This read me file accompanies the dataset associated with the manuscript: “*Cryptic genetic diversity and cytonuclear discordance characterize contact among Canada jay (Perisoreus canadensis) morphotypes in western North America.”*

The file “BJLS-6492\_dryad data file” contains microsatellite genotyping data. Below is a description of the columns found in this file.

*Order no:* shows the sample number

*ID:* gives the unique name for each sample

*Allopatric?:* indicates whether the sample was collected from a population where two or more morphotypes come into contact.

Pop name: indicates which unique population each sample cam from

Location: lists the location where samples were collected from.

Museum Accession number: This number lists the unique museum accession number for all samples that were obtained from museums.

Morph: This column lists the morphotype Boreal, Pacific, or Rocky Mountain that each bird was assigned to. For those birds that we could not assign a morphotype to, we labeled the individual as an intergrade

mtDNA clade: This lists the mtDNA clade that each bird was assigned to.

The next two unnamed columns are duplicates of columns 1 and 2.

The next 24 columns list the microsatellite genotyping data. All fields that contain -9 represent missing data.

Finally the last two columns list the latitude and longitude for each sample.