Sprint plan # 2

Contextproject: Programming life

Group: 2

User Story	Task	Task Assigned To	Estimated Effort per Task (1-5)
As a team we want to deliver our deliverables.	 deliver sprintplan3.pdf deliver sprintreflection2.pdf deliver productplanning.pdf deliver final productvision.pdf 	1. Boot 2. Boot 3. Nieuwdorp 4. Boot	1. 1 2. 1 3. 1 4. 1
As a team we want to create and maintain our documents.	Review and redact the product vision based on received feedback. Review and redact the arch. design based on received feedback. Create a product planning	1. Nieuwdorp 2. Hommes 3. Nieuwdorp	1. 4 2. 3 3. 3
As a user I want to be able to see the genome graph with edges between the nodes.	1. Draw edges between the nodes.	1. Vennik	1. 3
As a user I want to have a clear & efficient view of the genome graph	1. Use an algorithm to filter out the crossed lines. 2. Show % of ATCG per node on high-level view 3. Enlarge the scrollbar for bigger graphs 4. Research if a dropdownmenu is better than the menu used for nodes and edges loading 5. If so, implement a dropdown menu	1. Vennik 2. Nieuwdorp 3. Hommes 4. Hommes 5. Hommes	1. 5 2. 4 3. 3 4. 2 5. 3
As a user I want to see the DNA-sequence that belongs to each node.	Display DNA-sequence on node.	1. Vennik	1. 3

As a user I want to see a detailed view of the phylogenetic tree.	Show the names as nodes Add edges between nodes	1. Boot 2. Boot	1. 2 2. 2
As a user I want to be able to navigate between the phylogenetic tree and the genome graph.	Create a link from the phylogenetic tree to the genome graph Create a link from the genome graph to the phylogenetic tree	1. Boot 2. Boot	1. 2 2. 2
As a user I want to simply load all relevant files in a certain directory.	Create a workspace for data files Create a scanner for the files in the workspace Create a select menu for selecting which node, edge and nwk files to use for displaying	Oolbekkink Oolbekkink Oolbekkink	1. 3 2. 3 3. 3