

# Open Canvas

project : Phinch (Holly Bik)

Project Execution

<b>Problem</b> Difficult to visualize large -Omic datasets  Biological Data Viz tools require computational expertise & lots of manual effort  Lack of interactive data viz tools, bad UX/UIs	<b>Solution</b> <ul style="list-style-type: none"> <li>- Scalable framework</li> <li>- Low activation energy (plug &amp; play)</li> <li>- Emphasize novelty, interactivity</li> <li>- Work with designers on UX/UI</li> </ul>	<b>Unique Value Proposition</b>  Quick and easy framework for sharing/storing large datasets and exporting publication quality graphics  Customized, interactive exploration of -Omic datasets without needing any prior computational training  Exciting educational and outreach tool - teachers and citizen scientists can work with real genomic datasets	
	<b>Key Metrics</b>  Web Traffic/Analytics # of Downloads Publication citations GitHub pull requests Social Media mentions	<b>User Profiles</b> Target audience and early adopters Research Scientists Citizen Scientists Teachers Data Journalists	<b>User Channels</b>  Conference Talks Workshops Twitter/Web Word of Mouth Journal Article
<b>Resources Required</b>  Funding - (existing Sloan Foundation grant) Long-term Funding (new grant proposals) End Users to test and uptake product Data Viz Expertise (Pitch Interactive) Community Contributions to Code		<b>Contributor Profiles</b> Contribution types and ideal contributors Computer Scientists (Academics) Freelance Programmers with biological interest/expertise Postdocs, Grad Students, other research Trainees End Users (bug reports, data viz requests)	<b>Contributor Channels</b>  Github Slack Social Media Academic Journals Hackathons/Sprints

Product

Community

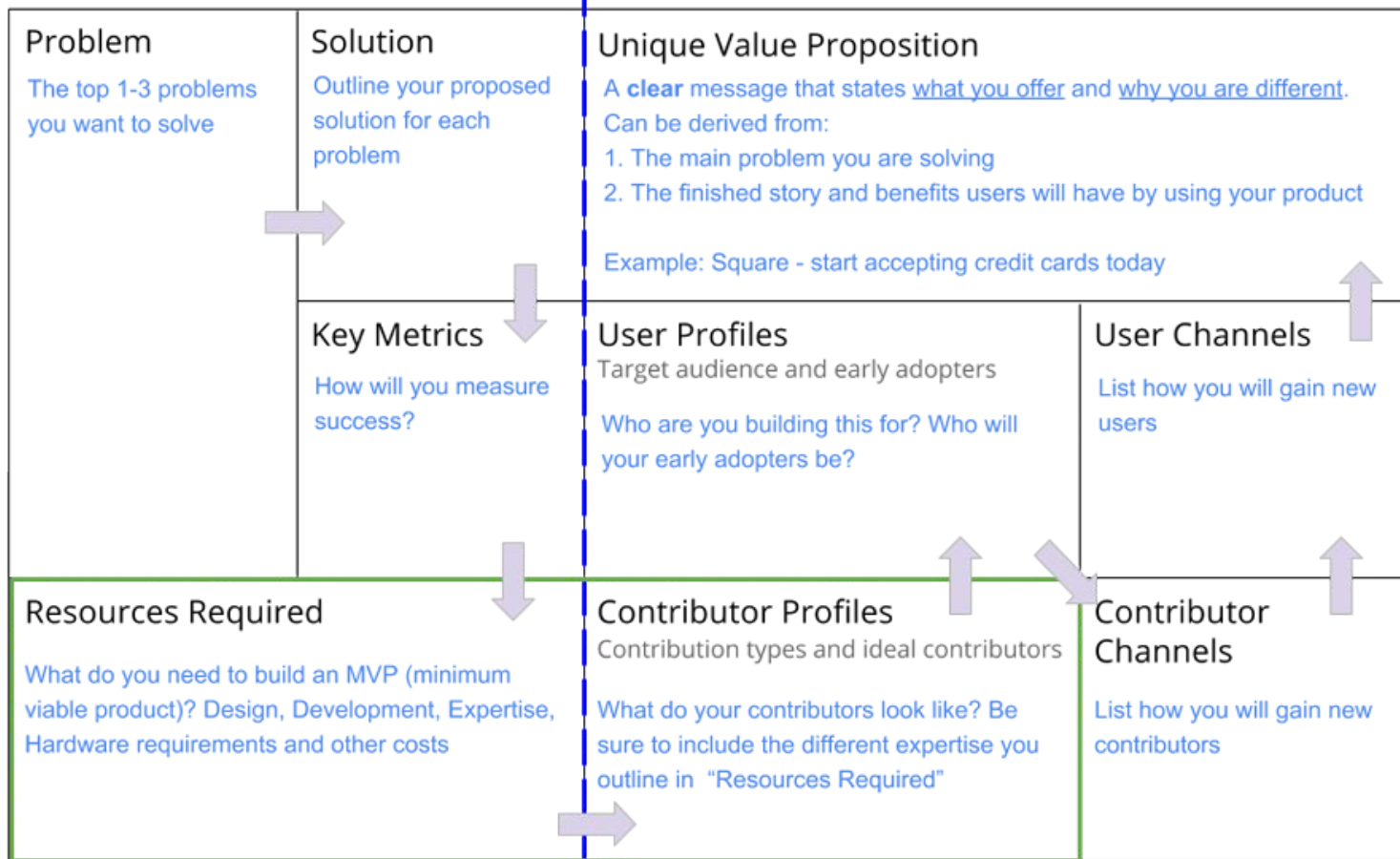
See next slide for instructions!

# Open Canvas

project : Title

Make your  
own canvas:  
Go to **File** >  
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Project Execution



Product

Community

Your users and contributors will likely overlap