Supplementary Table 2 - List of Passeriformes and Psittaciformes species analyzed with *Gallus gallus* (GGA) chromosome painting or BACs clones corresponding to GGA1-10.

| Species | Family | 2n[1] | Fusions | Fissions | References |
|---|---|---|---|---|---|
| | | | **Passeriformes** | | |
| *Conopophaga lineata* | Conopophagidae | 78 | GGA2seg[2]/micro[3] | GGA1=CLI2 and 7<br>GGA2=CLI1 and 5 | [1] |
| *Pica pica* | Corvidae | 76 | - | GGA1=PPI3 and 4 | [2] |
| *Garrulus glandarius* | Corvidae | 78 | - | GGA1=GGL2 and 4 | [2] |
| *Taeniopygia guttata* | Estrildidae | 80 | - | GGA1=TGU2 and 5 | [3] |
| *Serinus canaria* | Fringillidae | 80 | - | GGA1=SCA2 and 5 | [3] |
| *Fringilla coelebs* | Fringillidae | 80 | - | GGA1=FCO3 and 4 | [4] |
| *Synallaxis frontalis* | Furnariidae | 82 | - | GGA1=SFR1 and 5<br>GGA2=SFR3 and 7 | [5] |
| *Glyphorynchus spirurus* | Furnariidae | 80 | - | GGA1=GSP3 and 4<br>GGA2=GSP2 and 5 | [6] |
| *Progne chalybea* | Hirundinidae | 76 | - | GGA1=PCH2 and 5 | [7] |
| *Progne tapera* | Hirundinidae | 76 | - | GGA1=PTA2 and 5 | [7] |
| *Pygochelidon cyanoleuca* | Hirundinidae | 76 | GGA6/micro | GGA1=PCY2 and 5 | [7] |
| *Zonotrichia capensis* | Passerellidae | 80 | - | GGA1=ZCA2 and 5 | [8] |
| *Parus major* | Paridae | 80 | - | GGA1=PMA2 and 4 | [2] |
| *Periparus ater* | Paridae | 80 | - | GGA1=PAT2 and 4 | [2] |
| *Sitta europaea* | Sittidae | 80 | GGA8/9=SEU7<br>GGA5/10=SEU6 | GGA1= SEU2 and 5<br>GGA5=SEU6q and 10 | [2] |
| *Sylvia atricapilla* | Sylviidae | 74 | GGA4q/GGA4p=SAT5 | GGA1=SAT2 and 4 | [2] |
| *Saltator aurantiirostris* | Thraupidae | 80 | - | GGA1=SAU2 and 5 | [9] |
| *Saltator similis* | Thraupidae | 80 | - | GGA1=SSI2 and 5 | [9] |
| *Sicalis flaveola* | Thraupidae | 80 | - | GGA1=SFL2 and 4 | Present study |
| *Turdus rufiventris* | Turdidae | 78 | - | GGA1=TRU2 and 5 | [10] |
| *Turdus albicollis* | Turdidae | 78 | - | GGA1=TAL2 and 5 | [10] |
| *Turdus iliacus* | Turdidae | 80 | - | GGA1=TIL2 and 5 | [4] |

| | | | | | |
|---|---|---|---|---|---|
| *Turdus merula* | Turdidae | 80 | - | GGA1=TME2 and 5 | [11] |
| *Elaenia spectabilis* | Tyrannidae | 80 | - | GGA1=ESP2 and 5 | [12] |
| *Pitangus sulphuratus* | Tyrannidae | 80 | - | GGA1=PSU3 and 5 | [13] |
| *Serpophaga subcristata* | Tyrannidae | 82 | - | GGA1=SSU3 and 5 | [13] |
| *Satrapa icterophrys* | Tyrannidae | 82 | - | GGA1=SIC2 and 5 | [13] |
| **Psittaciformes** | | | | | |
| *Nymphicus hollandicus* | Cacatuidae | 72 | GGA6/7=NHO5<br>GGA4/8/9=NHO4 | GGA1=NHO3 and 6<br>GGA9=NHO4p and 10 | [14] |
| *Ara chloropterus* | Psittacidae | 70 | GGA1seg/4q=ACH1<br>GGA6/7=ACH6<br>GGA8/9=ACH7 | GAA1=ACH1q and 4<br>GGA2=ACH2 and 11 | [15] |
| *Anodorhynchus hyacinthinus* | Psittacidae | 70 | GGA1seg/4q=AHY1<br>GGA6/7=AHY6<br>GGA8/9=AHY7 | GGA1=AHY1q and 4 | [15] |
| *Psittacus erithacus* | Psittacidae | 62-64 | GGA1seg/4=PER1<br>GGA6/7=PER6<br>GGA8/9=PER7 | GGA1=PER1q and 4 | [16] |
| *Ara macaw* | Psittacidae | 62-64 | GGA1seg/4q=AMA1<br>GGA6/7=AMA6<br>GGA8/9=AMA7 | GGA1=AMA1, 4, and 9 | [17] |
| *Pyrrhura frontalis* | Psittacidae | 70 | GGA1q/4=PFR1<br>GGA6/7=PFR6 | GGA1=PFR1q and 4 | [18] |
| *Amazona aestiva* | Psittacidae | 70 | GGA6/7=AAE7 | GGA1=AAE2 and 5<br>GGA2=AAE1 and 12 | [18] |
| *Agapornis roseicollis* | Psittaculidae | 48 | GGA6/7=ARO6<br>GGA1/4=ARO4<br>GGA8/9=ARO5<br>GGA2/9=ARO9 | GGA1=ARO3 and 4q<br>GGA2=ARO2 and 9q<br>GGA9=ARO5q and 9q | [14] |
| *Melopsittacus undulatus* | Psittaculidae | 62 | GGA5/6/7=MUN4<br>GGA4/8/9=MUN5 | GGA1=MUN3 and 6<br>GGA6=MUN4p and 8p | [14] |

[1] Diploid number, [2] segment, and [3] microchromosome. The data of *Myiopsitta monachus* (Psittaciformes) were not compared here because the exact fusions involving chromosomes GGA10, 11, and 12 were not identified, since a flow sorted peak including the chromosomes GGA10, 11, and 12 was used [19]. The data of *Willisornis vidua* (Passeriformes) were not compared here because the authors reported several interchromosomal rearrangements in this species, however, the exact rearrangements were not described [20].

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