Abstraction in Technical Computing

by

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Abstract

Array-based programming environments are popular for scientific and technical computing. These systems consist of built-in function libraries paired with high-level languages for interaction. Although the libraries perform well, it is widely believed that scripting in these languages is necessarily slow, and that only heroic feats of engineering can at best partially ameliorate this problem.

This thesis argues that what is really needed is a more coherent structure for this functionality. To find one, we must ask what technical computing is really about. This thesis suggests that this kind of programming is characterized by an emphasis on operator complexity and code specialization, and that a language can be designed to better fit these requirements.

The key idea is to integrate code *selection* with code *specialization*, using generic functions and data-flow type inference. Systems like these can suffer from inefficient compilation, or from uncertainty about what exactly to specialize on. We show that dispatch on structured type tags helps address these problems. The resulting language, Julia, achieves a Quine-style "explication by elimination" of many of the productive features technical computing users expect.

Thesis Supervisor: Alan Edelman

Title: Professor

Acknowledgments

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Chapter 1

Introduction

Scientific computing has evolved from being essentially the only kind of computing that existed, to being a small part of how and why computers are used. Over this period of time, programming tools have advanced in terms of the abstractions and generalizations they are capable of.

Science as a whole evolves through the incorporation of specialized bits of knowledge into more powerful general theories. There is a parallel in the world of programming languages: special-purpose languages have been subsumed by more powerful and general languages. How has this trend affected scientific computing? The surprising answer is: not as much as we would like. We see scientists and technical users generally either turning away from the best new programming languages, or else pushing them to their limits in unusual ways.

In fact, we have discovered at least 49 programming languages designed for technical computing since Fortran (figure 1-1). This high number begs for an explanation. We propose that technical users crave the flexibility to pick notation for their problems, and language design is where you go when you need this level of flexibility. Language design is associated with the highest level of abstraction. The desire for scientists to rework their code is not so surprising when one reflects on the tradition of inventing new notation in mathematics and science.

The need for language-level flexibility is corroborated by the ways that the technical

Productivity				Performance
Matlab	Maple	Mathematica	SciPy	Fortress
SciLab	IDL	R	Octave	Chapel
S-PLUS	SAS	J	APL	X10
Maxima	Mathcad	Axiom	Sage	SAC
Lush	Ch	LabView	O-Matrix	ZPL
PV-WAVE	Igor Pro	OriginLab	FreeMat	
Yorick	GAUSS	MuPad	Genius	
SciRuby	Ox	Stata	JLab	
Magma	Euler	Rlab	Speakeasy	
GDL	Nickle	gretl	ana	
Torch7	A+	PDL	Nial	

Figure 1-1: 49 technical computing languages classified as primarily designed for productivity or performance

computing domain uses general purpose languages. Effective scientific libraries extensively employ polymorphism, custom operators, and compile time abstraction. Code generation approaches (writing a program that writes the needed program) are unusually common. Most of these techniques are presently fairly difficult to use, and so programmers working at this level give up the notational conveniences of the purpose-built languages above.

This thesis addresses the essential challenges created by this state of affairs. We argue that the situation can be improved considerably by abstracting, with a new language.

Compiler techniques, library design, high-performance computational kernels, new algorithms, and approaches to parallelism are all important to technical computing. However these sorts of technologies can usually be applied to multiple languages, as has happened in the C and Fortran language families. However in conjunction with a new language, these technologies can flourish. Indeed, we believe that language design has reached a level of importance that even (perhaps temporarily) supersedes the traditional focus of cache utilization and the use of matrix operations.

In brief, our idea is to *integrate code selection and specialization*. The aforementioned flexibility requirement can be explained as a sufficiently powerful means for *selecting* which piece of code to run. This notion subsumes method dispatch, function overloading, and

Mainstream PL	Technical computing
classes, single dispatch	complex operators
separate compilation	performance, inlining
parametric polymorphism	ad-hoc polymorphism, extensibility
static checking	experimental computing
modularity, encapsulation	large vocabularies
eliminating tags	self-describing data, acceptance of tags
data hiding	explicit memory layout

Figure 1-2: Priorities of mainstream object-oriented and functional programming language research and implementation compared to those of the technical computing domain.

potentially even branches. When such mechanisms are used in technical computing, there is nearly always a corresponding need to *specialize* code for specific cases to obtain performance. Polymorphic languages sometimes support forms of specialization, but often only through a designated mechanism (e.g. templates), or deployed rather conservatively out of resource concerns. We intend to show that an equilibrium point in the design space can be found by combining selection and specialization into a single dynamic multiple dispatch mechanism. The specific design is identified via subtyping theories and closure under data-flow operations.

1.1 The technical computing problem

Figure 1-2 compares the general design priorities of mainstream programming languages to those of technical computing languages. The priorities in each row are not necessarily opposites or even mutually exclusive, but rather are a matter of emphasis. It is certainly possible to have both parametric and ad-hoc polymorphism within the same language, but syntax, recommended idioms, and the design of the standard library will tend to emphasize one or the other. Of course, the features on the left side can also be useful for technical computing; we exaggerate to help make the point.

It is striking how different these priorities are. We believe these technical factors have

contributed significantly to the persistence of specialized environments in this area. Imagine you want just the features on the left. Then there are many good languages available: Haskell, ML, Java, C#, perhaps C++. Getting everything on the right, by comparison, always seems to be awkward. The most popular approach is to use multiple languages, as in e.g. NumPy [57], with a high-level productivity language layered on top of a large library written in lower-level languages. Seeking a similar tradeoff, others have gone as far as writing a C++ interpreter [58].

1.2 Solution space

It is clear that any future scientific computing language will need to be able to match the performance of C, C++, and Fortran. To do that, it is almost certain that speculative optimizations such as tracing [24] will not be sufficient — the language will need to be able to *prove* facts about types, or at least let the user request specific types. It is also clear that such a language must strive for maximum convenience, or else the split between performance languages and productivity languages will persist.

It is fair to say that two approaches to this problem are being tried: one is to design better statically-typed languages, and the other is to apply program analysis techniques to dynamically typed languages. Static type systems are getting close to achieving the desired level of flexibility (as in Fortress [5] or Polyduce [14], for instance), but it is still too early to call a winner between these two approaches (if, indeed, there even needs to be a winner).

Here we will use a dynamically typed approach, with approximate type inference (for an excellent discussion of the many meanings of the word "type" see [32]). There are several reasons for this. First, we want to emphasize that insufficient static checking is most likely not the current limiting factor in scientific computing productivity. Second, some idioms in this domain appear to be inherently, or at least naturally, dynamically typed (as we will explore in later chapters). Third, there has been a sense that "dynamic" or "scripting" language users do not want to hear about types: they are thought of as a distraction and

Put in a ref to these chapters by number please.

associated with verbosity and nuisance compiler errors. We hope to contribute an example of a language where types are useful and not burdensome.

1.2.1 The excess power problem

We claim that the amount of information that can be statically inferred exceeds most dynamic languages' capacity to exploit it. Much work has been done on program analysis and optimization techniques for dynamically typed languages. When static analyses (often incorporating run time information) are applied to dynamically typed programs, it is typically possible to recover a significant amount of type information (TODO cite). What, then, can one do with this information? If the goal is performance, various partial evaluations can be done: generating code without type checks, removing branches, type-specializing the storage of variables, and compile time method lookup are all valuable and yield large real-world gains.

For example, if method calls are dispatched on the first argument, but the types of all arguments can be inferred, some power has been "left on the table" — we could have had multi-methods for little extra cost. In fact, method-at-a-time JIT compilers (TODO cite) can specialize method bodies on all arguments, and might use multiple dispatch *internally* to select implementations at run time (TODO cite a system that did this). This argument does not apply equally to statically-typed languages, since they cannot simply "switch" their functions to generic functions without significant consequences for type checking.

This "excess power" problem applies to data structures as well. For example, static or run time analysis might reveal that a certain array can be represented as a native Int32 array [8]. If this information is not reflected in the source language, then certain uses like passing data to native code become unnecessarily more complicated. And if one is going to implement homogeneous arrays anyway, why not let programmers request them?

Some levels of performance are difficult to reach with implicitly specialized code and data. Given the knowledge that an array contains only Int32 data, we might want to go beyond essential optimizations like storing intermediate values in registers, and actually use different

cite

cite

algorithms. For example, in Miller-Rabin primality testing, checking three "witness" values suffices for all 32-bit arguments, but up to seven values might be needed for 64-bit arguments .(TODO cite) In cryptographic applications, exploiting this difference in an inner loop could cite bring significant benefits.

1.2.2 The divergence problem

Another problem that occurs when analyzing programs with complex type behavior is divergence: the analysis is likely to infer an overly-large result from failing to eliminate enough possible behaviors. Narrowing the inferred types requires some extra source of type information. Multi-method signatures work well for this purpose.

However the divergence problem also places a limit on what kinds of dispatch specifications are useful to program analyses: some sets of values tend to be more robust under execution than others.

Would love an example if possible

1.2.3 Staged programming

In demanding applications, selecting the right algorithm might not be enough, and we might need to automatically *generate* code to handle different situations. While these cases are relatively rare in most kinds of programming, they are remarkably common in technical computing. Code generation, also known as *staged programming*, raises several complexities:

- What is the input to the code generator?
- When and how is code generation invoked?
- How is the generated code incorporated into an application?

These questions lead to practical problems like ad-hoc custom syntax, extra build steps, and excessive development effort (writing parsers and code generators might take more work than the core problem). If code needs to be generated based on run time information,

then a staged library might need to implement its own dispatch mechanism for invoking its generated code.

We find that our particular style of dynamic multiple dispatch provides an unusually good substrate for staged programming. The reason is that the language, though semantically dynamically typed, has a canonical (and not optional!) type analysis stage. This is a natural place to hook in code generators: semantic (not just syntactic) information is available, but machine code has not been generated yet. In general, this semantic information can also include run time (and always, at least, approximate run time) information. Since the types used by dispatch have rich, nested structure, they convey enough information to drive code generation for a significant class of problems. In this framework code generators are invoked automatically by the compiler, and the results can be called directly or even inlined into user code. Our approach is closely related to exotypes [19].

1.3 Contributions of this thesis

Our first contribution is a discussion of the nature of technical computing that suggests which language-level abstractions might best support real use cases. Simply put technical computing has by and large happened without sufficient probing and analysis as to what it is. Our looking at the space in this manner has lead to a novel design. The success of this design is resonating with users worldwide.

Based on the theory that technical computing is characterized by complex operators and particular combinations of binding time behavior.

To our knowledge, the first technical computing system written in a single language. Past systems have applied specialization to a set of user-level routines. We show that it can be used for an entire system.

Second, we introduce the idea of integrated code selection and specialization.

Using data flow types to drive the design leads to novel dispatch features: - diagonal dispatch - unifying variadic methods with regular types

Easy polymorphism

- library writers can decide what to put at the type level without affecting users - a single mechanism to cover overloading, dispatch, and specialization - specialize by default (makes static polymorphism less of a black art)

Evan Miller expressed this well as "getting disparate groups of programmers to code in the same language...With Julia, it's the domain experts and the speed freaks." [42]

Subtyping algorithm

Library design issues

James Morris eloquently observed that "One of the most fruitful techniques of language analysis is explication through elimination. The basic idea is that one explains a linguistic feature by showing how one could do without it." [44] An additional contribution of this thesis is the application of this approach to features of technical computing environments that have not been subject to such analysis before.

Chapter 2

Problems in technical computing

The original numerical computing language was Fortran, short for "Formula Translating System", released in 1957. Since those early days, scientists have dreamed of writing high-level, generic formulas and having them translated automatically into low-level, efficient machine code, tailored to the particular data types they need to apply the formulas to. Fortran made historic strides towards realization of this dream, and its dominance in so many areas of high-performance computing is a testament to its remarkable success.

The landscape of computing has changed dramatically over the years. Modern scientific computing environments such as MATLAB [39], R [30], Mathematica [36], Octave [45], Python (with NumPy) [57], and SciLab [27] have grown in popularity and fall under the general category known as dynamic languages or dynamically typed languages. In these programming languages, programmers write simple, high-level code without any mention of types like int, float or double that pervade statically typed languages such as C and Fortran.

How should a computer scientist approach this space? We might try to maximize performance. Unfortunately, is all to easy to make unfounded assumptions about what users want. But instead we should study the real world, see what is happening and figure out how to steer it in a better direction.

2.1 What is a technical computing environment?

This question has not really been answered.

Views of this are strongly shaped by what systems happen to exist, and what people were exposed to as they learned to program.

Technical computing software has been designed haphazardly. Each system has evolved as a pile of features taken without what we consider a sufficiently critical argument.

Some languages provide a "convenient" experience that is qualitatively different from "inconvenient" languages. We believe this can be made somewhat precise. A large part of "convenience" is the reduction of the amount you need to know about any given piece of functionality in order to use it. This leads languages to adopt various forms of loose coupling, automation, and elision of software engineering distinctions that are considered important in other languages.

These systems are function-oriented, typically providing a rather large number of functions and a much smaller number of data types. These functions have a particular character one might describe as "manifest": you just call them, interactively, and see what they do. This notion includes the following features:

- 1. Performing fairly large tasks in one function
- 2. Minimal "set up" work to obtain suitable arguments
- 3. No surrounding declarations required
- 4. Permissiveness in accepting many data types and attempting to automatically handle as many cases as possible
- 5. Providing multiple related algorithms or behaviors under a single name

Language design choices affect the ability to provide this user experience (though the first two items are also related to library design). Informally, in order to provide the desired experience a language needs to be able to assign a meaning to a brief and isolated piece of code such as sin(x). This leads directly to making declarations and annotations optional, eliding administrative tasks like memory management, and leaving information implicit (for example

the definition scopes and types of the identifiers sin and x). These characteristics are strongly associated with the Lisp tradition of dynamically typed, garbage collected languages with interactive REPLs.

However, there are subtle cultural differences. A case in point is the MATLAB mldivide, or backslash, operator [38]. By writing only A\B, the user can solve square, over- or underdetermined linear systems that are dense or sparse, for multiple data types. The arguments can even be scalars, in which case simple division is performed. In short, a large amount of linear algebra software is accessed via a single character! This contrasts with the software engineering tradition, where clarifying programmer intent would likely be considered more important. Even the Lisp tradition, which originated most of the convenience features enumerated above, has sought to separate functionality into small pieces. For example Scheme provides separate functions list-ref and vector-ref [3] for indexing lists and vectors.

2.1.1 Case study: Vandermonde matrices

To get a sense of how current technical computing environments work, it is helpful to look at a full implementation example. NumPy [57] provides a function for generating Vandermonde matrices which, despite its simplicity, demonstrates many of the essential language characteristics this thesis addresses. The user-level vander function is implemented in Python (here lightly edited for presentation):

```
def vander(x, N):
    x = asarray(x)
    if x.ndim != 1:
        raise ValueError("x must be a one-dimensional array or sequence.")

v = empty((len(x), N), dtype=promote_types(x.dtype, int))

if N > 0:
    v[:, 0] = 1

if N > 1:
    v[:, 1:] = x[:, None]
    multiply.accumulate(v[:, 1:], out=v[:, 1:], axis=1)
```

return v

This code has many interesting features. Its overall structure consists of normalizing and checking arguments, allocating output, and then calling the lower-level kernel multiply.accumulate (written in C) to run the inner loop. Even though most of the work is done in C, notice why this part of the code is written in Python. vander accepts nearly anything as its first argument, relying on asarray to convert it to an array. There is no need to write down requirements on argument x, allowing the set of supported inputs to grow easily over time. Next, an output array of the correct size is allocated. The key feature of this line of code is that the data type to use is *computed* using the function promote_types. This kind of behavior is difficult to achieve in statically-typed languages.

The call to multiply accumulate invokes a C function called PyUFunc_Accumulate, which is over 300 lines long. The job of this function is to look up an appropriate computational kernel to use given the types of the input arguments and the operation being performed (multiply). In other words, it performs dynamic dispatch. This is notable because Python itself is already based on dynamic dispatch. However, Python's class-based dispatch system is not particularly helpful in this case, since the code needs to dispatch on multiple values. Therefore NumPy's C code implements a custom dispatch table. The targets of this dispatch are many compact, specialized loops generated by running C code through a custom preprocessor.

The mechanism is complicated, but the results are appealing: NumPy code has full runtime flexibility, yet can still achieve good inner-loop performance. Notice that this result is obtained through a clever, if painstaking, mixture of binding times. Python performs some late-bound operations, then calls an early-bound C routine, which performs its own late-bound dispatch to a loop of early-bound operations.

A central message of this thesis is that this kind of program behavior is useful, highly characteristic of technical computing, and can be provided automatically at the language level. Here is how the above dispatch scheme might be implemented in Julia (mirroring the

structure of the Python code as much as possible to facilitate comparison):

```
function vander(x, N::Int)
 x = convert(AbstractVector, x)
 M = length(x)
  v = Array(promote_type(eltype(x),Int), M, N)
    v[:, 1] = 1
  end
  if N > 1
    for i = 2:N; v[:, i] = x; end
    accumulate(MulFun(), v, v)
  return v
end
function accumulate(op, input, output)
 M, N = size(input)
 for i = 2:N
    for j = 1:M
      output[j,i] = op(input[j,i], input[j,i-1])
  end
end
```

This code implements the entire vander computation (with some details left out, for example it does not provide the axis argument of multiply.accumulate). The C component is replaced by a general type-based dispatch system that handles specializing and then invoking an efficient implementation of accumulate for each combination of argument types.

2.2 Why dynamic typing?

Mathematical abstractions often frustrate our attempts to represent them on a computer. For example, mathematicians can move instinctively between isomorphic objects such as scalars and 1-by-1 matrices, but most programming languages would prefer to represent scalars and matrices quite differently.

2.2.1 Mismatches with mathematical abstractions

Programs in general, deal with values of widely varying disjoint types: functions, numbers, lists, network sockets, etc. Type systems are good at sorting out values of these different types. However, in mathematical code most values are numbers or number-like. Numerical properties (such as positive, negative, even, odd, greater than one, etc.) are what matter, and these are highly dynamic.

The classic example is square root (sqrt), whose result is complex for negative arguments. Including a number's sign in its type is a possibility, but this quickly gets out of hand. When computing eigenvalues, for example, the key property is matrix symmetry. Linear algebra provides many more examples where algorithm and data type changes arise from subtle properties. These will be discussed in more detail in section 5.4.

We must immediately acknowledge that static type systems provide mechanisms for dealing with "dynamic" variations like these. However, these mechanisms require deciding which distinctions will be made at the type level, and once the choice is made we are restricted in how objects of different types can be used. In the domain of linear algebra, for example, it would be useful to have matrices that are considered different types for some purposes (e.g. having different representations), but not others (e.g. different types of matrices can be returned from the same function).

2.2.2 I/O and data-oriented programming

Inevitably there is a need to refer to a datatype at run time. A good example is file I/O, where you might want to say "read n double precision numbers from this file". In static languages the syntax and identifiers used to specify data types for these purposes are usually different from those used to specify ordinary types. A typical example is a constant like MPI_DOUBLE [55], which is used to indicate the data type of an MPI message. Many libraries, especially those for tasks like interacting with databases and file formats, have their own set of such tags. Using more than one of these libraries together can result in lots of boilerplate code to map between tag systems.

Polymorphic languages provide better interfaces in these cases. For example, C++ overloading can be used to provide the correct tag automatically based on the type of a function argument. However, these techniques only work for statically-known types. Code like that in the vander example that determines types using arbitrary computation is excluded. Dynamic typing provides a complete solution. Indeed, I/O examples are often used to motivate dynamic typing [2]. With dynamic typing, a single set of names can be used to refer to all types regardless of binding time, and the user may write whatever code is convenient to handle classifying data moving in and out of a program.

2.3 Code generation

- tensor contraction engine
- Firedrake, pyop2 (vs. c++ libmesh)
- JuMP vs. python puLP and pyomo (want to be able to pass functions, not C++ code as strings)
- FFTW
- pochoir

2.3.1 A compiler for every problem

[4]

2.4 Tradeoffs in current designs

"Ducking" the issue of typing is not really possible.

NumPy essentially adds an extra type system to Python (the dtype class). For a long time, there was no existing compiler that was aware of this type system. One was finally

implemented [46], but there are also other Python extensions and JITs with their own incompatible type systems.

Dictionaries for everything (python, js) is the wrong default. Almost every type somebody wants has a fixed number of fields with fixed types.

This phenomenon is increasingly noticed in other domains, particularly JavaScript and web programming. Modern JavaScript implementations are quite fast, but Google's Dart language is based on the premise that we could have a web language that is even faster, and offers more productivity as well. How so? Because Dart's designers observed that JavaScript programmers in practice often write code that could be defined using traditional OO classes, but the language does not support them.

Python is often described as a good glue language. This means it is effectively used as an interface standard, a kind of extended C ABI that makes it easier for libraries to interoperate, and easier for users to access those libraries. Something as straightforward as providing a standard N-dimensional numeric array class (NumPy), which does not exist at the level of C, goes a long way.

However, we wish to point out that in this picture, Python is not doing as much work as it might first appear. Python does not make it easier to implement the functionality inside NumPy, or other "under the hood" scientific libraries. In many cases it creates more work, through the need to write wrappers and interfaces for native code.

Merely being "dynamic" (e.g. Python) should not be considered the gold standard of flexibility. Although these systems permit tricks that can solve otherwise difficult programming problems, this is not always the kind of flexibility that is needed. When faced with the need to describe many functions with elaborate behavior and many cases, one does not primarily need permissiveness, but rather powerful and descriptive organizing principles.

2.4.1 Vectorization

The attitude of vectorization proponents has been somewhat famously summarized as "Life is too short to spend writing for loops." [1] The vectorized style has the following touted

benefits:

- 1. One may write sin(x) to compute the sine of all elements of x, which is much more compact than writing a loop.
- 2. Most of execution time can be spent inside an expertly-written library, taking advantage of special hardware (e.g. SIMD units or stream processors like GPUs) and parallelism implicitly.
- 3. The performance of the high-level language being used becomes less relevant.
- 4. The compiler can optimize across whole-array operations, potentially drastically rearranging computations in a way that would be difficult to do by examining one scalar operation at a time.

Under closer inspection, this argument begins to break down.

An equivocation takes place between being *able* to write a loop as sin(x) and being required to. The former is all we really want, and it is possible in any language that supports some form of function overloading. There is no inherent reason that a high-performance language like C++ or Fortran cannot also support more compact, high-level notation.

Studies reveal that many real-world applications written in vectorization languages actually do not spend most of their time in libraries. [43] It is possible for an interpreter to be so slow that it cancels the advantage of optimized kernels for a range of realistic data sizes.

The final reason actually is compelling. However at a certain point a "vectorization bottleneck" is reached, where performance is limited by expressing computations in terms of in-memory arrays. Figure 2-1 compares the operation rate of three different algorithms for evaluating the polynomial $x^2 + x - 1$ over double precision vectors of varying size. The solid line describes a naive algorithm, where each operation allocates a new array and fills it with one intermediate result (a total of three arrays are needed for the full evaluation). The dotted line algorithm is similar in spirit, except the three operations are "fused" into a single loop. Only one pass is made over the data and one new array is allocated. Naturally

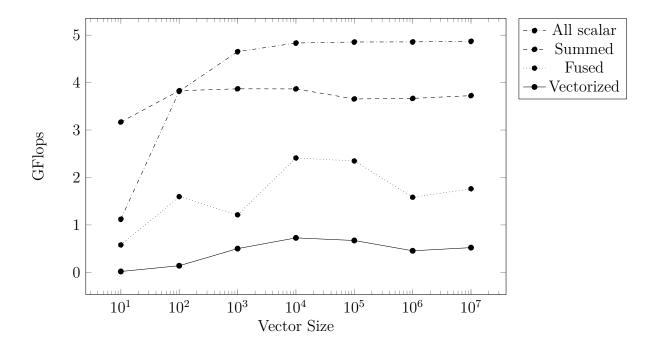


Figure 2-1: Performance of evaluating $x^2 + x - 1$ over a double precision vector of length n. Operation repeated $10^7/n$ times. (Core i7-3517U 1.9GHz, 4MB cache, 8GB RAM)

performance is much better in this case, and advanced array programming environments like APL [22] and ZPL [35] implement this optimization.

However, this level of performance is not impressive compared to the dashed line, which describes the operation rate for computing the same results but summing them instead of storing them. Finally, the dash-dotted line uses no memory at all, instead computing input data points from a linear range. Of course a real use case might require data to be read from memory or stored, so this comparison is unfair (though not entirely, since many numerical environments have represented even simple objects like linear ranges using dense vectors [37]). Still the point should be clear: vectors are not necessarily a good abstraction for high performance. The observation that memory is a bottleneck is nothing new, but for our purposes the important point is that languages that only support vectorized syntax cannot express the code rearrangements and alternate algorithms that can be necessary to get the best performance.

The performance model resulting from vectorized syntax can be difficult to understand, since it is difficult for users to predict which optimizations will be applied. A good example was discussed on the julia-users mailing list [40]. Jason Merrill discovered a Mathematica [36] list discussion of the fastest way to count how many values fall within a given range. Of many proposed implementations, the fastest turned out to be

Total@Unitize@Clip[data,min,max,0,0]

Arguably, it is difficult to know that this will run faster than, for example,

Count[data, x_/; min<=x<=max]</pre>

2.4.2 Data representation

Two major approaches: static language: complex and rigid but efficient memory layout dyn language: simple but flexible (all pointers)

It's reasonable to guess that these would trade off about equally; if you want run-time flexibility then it's not worth using complex specialized data representations. But this turns out not to be true! It's worth using efficient memory layouts even if the layout is not known in advance, and you need an extra dispatch to handle accessing an array (storage strategies).

2.4.3 What needs to be built in?

On"built-in", why built-in is bad.

Built-in-ness often conflates two aspects:

- 1. A feature being readily available and agreed-on by all language users
- 2. A feature tightly coupled to the rest of the system
- (2) implies (1), but not the other way around. (2) is the only technically interesting item, since the other can be addressed e.g. just by including a library in the standard software

distribution. Many technical computing languages have done a large amount of (2) while justifying it with point (1).

While a large part of our motivation is to move more decisions and functionality into libraries, it is equally important to identify what *must* be part of a language for the system to be successful. We believe that large amounts of functionality can be provided by add-ons, but that certain key features cannot be. Past failures to properly classify features this way have caused a lot of undue pain.

First, performance cannot be an add-on. If some users have a fast version of a language and others have a slow version (with the difference being an order of magnitude or more), library writers cannot be sure whether users will find their code fast enough to be useful. How are we to teach people to program in the language?

Psychologically, it may be difficult to accept a "non standard" extension that changes a language so fundamentally. There is a nagging, though perhaps totally unfounded, perception that something subtle may break. If indeed a bug arises due to the use of such an extension, a user is likely to conclude that the extension is dangerous or broken and stop using it. If, on the other hand, a bug arises due to a language's standard optimizing compiler, the user will simply file a bug report, then find a way to work around the problem.

Adding a JIT compiler to a language also requires acceptance of detriments like compilation pauses and pages with RWX permissions. In some cases this may lead to use of the extension being disallowed, perhaps for security reasons.

Type systems similarly fail when provided as optional extensions. Library writers face the same kinds of problems as with performance add-ons. Should I use type annotations in my library?

Dynamic dispatch mechanisms also make especially poor add-ons. Of course, every program makes decisions at run time, and so implements its own "dispatch" to some extent. But these behaviors are inextensible; if language users do not agree on a reusable dispatch framework their code will not be composable.

2.5 Social phenomena

Programming languages are observed to have strong network effects, and the difficulty of getting new languages adopted is well known. However based on [41] we believe this doesn't have to be the case. The formula of improving or redesigning general-purposes languages to be more appealing to domain experts might solve the problem. That way the new system has immediate appeal for at least some users, without the worry that a different tool will be needed as soon as requirements change slightly.

Barriers to contributing.

decisions about what to reflect at the type level should be less consequential somebody might write a c++ library seeking max performance and so might make array rank a static value. (we will discuss this further in section 5.1.3) somebody else who wants dynamic flexibility might have to write an entirely separate package.

Chapter 3

Available technology

3.1 The language design space

It is helpful to begin with a rather coarse classification of programming languages, according to how expressive their type systems are, and whether their type systems are dynamic or static:

	More types	Fewer types	
Dynamic Dylan, Julia		Scheme, Python, MATLAB	
Static	ML, Haskell, Scala	Fortran, Pascal, C	

The lower right corner tends to contain older languages. The upper right corner contains many popular "dynamic" languages. The lower left corner contains many modern languages resulting from research on static type systems. We are most interested in the upper left corner, which is notable for being rather sparsely populated. It has been generally believed that dynamic languages do not "need" types, or that there is no point in talking about types if they are not going to be checked at compile time. These views have some merit, but as a result the top-left corner of this design space has been under-explored.

3.1.1 Classes vs. functions

Technical computing is function oriented.

in C++ to "extend" an existing type you have to use another mechanism, overloading. however then you also need to accept different semantics, static resolution.

3.1.2 Separate compilation vs. performance

Writing the signature of a generic method that needs to be separately compiled, as in Java for example, can be a difficult exercise. The programmer must use the type system to write down sufficient conditions on all arguments. (TODO example from [25]) If, however, we are going to specialize the method, the compiler can analyze it using actual types from call sites, and see for itself whether the method works in each case (this is how C++ templates are type-checked; they are analyzed again for each instantiation).

It is quite interesting that performance and ease-of-use pull this design decision in the same direction.

3.1.3 Parametric vs. ad-hoc polymorphism

Can we have a definition of ad-hoc polymorphism and talk about its relation to multiple dispatch? and point to a technical computing example?

The idea of specialization unites parametric and ad-hoc polymorphism. Beginning with a parametrically-polymorphic function, one can imagine a compiler specializing it for various cases, i.e. certain concrete argument values. These specializations could be stored in a lookup table, for use either at compile time or at run time.

Next we can imagine that the compiler might not optimally specialize certain definitions, and that the programmer would like to provide hints or hand-written implementations to speed up special cases. For example, imagine a function that traverses a generic array. A compiler-generated specialization might inline a specific array types's indexing operations, but a human might further realize that the loop order should be switched for certain arrays

types based on their storage order.

However, if we are going to let a human specialize a function for performance, we might as well allow them to specialize it for some other reason, including entirely different behavior. But at this point separate ad-hoc polymorphism is no longer necessary; we are using a single overloading feature for everything.

Parametric polymorphism describes code that works for any object precisely because it does not do anything meaningful to the object, for example the identity function. In contrast, programming with tagged data (e.g. symbolic expression systems, XML) permits code to work for any object because every object has the same structure, allowing meaningful operations.

In theory, a parametrically-polymorphic function works on all data types. In practice, this can be achieved by forcing a uniform representation of data such as pointers, which can be handled by the same code regardless of what kind of data is pointed to. However this kind of representation is not the most efficient, and for good performance specialized code for different types must be generated.

3.1.4 Static checking vs. flexibility

3.1.5 Modularity vs. large vocabularies

In the context of software engineering, modularity is a primary concern. To build large systems, separate components must be isolated to some degree. Reasons for this include simple concerns like avoiding name conflicts, and a desire to separate interface from implementation to allow a component to change without affecting the rest of the system.

Modularity is sometimes taken to an extreme, and one will see fully qualified names like org.jboss.annotation.ejb.cache.simple.CacheConfig (selected from the JBOSS Java APIs).

Technical computing languages have often avoided and discouraged such designs. For example MATLAB for most of its history supported only a single namespace, which comes

pre-populated with thousands of functions.

3.1.6 Eliminating vs. embracing tagged data

3.1.7 Data hiding vs. explicit memory layout

Examples of CSC and CSR sparse representation. In one sense this is a perfect example of an interface with multiple implementations, and therefore a good use case for object-oriented programming. However in the technical computing world, *hiding* this difference in representation is not usually considered desirable. Clearly a sparse matrix class cannot contain all functions of matrices that users might want to compute. Yet when new functions are added, the programmer needs and wants to exploit representation details (CSC or CSR) for performance. The performance differences involved here are quite significant (TODO cite).

The loss of encapsulation due to multi-methods weighed in [9] is less of a problem for technical computing, and in some cases even advantageous.

3.1.8 Dynamic dispatch is key

Comment: Really nice!

It would be unpleasant if every piece of every program we wrote were forced to do only one specific task. Every time we wanted to do something slightly different, we'd have to write a different program. But if a language allows the same program element to do different things at different times, we can write whole classes of programs at once. This kind of capability is one of the main reasons *object-oriented* programming is popular: it provides a way to automatically select different behaviors according to some structured criteria.

In class-based OO there is essentially *no way* to create an operation that dispatches on existing types (the expression problem [59]). This clearly does not match technical computing, where most programs deal with the same few types (e.g. number, array), and might sensibly want to write new operations that dispatch on them.

We use the non-standard term "criteria" deliberately, in order to clarify our point of view, which is independent of any particular object system.

The sophistication of the available "selection criteria" account for a large part of the perceived "power" or leverage provided by a language. In fact it is possible to illustrate a hierarchy of such mechanisms. As an example, consider a simple simulation, and how it can be written under a series of increasingly powerful paradigms. First, written-out imperative code:

```
while running
  for a in animals
    b = nearby_animal(a)
    if a isa Rabbit
        if b isa Wolf then run(a)
        if b isa Rabbit then mate(a,b)
    else if a isa Wolf
        if b isa Rabbit then eat(a,b)
        if b isa Wolf then follow(a,b)
        end
    end
end
```

We can see how this would get tedious as we add more kinds of animals and more behaviors. Another problem is that the animal behavior is implemented directly inside the control loop, so it is hard to see what parts are simulation control logic and what parts are animal behavior. Adding a simple object system leads to a nicer implementation ¹:

```
class Rabbit
   method encounter(b)
      if b isa Wolf then run()
      if b isa Rabbit then mate(b)
   end
end
```

¹A perennial problem with simple examples is that better implementations often make the code longer.

```
class Wolf
   method encounter(b)
      if b isa Rabbit then eat(b)
      if b isa Wolf then follow(b)
   end
end

while running
   for a in animals
      b = nearby_animal(a)
      a.encounter(b)
   end
end
```

Here all of the simulation's animal behavior has been essentially compressed into a single program point: a.encounter(b) leads to all of the behavior by selecting an implementation based on the first argument, a. This kind of criterion is essentially indexed lookup; we can imagine that a is simply an integer index into a table of operations.

The next enhancement to "selection criteria" adds a hierarchy of behaviors, to provide further opportunities to avoid repetition:

has <: been defined yet?

```
abstract class Animal
   method nearby()
        # search within some radius
   end
end

class Rabbit <: Animal
   method encounter(b: Animal)
        if b isa Wolf then run()
        if b isa Rabbit then mate(b)
   end
end

class Wolf <: Animal
  method encounter(b: Animal)
   if b isa Rabbit then eat(b)</pre>
```

```
if b isa Wolf then follow(b)
end
end
while running
  for a in animals
    b = a.nearby()
    a.encounter(b)
end
end
```

We are still essentially doing table lookup, but the tables have more structure: every Animal has the nearby method, and can inherit a general-purpose implementation.

This brings us roughly to the level of most popular object-oriented languages. But in this example still more can be done. Notice that in the first step to objects we replaced one level of if statements with method lookup. However, inside of these methods a structured set of if statements remains. We can replace these by adding another level of dispatch.

```
class Rabbit <: Animal
   method encounter(b: Wolf) = run()
   method encounter(b: Rabbit) = mate(b)
end

class Wolf <: Animal
   method encounter(b: Rabbit) = eat(b)
   method encounter(b: Wolf) = follow(b)
end</pre>
```

We now have a *double dispatch* system, where a method call uses two lookups, first on the first argument and then on the second argument.

This syntax might be considered a bit nicer, but the design clearly begs a question: why is n = 2 special? It isn't, and we could clearly consider even more method arguments as part of dispatch. But at that point, why is the first argument special? Why separate methods in a special way based on the first argument? It seems arbitrary, and indeed we can remove the special treatment:

```
abstract class Animal
end
class Rabbit <: Animal
end
class Wolf <: Animal
nearby(a: Animal) = # search
encounter(a: Rabbit, b: Wolf) = run(a)
encounter(a: Rabbit, b: Rabbit) = mate(a,b)
encounter(a: Wolf, b: Rabbit) = eat(a, b)
encounter(a: Wolf, b: Wolf) = follow(a, b)
while running
    for a in animals
        b = nearby(a)
        encounter(a, b)
    end
end
```

Here we made two major changes: the methods have been moved "outside" of any classes, and all arguments are listed explicitly. This change has fairly significant implications. Since methods no longer need to be "inside" classes, there is no syntactic limit to where definitions may appear. Now it is easier to add new methods after a class has been defined. Methods also now naturally operate on combinations of objects, not single objects.

The shift to thinking about combinations of objects is fairly revolutionary. Many interesting properties only apply to combinations of objects, and not individuals. We are also now free to think of more exotic kinds of combinations.

We can define a method on any number of objects:

```
encounter(ws: Wolf...) = pack(ws)
```

We can also do diagonal dispatch:

```
encounter\{T <: Animal\}(a: T, b: T) = mate(a, b)
```

3.2 Multiple dispatch

A helpful way to classify languages with some kind of generic programming support is to look at which language constructs are generic. For example, in C++ the syntax object.method(x) is generic: a programmer can get it to do different things by supplying different values for object. The C++ syntax f(a,b) or a+b is usually not generic, but can be overloaded by supplying definitions for different argument types. However, this overloading only uses compile time types (no run time information), and so is essentially a form of renaming — it can be implemented by renaming each definition with a unique name, and replacing overloaded calls with an appropriate name based on compile time types. This is a much weaker form of generic programming, since different behaviors cannot be obtained by passing different values at run time.

For this reason, C++ programs are not fully generic: it is difficult to substitute new behaviors for every part of a program. Some languages take generic programming much further. For example, in MATLAB, *every* function is effectively generic, and can be overloaded by new classes. This enables a programmer to "intercept" every function call (and therefore, essentially everything a program does)

3.2.1 Example: lattices

This example will illustrate possible benefits of multiple dispatch and dynamic typing for mathematical computing. The benefits are not absolute, but notational and semantic: they involve code size and clarity, and the extent to which the entities provided by the language match a mental model of the domain.

A lattice is an algebraic structure where some pairs of elements satisfy a reflexive, antisymmetric, and transitive relation \leq . For purposes of this example, we will consider lattices that have a greatest, or top, element (\top) , and a least, or bottom element (\bot) . When working with lattices one often wants to compute a least upper bound, or join (\sqcup) , or a greatest lower bound, or meet (\sqcap) . Several interesting concerns arise when modeling lattices in a programming language. First, the structure is very general, and so admits implementations for many different kinds of elements. We want to write code using the operators \leq , \sqcup , and \sqcap , and have it apply to any kind of lattice. Therefore some kind of overloading or object-oriented programming is desirable. Second, some properties apply to all lattices, and we would like to avoid implementing them repeatedly.

Using "duck typing", the problem of modeling an abstraction like lattices disappears almost entirely. One may simply define methods for \leq , \sqcup , and \sqcap at any time, for any type, and that type will function as a lattice. That is certainly convenient, but it also fails to provide any reusable functionality for those defining lattices.

Figure 3.2.1 shows a small Julia library for lattices. It defines an abstract class LatticeElement that may be subclassed by objects that will be used primarily as elements of some lattice.

The library also provides standard LatticeElement provides some useful default method definitions.

Service

Join of two incomparables in general lattices does not have to be top

sentence

3.2.2 Symbolic programming

There has always been an essential divide between "numeric" and "symbolic" languages in the world of technical computing. To many people the distinction is fundamental, and we should happily live with both kinds of languages. But if one insists on an answer as to which approach is the right one, then the answer is: symbolic.

Love it!

Systems based on symbolic rewrite rules arguably occupy a further tier of dispatch sophistication. In these systems, you can dispatch on essentially anything, including arbitrary values and structures. These systems are typically powerful enough to concisely define the kinds of behaviors we are interested in.

However, symbolic programming lacks data abstraction: the concrete representations of values are exposed to the dispatch system (e.g. there is no difference between being a list and being something represented as a list).

Wish you would say more!

what does === mean here?

```
abstract LatticeElement
<=(x::LatticeElement, y::LatticeElement) = x===y
==(x::LatticeElement, y::LatticeElement) = x<=y && y<=x
< (x::LatticeElement, y::LatticeElement) = x<=y && !(y<=x)

immutable TopElement <: LatticeElement; end
immutable BotElement <: LatticeElement; end

const T = TopElement()
const L = BotElement()
<=(::BotElement, ::TopElement) = true
<=(::BotElement, ::LatticeElement) = true
<=(::LatticeElement, ::TopElement) = true
<=(::LatticeElement, ::TopElement) = true</pre>
```

Figure 3-1: A small Julia library for lattices

3.2.3 Predicate dispatch

Predicate dispatch is a powerful object-oriented mechanism that allows methods to be selected based on arbitrary predicates [21]. It is, in some sense, the most powerful possible dispatch system, since any computation may be done as part of method selection. Since a predicate denotes a set (the set of values for which it is true), it also denotes a set-theoretic type. Some type systems of this kind, notably that of Common Lisp [56], have actually included predicates as types. However, such a type system is obviously undecidable, since it requires computing the predicates themselves or, even worse, computing predicate implication.²

For a language that is willing to do run time type checks anyway, the undecidability of predicate dispatch is not a problem. Interestingly, it can also pose no problem for static type systems that wish to prove that every call site has an applicable method. Even without evaluating predicates, one can prove that the available methods are exhaustive (e.g. methods for both p and $\neg p$ exist). In contrast, and most relevant to this thesis, predicate types are a problem for code specialization. Static method lookup would require evaluating the predicates, and optimal code generation would require understanding something about what the predicates mean. One approach would be to include a list of satisfied predicates in a type. However, to the extent such a system is decidable, it is essentially equivalent to multiple inheritance. Another approach would be to separate predicates into a second "level" of the type system. The compiler would combine methods with the same "first level" type, and then generate branches to evaluate the predicates. Such a system would be very useful, and could be combined with a language like Julia or, indeed, most object-oriented languages. However this comes at the expense of conceding that predicates are not "real" types.

In considering the problems of predicate dispatch for code specialization, we seem to

²Many type systems involving bounded quantification, such as system $F_{<:}$, are already undecidable [47]. However, they seem to terminate for most practical programs, and also admit minor variations that yield decidable systems [15]. It is fair to say they are "just barely" undecidable, while predicates are "very" undecidable.

be up against a fundamental obstacle: some sets of values are simply more robust under evaluation than others. Programs that map integers to integers abound, but programs that map, say, even integers to even integers are rare to the point of irrelevance.

3.3 Domain theory

In the late 1960s Dana Scott and Christopher Strachey asked how to assign meanings to programs, which otherwise just appear to be lists of symbols [53]. For example, given a program computing the factorial function, we want a process by which we can assign the meaning "factorial" to it. This led to the invention of domain theory, which can be interpreted as modeling the behavior of a program without running it. A "domain" in this theory is a partial order of sets of values that a program might manipulate. Domain theory models computation as follows: a program starts with no information, the lowest point in the partial order ("bottom"). Computation steps accumulate information, gradually moving higher through the order. The advantage of this model, in essence, is that it provides a way to think about the meaning of a program without running the program. Even the "result" of a non-terminating program has a representation — the bottom element. Other elements of the partial order might refer to intermediate results.

The idea of "running a program without running it" is of great interest in compiler implementation. A compiler would like to discover as much information as possible about a program without running it, since running it might take a long time, or produce side effects that the programmer does not want yet, or, of course, not terminate. The general recipe for doing this is to design a partial order (lattice) that captures program properties of interest, and then describe all primitive operations in the language based on how they transform elements of this lattice. Then an input program can be executed, in an approximate sense, until it converges to a fixed point. For example, given a program that outputs an integer, we might decide that we only care whether this integer is even or odd. Then our posets are the even and odd integers, and we will classify operations in the program according

to whether they are evenness-preserving, evenness-inverting, always even, always odd, of uncertain evenness, etc. Most modern compilers use a technique like this (sometimes called abstract interpretation [16]) to semi-decide interesting properties like whether a variable is a constant, whether a variable might be used before it is initialized, whether a variable's value is never used, and so on.

Given the general undecidability of questions about programs, analyses like these are conservative in nature. For example, if our goal is to warn programmers about use of uninitialized variables, we want to be "better safe than sorry" and print a warning if such a use might occur. Such a use corresponds to any lattice element greater than or equal to the element representing "uninitialized". Such conservative analyses are the essence of compiler transformations for performance (optimizations): we only want to perform an optimization if the program property it relies on holds for sure, and not if there is any uncertainty.

The generality of this approach makes it a kind of "ur type system": we can discover a large variety of program properties as long as we are willing to accept some uncertainty. Of course, many programmers and language designers prefer to maximize safety, leading to different approaches that trade away some precision (e.g. syntactic type systems such as those in the ML language family [28], [50]). But if static guarantees are not the priority, or if a language considers itself unconcerned with "types", then we can take the domain-theoretic model as our type system. In some sense, it is our type system whether we like it or not (as implied by [52])!

3.3.1 Monotonicity

When this kind of type system is used, it usually becomes part of the meta-theory of the language: values are "lifted" into an abstract domain, and abstract values are manipulated only by a compiler or other program analysis. One good reason for this is that the abstract values (which we will now interchangeably call "types") must meet certain requirements in order to be useful. For an abstract interpretation to terminate, its lattice must be of finite height, and all functions of lattice elements must be monotonic (given lattice elements a and

b, we must have $a \subseteq b$ implies $f(a) \subseteq f(b)$ for all f).

Functions of types are not monotonic. isa(2,3) is false, but isa(2,Int) is true. This provides one way to separate "real types" from set-valued terms: if there is a way to interpret execution over types such that all functions are monotonic in the (alleged) type lattice, then we are really dealing with types. One way to do this is to specify a genuinely multi-valued term language like λ_{\aleph} [26]. In this approach a term like isa(2,Int) is interpreted as a union of all results obtained by substituting all possible single values for each sub-term. One cannot help but be impressed by such a language, though it seems prohibitively difficult to actually implement. We opt instead for a simpler approach where Int is interpreted as Type{Int} (in a near-borrowing of notation from [26] and [10]), and multi-valued terms exist only statically within a separate maximum-fixed-point evaluator.

3.4 Subtyping

Something about semantic subtyping and type systems for processing XML. XML at first seems unrelated to numerical computing, and indeed it was quite a while before we discovered these papers and noticed the overlap. However if one substitutes "symbolic expression" for "XML document", the similarity becomes clearer.

Chapter 4

The Julia approach

4.1 Core calculus and data model

Julia is based on an untyped lambda calculus augmented with generic functions, tagged data values, and mutable cells.

```
e ::= x
                                                                                   (variable)
        0 1 1 ...
                                                                                  (constant)
        \mid \text{new}(e_{tag}, e_1, e_2, \cdots)
                                                                       (data constructor)
                                                                               (projection)
         | e_1.e_2 = e_3
                                                                              (assignment)
         \mid if e_1 \ e_2 \ e_3
                                                                              (conditional)
         | e_1(e_2) |
                                                                              (application)
                                                                               (sequencing)
         function x_{name} e_{type} (x_1, x_2, \cdots) e_{body}
                                                                      (method definition)
```

The new construct resembles the Dynamic constructor in past work on dynamic typing [2]. The main difference is that the tag is determined by evaluating an expression. Why? The flippant answer is that we are already giving up decidable type checking, so we might as well. But what this really means is that constructing types is part of programming. This can be used to request specific kinds of objects from an API, or used as an implementation detail

to tell the compiler which object properties to track. In practice, new is always wrapped in a constructor function, so the programmer defining a data type is the only one who actually instantiates it. In fact, new for *user-defined* data types is syntactically available only within the code block that defines the type. This provides some measure of data abstraction, since it is not possible for user code to construct arbitrary instances. However, tags themselves (the first argument to new) can be constructed anywhere using nested applications of the syntax $T\{\cdots\}$. Tags are a specific subset of data values whose tag is the built-in value Tag. Once so constructed, a tag can be passed to a constructor (which is just a generic function) to obtain an instance if desired.

Constants are pre-built tagged values.

Types are a superset of tags that includes values generated by the special tags Abstract, Union, and UnionAll, plus the special values Any and Bottom:

```
type ::= \texttt{Bottom} \mid abstract \mid var \\ \mid \texttt{Union} \ type \ type \\ \mid \texttt{UnionAll} \ type <: var <: type \ type \\ \mid \texttt{Tag} \ x_{name} \ abstract_{super} \ value * \\ abstract ::= \texttt{Any} \mid \texttt{Abstract} \ x_{name} \ abstract_{super} \ value * \\ \mid \texttt{Abstract} \ \texttt{Tuple} \ \texttt{Any} \ type * \ type . . . .
```

The last item is the special abstract varargs tuple type.

The language implicitly maps tags to data descriptions, and ensures that the data part of a tagged value always conforms to the tag's description. Mappings from tags to data descriptions are established by special type declaration syntax. Data descriptions have the following grammar:

$$data ::= bit^n \mid ref \mid data*$$

where bit^n represents a string of n bits, and ref represents a reference to a tagged data value. Data may be declared mutable, in which case its representation is implicitly wrapped in a mutable cell. A built-in primitive equality operator === is provided, based on egal [6]

(mutable objects are compared by address, and immutable objects are compared by directly comparing both the tag and data parts bit-for-bit, and recurring through references to other immutable objects).

Functions are generally applied to more than one argument. In the application syntax $e_1(e_2)$, e_2 is an implicitly-constructed tuple of all arguments. e_1 must evaluate to a generic function, and its most specific method matching the tag of argument e_2 is called.

We use the keyword function for method definitions for the sake of familiarity, though method is arguably more appropriate. Method definitions subsume lambda expressions. Each method definition modifies a generic function named by the argument x_{name} . The function to extend is specified by name rather than by value in order to make it easier to syntactically restrict where functions can be extended. This, in turn, allows the language to specify when new method definitions take effect, providing useful windows of time within which methods do not change, allowing programs to be optimized more effectively (and hopefully discouraging abusive and confusing run time method replacements).

The signature, or specializer, of a method is obtained by evaluating e_{type} , which must result in a type value as defined above. A method has n formal argument names x_i . The specializer must be a subtype of the varargs tuple type of length n. When a method is called, its formal argument names are bound to corresponding elements of the argument tuple. If the specializer is a vararg type, then the last argument name is bound to a tuple of all trailing arguments.

The equivalent of ordinary lambda expressions can be obtained by introducing a unique local name and defining a single method on it.

Mutable lexical bindings are provided by the usual translation to operations on mutable cells.

4.2 Type system

Our goal is to design a type system for describing method applicability, and (similarly) for describing classes of values for which to specialize code. Set-theoretic types are a natural basis for such a system. A set-theoretic type is a symbolic expression that denotes a set of values. In our case, these correspond to the sets of values methods are intended to apply to, or the sets of values supported by compiler-generated method specializations. Since set theory is widely understood, the use of such types tends to be intuitive.

These types are less coupled to the languages they are used with, since one may design a value domain and set relations within it without yet considering how types relate to program terms [23] [12]. Since our goals only include performance and expressiveness, we simply skip the later steps for now, and do not consider in detail possible approaches to type checking. A good slogan for this attitude might be "evaluate softly and carry a big subtype relation!"

To avoid the dual traps of "excess power" and divergence, the system we use must have a decidable subtype relation, and must be closed under data-flow operations (meet, join, and widen). It must also lend itself to a reasonable definition of specificity, so that methods can be ordered automatically (a necessary property for composability). These requirements are fairly strict. Beginning with the simplest possible system, we added features as needed either to satisfy the aforementioned closure properties, or to allow us to write method definitions that seemed particularly useful (as it turns out, these two considerations lead to essentially the same features).

We will define our types by formally describing their denotations as sets. We use the notation [T] for the set denotation of type expression T. Concrete language syntax and terminal symbols of the type expression grammar are written in typewriter font, and metasymbols are written in mathematical italic. First there is a universal type Any, an empty type Bottom, and a partial order \leq :

where \mathcal{D} represents the domain of all values.

Next we add data objects with structured tags. The tag of a value is accessed with typeof(x). Each tag consists of a declared type name and some number of sub-expressions, written as Name $\{E_1, \dots, E_n\}$. The center dots (\dots) are meta-syntactic and represent a sequence of expressions. Tag types may have declared supertypes (written as super(T)). Any type used as a supertype must be declared as abstract, meaning it cannot have direct instances.

$$[\![\operatorname{Name} \{ \, \cdots \, \}]\!] = \{ x \mid \operatorname{typeof} (x) = \operatorname{Name} \{ \, \cdots \, \} \}$$

$$[\![\operatorname{Abstract} \{ \, \cdots \, \}]\!] = \bigcup_{\operatorname{super} (T) = \operatorname{Abstract} \{ \cdots \}} [\![T]\!]$$

These types closely resemble the classes of an object-oriented language with generic (parametric) types, invariant type parameters, and no concrete inheritance. We prefer parametric invariance partly for reasons that have been addressed in the literature [17]. Invariance preserves the property that the only subtypes of a concrete type are Bottom and itself. This is important given how we map types to data representations: an Array{Int} cannot also be an Array{Any}, since those types imply different representations. If we tried to use covariance despite this, there would have to be some other notion of which type a value really had, which would be unsatisfyingly complex (tuples are a special case where covariance works, because each component type need only refer to single value, so there is no need for concrete tuple types with non-concrete parameters). We also find that most uses of covariance are more flexibly handled by union type connectives (introduced below).

Next we add conventional product (tuple) types, which are used to represent the arguments to methods. These are almost identical to the nominal types defined above, but are different in two ways: they are *covariant* in their parameters, and permit a special form ending in three dots (...) that denotes any number of trailing elements:

$$\llbracket \mathrm{Tuple}\{P_1,\cdots,P_n\} \rrbracket = \prod_{1 \leq i \leq n} \llbracket P_i \rrbracket$$

$$\llbracket \mathrm{Tuple}\{\cdots,P_n\ldots\} \rrbracket, n \geq 1 = \bigcup_{i \geq n-1} \llbracket \mathrm{Tuple}\{\cdots,P_n^i\} \rrbracket$$

 P_n^i represents i repetitions of the final element P_n of the type expression.

The abstract tuple types ending in . . . correspond to variadic methods, which provide convenient interfaces for tasks like concatenating any number of arrays. Multiple dispatch has been formulated as dispatch on tuple types before [34]. This formulation has the advantage that *any* type that is a subtype of a tuple type can be used to express the signature of a method. It also makes the system simpler and more reflective, since subtype queries can be used to ask questions about methods.

The types introduced so far would be perfectly sufficient for many programs, and are roughly equal in power to several multiple dispatch systems that have been designed before. However, these types are not closed under data-flow operations. For example, when the two branches of a conditional expression yield different types, a program analysis must compute the union of those types to derive the type of the conditional. The above types are not closed under set union. We therefore add the following type connective:

$$[\![\operatorname{Union}\{A,B\}]\!] = [\![A]\!] \cup [\![B]\!]$$

As if by coincidence, Union types are also tremendously useful for expressing method dispatch. For example, if a certain method applies to all 32-bit integers regardless of whether they are signed or unsigned, it can be specialized for Union{Int32,UInt32}.

Union types are easy to understand, but complicate the type system considerably. To see this, notice that they provide an unlimited number of ways to rewrite any type. For example a type T can always be rewritten as Union{T,Bottom}, or Union{Bottom,Union{T,Bottom}}, etc. Any code that processes types must "understand" these equivalences. Covariant constructors (tuples in our case) also distribute over Union types, providing even more ways to rewrite types:

$$Tuple\{Union\{A,B\},C\} = Union\{Tuple\{A,C\},Tuple\{B,C\}\}$$

This is one of a few reasons that union types are often considered undesirable. When used with type inference, such types can grow without bound, possibly leading to slow or even non-terminating compilation. Their occurrence also typically corresponds to cases that would fail most static type checkers. Yet from the perspectives of both data-flow analysis and method specialization, they are perfectly natural and even essential [16] [29] [54].

The next problem we need to solve arises from data-flow analysis of the new construct. When a type constructor C is applied to a type S that is known only approximately at compile time, the type $C\{S\}$ does not correctly represent the result. The correct result would be the union of all types $C\{T\}$ where $T \leq S$. Interestingly, there is again a corresponding need for such types in method dispatch. Often one has, for example, a method that applies to arrays of any kind of integer (Array{Int32}, Array{Int64}, etc.). These cases can be expressed using a UnionAll connective, which denotes an iterated union of a type expression for all values of a parameter in a specified range:

$$[\![\mathtt{UnionAll} \ L \mathord{<\!:} \mathtt{T} \mathord{<\!:} U \ A]\!] = \bigcup_{L \leq T \leq U} [\![A[T/\mathtt{T}]]\!]$$

This is equivalent to an existential type [11]; for each concrete subtype of it there exists a corresponding T. Anecdotally, programmers often find existential types confusing. We prefer the union interpretation because we are describing sets of values; the notion of "there exists" can be semantically misleading since it sounds like only a single T value might be involved.

4.2.1 Examples

UnionAll types are quite expressive. In combination with nominal types they can describe groups of containers such as UnionAll T<:Number Array{Array{T}} (all arrays of arrays of some kind of number) or Array{UnionAll T<:Number Array{T}} (an array of arrays of potentially different types of number).

In combination with tuple types, UnionAll types provide powerful method dispatch specifications. For example UnionAll T Tuple{Array{T},Int,T} matches three arguments: an array, an integer, and a value that is an instance of the array's element type. This is a natural signature for a method that assigns a value to a given index within an array.

4.2.2 Type constructors

It is important for any proposed high-level technical computing language to be simple and approachable, since otherwise the value over established powerful-but-complex languages like C++ is less clear. In particular, type parameters raise usability concerns. Needing to write parameters along with every type is verbose, and requires users to know more about the type system and to know more details of particular types (how many parameters they have and what each one means). Furthermore, in many contexts type parameters are not directly relevant. For example, a large amount of code operates on Arrays of any element type, and in these cases it should be possible to ignore type parameters.

Consider Array{T}, the type of arrays with element type T. In most languages with parametric types, the identifier Array would refer to a type constructor, i.e. a type of a different *kind* than ordinary types like Int or Array{Int}. Instead, we find it intuitive and appealing for Array to refer to any kind of array, so that a declaration such as x::Array simply asserts x to be some kind of array. In other words,

$${\tt Array} = {\tt UnionAll} \ {\tt T} \ {\tt Array}'\{{\tt T}\}$$

where Array' refers to a hidden, internal type constructor. The { } syntax can then be used to instantiate a UnionAll type at a particular parameter value.

$$\frac{{}^{B}X^{L}, \Gamma \vdash T \leq S}{\Gamma \vdash \exists {}^{B}X.T \leq S} \qquad \frac{{}^{B}X^{R}, \Gamma \vdash T \leq S}{\Gamma \vdash T \leq \exists {}^{B}X.S} \qquad \overline{\Gamma \vdash X \leq X}$$

$$\frac{{}^{B}X^{L}, \Gamma \vdash B \leq T}{{}^{B}X^{L}, \Gamma \vdash X \leq T} \qquad \frac{{}^{A}X^{L}, \Gamma \vdash T \leq A}{{}^{A}X^{L}, \Gamma \vdash T \leq X}$$

$$\frac{{}^{B}X^{L}, {}^{A}Y^{L}, \Gamma \vdash B \leq Y \lor X \leq A}{{}^{B}X^{L}, {}^{A}Y^{L}, \Gamma \vdash X \leq Y} \qquad \frac{{}^{B}X^{R}, Y^{R}, \Gamma \vdash B \leq A}{{}^{B}X^{R}, Y^{R}, \Gamma \vdash X \leq Y}$$

$$\frac{{}^{B}X^{R}, Y^{R}, \Gamma \vdash A \leq T}{{}^{A}X^{R}, Y^{R}, \Gamma \vdash X \leq Y} \qquad \frac{{}^{B}X^{R}, Y^{R}, \Gamma \vdash T \leq B}{{}^{A}U^{T}X^{R}, \Gamma \vdash X \leq Y}$$

Figure 4-1: Subtyping algorithm for UnionAll (\exists) and variables. X and Y are variables, A, B, T, and S are types. B_AX means X has lower bound A and upper bound B.

4.2.3 Associated types and type computation

Multiple dispatch systems have often only dispatched on the classes of arguments. This makes it appear necessary to introduce a separate template system to handle other kinds of parameterization. Our type system makes it easy to match the expressive features of templates using dispatch.

4.2.4 Subtyping

Deciding subtyping for base types is straightforward: Bottom is a subtype of everything, everything is a subtype of Any, and tuple types are compared component-wise. The invariant parameters of tag types are compared in both directions: to check $A\{B\} \leq A\{C\}$, check $B \leq C$ and then $C \leq B$. In fact, the algorithm depends on these checks being done in this order, as we will see in a moment.

Checking union types is a bit harder. When a union $A \cup B$ occurs in the algorithm, we need to non-deterministically replace it with either A or B. The rule is that for all such choices on the left of \leq , there must exist a set of choices on the right such that the rest of the algorithm

accepts. This can be implemented by keeping a stack of decision points, and looping over all possibilities with an outer for-all loop and an inner there-exists loop. We speak of "decision points" instead of individual unions, since in a type like Tuple{Union{A,B}...} a single union might be compared many times.

The algorithm for UnionAll and variables is shown in figure 4-1. The first row says to handle a UnionAll by extending the environment with its variable, marked according to which side of \leq it came from, and then recurring into the body. In analogy to union types, we need to check that for all variable values on the left, there exists a value on the right such that the relation holds. The for-all side is relatively easy to implement, since we can just use a variable's bounds as proxies for its value (this is shown in the second row of the figure). We implement the there-exists side by narrowing a variable's bounds (raising the lower bound and lowering the upper bound). The relation holds as long as the bounds remain consistent (i.e. lower bound \leq upper bound). The algorithm assumes that all input types are well-formed, which includes variable lower bounds always being less than or equal to upper bounds.

However, at this point (starting in the third row, second column) the algorithm appears asymmetric. This is a result of exploiting the lack of contravariant constructors. No contravariance means that every time a right-side variable appears on the *left* side of a comparison, it must be because it occurs in invariant position, and the steps outlined in the first paragraph of this section have "flipped" the order (comparing both $B \le C$ and $C \le B$). This explains the odd rule for comparing two right-side variables: this case can only occur with differently-nested UnionAlls and invariant constructors, in which case the relation holds only if all involved bounds are equal. By symmetry, one would expect the bottom-left rule in the figure to update X's upper bound to $B \cap T$. But because of invariance, $T \le B$ has already been checked by the bottom-right rule in the figure. Therefore $B \cap T = T$. This is the reason the "forward" direction of the comparison needs to be checked first: otherwise, we would have updated B to equal T already and the $T \le B$ comparison would become vacuous. Alternatively, we could actually compute $B \cap T$. However there is reason to suspect

that serious trouble lies that way. We would need to either add intersection types to the system, or compute a meet without them. Either way, the algorithm would become much more complex and, judging by past results, very likely undecidable.

These subtyping rules are very likely Π_2^P -hard. Checking a subtype relation with unions requires checking that for all choices on the left, there exists a choice on the right that makes the relation hold. This matches the quantifier structure of 2-TQBF problems of the form $\forall x_i.\exists y_i.F$ where F is a boolean formula. If the formula is rewritten in conjunctive normal form, it corresponds to subtype checking between two tuple types, where the relation must hold for each pair of corresponding types. Now use a type $\mathbb{N}\{x\}$ to represent $\neg x$. The clause $(x_i \lor y_i)$ can be translated to $x_i <: Union\{\mathbb{N}\{y_i\}, True\}$ (where the x_i and y_i are type variables bound by UnionAll on the left and right, respectively). We have not worked out the details, but this sketch is reasonably convincing. Π_2^P is only the most obvious reduction to try; it is possible our system equals PSPACE or even greater, as has often been the case for subtyping systems like ours.

4.2.5 Type system variants

Our design criteria do not identify a unique type system. Many variants are possible. The following features would probably be fairly straightforward to add:

- Structurally-subtyped records
- μ -recursive types (regular trees)
- Regular types (allowing . . . in more places)

The following features would be difficult to add, or possibly break decidability of subtyping:

- Arrow types
- Negations
- Intersections, multiple inheritance
- Universal quantifiers

4.3 Dispatch mechanism

4.3.1 Ambiguities

4.3.2 Constructors

4.4 Higher-order programming

Generic functions are first-class objects, and so can be passed as arguments just as in any dynamically typed language with first-class functions. However, assigning useful type tags to generic functions and deciding how they should dispatch is not so simple. Past work has often described the types of generic functions using the "intersection of arrows" formalism [51] [20] [11] [13]. Since an ordinary function has an arrow type $A \to B$ describing how it maps arguments A to results B, a function with multiple definitions can natually be considered to have multiple such types. For example, a sin function with the following two definitions:

```
sin(x::Float64) = \# compute sine of x in double precision <math>sin(v::Vector) = map(sin, v)
```

could have the type (Float64 \rightarrow Float64) \cap (Vector \rightarrow Vector). The intuition is that this sin function can be used both where a Float64 \rightarrow Float64 function is expected and where a Vector \rightarrow Vector function is expected, and therefore its type is the intersection of these types.

This approach is effective for statically checking uses of generic functions: anywhere a function goes, we must keep track of which arrow types it "contains" in order to be sure that at least one matches every call site and allows the surrounding code to type check. However, despite the naturalness of this typing of generic functions, this formulation is quite problematic for dispatch and code specialization (not to mention that it might make subtyping undecidable).

4.4.1 Problems for code selection

Consider what happens when we try to define an integration function:

```
# 1-d integration of a real-valued function
integrate(f::Float64->Float64, x0, x1)

# multi-dimensional integration of a vector-valued function
integrate(f::Vector->Vector, v0, v1)
```

The -> is not real Julia syntax, but is assumed for the sake of this example. Here we wish to select a different integration routine based on what kind of function is to be integrated. However, these definitions are ambiguous with respect to the sin function defined above. Of course, the potential for method ambiguities existed already. However this sort of ambiguity is introduced non-locally — it cannot be detected when the integrate methods are defined.

Such a non-local introduction of ambiguity is a special case of the general problem that a generic function's type would change depending on what definitions have been added, which depends e.g. on which libraries have been loaded. This does not feel like the right abstraction: type tags are supposed to form a "ground truth" about objects against which program behavior can be selected. Though generic functions change with the addition of methods, it would be more satisfying for their types to somehow reflect an intrinsic, unchanging property.

An additional minor problem with the intersection of arrows interpretation is that we have found, in practice, that Julia methods often have a large number of definitions. For example, the + function in Julia v0.3.4 has 117 definitions, and in a more recent development version with more functionality, it has 150 methods. An intersection of 150 types would be unwieldy, even if only used when debugging the compiler.

A slightly different approach we might try would be to immitate the types of higher-order functions in traditional statically-typed functional languages. For example, we might wish to write map as follows:

```
map{A,B}(f::A\rightarrow B, x::List{A}) = isempty(x) ? List{B}() : List{B}(f(head(x)), map(f, tail(x)))
```

The idea is for the first argument to match any function, and not use the arrow type for

dispatch, thereby avoiding ambiguity problems. Instead, immediately after method selection, values for A and B would be determined using the element type of x and the table of definitions of f.

Unfortunately it is not clear how exactly B should be determined. We could require return type declarations on every method, but this would adversely affect usability (such declarations would also be helpful if we wanted to dispatch on arrow types, though they would not solve the ambiguity problem). Or we could use type inference of f on argument type A. This would not work very well, since the result would depend on partly-arbitrary heuristics. Such heuristics are fine for analyzing a program, but are not appropriate for determining the value of a user-visible variable, as this would make program behavior unpredictable.

4.4.2 Problems for code specialization

For code specialization to be effective, it must eliminate as many irrelevant cases as possible. Intersection types seem to be naturally opposed to this process, since they have the ability to generate infinite descending chains of ever-more-specific function types by tacking on more terms with \cap . There would be no such thing as a maximally-specific function type. In particular, it would be hard to express that a function has exactly one definition, which is an especially important case for optimizing code.

For example, say we have a definition f(g::String->String), and a function h with a single Int \rightarrow Int definition. Naturally, f is not applicable to h. However, given the call site f(h), we are forced to conclude that f might be called with a function of type $(Int \rightarrow Int) \cap (String \rightarrow String)$, since in general Int \rightarrow Int might be only an approximation of the true type of the argument.

The other major concern when specializing code is whether, having generated code for a certain type, we would be able to reuse that code often enough for the effort to be worthwhile. In the case of arrow types, this equates to asking how often generic functions share the same set of signatures. This question can be answered empirically. Studying the Julia Base library as of this writing, there are 1059 generic functions. We examined all 560211 pairs of

	matching pairs	GFs with matches	mean
arguments only	1831 (0.327%)	329	1.73
arguments and return types	241 (0.043%)	85	0.23

Figure 4-2: Number and percentage of pairs of functions with matching arguments, or matching arguments and return types. The second column gives the number of functions that have matches. The third column gives the mean number of matches per function.

functions; summary statistics are shown in figure 4-2. Overall, it is rare for functions to share type signatures. Many of the 85 functions with matches (meaning there exists some other function with the same type) are predicates, which all have types similar to Any \rightarrow Bool. The mean of 0.23 means that if we pick a function uniformly at random, on average 0.23 other functions will match it. The return types compared here depend on our heuristic type inference algorithm, so it useful to exclude them in order to get an upper bound. If we do that, and only consider arguments, the mean number of matches rises to 1.73.

The specific example of the sin and cos functions provides some intuition for why there are so few matches. One would guess that the type behavior of these functions would be identical, however the above evaluation showed this not to be the case. The reason is that the functions have definitions to make them operate elementwise on both dense and sparse arrays. sin maps zero to zero, but cos maps zero to one, so sin of a sparse array gives a sparse array, but cos of a sparse array gives a dense array. This is indicative of the general "messiness" of convenient real-world libraries for technical computing.

4.4.3 Possible solutions

The general lack of sharing of generic function types suggests the first possible solution: give each generic function a new type that is uniquely associated with it. For example, the type of sin would be GenericFunction{sin}. This type merely identifies the function in question, and says nothing more about its behavior. It is easy to read, and easily specific

enough to avoid ambiguity and specialization problems. It does *not* immediately solve the above problem of determining the result type of map. However there are corresponding performance benefits, since specializing code for a specific function argument naturally lends itself to inlining.

Another approach that is especially relevant to technical computing is to use nominal function types. In mathematics, the argument and return types of a function are often among its least interesting properties. In some domains, for example, all functions can implicitly be assumed $\mathbb{R} \to \mathbb{R}$, and the interesting property might be what order of integrable singularity is present (see section 5.5 for an application), or what dimension of linear operator the function represents. The idea of nominal function types is to describe the properties of interest using a data object, and then allow that data object to be treated as a function, i.e. "called". Some object-oriented languages call such an object a functor.

Julia accomodates this approach with a small adjustment to the evaluation rules: in the application syntax $e_1(e_2)$ when e_1 is not a function, evaluate $call(e_1, e_2)$ instead, where call is a particular generic function known to the system.

As an example, we can define a type for polynomials of order N over ring R:

```
immutable Polynomial{N,R}
    coeffs:Vector{R}
end

call{N}(p::Polynomial{N}, x) = sum([p.coeffs[i+1]*x^i for i=0:N])
```

Now it is possible to use a Polynomial just like a function, while also dispatching methods on the relevant properties of order and ring (or ignoring them if you prefer). It is also easy to examine the polynomial's coefficients, in contrast to functions, which are usually opaque.

It is even possible to define a "nominal arrow" type, which uses this mechanism to impose a classification on functions based on argument and return types:

```
immutable Arrow{A,B}
    f
end

call{A,B}(a::Arrow{A,B}, x::A) = a.f(x)::B
```

```
function map(f, A::Array)
    isempty(A) && return Array(Bottom, O)
    el = f(A[1]); T = typeof(el)
    dest = Array(T, length(A))
    dest[1] = el
    for i = 2:length(A)
        el = f(A[i]); S = typeof(el)
        if !(S <: T)
            T = typejoin(T, S)
            new = similar(dest, T)
            copy!(new, 1, dest, 1, i-1)
            dest = new
        end
        dest[i] = el::T
    end
    return dest
end
```

Figure 4-3:

Calling an Arrow will yield a no-method error if the argument type fails to match, and a type error if the return type fails to match.

It is worth examining the differences and tradeoffs between nominal functions and "real" arrow types.

4.4.4 Implementing map

So, given typed containers and no arrow types, how do you implement map? The answer is that the type of container to return must also be computed. This should not be too surprising, since it is implied by the definition of new at the beginning of the chapter: each value is created by computing a type part and a content part.

A possible implementation of map for arrays is shown in figure 4-3. The basic strategy is to try calling the argument function f first, see what it returns, and then optimistically assume that it will always return values of the same type. This assumption is checked on each iteration. If f returns a value that does not fit in the current output array dest, we re-

allocate the output array to a larger type and continue. The primitive typejoin computes a union-free join of types. For uses like map, this does not need to be a true least upper bound; any reasonable upper bound will do.

This implementation works well because it completely ignores the question of whether the compiler understands the type behavior of f. However, it cooperates with the compiler by making the optimistic assumption that f is well-behaved. This is best illuminated by considering three cases. In the first case, imagine f always returns Int, and that the compiler is able to infer that fact. Then the test !(S <: T) disappears by specialization, and the code reduces to the same simple and efficient loop we might write by hand. In the second case, imagine f always returns Int, but that the compiler is not able to infer this. Then we incur the cost of an extra type check on each iteration, but we return the same efficiently-stored Int-specialized Array (Array{Int}). This leads to significant memory savings, and allows subsequent operations on the returned array to be more efficient. The third case occurs when f returns objects of different types. Then the code does not do anything particularly efficient, but is not much worse than a typical dynamically typed language manipulating heterogeneous arrays.

Overall, the resulting behavior is quite similar to a dynamic language that uses *storage* strategies [8] for its collections. The main difference is that the behavior is implemented at the library level, rather than inside the runtime system. This can be a good thing, since one might want a map that works differently. For example, replacing typejoin with promote_type would collect results of different numeric types into a homogeneous array of a single larger numeric type. Other applications might not want to use typed arrays at all, in which case map can be much simpler and always return an Array{Any}.

It must be admitted that this map is more complex than what we are used to in typical functional languages. However, this is at least partly due to map itself being more complex, and amenable to more different interpretations, than what is usually studied in the context of those languages.

- 4.5 Performance model
- 4.5.1 Type inference
- 4.5.2 Specialization

Chapter 5

Case studies

5.1 Explication through elimination

This section will illustrate how we implement key features of technical computing systems using our methodology.

5.1.1 Conversion and comparison

Type conversion provides a classic example of a binary method. Multiple dispatch allows us to avoid deciding whether the converted-to type or the converted-from type is responsible for defining the operation. Defining a specific conversion is straightforward, and might look like this in Julia syntax:

```
convert(::Type{Float64}, x::Int32) = ...
```

A call to this method would look like convert (Float64, 1).

Using conversion in generic code requires more sophisticated definitions. For example we might need to convert one value to the type of another, by writing convert(typeof(y), x). What set of definitions must exist to make that call work in all reasonable cases? Clearly we cannot explicitly write all $O(n^2)$ possibilities. We need abstract definitions that cover many points in the dispatch matrix. One such family of points is particularly important:

those that describe converting a value to a type it is already an instance of. In our system this can be handled by a single definition that performs "triangular" dispatch:

```
convert{T,S<:T}(::Type{T}, x::S) = x
```

"Triangular" refers to the rough shape of the dispatch matrix covered by such a definition: for all types T in the first argument slot, match values of any type less than it in the second argument slot.

A similar trick is useful for defining equivalence relations. It is most likely unavoidable for programming languages to need multiple notions of equality. Two in particular are natural: an *intensional* notion that equates objects that look identical, and an *extensional* notion that equates objects that mean the same thing for some standard set of purposes. Intensional equality (=== in Julia, mentioned in section 4.1) lends itself to being implemented once by the language implementer, since it can work by directly comparing the representations of objects. Extensional equality (== in Julia), on the other hand, must be extensible to user-defined data types. The latter function must call the former in order to have any basis for its comparisons.

As with conversion, we would like to provide default definitions for == that cover families of cases. Numbers are a reasonable domain to pick, since all numbers should be equality-comparable to each other. We might try

```
==(x::Number, y::Number) = x === y
```

meaning that number comparison can simply fall back to intensional equality. However this definition is rather dubious. It gets the wrong answer every time the arguments are different representations (e.g. integer and floating point) of the same quantity. We might hope that its behavior will be "patched up" later by more specific definitions for various concrete number types, but it still covers a dangerous amount of dispatch space. If later definitions somehow miss a particular combination of number types, we could get a silent wrong answer instead of an error. (Note that statically checking method exhaustiveness is no help here.)

"Diagonal" dispatch lets us improve the definition:

```
==\{T <: Number\}(x::T, y::T) = x === y
```

Now === will only be used on arguments of the same type, making it far more likely to give the right answer. Even better, any case where it does not give the right answer can be fixed with a single definition, i.e. ==(x::S, y::S) for some concrete type S. The more general (Number, Number) case is left open, and in the next section we will take advantage of this to implement "automatic" type promotion.

5.1.2 Numeric types and embeddings

We might prefer "number" to be a single, concrete concept, but the history of mathematics has seen the concept extended many times, from integers to rationals to reals, and then to complex, quaternion, and more. These constructions tend to follow a pattern: a new set of numbers is constructed around a subset isomorphic to an existing set of numbers. For example, the reals are isomorphic to the complex numbers with zero imaginary part.

Human beings happen to be good at equating and moving between isomorphic sets, so it is easy to imagine that the reals and complexes with zero imaginary part are one and the same. But a computer forces us to be specific, and admit that a real number is not complex, and a complex number is not real. And yet the close relationship between them is too compelling not to model in a computer somehow. Here we have a numerical analog to the famous "circle and ellipse" problem in object-oriented programming: the set of circles is isomorphic to the set of ellipses with equal axes, yet neither "is a" relationship in a class hierarchy seems fully correct. An ellipse is not a circle, and in general a circle cannot serve as an ellipse (for example, the set of circles is not closed under the same operations that the set of ellipses is, so a program written for ellipses might not work on circles). This problem implies that a single built-in type hierarchy is not sufficient: we want to model custom *kinds* of relationships between types (e.g. "can be embedded in" in addition to "is a").

Current approaches

Numbers tend to be among the most complex features of a language. Numeric types usually need to be a special case: in a typical language with built-in numeric types, describing their behavior is beyond the expressive power of the language itself. For example, in C arithmetic operators like + accept multiple types of arguments (ints and floats), but no user-defined C function can do this (this situation is of course improved in C++). In Python, a special arrangement is made for + to call either an __add__ or __radd__ method, effectively providing double-dispatch for arithmetic in a language that is idiomatically single-dispatch.

Implementing type embeddings

Most functions are naturally implemented in the value domain, but some are actually easier to implement in the type domain. One reason is that there is a bottom element, which most data types lack.

It has been suggested on theoretical grounds [49] that generic binary operators should have "key variants" where the arguments are of the same type.

```
+(x::Number, y::Number) = +(promote(x,y)...)
+{T<:Number}(x::T, y::T) = no_op_err("+", T)
```

A full-featured promotion operator is a tall order. We would like

- each combination of types only needs to be defined in one order
- it falls back to join
- must prevent promotion above a certain point to avoid circularity in promoting fallback definitions of operators.

```
promote{T,S}(x::T, y::S) =
        (convert(promote_type(T,S),x), convert(promote_type(T,S),y))
promote_type{T,S}(::Type{T}, ::Type{S}) =
```

```
promote_result(T, S, promote_rule(T,S), promote_rule(S,T))
promote rule(T, S) = Bottom
promote_result(t,s,T,S) = promote_type(T,S)
# If no promote_rule is defined, both directions give Bottom. In that
# case use typejoin on the original types instead.
promote_result{T,S}(::Type{T},::Type{S},::Type{Bottom},::Type{Bottom}) = typejo
# Because of the promoting fallback definitions for Number, we need
# a special case for undefined promote_rule on numeric types.
# Otherwise, typejoin(T,S) is called (returning Number) so no conversion
# happens, and +(promote(x,y)...) is called again, causing a stack
# overflow.
promote_result{T<:Number,S<:Number}(::Type{T},::Type{S},::Type{Bottom},::Type{I</pre>
    promote_to_super(T, S, typejoin(T,S))
\# promote numeric types T and S to typejoin(T,S) if T<:S or S<:T
# for example this makes promote_type(Integer, Real) == Real without
# promoting arbitrary pairs of numeric types to Number.
promote_to_super{T<: Number</pre>
                                      \{(::Type\{T\}, ::Type\{T\}, ::Type\{T\}) = T\}
promote_to_super{T<:Number,S<:Number}(::Type{T}, ::Type{S}, ::Type{T}) = T</pre>
promote_to_super\{T <: Number, S <: Number\}(::Type\{T\}, ::Type\{S\}, ::Type\{S\}) = S
promote_to_super{T<:Number,S<:Number}(::Type{T}, ::Type{S}, ::Type) =</pre>
    error("no promotion exists for ", T, " and ", S)
```

Application to ranges

Ranges illustrate an interesting application of type promotion. A range data type, notated a:s:b, represents a sequence of values starting at a and ending at b, with a distance of s between elements (internally, this notation is translated to colon(a, s, b)). Ranges seem simple enough, but a reliable, efficient, and generic implementation is difficult to achieve. We propose the following requirements:

- The start and stop values can be passed as different types, but internally should be of the same type.
- Ranges should work with ordinal types, not just numbers (examples include characters, pointers, and calendar dates).

• If any of the arguments is a floating-point number, a special FloatRange type designed to cope well with roundoff is returned.

In the case of ordinal types, the step value is naturally of a different type than the elements of the range. For example, one may add 1 to a character to get the "next" encoded character, but it does not make sense to add two characters.

It turns out that the desired behavior can be achieved with six definitions:

First, given three floats of the same type we can construct a FloatRange right away:

```
colon{T<:FloatingPoint}(start::T, step::T, stop::T) = FloatRange{T}(start, step, stop)</pre>
```

Next, if a and b are of the same type and there are no floats, we can construct a general range:

```
colon{T}(start::T, step, stop::T) = StepRange(start, step, stop)
```

Now there is a problem to fix: if the first and last arguments are of some non-floatingpoint numeric type, but the step is floating point, we want to promote all arguments to a common floating point type. We must also do this if the first and last arguments are floats, but the step is some other kind of number:

```
colon{T<:Real}(a::T, s::FloatingPoint, b::T) = colon(promote(a,s,b)...)
colon{T<:FloatingPoint}(a::T, s::Real, b::T) = colon(promote(a,s,b)...)</pre>
```

These two definitions are correct, but ambiguous: if the step is a float of a different type than a and b both definitions are equally applicable. We can add the following disambiguating definition:

```
colon{T<:FloatingPoint}(a::T, s::FloatingPoint, b::T) = colon(promote(a,s,b)...)</pre>
```

All of these five definitions require a and b to be of the same type. If they are not, we must promote just those two arguments, and leave the step alone (in case we are dealing with ordinal types):

 $colon{A,B}(a::A, s, b::B) = colon(convert(promote_type(A,B),a), s, convert(promote_type(A,B),a))$

This example shows that it is not always sufficient to have a built-in set of "promoting operators". Library functions like this colon need more control.

Diversity of number and number-like types in practice

Originally, our reasons for implementing all numeric types at the library level were not entirely practical. We had a principled opposition to including such definitions in a compiler, and guessed that being able to define numeric types would help ensure the language was "powerful enough". However, defining numeric and number-like types and their interactions turns out to be surprisingly useful. Once such types are easy to obtain, people find more and more uses for them.

Even among basic types that might reasonably be "built in", there is enough complexity to require an organizational strategy. We might have

- Ordinal types Pointer, Char
- Integer types Bool, Int8, Int16, Int32, Int64, Int128, UInt8, UInt16, UInt32, UInt64, UInt128, BigInt
- Floating point types Float16, Float32, Float64, Float80*, Float128*, BigFloat, DoubleDouble
- Extensions Rational, Complex, Quaternion

Types with * have not been implemented yet, but the rest have. In external packages, people have implemented types for interval arithmetic, dual and hyper-dual numbers for computing first and second derivatives, and finite fields.

Some applications benefit in performance from fixed-point arithmetic. This has been implemented in a package as Fixed32{b}, where the number of fraction bits is a parameter.

A problem in the design of image processing libraries was solved by defining a new kind of fixed-point type [31]. The problem is that image scientists often want to work with fractional

pixel values in the interval [0, 1], but most graphics libraries (and memory efficiency concerns) require 8-bit integer pixel components with values in the interval [0, 255]. The solution is a Ufixed8 type that uses an unsigned 8-bit integer as its representation, but behaves like a fraction over 255.

Many real-world quantities are not numbers exactly, but benefit from the same mechanisms in their implementation. Examples include colors (which form a vector space, and where many different useful bases have been standardized), physical units, and DNA nucleotides. Date and time arithmetic is especially intricate and irregular (more on this in section 5.6).

5.1.3 Multidimensional array indexing

One-dimensional arrays are a simple and essential data structure found in most programming languages. The multi-dimensional arrays required in scientific computing, however, are a different beast entirely. Allowing any number of dimensions entails a significant increase in complexity. Why? The essential reason is that core properties of the data structure no longer fit in a constant amount of space. The space needed to store the sizes of the dimensions (the array shape) is proportional to the number of dimensions. This does not seem so bad, but becomes a large problem due to three additional facts.

First, code that operates on the dimension sizes needs to be highly efficient. Typically the overhead of a loop is unacceptable, and such code needs to be fully unrolled. Second, in some code the number of dimensions is a *dynamic* property — it is only known at run time. Third, programs may wish to treat arrays with different numbers of dimensions very differently. A vector (1d) might have rather different behaviors than a matrix (2d) (for example, to compute a norm). This kind of behavior makes the number of dimensions a crucial part of program semantics, preventing it from remaining a compiler implementation detail.

These facts pull in different directions. The first fact asks for static analysis. The second fact asks for run time flexibility. The third fact asks for dimensionality to be part of the

type system, but partly determined at run time (for example, via virtual method dispatch). Current approaches choose a compromise. In some systems, the number of dimensions has a strict limit (e.g. 3 or 4), so that separate classes for each case may be written out in full. Other systems choose flexibility, and just accept that most or all operations will be dynamically dispatched. Other systems might provide flexibility only at compile time, for example a template library where the number of dimensions must be statically known.

Whatever decision is made, rules must be defined for how various operators act on dimensions. For now we will focus on indexing, since selecting parts of arrays has particularly rich behavior with respect to dimensionality. For example, if a single row or column of a matrix is selected, does the result have one or two dimensions? Array implementations prefer to invoke general rules to answer such questions. Such a rule might say "dimensions indexed with scalars are dropped", or "trailing dimensions of size one are dropped", or "the rank of the result is the sum of the ranks of the indexes" (as in APL).

Our goal here is a bit unusual: we are not concerned with which rules might work best, but merely with how they can be specified, so that domain experts can experiment. In fact different domains want different things. Working with images, each dimension might be quite different, e.g. representing time, space, or color, so you don't want to drop or rearrange dimensions very often.

How are such rules implemented? For a language with built-in multidimensional arrays, the compiler will analyze indexing expressions and determine an answer using hard-coded logic. However, this approach is not satisfying: we would rather implement the behavior in libraries, so that different kinds of arrays may be defined, or so that rules of similar complexity may be defined for other kinds of objects. But these kinds of rules are unusually difficult to implement in libraries. If a library writes out its indexing logic using imperative code, the host language compiler is not likely to be able to analyze it. Using compile time abstracton (templates) would provide better performance, but such libraries tend to be difficult to write (and read), and the full complement of indexing behavior expected by technical users strains the capabilities of such systems.

Our dispatch mechanism permits a novel solution [7]. If a multiple dispatch system supports variadic functions and argument "splicing" (the ability to pass a structure of n values as n separate arguments to a function), then indexing behavior can be defined as method signatures.

This solution is still a compromise among the factors outlined above, but it is a new compromise that provides a modest, but definite, increment of power.

Below we define a function index_shape that computes the shape of a result array given a series of index arguments. We show three versions, each implementing a different rule that users in different domains might want:

```
# drop dimensions indexed with scalars
index_shape() = ()
index_shape(i::Real, I...) = index_shape(I...)
index_shape(i, I...) = tuple(length(i), index_shape(I...)...)
```

```
# drop trailing dimensions indexed with scalars
index_shape(i::Real...) = ()
index_shape(i, I...) = tuple(length(i), index_shape(I...)...)
# rank summing (APL)
index_shape() = ()
index_shape(i, I...) = tuple(size(i)..., index_shape(I...)...)
```

Inferring the length of the result of index_shape is sufficient to infer the rank of the result array.

These definitions are concise, easy to write, and possible for a compiler to understand fully using straightforward techniques.

The result type is determined using only dataflow type inference, plus a rule for splicing an immediate container (the type of f((a,b)...) is the type of f(a,b)). Argument list destructuring takes place inside the type intersection operator used to combine argument types with method signatures.

This approach does not depend on any heuristics. Each call to index_shape simply requires one recursive invocation of type inference. This process reaches the base case () for these definitions, since each recursive call handles a shorter argument list (for less-well-behaved definitions, we might end up invoking a widening operator instead).

5.2 Units

5.3 Array views

tim holy in issue 8839:

"without staged functions in my initial post in 8235. The take-home message: generating all methods through dimension 8 resulted in more than 5000 separate methods, and required over 4 minutes of parsing & lowering time (i.e., a 4-minute delay while compiling julia). By

comparison, the staged function implementation loads in a snap, and of course can go even beyond 8 dimensions."

5.4 Numerical linear algebra

The crucial roles of code selection and code specialization in technical computing are captured well in the linear algebra software engineer's dilemma, described in the context of LAPACK and ScaLAPACK by Demmel and Dongarra, et.al. [18]:

Multiple dispatch on special matrices

29 LAPACK types via composition of 9 types, issue 8240

QR factorization: want to support it on all input types, but many types (integer, rational) are not closed under the needed operations. compare to eigen: in C++, if you want a qrfact of an integer matrix, you might get a compile time error, or it might work but all the values will be truncated when stored. this is a big generic programming challenge. you can't expect types to have a "type to use for QR fact" trait.

great computes the type of the result needed:

```
function qrfact{T}(A::StridedMatrix{T}; pivot=false)
   S = typeof(one(T)/norm(one(T)))
   if S != T
        qrfact!(convert(AbstractMatrix{S},A), pivot=pivot)
   else
        qrfact!(copy(A),pivot=pivot))
   end
end
```

This code is not statically typeable, and yet with specialization a compiler could in fact determine the type of each call site. It just happens to be convenient to specify this behavior with a branch.

5.5 Boundary-element method

There are lots of general packages for FEM problems, but it is much more difficult to create such a package for BEM problems. The method requires integrating functions with singularities, many times in the inner loop of code that builds the problem matrix. Integrating such functions numerically on each iteration is much too slow. As a result, many special-purpose implementations have been written by hand for different problems.

Some recent work [48] managed a more general solution, using Mathematica to generate C++ code for different cases. This worked well, but was difficult to implement and the resulting system is difficult to use. We see the familiar pattern of using multiple languages and code-generation techniques, with coordination of the overall process done either manually or with ad-hoc scripts. To polish the implementation for use as a practical library, a likely next step would be to add a Python interface, adding yet another layer of complexity.

Thanks to Professor Steven Johnson for providing the following section describing the problem.

5.5.1 Galerkin matrix assembly for singular kernels

A typical problem in computational science is to form a discrete approximation of some infinite-dimensional linear operator \mathcal{L} with some finite set of basis functions $\{b_m\}$ via a Galerkin approach [refs], which leads to a matrix L with entries $L_{mn} = \langle b_m, \mathcal{L}b_n \rangle = \langle b_m, b_n \rangle_{\mathcal{L}}$ where $\langle \cdot, \cdot \rangle$ denotes some inner product (e.g. $\langle u, v \rangle = \int uv$ is typical) and $\langle \cdot, \cdot \rangle_{\mathcal{L}}$ is the bilinear form of the problem. Computing these matrix elements is known as the matrix assembly step, and its performance is a crucial concern for solving partial differential equations (PDEs) and integral equations (IEs).

The "easy" case: nonsingular assembly

For example, in the finite-element method (FEM) [refs], the basis functions b_m are typically low-order polynomials defined piecewise over geometric elements (typically triangles or tetrahedra), and \mathcal{L} is typically a differential operator like $-\nabla \cdot c(x)\nabla$ for some coefficients c(x), which leads to a bilinear form $\langle b_m, b_n \rangle_{\mathcal{L}} = \int \nabla b_m \cdot c(x) \nabla b_n$ (after integration by parts). Because the basis functions are localized and \mathcal{L} consists of local operations, the matrix L is sparse and L_{mn} need only be computed for m and n corresponding to neighboring elements. Moreover, these integrals are straightforward to evaluate by standard cubature schemes because the integrands are nonsingular: they typically have no divergences or discontinuities. In particular, because the functions b_m and c are usually smooth within a single element, one can use a fixed low-order cubature rule: you evaluate the integrand at a handful of precomputed points within each element, multiply by precomputed weights, and sum to obtain the approximate integral.

Even so, the basis functions and the coefficient function c(x) may need to be evaluated tens of millions of times for even a moderate-size mesh in three dimensions, so production FEM implementations in traditional high-level dynamic languages such as Matlab and Python are forced to offload matrix assembly to external C and C++ code. For example, the popular FEniCS [ref] and Firedrake [ref] FEM packages for Python both implement domain-specific compilers: a symbolic expression for the bilinear form is combined with fragments of user-specified C++ code to define functions like c(x), compiled to C++ code, and then compiled to object code which is dynamically loaded. In Julia, we believe this could be simplified considerably because functions like c(x) could be defined directly in Julia and code generation/compilation could be performed entirely within Julia without a C++ intermediary. Indeed, preliminary experiments with pure Julia FEM implementations [ref http://www.codeproject.com/Articles/579983/Finite-Element-programming-in-Julia] have demonstrated performance comparable to sophisticated solutions like FEniCS in Python and FreeFem++ in C++ [ref]

Singular assembly for integral operators

A much more challenging case of Galerkin matrix assembly arises for singular integral operators \mathcal{L} , which act by convolving their operand against a singular "kernel" function K(x): $u = \mathcal{L}v$ means that $u(x) = \int K(x-x')v(x')dx'$. For example, in electrostatics and other Poisson problems, the kernel is K(x) = 1/|x| in three dimensions and $\ln |x|$ in two dimensions, while in scalar Helmholtz (wave) problems it is $e^{ik|x|}/|x|$ in three dimensions and a Hankel function $H_0^{(1)}(k|x|)$ in two dimensions. Formally, Galerkin discretizations lead to matrix assembly problems similar to those above: $L_{mn} =: \langle b_m, \mathcal{L}b_n \rangle = \int b_m(x)K(x-x')b_n(x')dx\,dx'$. However, there are several important differences from FEM:

- The kernel K(x) nearly always diverges for |x| = 0, which means that generic cubature schemes are either unacceptably inaccurate (for low-order schemes) or unacceptably costly (for adaptive high-order schemes, which require huge numbers of cubature points around the singularity), or both.
- Integral operators typically arise for *surface* integral equations (SIEs) [ref], and involve unknowns on a surface. The analogue of the FEM discretization is then a boundary element method (BEM) [ref], which discretizes a surface into elements (e.g. triangles), with basis functions that are low-order polynomials defiend piecewise in the elements. However, there are also volume integral equations (VIEs) which have FEM-like volumetric meshes and basis functions.
- The matrix L is typically dense, since K is long-range. For large problems, L is often stored and applied implicitly via fast-multipole methods [refs] and similar schemes, but even in this case the diagonal L_{mm} and the entries L_{mn} for adjacent elements must typically be computed explicitly. (Moreover, these are the integrals in which the K singularity is present.)

These difficulties are part of the reason why there is currently *no* truly "generic" BEM software, analogous to FEniCS for FEM: essentially all practical BEM code is written for a

specific integral kernel and a specific class of basis functions arising in a particular physical problem. Changing anything about the kernel or the basis—for example, going from two- to three-dimensional problems—is a major undertaking.

We believe that Julia should be an ideal platform on which to attack this problem:

- Multiple dispatch allows the cubature scheme to be selected at compile time based on the dimensionality, the degree of the singularity, the degree of the polynomial basis, and so on, and allows specialized schemes to be added easily for particular problems with no run time penalty.
- Staged functions allow computer algebra systems to be invoked at compile time to generate specialized cubature schemes for particular kernels. New developments in BEM integration schemes [48] have provided efficient cubature-generation algorithms of this sort, but it has not yet been practical to integrate them with run time code in a completely automated way.

A prototype implementation of this approach follows.

5.6 Dates

compare to python DateTime, compare code length

5.7 JUMP

Metaprogramming tools reused for symbolic algebra

5.8 Computational geometry

robust predicates (dispatch over points, lines) and Voronoi Delaunay.jl benchmarked against CGAL 5.9 Beating the incumbents

- erfinv and digamma using horner macro

- randn beating matlab

@evalpoly macro has separate cases for real and complex in order to automatically take advantage of a subtle algorithm [33]. A macro is perfect for generating the neccessary code, however it lacks the type information needed to select between the two cases. In Julia, it can generate a branch with a type check, and rely on the unused case being removed by automatic specialization. (A generic function with two definitions could be generated instead, but in high-performance programming the "force inlining" behavior of macros is welcome.)

If nothing else, demonstrates that removing glue code overhead is worthwhile.

grisu: 6kLOC to 1kLOC (PR 7291)

Chapter 6

Conclusion

A generation of dynamic languages have been designed by trying variants of the class-based object oriented paradigm. This process has been aided by the development of standard techniques (e.g. bytecode VMs) and reusable infrastructure such as code generators, garbage collectors, and whole VMs like the JVM and CLR.

It is possible to envision a future generation of languages that generalize this design to set-theoretic subtyping instead of just classes. This next generation will require its own new tools, such as partial evaluators (already under development in PyPy and Truffle). One can also imagine these future language designers wanting reusable program analyses, and tools for developing lattices and their operators.

It is interesting to observe that the data model of a language like Julia consists of two key relations: the subtype relation, which is relatively well understood and enjoys useful properties like transitivity, but also the typeof relation, which relates individual values to their types (i.e. the 'typeof' function). The typeof relation appears not to be transitive, and also has a degree of arbitrariness: a value is of a type merely because it is labeled as such, and because various bits of code conspire to ensure that this labeling makes sense according to various criteria.

We have speculated about whether future languages will be able to do away with this distinction. One approach is λ_{\aleph} . We have also speculated that this could be done using types

based on non-well-founded set theory, combining the subset-of and element-of relations using self-containing sets. We are not yet sure what a practical language based on this idea might look like.

There are several key aspects of performance programming that our design does not directly address, e.g. storage and in-place optimizations.

future work: product domains

Appendix A

Subtyping algorithm

```
abstract Ty
                                    type UnionT <: Ty</pre>
                                      a; b
type TypeName
                                     end
 super::Ty
  TypeName() = new()
                                    type Var
                                      lb; ub
end
                                     end
type TagT <: Ty</pre>
                                    type UnionAllT <: Ty</pre>
 name::TypeName
 params
                                      var::Var
 vararg::Bool
                                     end
## Any, Bottom, and Tuple
const AnyT = TagT(TypeName(), ()); AnyT.name.super = AnyT
type BottomTy <: Ty; end; const BottomT = BottomTy()</pre>
const TupleName = TypeName(); TupleName.super = AnyT
## type application
inst(t::TagT) = t
inst(t::UnionAllT, param) = subst(t.T, Dict{Any,Any}(t.var => param))
inst(t::UnionAllT, param, rest...) = inst(inst(t,param), rest...)
super(t::TagT) = t.name===TupleName ? AnyT : inst(t.name.super, t.params...)
extend(d::Dict, k, v) = (x = copy(d); x[k]=v; x)
                 env) = t
subst(t::TagT,
                  env) =
```

```
t == = AnyT ? t : TagT(t.name, map(x->subst(x,env), t.params), t.vararg)
subst(t::UnionT, env) = UnionT(subst(t.a,env), subst(t.b,env))
subst(t::Var,
                 env) = get(env, t, t)
function subst(t::UnionAllT, env)
 newVar = Var(subst(t.var.lb,env), subst(t.var.ub,env))
 UnionAllT(newVar, subst(t.T, extend(env, t.var, newVar)))
end
rename(t::UnionAllT) = let v = Var(t.var.lb, t.var.ub)
 UnionAllT(v, inst(t,v))
end
type Bounds
 lb; ub
         # current lower and upper bounds of a Var
 right::Bool # this Var is on the right-hand side of A <: B
type UnionState
                      # number of union decision points we're inside
 depth::Int
 more::Bool
                      # new union found; need to grow stack
 stack::Vector{Bool} # stack of decisions
 UnionState() = new(1,0,Bool[])
end
type Env
 vars::Dict{Var,Bounds}
 Lunions::UnionState
 Runions::UnionState
 Env() = new(Dict{Var,Bounds}(), UnionState(), UnionState())
issub(x, y)
                         = forall_exists_issub(x, y, Env(), false)
issub(x, y, env)
                        = (x === y)
issub(x::Ty, y::Ty, env) = (x === y) || x === BottomT
function forall_exists_issub(x, y, env, anyunions::Bool)
  for forall in false:anyunions
    if !isempty(env.Lunions.stack)
      env.Lunions.stack[end] = forall
    end
    !exists_issub(x, y, env, false) && return false
    if env.Lunions.more
      push!(env.Lunions.stack, false)
```

```
sub = forall_exists_issub(x, y, env, true)
      pop!(env.Lunions.stack)
      !sub && return false
    end end
  return true
end
function exists_issub(x, y, env, anyunions::Bool)
  for exists in false:anyunions
    if !isempty(env.Runions.stack)
      env.Runions.stack[end] = exists
    env.Lunions.depth = env.Runions.depth = 1
    env.Lunions.more = env.Runions.more = false
    found = issub(x, y, env)
    if env.Lunions.more
      return true # return up to forall_exists_issub
    if env.Runions.more
      push!(env.Runions.stack, false)
      found = exists_issub(x, y, env, true)
      pop!(env.Runions.stack)
    end
    found && return true
  end
  return false
end
function issub_union(t, u::UnionT, env, R, state::UnionState)
  if state.depth > length(state.stack)
    state.more = true
    return true
  ui = state.stack[state.depth]; state.depth += 1
  choice = u.(1+ui)
  return R ? issub(t, choice, env) : issub(choice, t, env)
end
issub(a::UnionT, b::UnionT, env) = issub_union(a, b, env, true, env.Runions)
issub(a::UnionT, b::Ty, env)
                               = issub_union(b, a, env, false, env.Lunions)
                               = issub_union(a, b, env, true, env.Runions)
issub(a::Ty, b::UnionT, env)
# take apart unions before handling vars
issub(a::UnionT, b::Var, env) = issub_union(b, a, env, false, env.Lunions)
```

```
issub(a::Var, b::UnionT, env) = issub_union(a, b, env, true, env.Runions)
function issub(a::TagT, b::TagT, env)
  a === b && return true
 b === AnyT && return true
  a === AnyT && return false
  if a.name !== b.name
    return issub(super(a), b, env)
  end
  if a.name === TupleName
    va, vb = a.vararg, b.vararg
   la, lb = length(a.params), length(b.params)
    ai = bi = 1
    while ai <= la
      bi > 1b && return false
      !issub(a.params[ai], b.params[bi], env) && return false
      ai += 1
      if bi < lb || !vb
        bi += 1
      end end
    return (la==lb && va==vb) || (vb && (la >= (va ? lb : lb-1)))
  end
  for i = 1:length(a.params)
    ai, bi = a.params[i], b.params[i]
    (issub(ai, bi, env) && issub(bi, ai, env)) || return false
  return true
end
function join(a,b,env)
  (a===BottomT \mid | b===AnyT \mid | a === b) && return b
  (b===BottomT || a===AnyT) && return a
  UnionT(a,b)
end
issub(a::Ty, b::Var, env) = var_gt(b, a, env)
issub(a::Var, b::Ty, env) = var_lt(a, b, env)
function issub(a::Var, b::Var, env)
 a === b && return true
  aa = env.vars[a]; bb = env.vars[b]
  if aa.right
    bb.right && return issub(bb.ub, bb.lb, env)
    return var_lt(a, b, env)
  else
    if !bb.right # check âLĂa,b . a<:b
```

```
return issub(aa.ub, b, env) || issub(a, bb.lb, env)
    end
    return var_gt(b, a, env)
  end
end
function var_lt(b::Var, a::Union(Ty,Var), env)
 bb = env.vars[b]
  !bb.right && return issub(bb.ub, a, env) # check \forallb . b<:a
  !issub(bb.lb, a, env) && return false
  # for contravariance we would need to compute a meet here, but
  # because of invariance bb.ub \sqcap a == a here always.
 bb.ub = a # meet(bb.ub, a)
  return true
end
function var_gt(b::Var, a::Union(Ty,Var), env)
  bb = env.vars[b]
  !bb.right && return issub(a, bb.lb, env) # check \forallb . b>:a
  !issub(a, bb.ub, env) && return false
  bb.lb = join(bb.lb, a, env)
  return true
end
function issub_unionall(t::Ty, u::UnionAllT, env, R)
 haskey(env.vars, u.var) && (u = rename(u))
  env.vars[u.var] = Bounds(u.var.lb, u.var.ub, R)
  ans = R ? issub(t, u.T, env) : issub(u.T, t, env)
  delete!(env.vars, u.var)
  return ans
end
issub(a::UnionAllT, b::UnionAllT, env) = issub_unionall(a, b, env, true)
issub(a::UnionT, b::UnionAllT, env)
                                     = issub_unionall(a, b, env, true)
issub(a::UnionAllT, b::UnionT, env)
                                      = issub_unionall(b, a, env, false)
issub(a::Ty, b::UnionAllT, env)
                                       = issub_unionall(a, b, env, true)
issub(a::UnionAllT, b::Ty, env)
                                      = issub unionall(b, a, env, false)
```

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