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**1. Which professional category best describes you? Please select only one.**

- Bioinformatics professional
  Life Science Researcher  
 Educator
  Student  
 Healthcare professional
  Technician  
 Librarian / Information Specialist
  Computer Scientist / Software Developer  
 Other (please specify)

**2. Please pick one category that best describes your organization.**

- College or University
  Research Institute  
 Commercial / Industry
  Government  
 Non-profit Organization
  Hospital/health care organization  
 Other (please specify)

**3. How frequently do you use NCBI's Identical Protein Groups (IPG)?**

- Daily  
 Weekly  
 Monthly  
 Yearly  
 This is my first visit

**4. Do you want to know the source of the protein names in an IPG?**

- Yes  
 No

**5. What was your primary motivation for visiting IPG today?**

- Determine the taxonomic distribution for a set of identical proteins  
 Find an exemplar sequence for a set of identical proteins (e.g. "best" accession)  
 Find all sequence records representing a set of identical proteins  
 Find the corresponding nucleotide coordinates for a protein's coding sequence (CDS) in one or more organisms  
 Identify genome assemblies on which this protein is annotated  
 Find the best name for a protein  
 Look at a graphical display of a protein sequence  
 Find relevant literature  
 View a protein in its genomic context(s)  
 Evaluate the domain structure of a protein  
 Other (please specify)

**6. Did you find what you were looking for?**

- Yes  
 No

**7. Considering your *primary* motivation for visiting IPG today, how easy was it to find what you were looking for on the IPG web page ?**

- Very easy  
 Easy  
 Neither easy nor difficult  
 Difficult  
 Very difficult

**8. Considering your *primary* motivation for visiting IPG today, how easy was it to download/retrieve the data you were looking for?**

- Very easy  
 Easy  
 Neither easy nor difficult  
 Difficult  
 Very difficult  
 Not Applicable

**9. Considering your *primary* motivation for visiting IPG today, will you also use the NCBI Protein database to accomplish some part or all of this task?**

- Yes  
 No, IPG is sufficient  
 I'm not familiar with NCBI Protein

**10. Which of the following tasks do you perform with data from IPG? Please check all that apply.**

- Search for related proteins or IPGs (e.g. using BLAST or other sequence similarity search tool)  
 Perform a multiple alignment to other proteins (e.g. using COBALT, MAFFT, muscle, kalign, etc.)  
 Generate a taxonomic tree of organisms in which this protein has been identified  
 Evaluate the gene neighborhood in one or more taxa  
 Look for possible taxonomic mis-classifications (i.e. protein/taxonomic mismatch)  
 Other (please specify)

**11. In addition to the website, do you retrieve data from IPG using any of the following mechanisms?**

- Yes, using the E-Utilities / EDirect  
 Yes, using other computational approaches (e.g. custom scripts)  
 No, I only use the website

**12. Please outline the workflow that you use with the data you are seeking today at NCBI, including other NCBI and non-NCBI resources used.**

**13. If there was one thing you could change about the IPG pages, what would it be?**

**14. How likely is it that you would recommend the IPG pages to a friend or colleague?**

Not at all likely

Extremely likely

0	1	2	3	4	5	6	7	8	9	10
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**We welcome your input! Please contact us at [suggest@ncbi.nlm.nih.gov](mailto:suggest@ncbi.nlm.nih.gov) if you would like to provide additional feedback about the IPG resource.**

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Done

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