Genew

Human Gene Nomenclature Database

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What is Genew

- Genew is the only resource that provides data for all human genes which have approved symbols.
- It's managed by HUGO Gene Nomenclature Committee (HGNC)
- o First published in 1979
- An estimated total of 26,000-40,000 genes

Implementation

- o Microsoft Access 97 relational DBMS
- o 13 tables, over 170 fields
- search engine based on a Perl frontend querying a PostgreSQL database, data in text files exported from off-line database http://www.gene.ucl.ac.uk/cgi-bin/nomenclature/searchgenes.pl

Data flow

Source of data

- Data submissions (authors, journals, literature review, chromosome sequencing projects, gene families, homologs from other species)
- Other database files
 - o LocusLink
 - o SWISS-PROT
 - o The Genome Database GDB
 - Mouse Genome Database MGD (not exported)

Data flow

Output

- Online Genew Search Engine
 - o Search: gene symbol, gene name, aliases
 - Retrieved: gene symbol, gene name, aliases, cytogenetic location, OMIM (Online Mendelian Inheritance in Man) number, PMID (PubMed ID)
- Online text files

Other information

- LBLAST
- Confidentiality
- LocusLink updates
- Sequence analysis
- o http://www.gene.ucl.ac.uk/nomenclature/guidelines.html

Rules

- o Each gene record
 - Has a unique HGNC ID
 - Has a status...

Rules

- Status
 - Pending
 - Approved
 - Reserved
 - Symbol Withdrawn
 - Entry Withdrawn
 - Approved Non-Human

Rules

- Each gene symbol
 - Must be unique
 - Is a short-form representation (or abbreviation) of the descriptive gene name
 - should only contain Latin letters and Arabic numerals
 - should not contain punctuation
 - should not end in "G" for gene
 - not contain any reference to species, for example "H/h" for human









