| | ONLY | | | |
|---|---------------|--------------------|---------------|---|
| | AUSTIN | | | |
| # Task | WorkSize | Status | Assignments | Comments |
| 3 Limma Analysis | 8 | PARTIALLY COMPLETE | Fahmi, Austin | Complete per specifications; future work needed |
| 11 Enhancements to HeatMap | 3 | PARTIALLY COMPLETE | Adam | |
| 9 Identifying and Fixing Perf bottlenecks | 3 | PARTIALLY COMPLETE | Adam | |
| 18 Launcher (set diff memory profiles) | 4 | 5 Done | Adam | |
| 1 MOG latest version of Java | 3 | 5 Done | Austin, Fahmi | |
| 2 Filter Genes | 7 | 5 Done | Austin, Kamsi | |
| 6 Complex AND and OR queries | 5 | 5 Done | Austin | |
| 7 Extend Logging.Playback | 5 | 5 Done | Austin | |
| 10 Add Violin Plot to MOG | 5 | 5 Done | Kamsi | |
| 15 Display MOG cpu meory usage in taskbar | 1 | 5 Done | Kamsi, Austin | |
| 16 Add an option to add mean/median of values (Gene filter) | 3 | 5 Done | Austin | |
| 19 Setting up CI/CD | 1 | 5 Done | John | |
| 20 Docker build process | 3 | 5 Done | John | |
| 21 Package/bundle app frontend into standard OS executable format | 3 | 5 Done | John | |
| 13 Add help using animations, gifs | 1 | INCOMPLETE | John | Need clarification |
| 12 Extend MOG as general plotting | 0 | INCOMPLETE | - | Need clarification |
| 4 Add unit testing | 8 | INCOMPLETE | Kamsi | |
| 8 Enable hypothesis testing in boxplot charts | 0 | INCOMPLETE | Harsha | Need clarification |
| 22 Web Framework | 8 | INCOMPLETE | * | Need clarification |
| 14 Auto fetch standard gene metadata for organisms | 0 | INCOMPLETE | - | Need clarification |
| 17 Add set/unset plots to MOG | 0 | INCOMPLETE | - | Need clarification |
| | | | | |
| | 0 = unsure or | n/a | | |
| | | | | |
| Task ID | | 0 UNASSIGNED | | |
| | | 1 TODO | | |
| | | 2 Next Up | | |
| | | 3 Just Started | | |
| | | 4 In Progress | | |
| | | 0 PAUSED | | |
| | | 5 Done | | |