These cervus documents are based on a genepop file created by filtering dDocent output on January 25, 2016 (see [Evernote](https://www.evernote.com/shard/s5/nl/154400311/1c7f5131-bba9-4a2e-8c5c-91b63d3f5ee8/)). Non-APCL species were removed from the genepop and the file, **DP10g95.genepop** was moved to the PC and converted to cervus, **DP10g95\_converted.csv**.

The project is saved as **DP10g95\_project.crv**

An allele frequency analysis was run and saved as **DP10g95\_AF.alf**  and **DP10g95\_AF.txt**.

An identity analysis was run and saved as **DP10g95\_ID.csv**(blue icon) **and DP10g95\_ID.txt**.

The ID.csv file was imported into google docs and saved as **DP10g95\_ID.csv**(green icon).