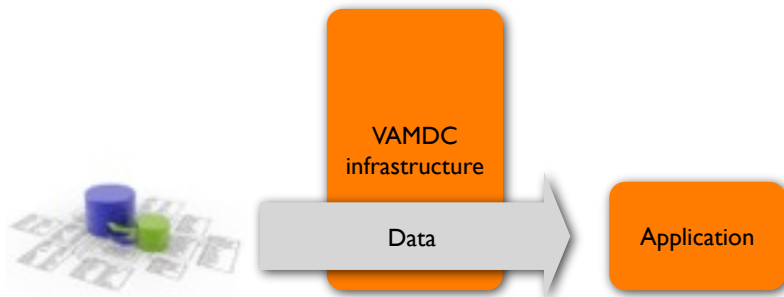


VAMDC technology

Guy Rixon
Innsbruck, February 2013



Plan A

Dump each database into a file and put on web.



Pro:

- “The simplest thing that could possibly work”
- Everything you can get has its own URL

Con:

- Data-sets too large (up to 10GB)
- No easy way to make data extracts

Plan B

Pre-compute all possible data extracts and dump on web



Pro:

- Selection now easy
- One URL for each possible extract

Con:

- Impossible to implement!

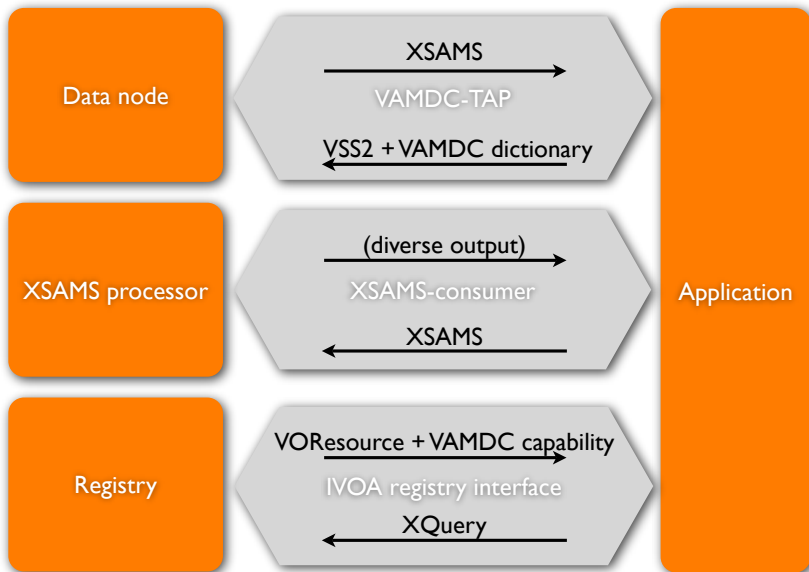
Plan C

Compute data extracts on demand
but
index them on the web as if pre-computed



- Pro:
 - Implementation now feasible
 - Still have a URL for every data-extract
- Con:
 - Some assembly required
 - Need to define standards for services, queries etc.

The core standards



Data URLs

The address of the service
(this one for CDMS)



[http://cdms.phl.uni-koeln.de:8090/DjCDMS/tap/sync?
REQUEST=doQuery&LANG=VSS2&FORMAT=XSAMS
&QUERY=SELECT+ALL+WHERE+MoleculeStoichiometricFormula+%3D+%27CO%27](http://cdms.phl.uni-koeln.de:8090/DjCDMS/tap/sync?REQUEST=doQuery&LANG=VSS2&FORMAT=XSAMS&QUERY=SELECT+ALL+WHERE+MoleculeStoichiometricFormula+%3D+%27CO%27)

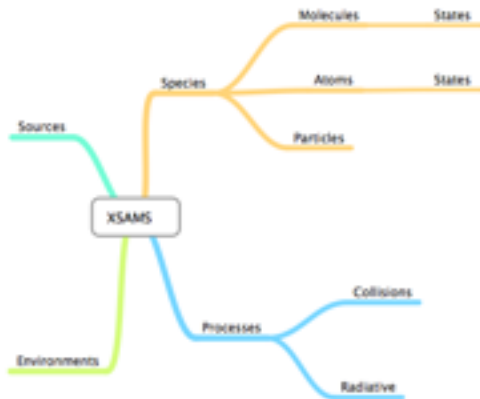
The data to be extracted

Call these with HTTP GET

XSAMS

- “XML schema for Atoms, Molecules & Solids”
- Developed by IAEA & VAMDC:
 - Proposed 2003, at IAEA DCN meeting
 - First versions by (IAEA, NIST, ORNL U. Pierre & Marie Curie, OPM, RFNC-VNIITF)
 - Subsequent development by VAMDC
- See <http://www-amdis.iaea.org/xsams/>
- See also <http://www.vamdc.org/documents/standards/#data-model>

XSAMS structure: top



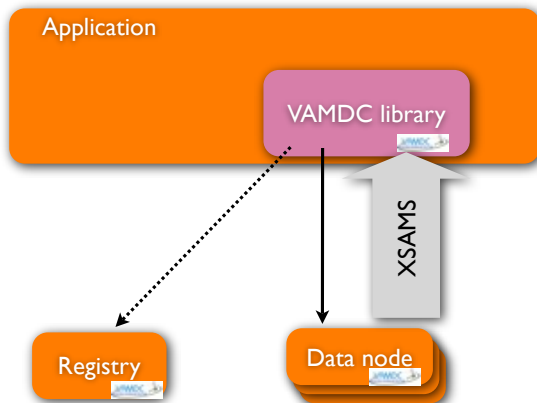
XSAMS structure: bottom

- All quantities have units
- All values can have associated uncertainties
- All values can have a source reference
- XML \Rightarrow no encoding issues for numbers

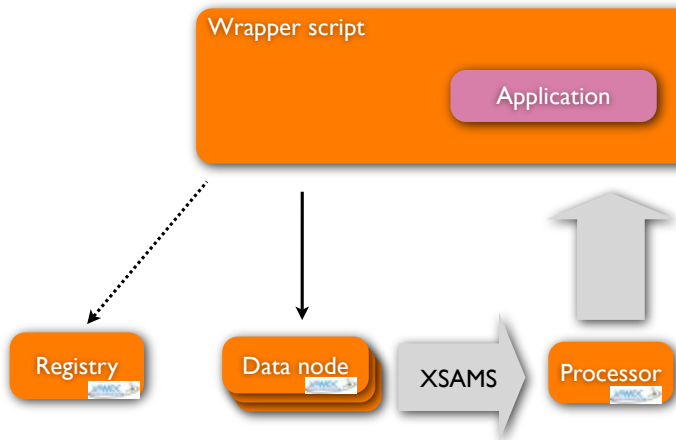
XSAMS for molecules

- “Case-by-case” XSAMS:
- Separate, additional schema for each class of molecule:
 - 1. Diatomic closed shell (dcs): CO, N₂, NO⁺
 - 2. Hund’s case (a) diatomics (hunda): NO, OH [for low J]
 - 3. Hund’s case (b) diatomics (hundb): O₂, OH [for high J]
 - 4. Closed-shell, linear triatomic molecules (1tcs): CO₂, HCN
 - ...etc up to at least 12 cases

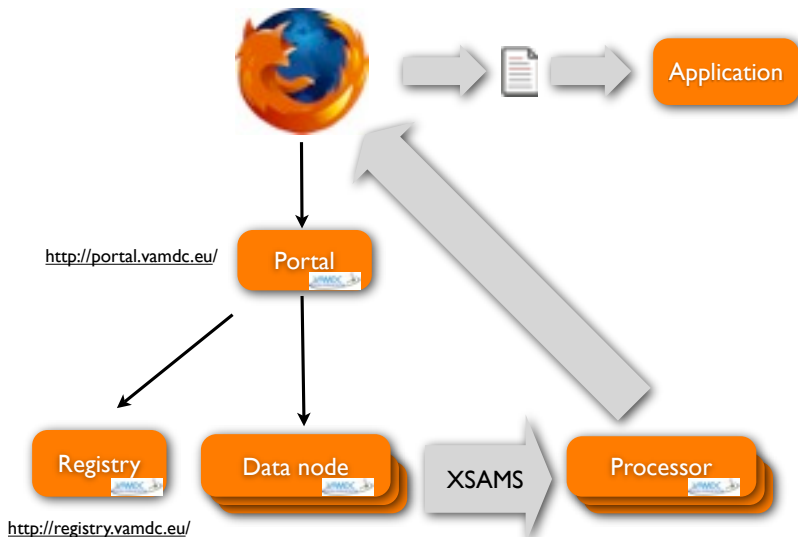
Adapted application



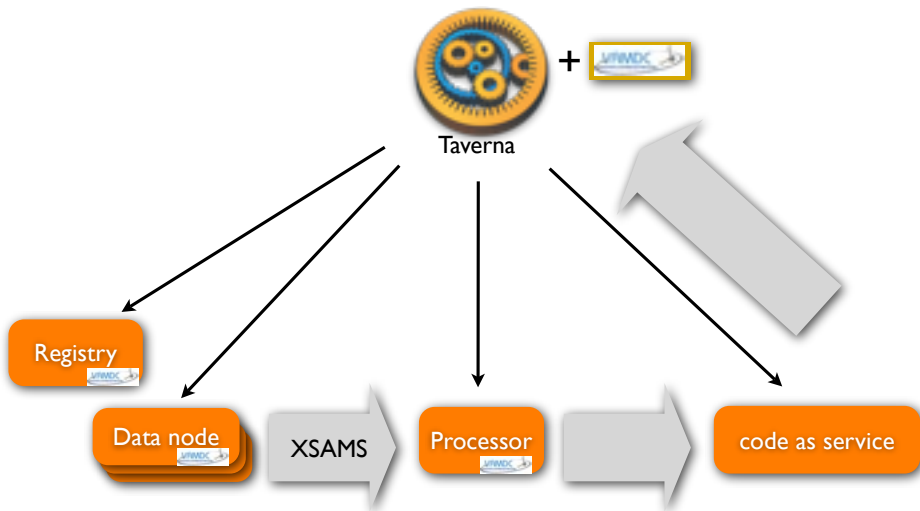
Wrapped application



Portal, nodes & processors



Taverna; code as service

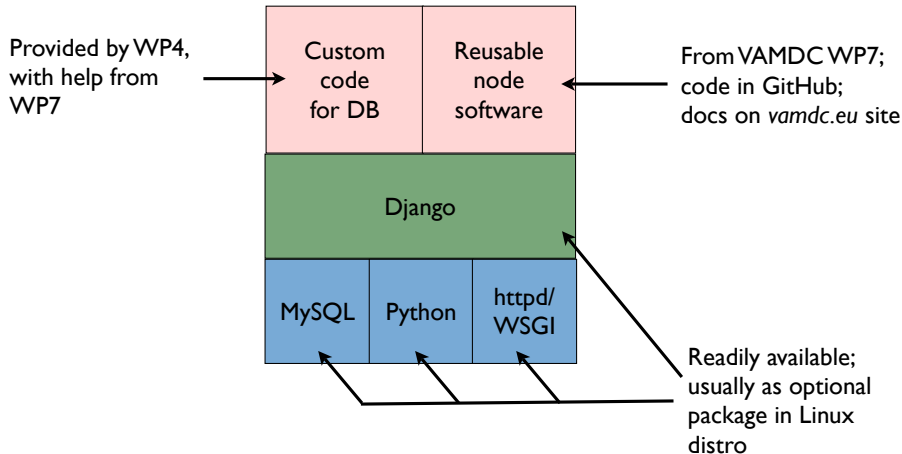


So how do I make a node?

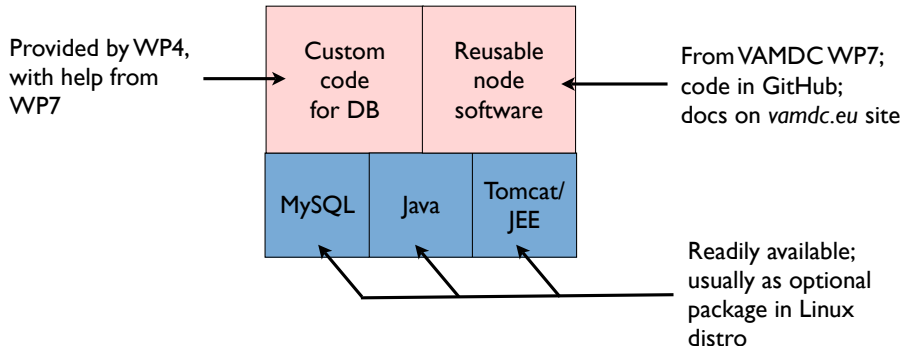
(And will it hurt?)

Node = database + web server + node software

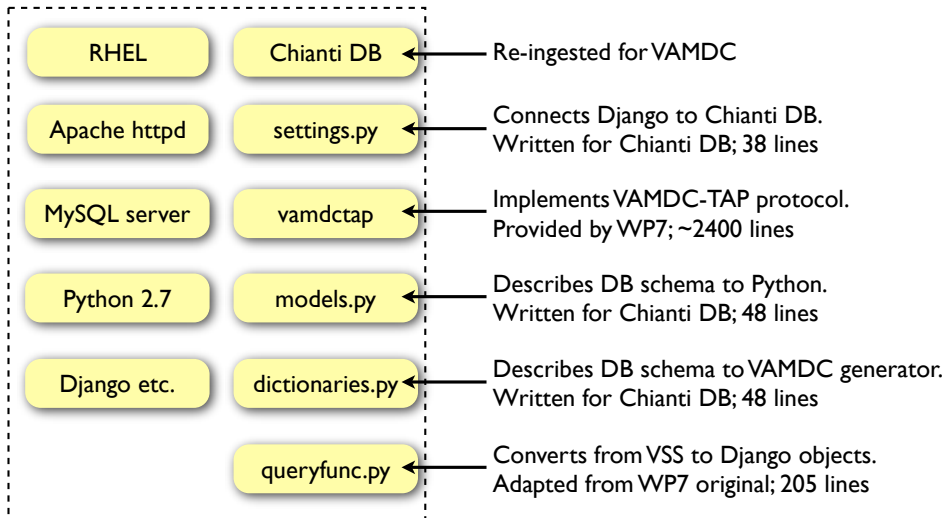
Node software in Python



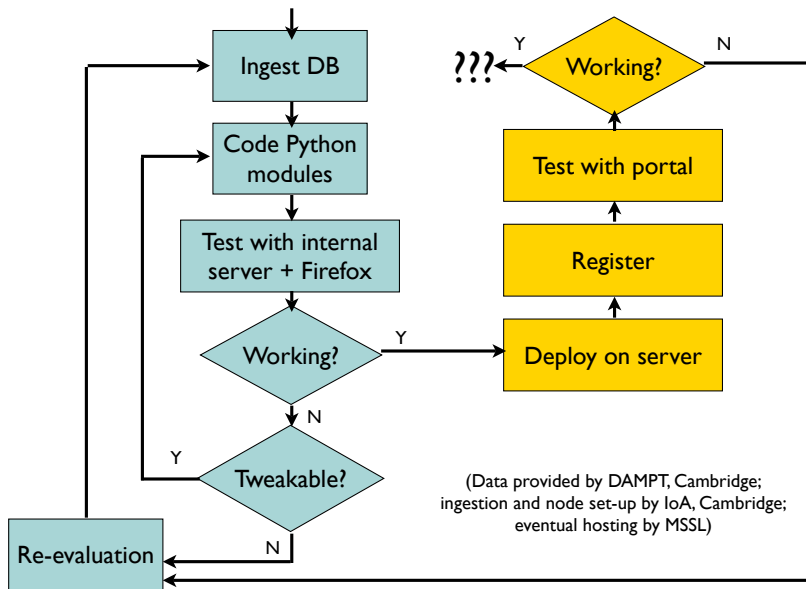
Node software in Java



Node example: Chianti



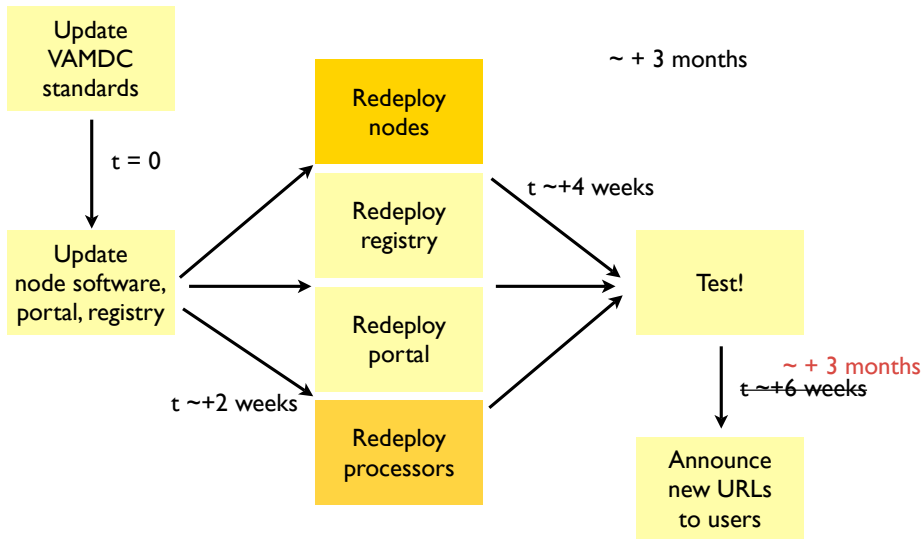
Chianti example (cont.)



System versions

- System version defined by standards version
- Three so far:
 - 11.05 (withdrawn)
 - 11.12 (current, released)
 - 12.07 (in preparation, to be released in 2013)
- Expect one new version per year from now on
- new standards \Rightarrow new deployments on new URLs

Annual updates of standards



Timeline of node registrations

