## data homogenization processing summary and QC check results: $JRB\_CZO$ notes included with key file:

| source   | Var_long                                    | var           | var_notes                           |
|----------|---|---------------|-------------------------------------|
| location | Google Directory                            |               | JRB_CZO                             |
| location | Network (e.g. LTER, CZO, DIRT, NutNet, etc) | network       | CZO                                 |
| location | Site code (e.g. LUQ) or name                | $site\_code$  | JRB-CZO                             |
| location | Location name                               | location_name | Redondo Peak catchments             |
| location | Elevation                                   | elevation     | varies by catchment                 |
| location | Mean Annual Precipitation                   | map           | depends on elevation                |
| location | Slope                                       | slope         | varies by catchment                 |
| location | aboveground biomass                         | agb           | varies by watershed, 134-230 MgC/ha |

## files processed:

| type             | filename                    |
|------------------|-----------------------------|
| provided data    | JRB_ZOB_Soil_Summary2       |
| homogenized data | JRB_ZOB_Soil_Summary2_HMGZD |

## variable conversion

| source   | var          | Var_long                                      | given_unit    | $target\_unit$ | factor | varNotes  |
|----------|--------------|---|---------------|----------------|--------|---|
| location | agb          | aboveground biomass                           | g m-2         | t ha-1         | 100    | converted   |
| location | map          | Mean Annual<br>Precipitation                  | mm            | mm             |        | NOT<br>converted  |
| profile  | clay         | Clay  | %             | percent        |        | NOT   |
|          |              |   | ~             |                |        | converted   |
| profile  | coarse_frac  | Coarse Fraction                               | %             | percent        |        | NOT<br>converted  |
| profile  | lyr_n_tot    | Bulk Layer Total                              | %             | percent        |        | NOT   |
| -        | v — —        | Nitrogen concentration                        |               | 1              |        | converted   |
| profile  | $lyr\_soc$   | Bulk Layer Organic                            | %             | percent        |        | NOT   |
|          |              | Carbon (CN analyzer) concentration, inorganic |               |                |        | converted   |
|          |              | C removed or not present                      |               |                |        |   |
| profile  | sand         | Sand  | %             | percent        |        | NOT   |
| profile  | silt         | Silt  | %             | noveont        |        | $\begin{array}{c} { m converted} \\ { m NOT} \end{array}$ |
| prome    | SHU          | SIII  | 70            | percent        |        | converted   |
| profile  | $layer\_bot$ | Layer Bottom                                  | $\mathrm{cm}$ | cm             |        | NOT   |
| Cl       | 1 .1         | т • 1 11                                      |               |                |        | converted   |
| profile  | layer_mid    | Layer middle                                  | cm            | cm             |        | NOT<br>converted  |
| profile  | layer_top    | Layer Top                                     | cm            | cm             |        | NOT   |
|          |              |   |               |                |        | converted   |

QC results: location data

| dataset | source   | var | error        |
|---------|----------|-----|--------------|
| JRB_CZO | location | agb | out of range |

QC results: profile data, data range

profile data range checks passed

QC results: profile data, data type

profile data type checks passed