Sprint Reflection — *Programming Life Team 1* — Week 4.4 — Sprint 3

Requirement/Story	Task	Actual time spent					Estimated time	Done	Notes
		Chak Shun	Justin	Mark	Marissa	Maarten	hours	y/n	
Enable us to interactively explore a sequence graph representing the genome architecture of multiple strains.	Orientate on visual aspects of nodes.	3,0					3	yes	
	Implement the visual aspects.	5,0					7	no	To be refined, basics are done.
	Implement visual interaction via the user interface.					1,0	4	no	
As a user, I want to be able to interact with the data.	Orientate on different layouts of the user interface.				2,0	2,0	3	yes	
	Improve the existing basic user interface.					10,0	5	yes	
Provide semantic zooming to enable useful visual interpretation at various zoom levels from whole-genome to individual mutations.	Refine semantic zoom level for amino acids.		6,0	6,0			5	no	Postponed since the customer meeting
	Implement semantic zoom level for joining small point mutations.		11,0		7,0		7	no	To be refined
As a user, after the first loading of the data, I want to be able to load the data more quickly.	Implement a new file format for storing importation generated graph data.			4,0			7	no	It is done, but since we have chosen to use a different data model it has to be changed again. It will be "yes" by tomorrow.