

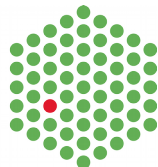
GIFTs Update

Ensembl Application Team Away Day Apr 2019

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EMBL-EBI



e!Ensembl

GIFTs

Genome Integration with FuncTion and Sequence

- Connect genome/proteome worlds for max impact
- Map from genome coords to protein residues
- Main obstacle: data synchronisation
- Solution: develop common infrastructure
 - track entities btw resources
 - synchronise content
 - share curational tasks
 - support mapping btw entities at nucleotide/AA precision

Goals

- Support:
 - DNA features ↔ proteins projections
 - common curation/analysis shared across resources
- System for agreed reference gene/transcript/protein sets from e! and UniProt
- Resource for genome integration with function and sequence

Deliverables

- GIFTs database (e!/Uniprot)
 - known genes, transcripts, proteins and their relationships
 - all previous transcript versions stored with relationships to genes
 - drives the mapping of UniProtKB sequences to genomes
 - support xrefs management and share curation interface
- Curation interface
 - view mappings
 - validate/edit automatic mappings

Deliverables (2)

- Protein-Genome Mapping Pipeline
 - DB remains in synch with e! and UniProt releases
 - automatically align proteins to reference genomes
 - store matches
 - <100% matches trigger further analysis and/or resolution through curation interface
- GIFTs Resource
 - present DB mappings as simple EMBL-EBI webpage
 - download of mapping files
 - project docs and use cases

Milestones

- Year 1 (2017)
 - DB schema designed, implemented, populated
 - API prototype to support protein-genome mapping pipeline
 - pipeline developed (use API and GIFTs DB)
 - commence work on curation interface
- Year 2 (2018)
 - develop curation interface (use API and GIFTs DB)
 - develop GIFTs website
 - develop automated method to query/combine e!/UniProt annotation → agreed reference genome/protein sets

Timeline

- May '18 (6 months to go!)
 - prototype schema designed
 - DB test server instance
 - *API backend skeleton (Django), no endpoints*
 - genebuild pipeline initial implementation (Perl)
 - front end work starts, mockup using precompiled JSON
- June '18
 - Web VM/virtual users request
 - several schema redesigns
 - back end API
 - prototype endpoint for bulk load gene/transcripts
 - endpoints to support front end build

Timeline

- July '18
 - new schema, more API endpoints (pipeline/front end)
 - loaded human/mouse
 - virtual users, no VMs yet!
 - plan B: OpenStack (can't reach DB, mirror)
 - 1st prototype interface
- August '18
 - several schema/front end API updates
 - 1st demo curation interface
 - finally, the VMs!
- September '18
 - several schema/front end API updates
 - first instances running dev/prod (HX/HH)
 - semi-automated deployment process
 - production DB VM request (HX/HH)
 - Matt leaves!

Timeline

- October '18
 - production DB VMs ready (HX)
 - pipeline tests alignment API
 - feedback on curation tool
- November '18
 - new requirement: unmapped entries
 - schema/API updates
 - authentication set up/tests
 - dev nightly builds
 - DB server monitoring/tuning

Timeline

- December '18
 - debugging web-prod URL mapping, Apache rewrite rules
 - tuning Apache (multi-process)
 - configurable deployment
 - pre-release out: <http://www-test.ebi.ac.uk/gifts>
 - bug-fixes
 - collaborative workflow simulation starts
- January '19
 - curators feedback
 - search issues (data/speed), bug-fixes
 - schema/API improvements
 - new requirement: mapped with unmapped

Timeline

- February '19
 - GIFTs demo at EBI service retreat
 - release/workflow process (Beth)
 - schema/API/front end updates
 - mapped with unmapped
 - search speed improvements
 - Load Balancer problems
 - VM kernel upgrade: build process disruption
- March '19
 - GIFTs presented to UniProt collaborators
 - feature requests:
 - review unmapped
 - changes to existing endpoints
 - REST based e! pipeline time out:
 - REST polling (task queue manager)
 - Mark steps in

To Be Completed

- API docs
- REST polling
 - Celery/Redis deployed
- review unmapped (front end)