

## EuPathDB 2016 Vocabulary Survey Results - In order of appearance

**Q1 - Please indicate your familiarity with these terms and concepts.**

<b>Question</b>	<b>not at all familiar</b>	<b>have heard of it</b>	<b>somewhat familiar</b>	<b>Very Familiar</b>
Gene ID	0.00%	3.85%	30.77%	65.38%
Annotation	3.85%	11.54%	15.38%	69.23%
Parasite Isolate	3.85%	23.08%	7.69%	65.38%
Genomic Segment	4.00%	20.00%	36.00%	40.00%
Motif	3.85%	7.69%	19.23%	69.23%
Genomic Sequence	0.00%	3.85%	3.85%	92.31%
EST	15.38%	15.38%	34.62%	34.62%
ORF	0.00%	0.00%	19.23%	80.77%
SNP	4.00%	8.00%	16.00%	72.00%
Non-synonymous	11.54%	15.38%	15.38%	57.69%
Allele Frequency	0.00%	26.92%	34.62%	38.46%
Proteomics	0.00%	3.85%	42.31%	53.85%
Mass Spec Peptide	7.69%	19.23%	42.31%	30.77%
Microarray	4.00%	12.00%	20.00%	64.00%
Expression Value	4.00%	20.00%	28.00%	48.00%
RNA-Seq	0.00%	11.54%	38.46%	50.00%
Alignment	3.85%	0.00%	11.54%	84.62%
eQTL	65.38%	19.23%	11.54%	3.85%
ChIP-Seq	11.54%	38.46%	34.62%	15.38%
Histone modification	7.69%	15.38%	53.85%	23.08%
Nucleosome Mapping	30.77%	38.46%	23.08%	7.69%
Synteny	15.38%	23.08%	23.08%	38.46%
Ortholog	3.85%	3.85%	30.77%	61.54%
Contig and Scaffold	3.85%	19.23%	34.62%	42.31%
GO Terms/Numbers	26.92%	15.38%	38.46%	19.23%
GO Term Enrichment	30.77%	30.77%	26.92%	11.54%
EC Number	53.85%	3.85%	19.23%	23.08%
Signal Peptide	3.85%	15.38%	34.62%	46.15%
BLAST	0.00%	3.85%	11.54%	84.62%
Genomic colocation	42.31%	11.54%	34.62%	11.54%
Mapping	3.85%	15.38%	53.85%	26.92%
Transcript	0.00%	0.00%	30.77%	69.23%

Protein Data Bank	3.85%	15.38%	42.31%	38.46%
Transmembrane Domains	3.85%	3.85%	38.46%	53.85%
Epitope	8.00%	4.00%	24.00%	64.00%
Antibody	0.00%	7.69%	11.54%	80.77%
Splice Site Junction	15.38%	15.38%	34.62%	34.62%
Comparative genomic hybridization	26.92%	42.31%	15.38%	15.38%
FASTA	3.85%	3.85%	19.23%	73.08%
GFF	65.38%	11.54%	11.54%	11.54%
BAM	57.69%	15.38%	11.54%	15.38%
Genome Browser	11.54%	23.08%	30.77%	34.62%
OrthoMCL	52.00%	28.00%	12.00%	8.00%
Paralog	7.69%	19.23%	26.92%	46.15%
CDS	16.00%	4.00%	28.00%	52.00%
PubMed Compound	34.62%	19.23%	23.08%	23.08%
Metabolic Pathway	3.85%	0.00%	34.62%	61.54%
Metabolomics	7.69%	19.23%	30.77%	42.31%
Substrate	3.85%	3.85%	23.08%	69.23%
KEGG Pathway	30.77%	26.92%	7.69%	34.62%
ChIP-chip	26.92%	34.62%	30.77%	7.69%
Protein Microarray	11.54%	42.31%	42.31%	3.85%
Antibody Microarray	11.54%	53.85%	30.77%	3.85%
Metadata	20.00%	40.00%	28.00%	12.00%
Ontology	19.23%	30.77%	30.77%	19.23%
ncRNA	34.62%	19.23%	19.23%	26.92%
Polysomal Profiling	53.85%	26.92%	11.54%	7.69%
Locus	3.85%	3.85%	30.77%	61.54%

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