**Supplemental Table 1. Breeds and Naming Convention for the 15 Samples Analyzed in this Study.**

The main text uses the abbreviation to refer to each breed. Genomes sequenced from pooled blood samples are specified in brackets in the “Name” column.

|  |  |  |  |
| --- | --- | --- | --- |
| **Study** | **Platform** | **Name** | **Abbreviation** |
| This study | Illumina NextSeq 500 | Ross 308 | Ross 308 |
| 1978 | 1978 |
| 1957 | 1957 |
| Fan *et al.*2013 (SRP022583) | Illumina Genome Analyzer II | Silkie | Silkie |
| Taiwan heritage L2 line | L2 |
| Rubin *et al*.*,*2010 (SRP001870) | SOLiD | Ross 308 (10 males) | CB1 |
| Undisclosed commercial broiler (10 females) | CB2 |
| Swedish layer (11 males) | WLA |
| Commercial White Leghorn (8 males) | WLB |
| High growth line (7 males, 4 females) | High |
| Low growth line (7 males, 4 females) | Low |
| Obese (10 males) | Obese |
| Rhode Island Red (8 males) | RIR |
| Swedish red jungle fowl (8 males) | RJFswe |
|  |  | Red jungle fowl | RJF |