

**APPENDIX B. External collaborators, partners and known users**

	<b>Collaborator</b>	<b>Form of collaboration</b>
Service Provider	EMBOSS MRC Roslin Franklin Centre, <a href="http://emboss.sourceforge.net/">http://emboss.sourceforge.net/</a>	Soaplab executes EMBOSS services
	BioMART Ewan Birney, <a href="http://www.ebi.ac.uk/biomart/">http://www.ebi.ac.uk/biomart/</a>	Taverna plugins to execute BioMART queries. BioMart is a generic data management system which offers a range of advanced query interfaces and administration tools.
	Blueprint <a href="http://www.blueprint.org/">http://www.blueprint.org/</a> Ian Robertson, Mount Sinai Hospital, Canada	Taverna plug-ins to execute SeqHound services. <a href="http://www.blueprint.org/seqhound/">http://www.blueprint.org/seqhound/</a>
Specialist Platform developer	BIRN <a href="http://www.nbirn.net/">http://www.nbirn.net/</a> Mark Ellisman, SDSC, USA	Joint work on workflow management and provenance and semantics. Interoperating with the Kepler workflow environment.
	SEEK & GEON <a href="http://www.geongrid.org/">http://www.geongrid.org/</a> <a href="http://seek.ecoinformatics.org">http://seek.ecoinformatics.org</a> Bertram Ludaescher, UC Davis, USA	Joint work on workflow management and provenance and semantics. Interoperating with the Kepler workflow environment.
	Virginia Bioinformatics Institute Bruno Sobral, <a href="https://www.vbi.vt.edu/">https://www.vbi.vt.edu/</a>	Taverna used with Pathport's Toolbus integration tool
	The Sanger Centre Thomas Down, <a href="http://www.efamily.org.uk/software/dasservers.shtml">http://www.efamily.org.uk/software/dasservers.shtml</a>	The DALEC joint collaboration links Distributed Annotation Service with Taverna, as part of the eFamily project
	BioMOBY <a href="http://www.biomoby.org">http://www.biomoby.org</a> Mark Wilkinson, University of British Columbia, iCAPTURE Centre, St. Paul's Hospital, Vancouver, BC	Joint work. myGrid developed Java components for BioMOBY; all BioMOBY service executed; merging myGrid semantic discovery and registry with BioMOBY Central. Using the myGrid ontologies and semantic discovery

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Life Science Application User	Roslin Institute <a href="http://www.ri.bbsrc.ac.uk/">http://www.ri.bbsrc.ac.uk/</a> , Dave Burt	Using Taverna for QTL analysis of chicken genome
	Animal Genomics University of Liverpool, Steve Kemp <a href="http://www.genomics.liv.ac.uk/animal/Archives1.html">http://www.genomics.liv.ac.uk/ animal/Archives1.html</a>	Using Taverna to hunt for trypanotolerance genes in African cattle
	Newcastle Centre for Integrated Systems Biology of Ageing.	Using Taverna workflows for the identification of genes involved in apoptosis
	Virtual Lab for eScience Scott Marshall & Marcos Roos Integrative Bioinformatics Unit , University of Amsterdam <a href="http://www.micro-array.nl/IBU/index.html">http://www.micro- array.nl/IBU/index.html</a>	Using Taverna for the Vle Virtual Laboratory project
	Unilever Centre for Molecular Informatics Peter Murray-Rust, University of Cambridge, UK <a href="http://wwmm.ch.cam.ac.uk/">http://wwmm.ch.cam.ac.uk/</a>	Using Taverna for workflows using the JUMBO library to analyse small molecules
	Larry Hunter Center for Computational Pharmacology, Aurora, CO 80045-0511 USA	Using Taverna and working with the Semantic technologies.
	Anders Lanzen, Svern Helge Grindhaug, Pal Puntervoli Computational Biology Unit University of Bergen, Norway	Developed the Taverna collaboration service for people to interact with workflows
	EMBL-EBI and Sanger Centre <a href="http://www.ebi.ac.uk">http://www.ebi.ac.uk</a>	e-Family and other projects. Using Taverna for automatic target selection for protein structure and function studies
	Purdue University Purdue, USA	Using Taverna for bioinformatics training
	Paolo Romano ( <a href="mailto:paolo.romano@istge.it">paolo.romano@istge.it</a> ) Bioinformatics and Structural Proteomics National Cancer Research Institute (IST) Largo Rosanna Benzi, 10, I- 16132, Genova, Italy <a href="http://www.nettab.org/promano/">http://www.nettab.org/proman o/</a>	Using Taverna for workflows, and have developed a workflow portal.
Peter Covitz CaBIG <a href="http://cabig.nci.nih.gov">http://cabig.nci.nih.gov</a>	Taverna can execute caBIG services. Joint discussions on information Models	
Commercial	Lexicon Genetics <a href="http://www.lexicon-genetics.com/">http://www.lexicon- genetics.com/</a>	Commercial BioTech who contributed to Taverna codebase
	BioTeam Cambridge Massachusetts. <a href="http://rt.bioteam.net:8080/metadot/index.pl?id=2324&amp;isa=Category&amp;op=show">http://rt.bioteam.net:8080/met adot/index.pl?id=2324&amp;isa=Ca tegrity&amp;op=show</a>	Commercial BioTech consultancy using Taverna
	Dennis Quan, Sean Martin T.J. Watson IBM	Haystack and BioDASH provider; worked closely with Dennis and published joint papers; myGrid used as an example in IBM's presentations at WWW204.

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Generic infrastructure developer	Yolanda Gil, Carl Kesselman ISI, USC California, USA	Joint work on workflows and agent technologies; funded through an e-Science sisters grant.
	EGEE <a href="http://public.eu-egee.org/">http://public.eu-egee.org/</a> Tristan Glatard & Johan Montagnat French National Center for Scientific Research	Using Taverna for image processing workflows.
	GryPhyn Ian Foster Argonne National Labs	Investigating the use of Soaplab and GRIMOIRES. Joint work on Provenance funded through a sisters grant with Moreau.