Hox

* dmela-hox-genes.csv : file provided by Gregg that contains gene families associated with the HOX genes.
* \*.gene\_type : output from MCScanX [Duplicate\_gene\_classifier](https://github.com/wyp1125/MCScanX#duplicate-gene-classifier)
* HOX\_FIND\_2.py : python script used to generate results presented in the 2023\_hox tab of chelicerate\_genomes.xlsx

Duppipe

Dot\_plots

* \*.png : dot plots produced using MCScanX dot\_plotter
* Control\_File\_Maker.py : python script used to make dot\_plotter control files from gff file input

Updated.tar.gz

* Compressed directory where MCScanX analyses were completed

Duppipe.tar.gz

* Compressed directory where DupPipe was ran