



INTELLECTUAL PROPERTY
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LESSONS LEARNED FROM THE ON-THE-JOB TRAINING IN SEQUENCE SEARCH AT THE SWISS PATENT OFFICE

Training Program



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Swiss Federal Institute of Intellectual Property (IPI)
(16-20 April 2018)

Funds-in-Trust Australia 2017-2018

Cooperation on Examination and Training Section, PCT
International Cooperation Division, WIPO

Objectives



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Develop competencies of the trainees for examining inventions involving protein and genetic sequences

Search for sequences using various public and commercial databases

Deal with technology-specific issues related to claim wording, or novelty and obviousness

Identify the source of the genetic material

Expected outcomes



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Develop competencies of the trainees for examining inventions involving protein and genetic sequences

Increase proficiency in using commercial and public databases

Accurately evaluate search requirements of any given claim so as to decide which type of prior arts to search for

Learn efficient search techniques and best practices that ensures thorough search

Assess novelty and inventiveness of sequence claims using the search products obtained

Pre-Training



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Gather sample cases



Training Method



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For each case:



Databases used



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Commercial

- SequenceBase (USGENE, WOGENE, GENESEQ)
- GenomeQuest
 - DNA, RNA & protein sequence search
- PatBase
- STNext
 - Full-text patent search

Open-access

- NCBI BLAST
 - Comparing sequences and calculating homology
- ENSEMBLE
 - Gene search and computing multiple alignments

Databases used



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<ul style="list-style-type: none">• Registry	<ul style="list-style-type: none">• Full-text search; search for sequences containing uncommon amino acids and metal complexes	<ul style="list-style-type: none">• Commercial
<ul style="list-style-type: none">• GQ-PAT (Collection)	<ul style="list-style-type: none">• Sequence search results include patent information as well as biological information	<ul style="list-style-type: none">• Commercial
<ul style="list-style-type: none">• PatSeq Finder	<ul style="list-style-type: none">• sequence similarity search tool	<ul style="list-style-type: none">• Open-access
<ul style="list-style-type: none">• GeneCards	<ul style="list-style-type: none">• Human gene sequence search tool that includes genomic, transcriptomic, proteomic, genetic, clinical and functional information	<ul style="list-style-type: none">• Open-access
<ul style="list-style-type: none">• UniProt	<ul style="list-style-type: none">• Protein sequences search which also gives the functional information	<ul style="list-style-type: none">• Open-access
<ul style="list-style-type: none">• iHOP	<ul style="list-style-type: none">• Searching literatures containing the specific gene name or the accession number	<ul style="list-style-type: none">• Open-access
<ul style="list-style-type: none">• EBI	<ul style="list-style-type: none">• Nucleotide sequence search, including assembly and functional information	<ul style="list-style-type: none">• Open-access
<ul style="list-style-type: none">• NCBI OMIM	<ul style="list-style-type: none">• Human gene search, with phenotype information and genetic disorders	<ul style="list-style-type: none">• Open-access
<ul style="list-style-type: none">• NCBI MeSH	<ul style="list-style-type: none">• Finding alternate words or synonyms for diseases and other life science-related terms	<ul style="list-style-type: none">• Open-access
<ul style="list-style-type: none">• NCBI Pubmed	<ul style="list-style-type: none">• Medical terminology search	<ul style="list-style-type: none">• Open-access

Databases used



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<ul style="list-style-type: none">• <u>Genesys</u>	<ul style="list-style-type: none">• Plant genetic resources search	<ul style="list-style-type: none">• Open-access
<ul style="list-style-type: none">• INN	<ul style="list-style-type: none">• Identification of pharmaceutical substances or active pharmaceutical ingredients (database of generic names)	<ul style="list-style-type: none">• Open-access
<ul style="list-style-type: none">• FDA	<ul style="list-style-type: none">• General information on drugs and other life science-related queries	<ul style="list-style-type: none">• Open-access
<ul style="list-style-type: none">• Clinical Trial	<ul style="list-style-type: none">• Searching drugs and their clinical trial stage, registration status and summary results	<ul style="list-style-type: none">• Open-access
<ul style="list-style-type: none">• Google Scholar	<ul style="list-style-type: none">• General search on academic literature	<ul style="list-style-type: none">• Open-access
<ul style="list-style-type: none">• Wikipedia	<ul style="list-style-type: none">• General/ background information search	<ul style="list-style-type: none">• Open-access

Outcomes



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- ❑ Wider understanding with regard the analysis and search of each case
- ❑ Optimized use of the databases and online platforms for sequence search
- ❑ “Manipulate” the features of each database to effect an inter-platform search
- ❑ Various online platforms are introduced for auxiliary search



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IPOPHL's Best Practices for Searching Inventions Related to Sequences

Search Method/Databases used



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- Keyword search
 - Google, Google Patents, Google Scholar



3-person search team

- Full-text search
 - EpoqueNet, Web of Science
 - Open-access patent databases

- Sequence search
 - STN, PatSeq, BLAST

- IPAS

- International work products



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THANK YOU