

Context project:	Programming Life				
Group:	3				
User Story	Task	Assigned To	Estimated effort (1-5) 5 = highest	Priority (1-5) 1 = highest	Actual effort (in hours)
<i>As a stakeholder, I need to have assurances of the build quality of the software</i>	Create a draft for the final report	Boris (All)	5	1	
	Finish the Emergent Architecture Design	Tom (All)	3	1	
	Fix MVC structure	Boris/Kasper	3	1	
	Fix checkstyle with intelij tool	Mathieu	4	3	
	Specify architecture for SIG	Sam	3	4	
<i>As a user, I want to view metadata of different genomes, so I can draw conclusions from my data</i>	Implement parsing of new annotation format (gff)	Tom	3	3	
	Embed all delivered metadata in nodes	Tom	3	3	
	Gene annotation is visual displayed	Boris	4	1	
	Mouse-over on a node can display all the information	Mathieu	3	2	
<i>As a user, I want the application to be stable, so I can work with it efficiently</i>	The application should not crash when loading files when the view is already loaded.	Kasper	3	1	
	The application should not crash when loading 671 strains	Mathieu	5	2	
<i>As a user, I want to know how to use the application, so I can work with it efficiently</i>	Pressing F1 gives a help menu	Kasper	1	3	
	Add the space bar binding to help	Kasper	1	5	
<i>As a developer, I want to know how users use the application, so I can improve the design</i>	Perform a user test with actual users	Tom (All)	5	1	
<i>As a user, I want to see already known mutation data that I can use to derive my own conclusions</i>	Known mutations parser	Tom	3	4	
	Show known mutations	Boris/Tom	3	2	
<i>As a user, I want to see what the application can do before I start using it, so I know what to expect for the application</i>	Make a video for the customer	Boris/Sam	2	1	

<i>As a user, I want to have access to all known information, so I can make an informed decision about the data</i>	All edges should be displayed	Sam	4	3	
<i>As a user, I want to know why a feature isn't working when I try to use it, so I know what to do to fix the error</i>	Display notification when spacebar is pressed and phylo tree is not loaded	Kasper	2	3	
<i>As a user, I want to move quickly through the data, so I can use the application efficiently</i>	Increase the zoom ratio	Kasper	1	2	
<i>As a user, I want to have a clear visual representation of the data, so I can use it to draw conclusions</i>	Fix spacing of nodes	Sam	5	2	
	Fix interestingness	Sam	4	2	
	Visually encode the size of node labels	Mathieu	2	5	
	Fix the empty edges in graph	Sam	3	5	
	Edge weight after filtering	Boris	2	3	
<i>As a user, I want to navigate through the graph, so I can search for interesting positions.</i>	Click on the bar to navigate through the graph	Boris	1	4	
<i>As a user, I want to quickly continue working where I left off, so I can use the application efficiently</i>	Fix recent files menu	Kasper	2	2	
<i>As a user, I want to zoom in on the graph, so I can get a more in-depth view of the graph</i>	Fix scaling factor for zoom	Boris	1	2	
The time estimations are done in the following scale:	1=15-60m	2=1-2h	3=2-4h	4=4-8h	5=8+h
	1	2	3	4	5