#### FacsXpert July 8, 2004 7<sup>th</sup> International Protégé Conference

- Builds protocols for studies with FACS instruments
- Uses a modified Protégé-based architecture that promotes runtime extensibility for the end-user

#### **FacsXpert**



- Built
  - At Stanford University's Herzenberg Lab
  - In close collaboration with Lee Herzenberg, James Tung, David Parks, Wayne Moore and other researchers in the Herzenberg Laboratory
- With help from
  - Mark Musen, Ray Fergerson, Monica Crubezy, Natasha Noy and the Protégé community

## FacsXpert



- Presented by
  - Stephen Meehan, software engineer Herzenberg Lab, Stanford University <u>meehan@darwin.stanford.edu</u>
  - Since 1992, Stephen has pursued a "dream" of a "software democracy" architecture where:
    - An application derives the majority of its behavior from runtime interpretation of its UML design model
    - A user can extend this model while the application runs

Thus the model is more than a design artifact, it is runtime government by the people ... <u>for</u> the people



#### **Scope of Presentation**

- 1. The FACS domain and need for runtime extensibility
- 2. FacsXpert's approach to data structure extensibility
- 3. FacsXpert's approach to data integrity extensibility
- 4. FacsXpert's approach to GUI extensibility



### **FACS** instruments

Count and characterize fluorescentlabeled cells in suspension to

- Monitor the progress of HIV infections by counting the number of CD4 T lymphocytes in blood from HIV-infected people
- Classify and stage tumors and to monitor bone marrow transplant survival
- Do the basic science and studies that underlie all these clinical advances

# FACS protocols specify what will be done in a FACS assay



#### For each tube in the assay

- Add up to 20 reagents, each linked to one of 12 fluorescent dyes
- Add one cell sample of up to 5 million cells
- Incubate and analyze with a multi-laser (Hi-D) FACS instrument
- Store the resulting data file (often about 5 megabytes)

One assay often has 60 or more tubes



## Sophisticated software is needed

- Must compute a feasible combination of staining reagents (which "label" cells) by taking into consideration:
  - Targeted species
  - Targeted cellular markers
  - Highly variable optical characteristics of fluorescent elements
  - Optical detection capabilities of a FACS instrument's configuration
  - Fluorescence interference between fluorescent elements
  - Purpose of cell labeling: to gate, dump or discover cell populations
  - Availability of inventory
  - Bio-chemical affinities between staining steps



#### Sophisticated software is needed

- Pipetting guide's
  - Reagent and specimen amounts must follow highly variable dilution recommendations
  - Organization must cater to pipetting technician work flow which differs from the planning work flow of the scientist
- Final protocol knowledge must interoperate with
  - Instrument software (offline as well as real time)
  - Analysis software
- GUI for decision making must be highly intuitive

Senior FACS scientists are often somewhere between computer naïve and totally computer-phobic

#### Highly extensible software is needed



- FACS vocabulary and "best practices" are
  - Minimally standardized
  - Rapidly changing
- From day to day, FACS scientists cannot predict
  - What questions they will be asking
  - What materials they will be working with
- Hence, their "research planning" software applications must be highly flexible and customizable

## cont.)

## Highly extensible software (cont.)

- THUS ... scientists (in addition to knowledge/software engineers) must be able to extend
  - Data structures (classes, attributes, etc)
  - Data integrity (rules that check knowledge inputs)
  - The GUI
  - ... without crashing the current application or future upgrades!!!
- Protégé provides the foundation for such a runtime evolvable system
- Commercial alternatives (e.g. Oracle, Rational Rose) suffer from higher financial cost, higher sys admin cost, closed source and design-time centricity

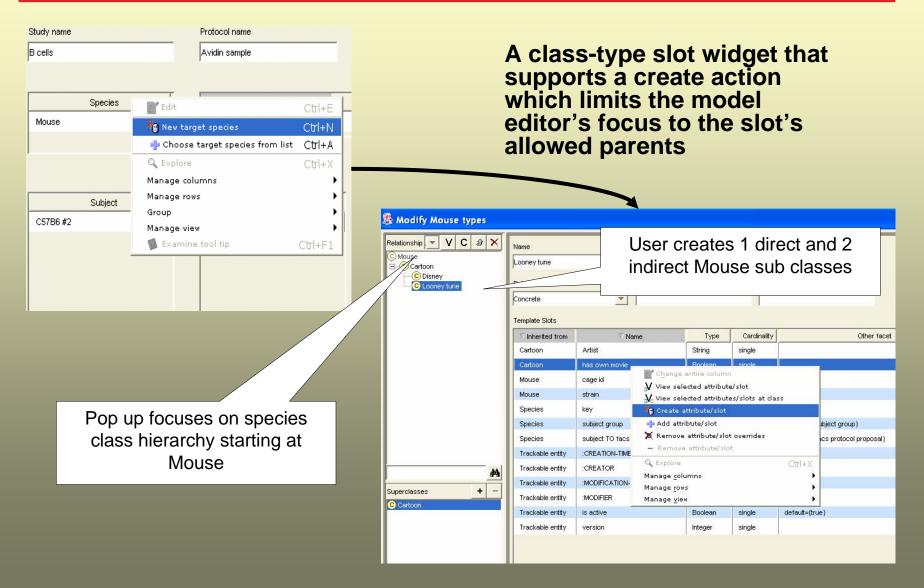


#### Data structure extensibility

- Protégé supports data structure extensibility through object-oriented sub classing
- However, the its model editor GUI overwhelms the scientist by exposing the whole model

FacsXpert required a model editor that only focuses on a specific part of the model at the specific time that it is relevant to the application

# *FacsXpert* solution for data structure extensibility



# FacsXpert solution for data structure extensibility



• A small # of programming idioms ready *FacsXpert* for such extensions; for example:

– Creating instances

```
myProject.createInstance( null,
    DisplayUtilities.pickCls( null,
        Collections.singletonList( designTimeLeafClass
)))
```

- Querying class type

designTimeLeafClass .equals( unknownClass ) || unknownClass.hasSuperclass( designTimeLeafClass )



#### Instantiating a new subject

#### 🖄 Instance: researcherSpecificKB 00038 The programming idiom detects Key Tracking the need to query for sub classes to the mouse class Strain Groups name Cage Id Protocol name Study name B cell sample Demo Demo Avidin sample Species √ FACS configure Aria 12 color Protocols proto Tiese End user's extended attributes Remove selected subject Ctrl+R Select Class Q Explore automatically appear A Eind/filter C Mouse Manage <u>c</u>olumns - Cartoon Manage rows - 🔘 Disney Group C Looney tune Manage <u>v</u>iew 🐔 Examine tool tij 4 Artist ġġ, Has Own Movie 🗸 ок 🗙 Cancel ۰. 🗸 Done



### Darwin "crash-proofs" data structure extensibility

We built a tool named Darwin that guards model evolution by handling model editions which break hard-coded expectations

- For any given class, Darwin prevents the "extinction" of one or more slot associations and one or more slot facets
- One can set 1 of 2 "watch dogs" to monitor changes to "endangered" model elements:
  - Golden retriever barks when user changes such an element and allows rollback (for engineers)
  - Doberman Pincher prevents user from making the change (for end-users)



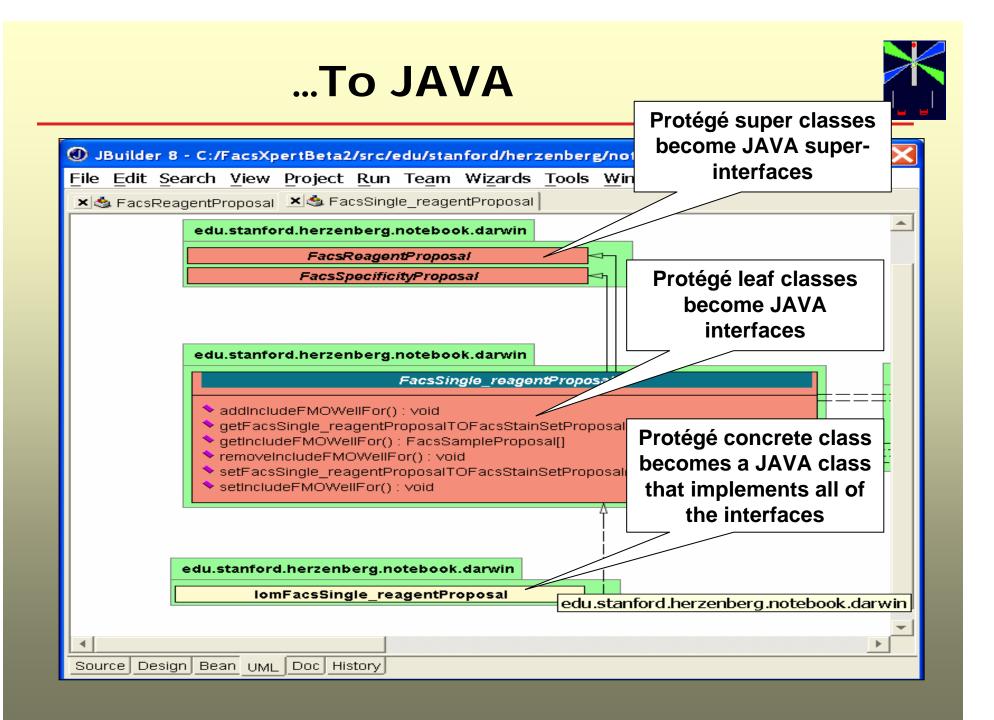
#### Darwin's JAVA generator

- Darwin generates JAVA modules that benefit client code by
  - Establishing a simple and consistent idiom for <u>type-</u> <u>safe</u> access to the <u>endangered</u> portions of the Protégé model
  - Adding JAVA compiler checking as a means of tracking model dependencies
- Darwin translates
  - All Protégé classes into JAVA interfaces hence supporting multiple inheritance
  - Concrete Protégé classes into JAVA classes that implement all associated JAVA interfaces

## From Protégé class...



🔘 Classes 🔝 Slots 📰 Forms 稭 FacsXpert 1.0 b	eta 2.0						
Relationship Superclass 🛛 🔽 💙 C 🕹 🗙	C Facs single-reagen	t proposal (type=:STANDARD-CLA	NSS)				
	Name	Documentation		Constr	aints V C + -		
E-C:SYSTEM-CLASS	Facs single-reagent	This is a p	roposal for				
E(C) Human agent	I dee ongie reagent i	FACS channe.	-				
ÈC Locator C Endeavor	Role	a single de	terminant.				
C Participation							
C Relation	Concrete	<b>_</b>					
C Ontology version	Template Slots						
C Vial shipment	$\nabla$ Inherited from	⊽ Name	Туре	Cardinality	Other fac		
🗄 🗠 🖸 Trackable entity	Facs specificity	assigned handle	Instance	single	classes={Molecule}		
i⊟© Research workflow element i⊟© Proposal	Facs specificity	assigned optical detector	String	single			
	Facs specificity	chosen fluorescent element	Instance	single	classes={Fluorescent element}		
	Facs specificity	chosen handle	Instance	single	classes={Molecule}		
E Facs volume and repeat proposal				-			
C Facs sample proposal	Facs specificity	determinant	Instance	required si	classes={Determinant}		
	Facs specificity	facs lot and dilution proposal	Instance	multiple	classes={Facs lot and dilution prop		
C Facs dilution and stain step propos∉	Facs specificity	one step recommended titration	Symbol	single	allowed-values={high,low,medium,		
C Facs cocktail volume proposal	Facs specificity	picked chemical reagent	Boolean	single	default={false}		
Facs plate layout proposal	F			single	allowed-values={high,low,medium,		
⊟⊸C Facs parameter proposal	Fa	ss "FACS single-re	single	allowed-values={high,low,medium,			
C Facs multi-reagent proposa	propos	al" has two direct	super	multiple	ality Other face classes={Molecule} classes={Fluorescent element} classes={Molecule} si classes={Determinant} classes={Determinant} classes={Facs lot and dilution prop allowed-values={high,low,medium, default={false} allowed-values={high,low,medium, allowed-values={high,low,medium, allowed-values={high,low,medium, allowed-values={high,low,medium, allowed-values={high,low,medium, allowed-values={high,low,medium, default={true}}		
C Facs single-reagent propos	Tr	classes		single			
C Facs der proposal	Trackab	TEATOR	String	single			
	entity	:MODIFICATION-TIMESTAMP	String	single			
AA .	Trackable entity	:MODIFIER	String	single			
Superclasses -	Trackable entity	is active	Boolean	single	default={true}		
C Facs reagent proposal	Trackable entity	version	Integer	single			
C Facs specificity proposal		facs single-reagent proposal TO	Instance	required si	classes={Facs stain set proposal}		
		include FMO well for	Instance	multiple	classes={Facs sample proposal}		





## Data integrity extensibility

#### Primary goal

 Runtime extensibility of constraints that check knowledge inputs (hereafter <u>checks</u>). Known to the Protégé community as "knowledge acquisition"

#### Out of scope

- Runtime extensibility of constraints that
  - infer new knowledge
  - validate ontologies (consistency checking, disjointedness, semantic imbalance etc.)

#### In practical terms...

 Allow scientists to author and authorize checks while FacsXpert runs



#### Requirements

- A. Decouple check inquirers from check providers
- **B.** Support reuse for both of the above actors:
  - A. Inquirer: "Is anything wrong?"
  - **B.** Provider:
    - A. Domain independent
    - B. Parameter-izable check <u>or</u> check template; user completes parameters when "binding"
- C. Make checks model associated and oriented
- D. Integrate checks with Protégé forms & pick lists



#### Requirements

- E. Make checks scaleable
  - A. Focus on limited instance context
  - **B. Incremental execution**
- F. Stay in sync with model re-factoring
- G. Support stepwise debugging
- H. Represent constraints in model
  - A. Start at :CONSTRAINT super class
  - B. Support inquirer querying of characterizations such as: advisory, warning, error and fatal

#### Requirements

- I. Allow user to add checks while FacsXpert runs
- J. Support end user, ad hoc check authoring
- K. Allow computability of dependencies, conflicts & bottlenecks
  - A. Prevent conflicts between checks
  - B. Prevent conflicts between checks and model facets and onto-clean meta classes/slots
- L. Have reasoning agents that use computability (otherwise prior requirement is "pie in the sky")



## Data integrity extensibility

**Engineering options (in 2002)** 

- 1. Reuse an existing runtime computable grammar
  - ✓ Pal with EzPAL
  - ✓ JESS
  - ✓ OCL
- 2. Invent a new "non programmatic" grammar; implementation options included:
  - 1. Translate invented grammar to accepted grammar to reuse interpreter
  - 2. Interpret the invented grammar directly
- 3. New JAVA validity checking framework that promotes "parameterized check templates"

#### We took choice 3

... and met 7 of 12 requirements

- ✓ A. Decouple check inquirers from check providers
- ✓ B. Support reuse in both of the above actors
- ✓ C. Make checks model associated *and* oriented
  - 0 By sub classing Darwin
- ✓ D. Integrate checks with Protégé forms & pick lists
- ✓ E. Make checks scaleable
- ✓ F. Stay in sync with model re-factoring:
  - o by sub classing Darwin class
- ✓ G. Support step-wise debugging



### An example of validity checking

Optical detector class contains validity checks such as

- Number assigned for a FACS parameter must be
  - Between 1 and the number of colors for the parent FACS configuration
  - Unique amongst sibling optical detectors
- The laser configuration associated with a detector must also be associated with the FACS configuration parent



## Checking integrity for parameter number value

Ar 🔛	ria 12	color	(type=Facs	, name=re	agents_16789)							
Config Aria 13	uration na 2 color	ame		۱ 	Max colors 12	_	Tracking					
VVar ∠	configural velength 407 488 633	tions Krypto Diode HeNe				a configuration template erg settings.conf		signi	ifies that	m for op at the us ect paran	er has	entered
	l detector: ser config	_	∆ Band pass filter	Parameter r	name Parameter	rder						
Diod	le 488 le 488	,	780/60	Cy7PE Cy55PE	12		Cy55PE	(type=C		name=reagents_	<b> X</b>	
	ie 488 ie 488		665/30 610/20	Cy5PE TRPE	2		band pass	center (	(nm)	🔽 In use		
	ie 488 ie 488		575/25 525/50	PE FL	4		710/40	710	0	j∙ in use	Tracking	
HeN HeN	-		780/60 710/40	Cy7APC Cy55APC	6 7		⊽Wavelen			Key color Cy55PE	PMT Label	
НеМ Кгут	-		660/40 540/80	APC Alexa430	8		488 Di	ode 488				
Кгур Кгур			465/30 440/40	CasYel CasBlu	10 11		Facs Parameter of		parameter # not exceed the max	Dichroic × colors (12) for the F	Dichoric Type ACS configuration	



## **Checking integrity of Laser** configuration relationship

State (	type=Optical detect	or, name=reagents						
band pass 710/40	center (nm) range (nm 710 40	) 🔽 In use	Tracking					
		Key color	💈 laser pick list		X			
VVavelen 488 Dio	Laser 📫 Choose laser from l	Cy55PE	⊽ Laser	Wavelength				
	🔍 Explore	Ctrl+X	Argon	488	<u> </u>			
Facs	Manage <u>c</u> olumns	► lic	Diode	635				
Aria 12 co	Manage <u>r</u> ows <u>G</u> roup Manage view	> <sup>  C</sup>  >	Can not select item because     The laser configuration is not setup for the current FACS configuration ""Diode"" is incompatible with "["Krypton", "Diode 488", "HeNe"]"					
			Liloge 466	486				
			DYE	595				
			Helium neon 633	405				
			HeNe	633				
			Kr/Ar UV	355				
			Krypton	407	<b>•</b>			
disabled. A	plays invalid o n explanatory on the mouse	/ tool tip		<u>Pick</u>				

( over the disabled item.



#### What are our next steps?

#### Meet remaining requirements

- H. Represent constraints in model
- I. Allow user to add checks while FacsXpert runs
- J. Support end user, ad hoc check authoring
- K. Allow computability of dependencies, conflicts & bottlenecks
- L. Have reasoning agents that use computability, otherwise prior requirement is pie in the sky

#### Current approach is limited:

- H and I are easily achievable, but not J, K and L
- At best it will become hidden plumbing for approach that hits all 12 requirements



#### And then there's the GUI





FACS scientists and the Protégé GUI were like oil and water

- They were confused by the slot buttons with +, -, C,V,
   X ... and they balked at the dialogs
- They begged for a highly customizable Xcel-like widget (grid) for manipulating tables of data
- Then, after many iterations of developing this table widget, they ultimately decided that navigation and group-based modifications of items in the table had to be further simplified.

They wanted to access the table via a tree!



## From a table (linear) view to...

🕅 Pipetting pl	an									
<u>N</u>	1	🗸 Done	Sample properties	Year Sample pipetting	<b>5</b>	Stain set pipetting	Comfort %	Plate setup	🊧 Full table	🕵 Help
⊽ Tissue	⊽ Stain set	⊽ Subject	Cell sample total ml	Cells/ml X 10 <sup>n</sup> 6	Repeats		Include Change entire c Explore		nce minus one (FMO) is a separate tube	Stain step volume (ul)
is not Lymph node						Ma Ma	Find/filter mage columns mage rows	•		-
Lung	B stain	C57B6 #2	2.5	25	1		nage view	•	B Save this vie Open differe Copy	
Lung	B stain	C57B6 #3	2.5	25	1	25	V	-	Paste view	Ctrl+Insert
Lung	B stain	CD22KO #4	2.5	25	1	25	V			1 25.0
Lung	B stain	CD22KO #5	2.5	25	1	25	V			1 25.0
Lung	B stain	CFSE C57B6	10	25	1	50	<b>N</b>			1 50.0
Lymph node leg	B stain	C57B6 #2	2.5	25	1	25	N			1 25.0
Lymph node leg	B stain	C57B6 #3	2.5	25	1	25	ম			1 25.0
Lymph node leg	B stain	CD22KO #4	2.5	25	1	25	<b>N</b>			1 25.0
Lymph node leg	B stain	CD22KO <b>#</b> 5	2.5	25	1	25	<u>र</u>	CD5/Cychrome	, lgD/Cy7-PE	1 25.0
					a 45 of 60 tub		-			1 50.0



## ...To a tree (hierarchical) view

JUNE	🖌 Done	Samp propen	le 🛛 🔫 S ties p	ample ipetting	Stain set pipetting	Comfort %	Plate setup	🏘 Full table	🥵 Help
ee: Stain set, Tissue, Subj	ect 9 tree select	ions filter 8 d	of 60 tube pla	n rows					
root ■ B stain ■	▲ V Stain set	⊽ Subject	⊽ Tissue	Include in assay		ce minus one Change entire for all cells in the s	column	Stain volume (ul)	
C57B6 #3	B stain	C57B6 #2	Lung	ব		Hanage columns Manage rows		25.0	Ī
	B stain	CD22KO #4	Lymph node	ų		Group Manage view		25.0	
C57B6 #3 CD22KO #4 CD22KO #5	B stain	CD22KO #4	Lymph nod	ম			1	25.0	
	B stain	C57B6 #2	Spleen	N			1	25.0	
	B stain	C57B6 #3	Spleen	ম			1	25.0	
	B stain	CD22KO #4	Spleen	되			1	25.0	T
	B stain	CD22KO <b>#</b> 5	Spleen	<b>N</b>			1	25.0	
Mature B → ♥ Lung	B stain	CFSE C57B6	Spleen	Y				50.0	
C57B6 #2 C57B6 #3 C522KO #4 C522KO #5 C52E C57B6 CFSE C57B6									

## Xpert grid/table supports



#### User customizability

- Customize column: names, sizes, display order and sort order
- Filter rows

#### Persistence of user customizations

- Automatically remember and use the most recent customizations
- Allow the user to save customizations
- Allow the user to save and retrieve sets of customizations into property files

#### Tree-based read/write access

- Define a tree structure to summarize elements of the underlying table
- Select any combination of tree nodes
- See the column entries sieved by the node selections
- Modify all cells in a column for the sieved rows

### **GUI extensibility**



- Added pick list plug-in framework
- Added system menu plug-in and overrides
- Added numerous other slot widgets

