

SweFreq NG

<p>Product owner</p> <p>Johan Rung, SciLifeLab Data Office</p>	<p>Project lead</p> <p>Niclas J (Team: Johan V, Andreas K)</p>
<p>Goals / Success metrics</p> <ul style="list-style-type: none"> Establish a web resource that <ul style="list-style-type: none"> Allows download of several data set allele frequencies (after user identification and registration) Allows separate description of, and terms of use for, each dataset Provide identification by ELIXIR ID Allows browsing the frequency data in a graphical browser, both for GRCh37 and GRCh38 Provides GA4GH Beacon access to the frequency data (dependent on ELIXIR Beacon project) Is compatible with the SciLifeLab graphical profile Provide documentation on how datasets should be formatted 	<p>Out of scope</p> <ul style="list-style-type: none"> Allowing controlled access to the individual data Automated generation of frequency datasets from raw data Combination of more than one dataset
<p>Key stakeholders</p> <div> <div> <p>External</p> <ul style="list-style-type: none"> Sverker Lundin, SciLifeLab Nat. proj. coordination - C Per Kraulis, Data Office - C Hanna Kultima, Data Office - C Adam Ameer, SweGen - I Patrick Sullivan, Exac data - I </div> <div> <p>Internal</p> <ul style="list-style-type: none"> NBIS managers - I </div> </div>	

Related projects/activities <ul style="list-style-type: none"> ELIXIR Beacon project 	Estimated resources 1-2 PM ?? (Johan R is looking at getting funding from SciLifeLab for future maintenance)
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Data <i>Background/observations that has caused us to want to start a project/activity</i>	Insight <i>Insights gathered from the Data</i>	Belief <i>Beliefs grounded in the Insights</i>	Bet <i>Suggestions for addressing the Beliefs</i>
<p>SweFreq has been established as a platform to publish variant frequency data from human WGS datasets.</p> <p>The SciLifeLab national projects, as well as other Swedish research projects, will produce human WGS datasets from which they want to publish variant frequency data.</p>	<p>The SciLifeLab national programmes data sets should be made available to the research community.</p> <p>It is currently outside of the capability of the reference population project or Scilifelab centrally to provide a stable access to the allele frequency data.</p>	<p>As NBIS promotes Open Access, and want to maximize the impact of the resources spent on national platforms, a web resource to allow access to the reference population allele frequency data should be established.</p>	<p>Modify the established SweFreq web resource so that it allows access to several variant frequency data sets produced by the SciLifeLab national programmes, and other Swedish research projects.</p>