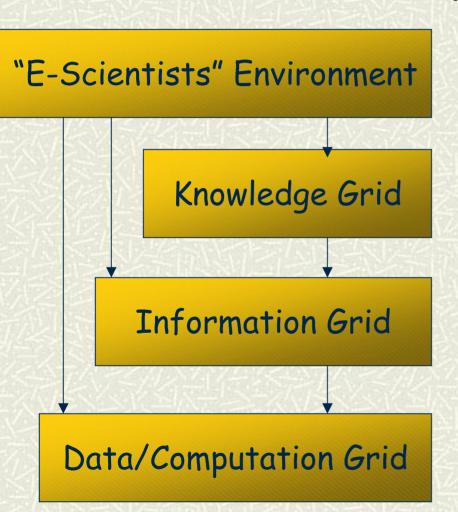
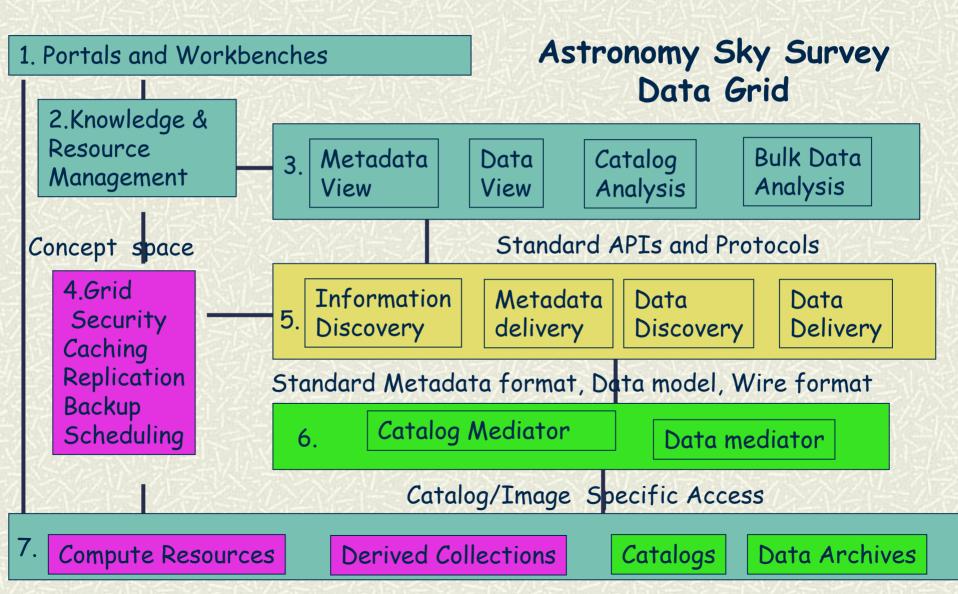
What is the Grid?



In silico experiments

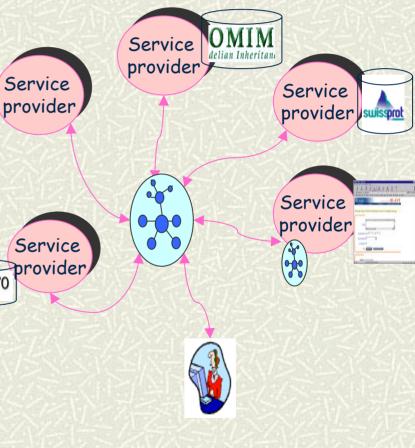
- Information harvesting & PSE
- Dynamically forming virtual organisations to solve problems.
- Describing, searching for and weaving resources: people. applications, db, content, instruments
- Orchestrating resources
- Support for scientific method: provenance, argumentation, opinion contextualisation etc

Grid is metadata based middleware



Grid is... dynamic marshalling of resources

- Needs describing the resources, mapping between resources.
- Resources include BIG databases - the instances will not be on the web, they will be inside applications and databases (there isn't much difference).
- Semantic web technologies for shared meaning (through ontologies) and shared models (e.g. exporting results through RDF and using inference over them).



Isn't information all computationally accessible already?

- Document publishing paradigm.
- Descriptive knowledge.
- Ontologies for controlling content already used.
- Evolving, nonpredictive schemas
- **#** XML is king.

ile <u>E</u> dit <u>V</u> iew	<u>Go</u> <u>C</u> ommunicator <u>H</u> elp
Database	InterPro
Accession	IPR000025; Metatonin_receptor (matches 22 proteins)
Name	Melatonin receptor
Туре	Family
Dates	08-OCT-1999 (created) 27-MAR-2000 (last modified)
Signatures	PR00657; MELATONINR (22 proteins)
Parent 🛈 [tree]	PR000276, Rhodopsin-like GPCR superfamily (3990 proteins)
<u>Children</u> (1) [tree]	PR002278; Melatonin 1 A receptor (12 proteins) PR002273; Melatonin 1 C receptor (5 proteins) PR002280; Melatonin-related 1X receptor (3 proteins)
Function 🕕	melatonin receptor (<u>GO:0008502</u>)
Component 🕕	membrane (<u>GO.0016020</u>)
<u>Abstract</u>	G-protein-coupled receptors (GPCRs) constitute a vast protein family that encompasses a wide range of functions (including various autocrine, paracrine and endocrine processes). They show considerable diversity at the sequence level, on the basis of which they can be separated into distinct groups. We use the term clan to describe the GPCRs, as they embrace a group of families for which there are indications of evolutionary relationship, but between which there is no statistically significant similarity in sequence [1]. The currently known clan members include the rhodopsin-like GPCRs, the sectim-like GPCRs, the sequence [1]. The currently known clan members include the rhodopsin-like GPCRs, the section-like GPCRs, the sequence [1]. The currently known clan members include the rhodopsin-like GPCRs, the sectim-like GPCRs, the sequence [1]. The currently known clan members include the rhodopsin-like GPCRs, the section-like GPCRs, the sequence [1]. The currently known clan members include the rhodopsin-like GPCRs, the section-like GPCRs, the sequence [1]. The currently known clan members include the rhodopsin-like GPCRs themselves represent a widespread protein family that includes hormone, neurotransmitter and light receptors, all of which transduce extracellular signals through interaction with guanine nucleotide-binding (G) proteins. Although their activating ligands vary widely in structure and character, the amino acid sequences of the receptors are very similar and are believed to adopt a common structural framework comprising 7 transmembrane (TM) helices [2, 3, 4]. Melatonin is secreted by the pineal gland during darkness [5]. It regulates a variety of neuroendocrine functions and is thought to play an essential role in circadian rhythms. Drugs that modify the action of melatonin, and hence influence circadian cycles, are of clinical interest (for example, in the treatment of jet-lag). Melatonin receptors are found in the retina, in the pars tuberalis of the pitulary, and in discrete areas of the brain. The
Examples	
References	1. Attwood T.K., Findlay J.B.C. <i>Fingerprinting G-protein-coupled receptors.</i> Protein Eng. 7: 195-203(1994). [<u>MEDLINE:94224751]</u> [PUB00004961]
-0-	Document Done

Grid is ... services, services, services

- The first generation of Grid was protocol based.
- Second generation is service based: Open Grid Service Architecture.
- Semantic Web description and annotation technologies core to service sophisticated service description and processing.



Descriptions => Automated discovery & search, selection, (imprecise) matching, composition & interoperation, invocation, execution monitoring

Reasoning is darn handy

The Semantic Grid is ...knowledge management

- Q: What ATPase superfamily proteins are found in mouse?
- 1. P21958 (from Swiss-Prot).
- 2. InterPro is a pattern database and could tell you if you had permission and paid.
- 3. Attwood's lab expertise is in nucleotide binding proteins
- 4. Jones published a new paper on this in Nature Genetics two weeks ago
- 5. Smith in your lab already asked this question...

Remarks

- **#** Semantic Web is a part of the Grid vision?
- Semantic Web technologies should be relevant for Grid metadata at all levels.
- There isn't one Grid, there are collections of Grids for communities - might be a more tractable model for the Semantic Web.
- Most facts will stay in databases. Metadata about the (scientific) process and facts could be in RDF.
- # E-Science (everyone?) loves XML and ignores RDF

Annotations sit in other (non RDF) databases.

Reliability, scalability, performance, explanation, longevity, evolution ...





