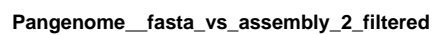
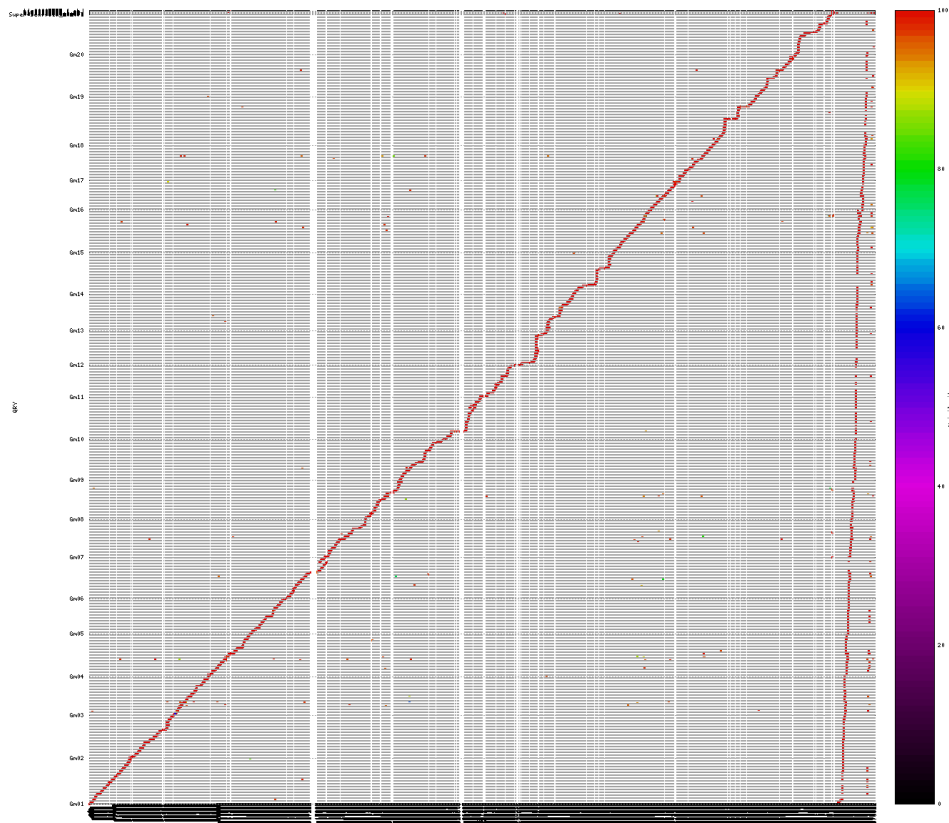


**Pangenome\_\_fasta\_vs\_assembly\_1\_filtered**





FEATURES - color spectrum with similarity scores between the 2 sequences is displayed to the right.

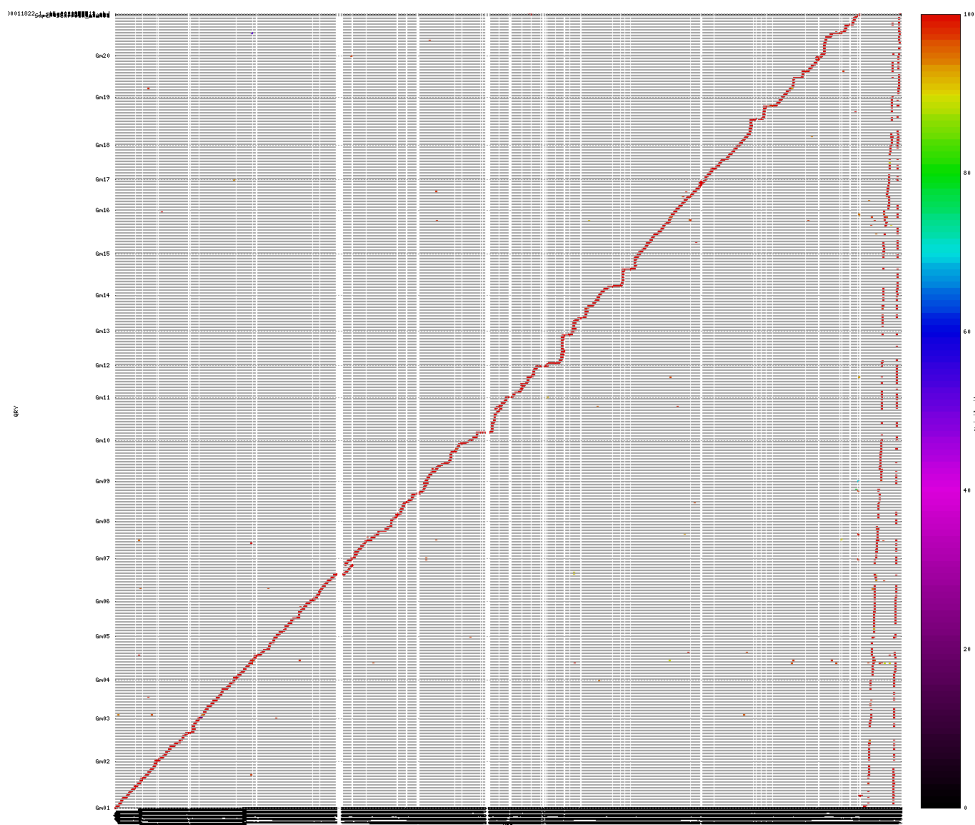
- filtered for one-to-one mapping between the 2 sequences
- y-axis is cleaned up

COMMAND LINE STATEMENTS: `cd <directory>`

`mummerplot -p <prefix name> --large --png --color --filter -y [1,1013761900] -t DOTPLOT`  
`pangenome__fasta_vs_assembly_2_bbmhFilter.delta`

RUNTIME ~ 1min

**Pangenome\_\_fasta\_vs\_assembly\_3\_filtered**



FEATURES - color spectrum with similarity scores between the 2 sequences is displayed to the right.

- filtered for one-to-one mapping between the 2 sequences
- y-axis is cleaned up

COMMAND LINE STATEMENTS: `cd <directory>`

`mummerplot -p <prefix name> --large --png --color --filter -y [1,1013761900] -t DOTPLOT`  
`pangenome__fasta_vs_assembly_3_bbmhFilter.delta`

RUNTIME ~ 1min