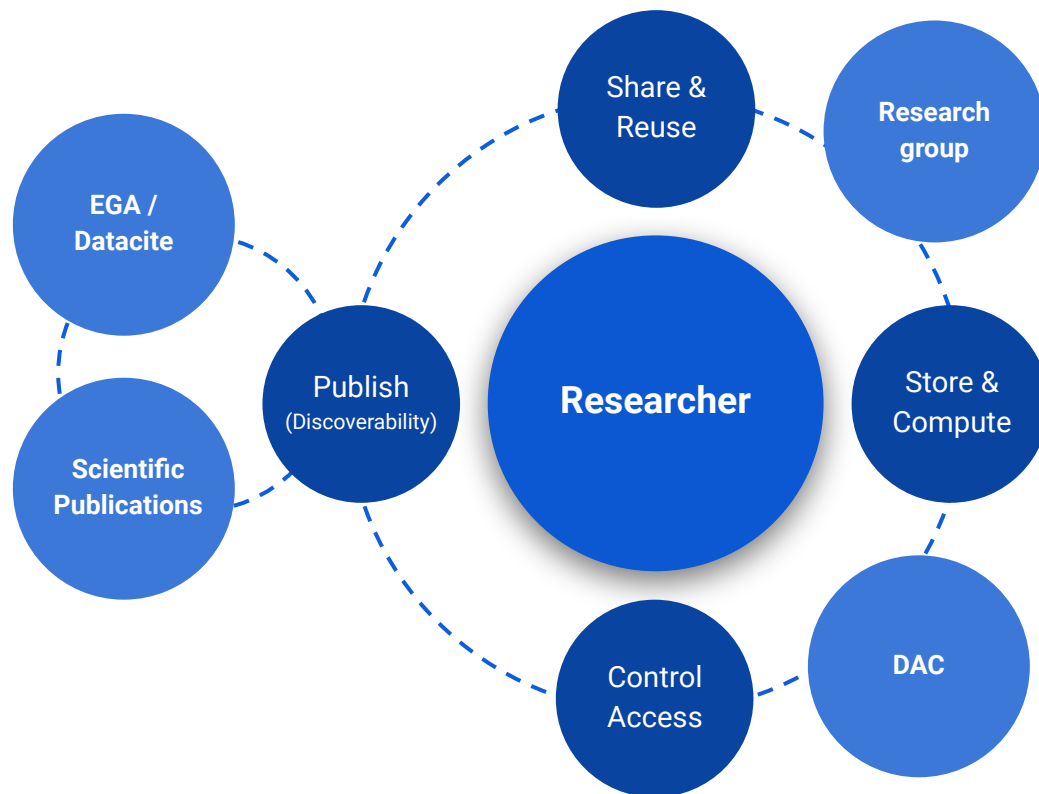


# Developing Technologies And Standards for Enabling Sensitive Data Archiving, Sharing and Reuse

Stefan Negru

CSC – IT CENTER FOR SCIENCE

# Understanding the landscape



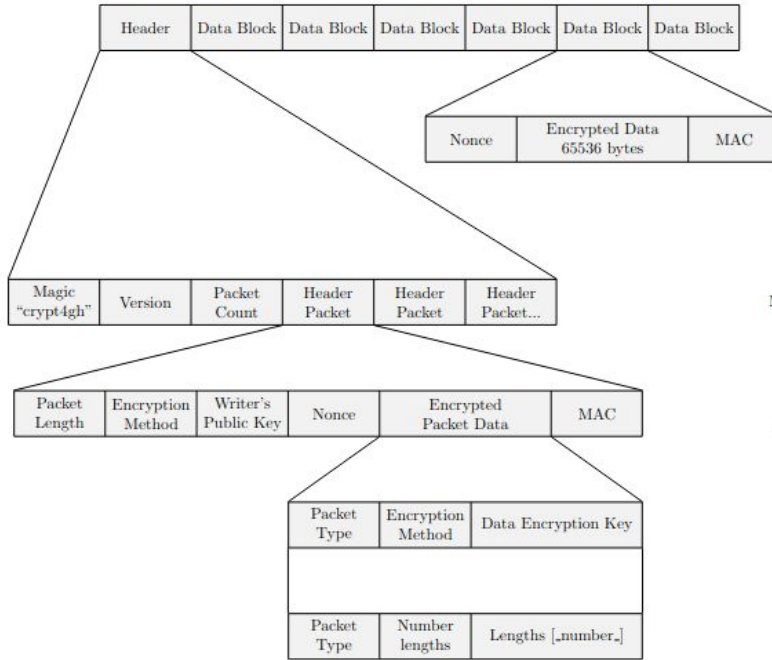
# Some principles to consider ...

- DARE (Data at Rest Encryption)
  - Facilitating data re-encryption
- FAIR (Findable Accessible Interoperable and Reusable)
  - Discoverable by other researchers
  - Reusable by other researchers
- Controlled Access to data - there is a DAC that controls access
  - Specify and encode permissions for the data, what can be used for and till when

# A few APIs/Standards

API/Standards Name	API/Standard Purpose	URL to Spec
htsget	A protocol for secure, efficient and reliable access to sequencing read and variation data	<a href="http://samtools.github.io/hts-specs/htsget.html">http://samtools.github.io/hts-specs/htsget.html</a>
DUO	Allows users to semantically tag genomic datasets with usage restrictions, allowing them to become automatically discoverable based on a health, clinical, or biomedical researcher's authorization level or intended use.	<a href="https://github.com/EBISPOT/DUO">https://github.com/EBISPOT/DUO</a>
Refget	Refget enables access to reference sequences using an identifier derived from the sequence itself.	<a href="http://samtools.github.io/hts-specs/refget.html">http://samtools.github.io/hts-specs/refget.html</a>
Crypt4GH	A file container specification enabling direct byte-level compatible random access to encrypted genetic data stored in community standards such as SAM/BAM/CRAM/VCF/BCF.	<a href="http://samtools.github.io/hts-specs/crypt4gh.pdf">http://samtools.github.io/hts-specs/crypt4gh.pdf</a>
Beacon	Discover genomic variants, individuals, and individuals	<a href="https://github.com/ga4gh-beacon/beacon-v2/">https://github.com/ga4gh-beacon/beacon-v2/</a>
Phenopackets	It merges the existing GA4GH metadata-schemas work with a more focused model from the phenopacket-reference-implementation.	<a href="https://phenopacket-schema.readthedocs.io/en/latest/">https://phenopacket-schema.readthedocs.io/en/latest/</a>
GA4GH Passports	GA4GH Passport specification aims to support data access policies within current and evolving data access governance systems.	<a href="https://github.com/ga4gh-duri/ga4gh-duri.github.io/blob/master/researcher_ids/ga4gh_passport_v1.md">https://github.com/ga4gh-duri/ga4gh-duri.github.io/blob/master/researcher_ids/ga4gh_passport_v1.md</a>
WES API	Workflow Execution Service API describes a standard programmatic way to run and manage workflows. Having this standard API supported by multiple execution engines will let people run the same workflow using various execution platforms running on various clouds/environments.	<a href="https://github.com/ga4gh/workflow-execution-service-schemas">https://github.com/ga4gh/workflow-execution-service-schemas</a>

# Encryption - Crypt4GH \*



**crypt4gh File**  
 Reader-specific encrypted header  
 Encrypted data in blocks

**Data Block**  
 Decryption key  $K_{data}$  is stored in  
 Data Encryption Parameters header packet

**File Header**  
 Magic number, version and packet count are unencrypted  
 Header packets individually encrypted for reader

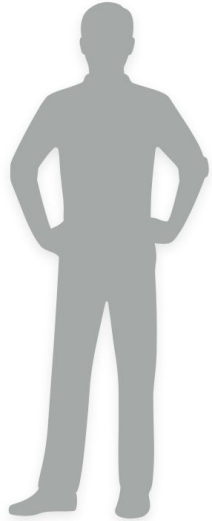
**Header Packet**  
 Packet data encrypted using key  $K_{shared}$  derived from  
 writer's public key ( $K_{pub}$ ) and reader's secret key ( $K_{sr}$ )

**Data Encryption Packet (plain-text)**  
 Stores  $K_{data}$

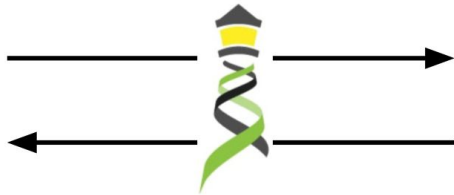
**Data Edit List Packet (plain-text)**  
 List of byte counts to alternately  
 exclude and include in output

\* specification & figure provided by  
<http://samtools.github.io/hts-specs/crypt4gh.pdf>

# Discovery - Beacon \*



“Do you have a ‘C’ at  
chromosome 13 at  
position 32,936,732?”



“Yes” (or “no”)

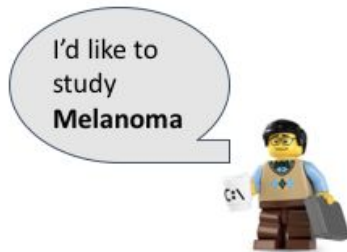
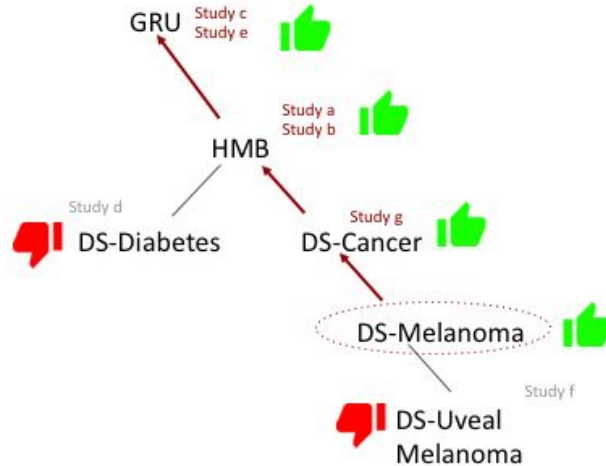


\* Figure provided by: <http://docs.genomebeacons.org/what-is-beacon-v2/>

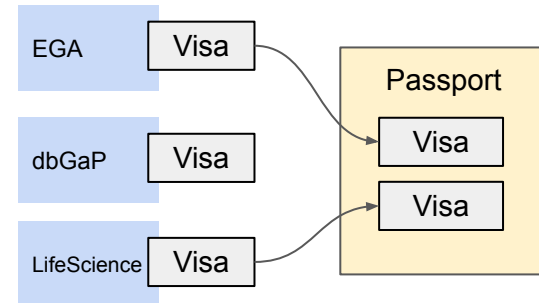
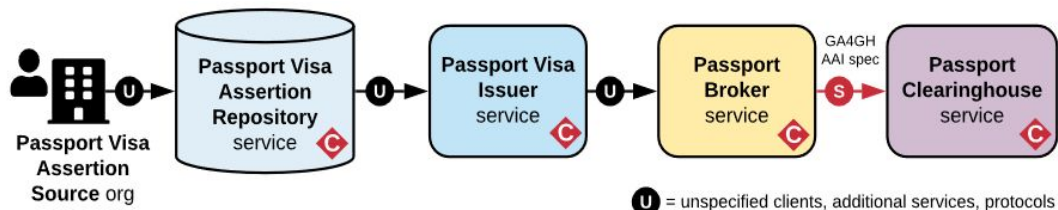
# Controlled Access - Data Use Ontology

Data Repository

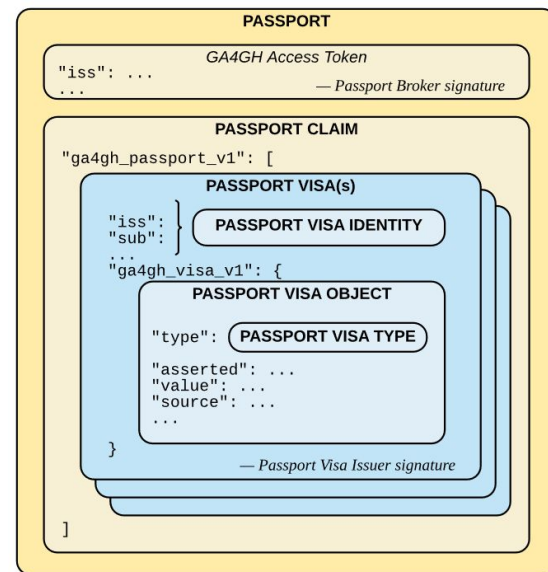
Dataset	DU restriction
Study a	HMB
Study b	HMB
Study c	GRU
Study d	DS- Diabetes
Study e	GRU
Study f	DS- Uveal Melanoma
Study g	Cancer



# Controlled Access - GA4GH Passport in brief

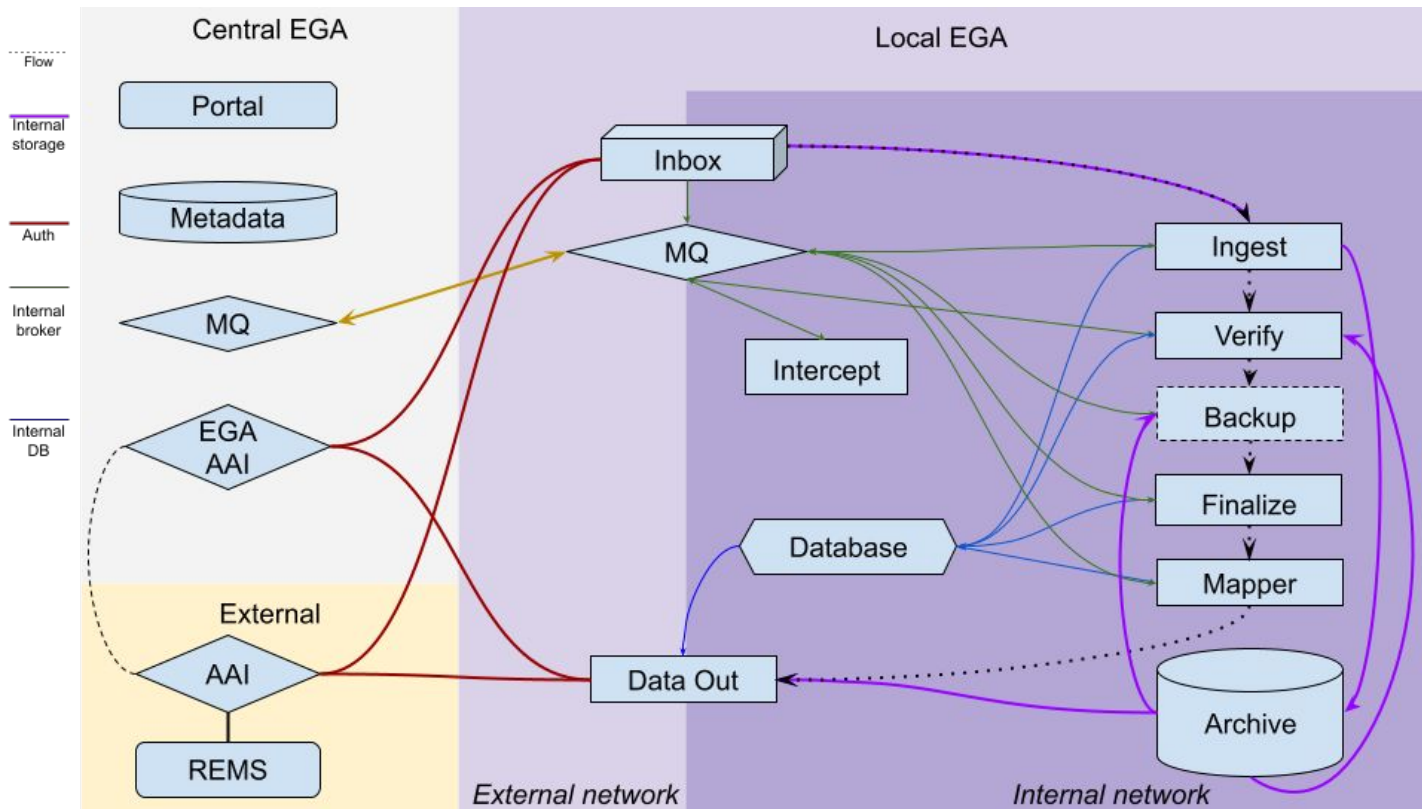


Visa type	Description
<b>AffiliationAndRole</b>	User's role within their institution - e.g. <a href="mailto:faculty@cam.ac.uk">faculty@cam.ac.uk</a> (eduPersonAffiliation)
<b>AcceptedTermsAndPolicies</b>	Acknowledged terms, policies, and conditions - e.g. attestations for registered access
<b>ResearcherStatus</b>	Bona fide researcher status - e.g. for registered access
<b>ControlledAccessGrants</b>	Permission to controlled access datasets - e.g. EGA, dbGaP
<b>LinkedIdentities</b>	Mapping of user identities - e.g. <a href="mailto:user@lifescience.org">user@lifescience.org</a> equal to <a href="mailto:username@csc.fi">username@csc.fi</a>

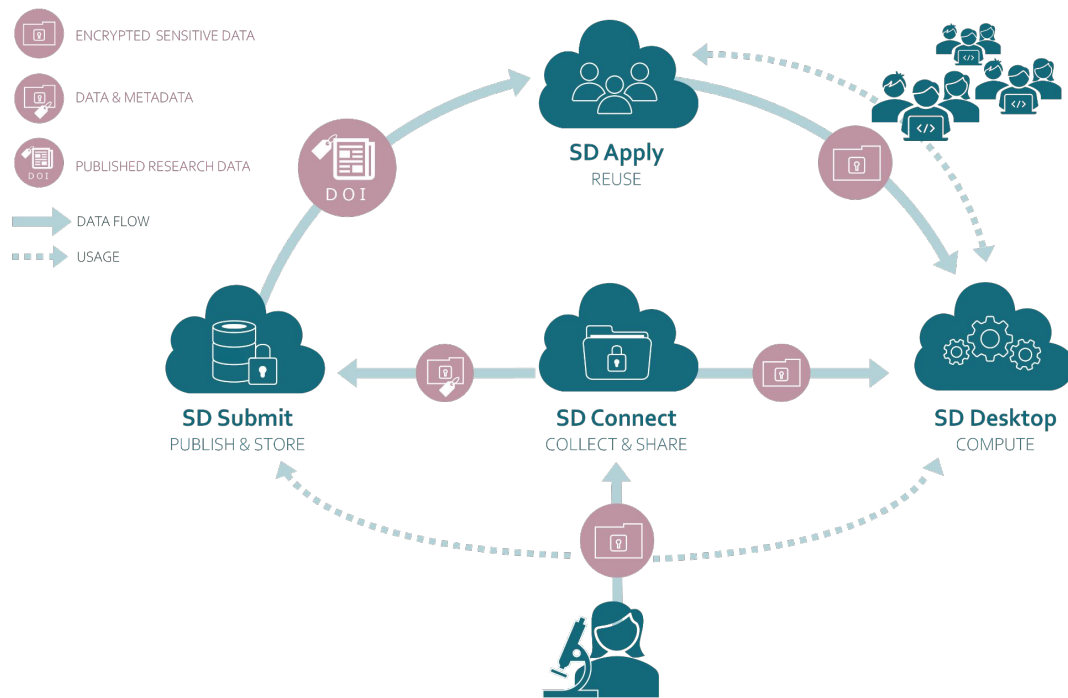




# Sensitive Data Archive - NelC Heilsa solution



# CSC - Sensitive Data Services for Research \*



\* more information at: <https://research.csc.fi/sensitive-data-services-for-research>

# Additional Resources

- Crypt4GH libraries
  - Python <https://github.com/EGA-archive/crypt4gh>
  - Go <https://github.com/neicnordic/crypt4gh>
  - Rust <https://github.com/EGA-archive/crypt4gh-rust>
  - C <https://github.com/silverdaz/crypt4gh>
  - Java <https://github.com/uio-bmi/crypt4gh>
  - Samtools <https://github.com/samtools/htslib-crypt4gh>
- NeIC Nordic:
  - <https://github.com/topics/neic-sda>
- Beacon
  - <https://github.com/CSCfi/beacon-python>
  - <https://github.com/CSCfi/beacon-network>
  - <https://github.com/EGA-archive/beacon2-ri-api>

Thank you!