

NCBI PROTEIN SEQUENCE INFORMATION FEEDBACK

If you have used our site before to retrieve protein information, please take a few moments to complete our survey. Your feedback will help us make the site more valuable. Your responses are strictly confidential.

1. As specifically as possible, what protein information are you looking for?

2. How many times a month do you visit this Web site for protein information?

- 20 or more
- 10 – 20
- 5 – 10
- 1 – 5

3. On a scale of 1 to 5, how would you rate your overall satisfaction with the protein information on this Web site?

- 5 – very satisfied
- 4 – satisfied
- 3 – neutral
- 2 – not very satisfied
- 1 – completely dissatisfied

4. Based on the categories of protein information available on this Web site, which is most useful to you? (Please rank order categories A through E using numbers 1-5, where 1 is most useful and important, 2 is less useful and important, etc.)

- CATEGORY A:**
Classification into protein families based on sequence similarity and function
(*e.g., SAM-dependent methyltransferases, Cytochrome P450, etc.*)
- CATEGORY B:**
Identification of protein domain architecture, with links to alignments and structures
(*e.g., SH3 domain, RAS domain, etc.*)
- CATEGORY C:**
Features derived from computational predictions
(*e.g., Transmembrane regions, Coiled-coil regions, etc.*)

- **CATEGORY D:**
Protein function based on experimental evidence from published literature
(e.g., *Integrin receptor signal signalling pathway. Nat Genet. 1993 Oct; 5(2):51-7.*)

- **CATEGORY E:**
Annotation of functional sites, processing pathways, and degradation sites based on evidence from published literature
(e.g., *Position of modified residues and/or active sites, Proteasome cleavage sites, etc.*)

5. Do you have additional comments or suggestions regarding NCBI's protein information Web pages?

Thank you for your assistance. We appreciate your time and ideas.

Note: Public reporting burden for this collection of information is estimated to average 5 minutes per response, including the time for reviewing instructions, searching existing data sources, gathering and maintaining the data needed, and completing and reviewing the collection of information. Potential respondents must be informed of the fact that an agency may not conduct or sponsor, and a person is not required to respond to, a collection of information unless it displays a currently valid OMB control number. Send comments regarding this burden estimate or any other aspect of this collection of information, including suggestions for reducing this burden, to: NIH, Project Clearance Branch, 6705 Rockledge Drive, MSC 7974, Bethesda, MD 20892-7974, ATTN: PRA (0925-0486). Do not return the completed form to this address.

Questions or Comments?
Write to the NCBI Service Desk