We are very interested in your comments and interests in our ontology workshop. Please fill out this survey, which will provide invaluable feedback to us in terms of developing useful workshops for users.

Would you class yourself as:	
☐ Bench scientist	Which ontology would you like to use in your
☐ PI/Project manager	project(s)?
☐ Bioinformaticist within a database resource	☐ Gene Ontology (GO)
☐ Bioinformaticist in a lab	Molecular function/Biological process/Cellular component
☐ Other (please specify)	☐ Plant Ontology (PO)
	☐ Trait Ontology (TO)
	☐ Anatomy and growth stage ontology for your
Do you commonly use biological databases for your	favorite organism  ☐ Others (please specify)
work? If so, which ones do you use most frequently?	Others (please speerry)
(Please cross out those that do not apply)	Did you find the introduction to the GO and PO in
□ NCBI (Entrez, GenBank, PubMed, RefSeq)	this workshop useful?
☐ EBI (http://www.ebi.ac.uk/)	☐ Very useful
☐ UniProt/SWISSPROT	☐ Mostly useful
☐ Gramene/TAIR/MaizeGDB/TIGR/GrainGenes	☐ Fairly useful
☐ GRIN/IRIS/ABRC (Plant germplasm stock	□ Not useful
repositories)	
☐ MGI/SGD/FlyBase/Dictybase.	Did you find the content of the GO&PO training
☐ RGD/TIGR/ZFIN/Wormbase/PSU Sanger	section useful?
☐ Others (please specify)	☐ Very useful
	☐ Mostly useful
	☐ Fairly useful
How did you learn about ontologies?	□ Not useful
☐ This workshop	W14111
By visiting model organism databases	Would you like to be invited to a more detailed
☐ Learning from colleagues	workshop about GO or PO? □ GO
Read an article	□ PO
☐ Visited posters and other workshops ☐ Others (places appoint)	
☐ Others (please specify)	Would you like to be contacted about contributing
	annotations to GO or PO?
What model organism(s) do you work with? (Please	□ GO
specify).	□ PO
SP	
	Do you have any additional comments and/or
Which dataset from your project would you like to	suggestions about ontologies in general and the
annotate with ontologies?	workshop?
☐ Genes/gene products (proteins and transcripts)	
ESTs/Microarrays	
□ Proteomics	
Mutant Phenotypes / QTL	
☐ FSTs (Flanking Sequence Tags/insertion/deletion	
mutant lines)	
☐ Genetic stocks/germplasms ☐ Diseases	
☐ Others (please specify)	
in Onicis (picase specify)	