAL = 2.0; // TRUE = Animation
public boolean TRACE;
integer F_ID,P_ID; // Counter fuer IIs
integer UPDATES; // Anzahl der Ereignisse
constant float RTI_delta = 0; //Delta fuer Attribut-Updates //5
public float SIM_TIME; // Simulationszeit
public float STEADY_STATE; // Steady-State-Zeitpunkt
public float TRACE_START; // Startzeit fuer Schreiben des Trace-Files
public float TRACE_STOP; // Stoppzeit fuer Schreiben des Trace-Files
}
```

SLX/Windows NT/PC

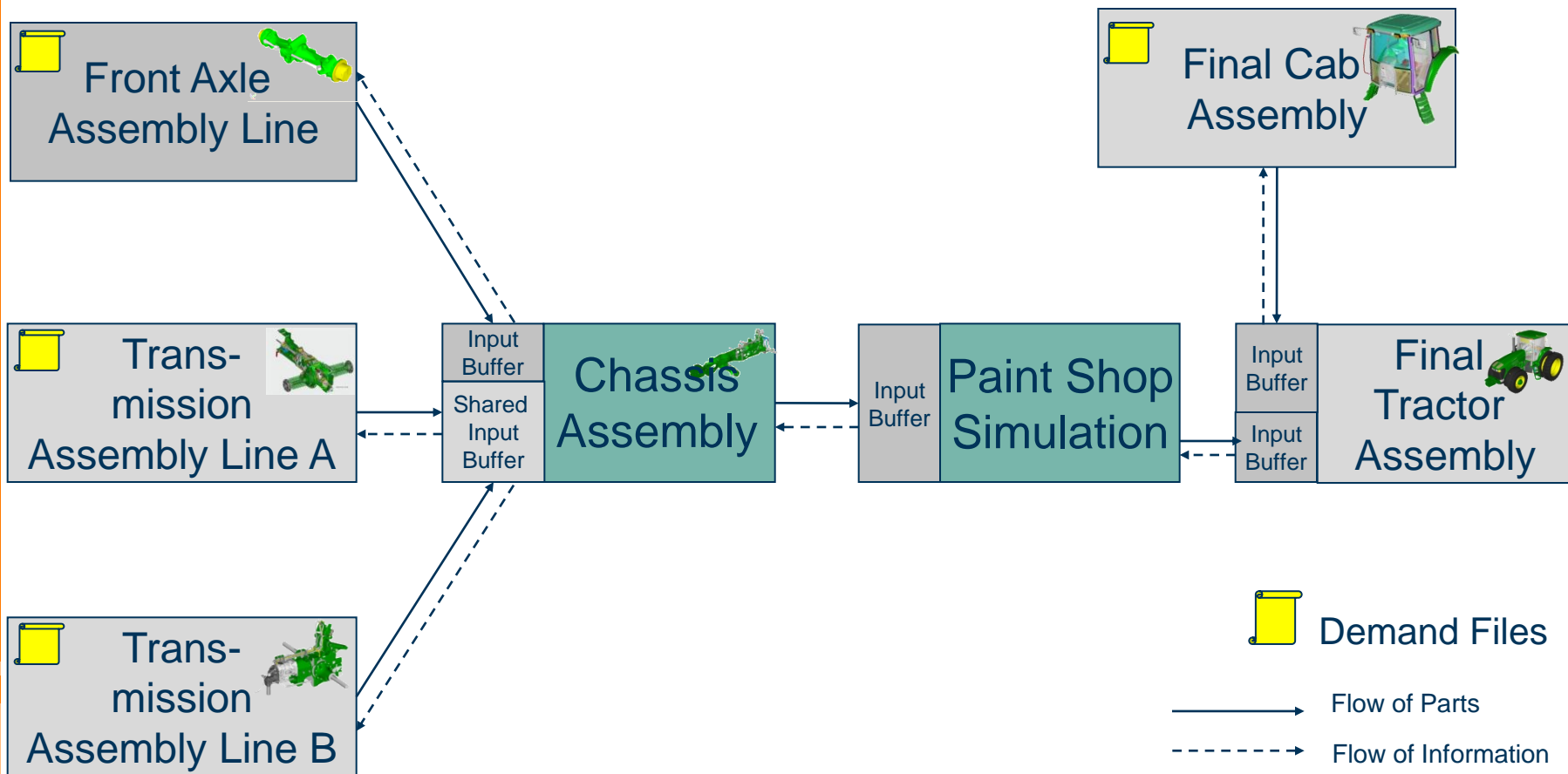


C++/SGI/Irix



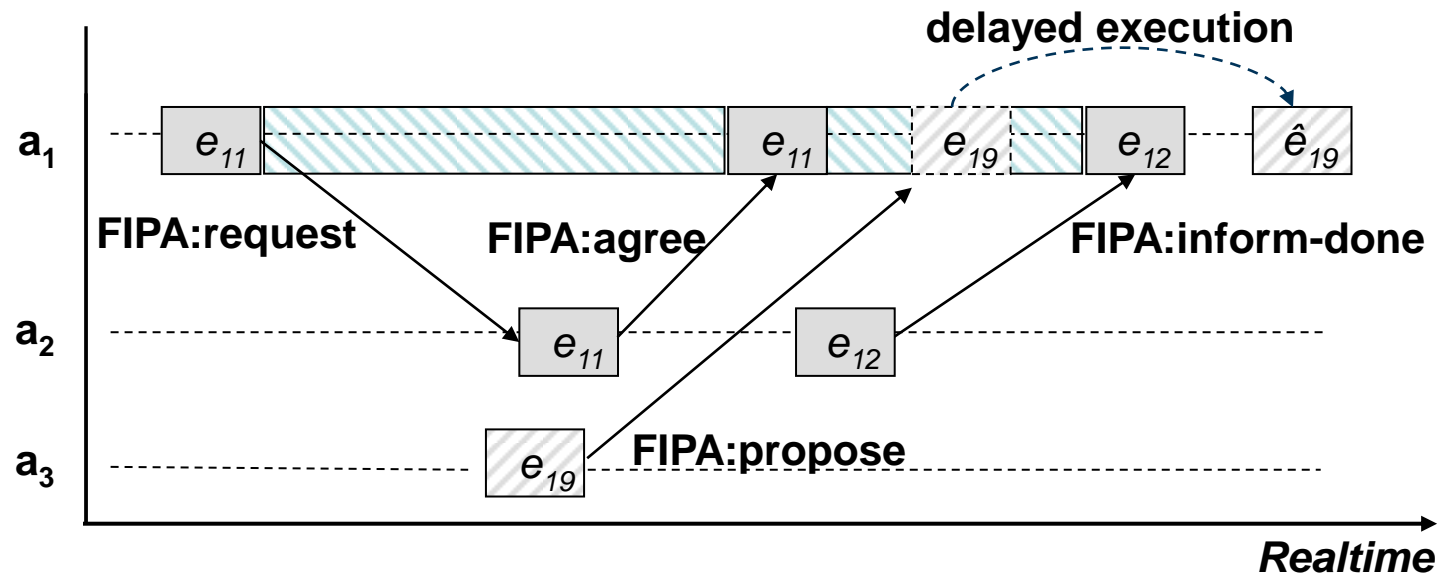
Case Study III – Digital Manufacturing

- Several scenarios combining DES-models



Case Study IV – Agent-based Simulation

- Test of synchronization algorithms for efficient distributed simulation of agent-based models
- Exploitation of properties of agent interaction patterns to reduce risk of rollbacks in time warp



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Lessons Learned

- Intelligent partitioning of models and efficient domain-specific synchronization algorithms are crucial for performance
 - For DES models, conservative synchronization is easily applicable (drawback: dependence on lookahead)
 - Optimistic synchronization promises better performance (if you can incorporate external knowledge)
 - Combined models need special consideration – do not try to distribute DE across models
- HLA is an industry strength DS standard – but there are potentially better solutions for massively parallel simulation requirements

Trends

- Efficient usage of GPU and multicore architectures
 - Huge potential, currently unused in COTS simulation packages
- Mobile devices
 - interface for life training exercises
 - New challenges: energy efficiency of algorithms, ...

Potential Impact for Computational Systems Biology

- Multicore architectures (including GPU) offer huge potential – but require suitable algorithms and modeling paradigms
 - How to compartmentalize models at molecular level so that they can be efficiently simulated on a given architecture?
- Usability of existing synchronization algorithms?
 - For time stepped, round-based models there may be simpler, better scalable synchronization algorithms
 - Properties of modeling paradigm and interaction patterns should be taken into account
- Usability of existing technological solutions and standards?
 - HLA with its current RTI implementations is not too well suited for massively parallel models
 - Potential adaptability for massively parallel applications (“lightweight parallel HLA”)

Questions?

- Thank you for your attention!