

2015 EuPathDB Vocabulary Survey Results

# Question	not at all familiar	have heard of it	somewhat familiar	Very Familiar	Total Responses	Mean
6 Genomic Sequence	0	0	4	24	28	3.86
29 BLAST	0	0	6	22	28	3.79
17 Alignment	0	1	5	22	28	3.75
39 FASTA	1	1	3	23	28	3.71
32 Transcript	1	1	5	21	28	3.64
1 Gene ID	0	1	10	17	28	3.57
33 Protein Data Bank	0	3	8	17	28	3.5
8 ORF	0	3	8	17	28	3.5
36 Antibody	1	4	5	18	28	3.43
58 Locus	0	5	7	16	28	3.39
9 SNP	0	5	7	16	28	3.39
16 RNA-Seq	0	2	14	12	28	3.36
47 Metabolic Pathway	1	5	5	17	28	3.36
5 Motif	0	3	12	13	28	3.36
12 Proteomics	0	3	12	13	28	3.36
23 Ortholog	0	4	12	12	28	3.29
34 Transmembrane Domains	3	3	6	16	28	3.25
49 Substrate	4	3	3	18	28	3.25
2 Annotation	1	4	11	11	27	3.19
3 Parasite Isolate	3	4	5	15	27	3.19
24 Contig and Scaffold	1	4	12	11	28	3.18
11 Allele Frequency	1	6	9	12	28	3.14
14 Microarray	0	6	13	9	28	3.11
10 Non-synonymous	3	6	7	12	28	3
42 Genome Browser	2	8	7	11	28	2.96
28 Signal Peptide	2	7	9	10	28	2.96
35 Epitope	2	9	5	12	28	2.96
31 Mapping	1	8	11	8	28	2.93
4 Genomic Segment	1	5	17	4	27	2.89
44 Paralog	5	4	8	10	27	2.85
45 CDS	6	6	3	13	28	2.82
13 Mass Spec Peptide	2	7	13	6	28	2.82
48 Metabolomics	4	6	10	8	28	2.79
20 Histone modification	3	9	9	7	28	2.71
37 Splice Site Junction	2	11	9	6	28	2.68
55 Ontology	4	10	6	8	28	2.64
54 Metadata	4	9	9	6	28	2.61
50 KEGG Pathway	7	6	6	9	28	2.61
22 Synteny	7	8	4	9	28	2.54
25 GO Terms/Numbers	6	9	7	6	28	2.46
15 Expression Value	5	11	6	6	28	2.46
7 EST	6	7	12	3	28	2.43
46 PubMed Compound	8	7	9	4	28	2.32
19 ChIP-Seq	6	11	7	4	28	2.32
56 ncRNA	10	7	6	5	28	2.21
52 Protein Microarray	7	12	7	2	28	2.14
30 Genomic colocation	9	8	9	2	28	2.14
51 ChIP-chip	10	8	7	3	28	2.11
53 Antibody Microarray	10	9	6	3	28	2.07
26 GO Term Enrichment	12	8	4	4	28	2
27 EC Number	15	2	7	4	28	2
38 Comparative genomic hybridization	11	8	8	1	28	1.96
43 OrthoMCL	16	7	1	4	28	1.75
21 Nucleosome Mapping	16	6	4	2	28	1.71
40 GFF	18	4	2	4	28	1.71
41 BAM	19	4	1	4	28	1.64
18 eQTL	20	6	0	2	28	1.43
57 Polysomal Profiling	20	5	3	0	28	1.39