SciBmad: A full-featured ecosystem for modern, differentiable accelerator physics simulations

Matt Signorelli, David Sagan

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Accelerator Software Wish-List



red Simulation Program Accelerato Physicist Actu **Dynamic Aperture Program** Lattice Design Program Control System Programs **IBS Simulation Programs** Etc.

□ Modularity!!

- Maximal code re-use, minimal reinventing the wheel
- **Plug-and-play** different optimizers, symplectic integrators, tracking methods, etc. with ease

Q Runs optimally on all architectures, with CPU & GPU parallelization

Easy to use and integrate with other programs/tools

□ *Fully differentiable* using automatic differentiation

• Fast, accurate calculation of gradients for optimizations and machine learning using **forward and backward differentiation**

□ Full featured accelerator software toolkit

• Linear and nonlinear tracking, nonlinear parametric normal forms including spin, Bmad's advanced lattice design tools (e.g. superposition, multipass), etc

Can we have all of this? Enter: SciBmad



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SciBmad: What it is (and isn't)



- SciBmad (formerly called Bmad-Julia) is NOT
 - × A rewrite of the current Bmad in a different programming language. No Fortran code in SciBmad
 - × An interface to the current Bmad. Lattice translation between the two will exist though!
 - × The end of the current Bmad. Maintenance development of the current Bmad will continue
- SciBmad is
 - ✓ Inspired by the experience (both good and bad) with developing the current Bmad
 - ✓ A new software ecosystem for modern, differentiable accelerator physics simulations
 - ✓ Written fully in the **julia** programming language
- By leveraging the julia programming language, SciBmad will achieve all points on the wish-list!





julia ? What's that?

- Julia is a high-level, HPC language that "walks like Python, runs like C"
 - As simple as Python, but as fast as C
- Adopts multiple dispatch and just-intime (JIT) compilation as central paradigms
 - Where types are inferable at compile-time, 1d. it will be compiled (using LLVM toolkit), else dynamically-dispatched with runtime type
- Features a powerful type system for highly-polymorphic code

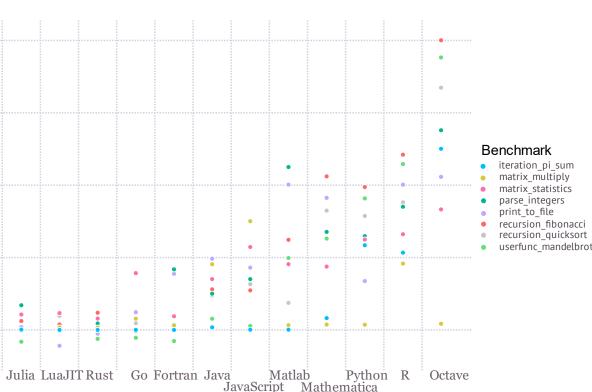


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The Power of Multiple Dispatch and JIT



- Universal polymorphism in Julia is easy and fast! Consider the function:
- Generally pass:

z0 = Vector{Float64}(...)
track_drift(z0, L)

• However, say we'd instead like to track a Taylor map of Truncated Power Series (TPS) defined in some other package. Just pass:

z0 = Vector{TPS64}(...) TI
track_drift(z0, L) JI

That's it! JIT compiled → fast!

- In fact, we can use types from *any* Julia (AD) package
 - E.g. Dual numbers and "tapes" in <u>ForwardDiff.jl</u>, <u>ReverseDiff.jl</u>, <u>Enzyme.jl</u>, <u>Zygote.jl</u>, etc etc.
 - For fun, we can even use <u>Symbolics.jl</u> (compiled!):
- Multiple dispatch and JIT compilation enable massive composability of packages
 - Plug-and-play (and differentiate) to your heart's desire!



function track drift(z0, L)

zf = similar(z0)

5

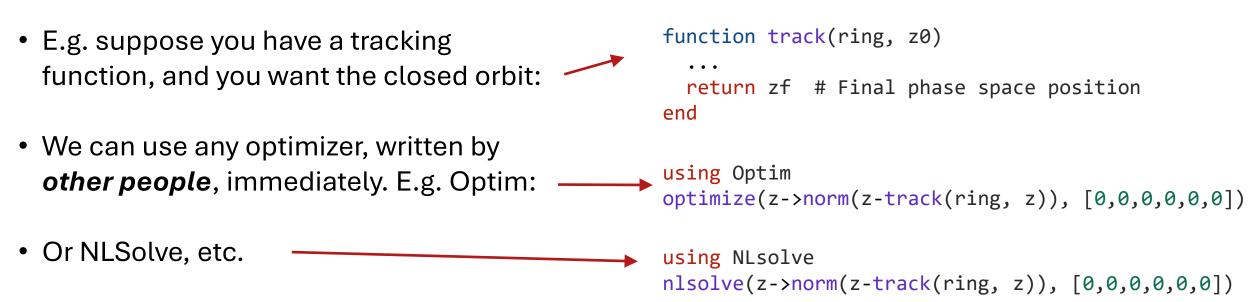
using Symbolics @variables z0[1:6] L # creates symbolic vars track_drift(z0, L)

zf[1] = z0[1]+z0[2]*L/(1.0+z0[6]) zf[2] = z0[2] zf[3] = z0[3]+z0[4]*L/(1.0+z0[6]) zf[4] = z0[4] $zf[5] = z0[5]-L*((z0[2]^2)+(z0[4]^2))/(1.0+z0[6])^2/2.0)$ zf[6] = z0[6]return zf end

SciBmad: Because we are lazy!



- We just showed how easy it is in Julia to use any 'number' type defined by any package
- In fact, we also can use *any optimizer, any (symplectic) integrator, plotting package, architecture-specific parallelization, etc.* written by other people in Julia **with minimal effort!**
 - SciBmad offloads the work from the accelerator physicists to other experts



• This will be JIT compiled + fast too!



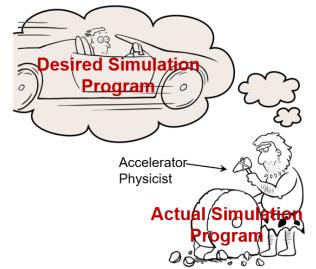
SciBmad: Because we are lazy!



Symplectic Integrators

Note that all symplectic integrators are fixed timestep only.

- SymplecticEuler: First order explicit symplectic integrator
- VelocityVerlet: 2nd order explicit symplectic integrator. Requires f_2(t,u) = v, i.e. a second order ODE
- VerletLeapfrog: 2nd order explicit symplectic integrator.
- PseudoVerletLeapfrog: 2nd order explicit symplectic integrator.
- McAte2: Optimized efficiency 2nd order explicit symplectic integrator.
- Ruth3: 3rd order explicit symplectic integrator.
- McAte3: Optimized efficiency 3rd order explicit symplectic integrator.
- CandyRoz4: 4th order explicit symplectic integrator.
- McAte4: 4th order explicit symplectic integrator. Requires quadratic kinetic energy.
- CalvoSanz4: Optimized efficiency 4th order explicit symplectic integrator.
- McAte42: 4th order explicit symplectic integrator. (Broken)
- McAte5: Optimized efficiency 5th order explicit symplectic integrator. Requires quadratic kinetic energy.
- Yoshida6: 6th order explicit symplectic integrator.
- KahanLi6: Optimized efficiency 6th order explicit symplectic integrator.



- Many forward/backward autodiff packages available
 - ForwardDiff.jl, ReverseDiff.jl, Enzyme.jl, Zygote.jl
 - Our ecosystem will be compatible with all such
- We also don't need to write any symplectic integrators <u>DifferentialEquations.jl</u> already has many *differentiable* ones:
- All of Julia's plotting packages at one's fingertips
 - Makie.jl, Plots.jl, PyPlot.jl, etc
- Powerful scientific ML tools: https://sciml.ai/
- GPU parallelization using <u>CUDA.jl</u> type CuArray (so long as structure-of-arrays used on CPU)
 - Universally polymorphic functions that work on both CPU and GPU!
- Lattice definition itself in the Julia programming language



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But I love Python and refuse to learn Julia!



- That's ok: we will have a full-featured Python interface to the (fast) SciBmad Julia ecosystem
- Gradients output for use with **PyTorch** and **Xopt** will be simple
- All other features will also be available, except the lattice definition will always be in Julia (with translators from other formats available)
- For Julia users, the experience will be fantastic. For Python users, it will be as good as the usual two-language experience: fast underlying library, with a full-featured Python wrapper for ease of use



Current SciBmad Ecosystem Status



Stable, officially-registered, ready for use:

• **<u>GTPSA.jl</u>**: Julia interface to L. Deniau's Generalised Truncated Power Series Algebra library

In development:

- AcceleratorLattice.jl: Accelerator lattice definition/manipulation
- NonlinearNormalForm.jl: Parametric nonlinear normal forms and analysis of DA maps including spin using Lie algebraic methods
- AtomicAndPhysicalConstants.jl: Atomic/subatomic particle properties and other physical constants for simulations

Starting development:

• **BeamTracking.jl:** CPU/GPU particle tracking methods/interfaces



AcceleratorLattice.jl



Accelerator lattice construction and manipulation is done using the Julia language itself:

```
using AcceleratorLattice
# Returns a FODO cell with specified quad strength
function FODO(k1)
 @eles begin
    qf = Quadrupole(L = 0.6, Kn1 = k1)
    d = Drift(L = 0.4)
    qd = Quadrupole(L = -0.6, Kn1 = -k1)
  end
  return BeamLine([qf, d, qd, d])
end
@ele begin0 = BeginningEle(pc_ref = 1e7, species_ref = Species("electron"))
# Construct a BeamLine using Julia functions!
my_beamline = BeamLine([begin0, FODO(0.36), FODO(0.30)])
my lat = Lat([my beamline])
```



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AcceleratorLattice.jl



Fully-featured lattice elements:

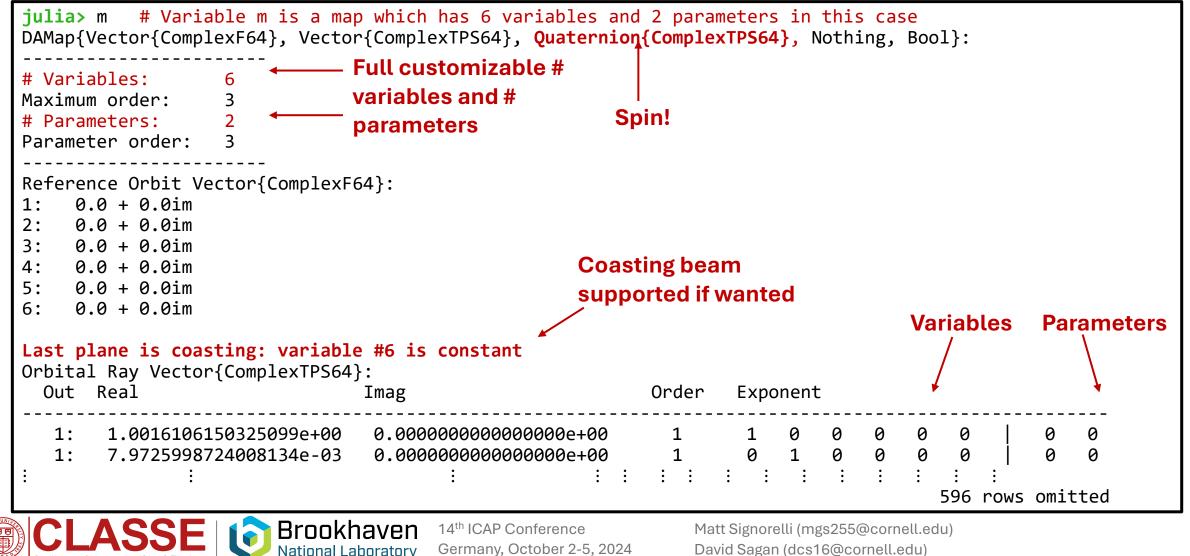
<pre>julia> show(lat["qf"][1])</pre>			
Ele: "qf" (b1>>2) Quadrupole			
AlignmentGroup:			
offset	[0.0, 0.0, 0.0] m	offset_tot	[0.0, 0.0, 0.0] m
x_rot	0 rad	x_rot_tot	0 rad
y_rot	0 rad	y_rot_tot	0 rad
tilt	0 rad	tilt_tot	0 rad
BMultipoleGroup:			
Order Integrated	Tilt (rad)		
1 false	0.0	0.34 Kn1	0.0 Ks1 (1/m^2)
	-0.0	11341179236737171 Bn1	-0.0 Bs1 (T/m^1)
FloorPositionGroup:			
r (r_floor)	[0.0, 0.0, 0.0] m		
q (q_floor)	1.0 + 0.0·i + 0.0·j + 0.0·k		
theta (theta_floor)	0.0 rad		
phi (phi_floor)	0.0 rad	psi (psi_floor)	0.0 rad
LengthGroup:			
L	0.6 m	orientation	1
S	0.0 m	s_downstream	0.6 m
ReferenceGroup:			
species_ref	<pre>Species("electron")</pre>	<pre>species_ref_exit</pre>	<pre>Species("electron")</pre>
pc_ref	1.0e7 eV	<pre>pc_ref_exit</pre>	1.0e7 eV
E_tot_ref	1.000000005e7 eV	E_tot_ref_exit	1.00000005e7 eV
Etc			



NonlinearNormalForm.jl



Real and complex parametric DAMaps including spin and coasting beam, for example:



NonlinearNormalForm.jl



All the following tools are implemented already:

julia> m_lin = cutord(m, 2); # extract the linear part in orbital

julia> m_nonlinear = inv(m_lin) • m; # remove the linear part

julia> F = log(m_nonlinear); # Get the Lie operator (including quaternion) generating nonlinear part

julia> m = m_lin • exp(F); # Reconstruct same map using Lie exponent and linear part separately

julia> a = normal(m); # Calculate the nonlinear (parametric) normalizing canonical transformation

julia> R_z = inv(a) o m o a; # Nonlinear amplitude-dependent rotation in regular phase space (x, px, ...)

julia> c = to_phasor(m); # Get the transform to phasors basis $\sqrt{(J)*exp(\pm im*\phi)}$

julia> R_J = inv(c) • R_z • c; # Nonlinear amplitude-dependent rotation in phasors basis

julia> a_spin, a0, a1, a2 = factorize(a); # Spin part, nonlinear parameter-dependent fixed point, a1, a2

julia> Σ = equilibrium_moments(m, a); # Calculate equilibrium sigma matrix when fluctuation-dissipation

julia> a = normal(m, m=[0; 1], m_spin=[-1]); # Leaving in a Q_y - Q_spin resonance



Conclusions



SciBmad is a modern accelerator physics ecosystem which will provide:

Modularity!!

- Maximal code re-use, minimal reinventing the wheel
- Plug-and-play different optimizers, symplectic integrators, tracking methods, etc. with ease

M Runs optimally on all architectures, with CPU & GPU parallelization

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• Fast, accurate calculation of gradients for optimizations and machine learning using forward and backward differentiation

Full featured accelerator software toolkit

- Linear and nonlinear tracking, nonlinear parametric normal forms including spin, Bmad's advanced lattice design tools (e.g. superposition, multipass), etc
- Goal: first accelerator simulations by end of year



Contributors and Thank You!



- Development of SciBmad is currently in full-gear, and would not be this far along without the help of many
 - Dan Abell (BeamTracking.jl and AtomicAndPhysicalConstants.jl)
 - J. Scott Berg
 - Oleksii Beznozov (BeamTracking.jl GPU)
 - Alex Coxe (AtomicAndPhysicalConstants.jl)
 - Laurent Deniau
 - Auralee Edelen
 - Etienne Forest (significant help with NonlinearNormalForm.jl)
 - Juan-Pablo Gonzalez
- Open to more collaborators!



- Georg Hoffstaetter de Torquat
- Gavin Hunsche (BeamTracking.jl)
- Lixing Li (AtomicAndPhysicalConstants.jl)
- Chris Mayes
- Ryan Roussel
- David Sagan (AcceleratorLattice.jl)
- Matt Signorelli (NonlinearNormalForm.jl and GTPSA.jl)
- Sophia Yang (BeamTracking.jl)



Thank you! Questions?



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Backup Slides



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C++: Single Dispatch

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```
-class A { };---
class B : public A { };
class C : public A { };
class Foo {
  virtual void my fun(A* arg1, A* arg2) { std::println("A,A"); }
  virtual void my fun(B* arg1, B* arg2) { std::println("B,B"); }
  virtual void my fun(C* arg1, B* arg2) { std::println("C,B"); }
  virtual void my fun(B* arg1, C* arg2) { std::println("B,C"); }
  virtual void my fun(C* arg1, C* arg2) { std::println("C,C"); }
};
void call my fun(A* arg1, A* arg2) {
   Foo *pFoo = new Foo;
  pFoo->my_fun(arg1, arg2); // SINGLE DISPATCH: prints "A,A"
}
int main() {
  A^* arg1 = new B();
  A^* arg2 = new C();
   call my fun(arg1, arg2); // prints "A,A"
  return 0;
}
                                         14<sup>th</sup> ICAP Conference
                          Brookhaven
```

 C++ only uses single dispatch, so this will print "A,A" even though both are B,C

Julia: Multiple Dispatch



```
abstract type A end
struct B <: A end
struct C <: A end</pre>
```

```
my_fun(arg1::A, arg2::A) = println("A,A")
my_fun(arg1::B, arg2::B) = println("B,B")
my_fun(arg1::C, arg2::B) = println("C,B")
my_fun(arg1::B, arg2::C) = println("B,C")
my_fun(arg1::C, arg2::C) = println("C,C")
```

```
function call_my_fun(arg1::A, arg2::A)
  return my_fun(arg1, arg2)
end
```

```
arg1 = B()
arg2 = C()
```

```
call_my_fun(arg1, arg2) # prints B,C
```

- Julia will dynamically-dispatch based on the runtime types
- Call to my_fun done by going to method lookup table using runtime types, and JITcompiling my_fun if not already compiled

