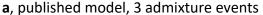
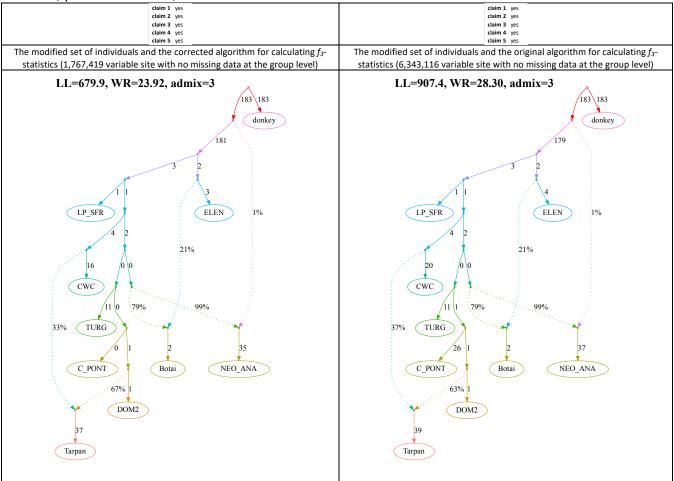
Figure 3–source data 7. Published admixture graphs from Librado *et al.* (2019) and alternative graphs found with *findGraphs* (10 populations, 3 to 5 admixture events) for the modified group composition and using the updated algorithm for calculating *f*-statistics. The graphs were also re-fitted on the original set of SNPs/individuals and using the original algorithm for calculating *f*-statistics. Selected alternative graphs found with *findGraphs* when more admixture events were allowed (from 6 to 9) are also shown. Model parameters (graph edges) that were inferred to be unidentifiable are plotted in red.

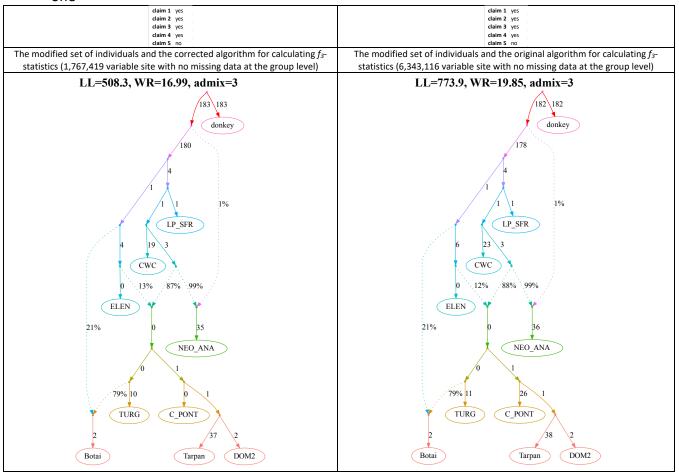




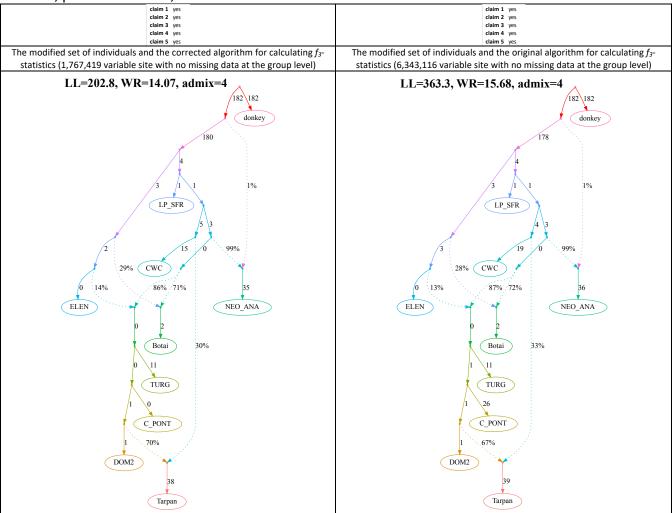
Claims by Librado et al. 2021 relying on the admixture graph:

- 1) NEO-ANA-related admixture is absent in DOM2;
- DOM2 and C-PONT are sister groups;
- there is no gene flow connecting the CWC and the cluster associated with Yamnaya horses and horses of the later Sintashta culture whose ancestry is maximized in the Western Steppe (DOM2, C-PONT, TURG);
- 4) there was a gene flow from a deep-branching ghost group to NEO-ANA;
- 5) Tarpan is a mixture of a CWC-related and a DOM2-related lineage.

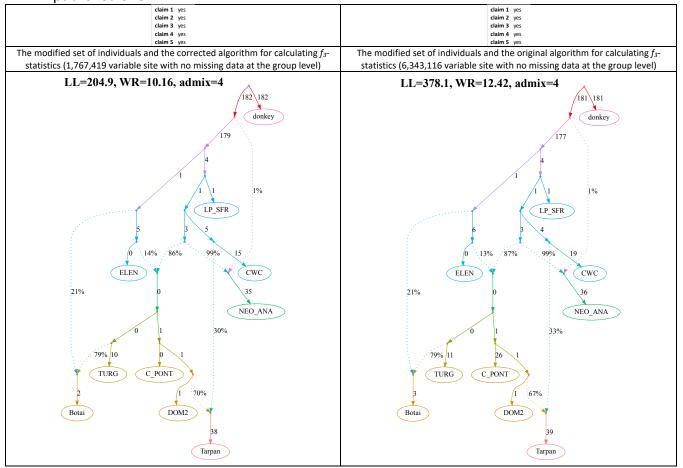
b, an alternative model with 3 admixture events fitting significantly better than the published one



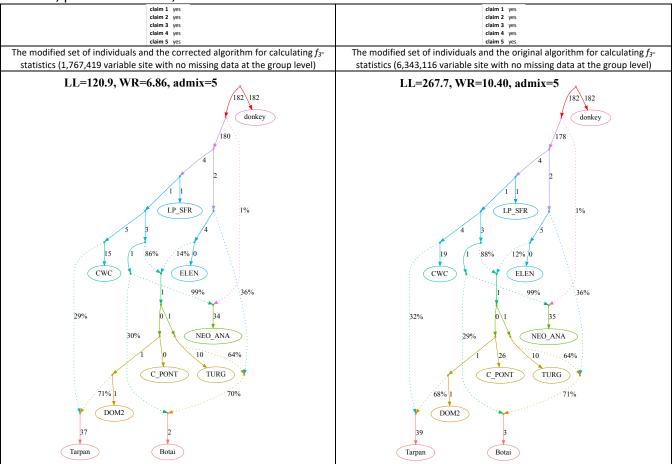
c, published model, 4 admixture events



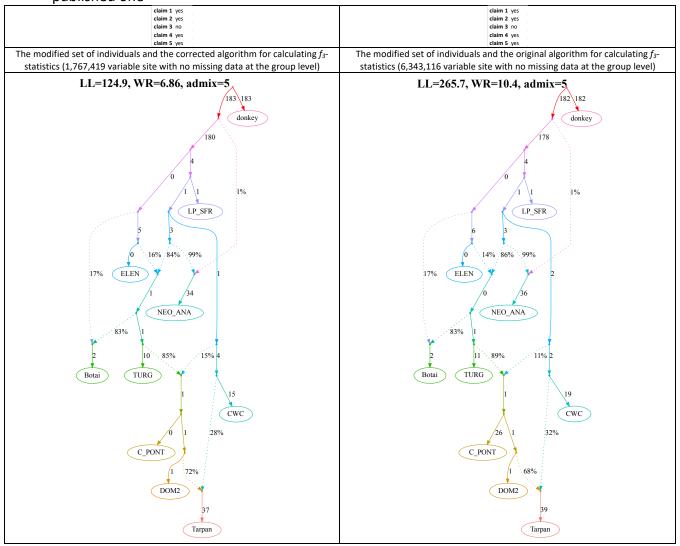
d, an alternative model with 4 admixture events fitting not significantly worse than the published one



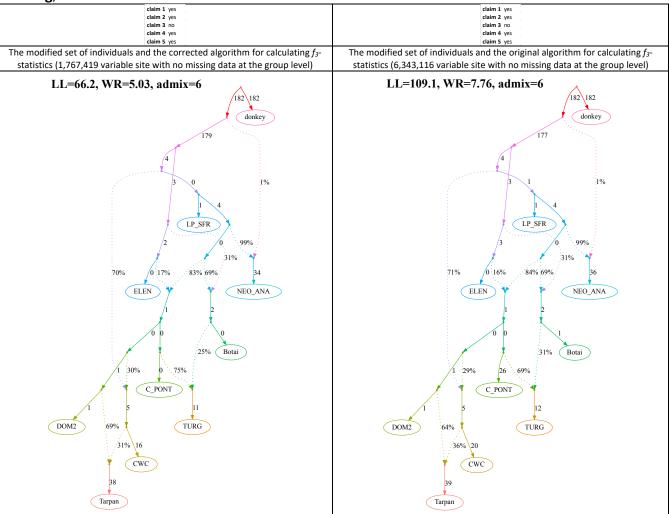
e, published model, 5 admixture events

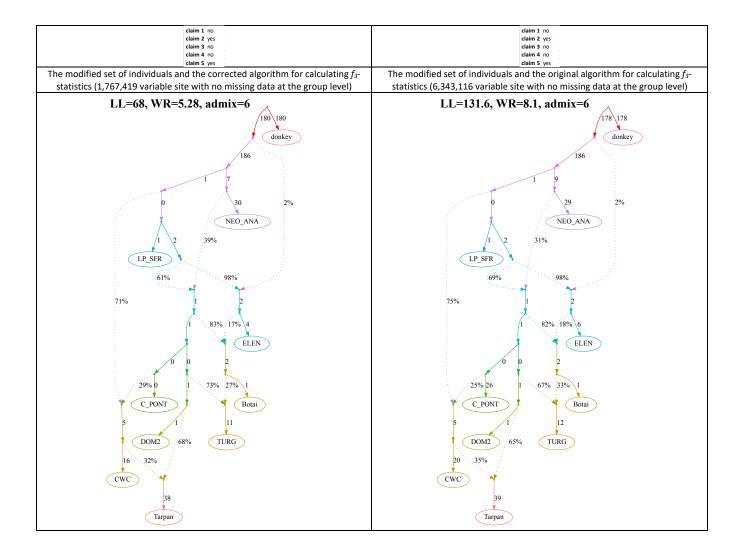


f, an alternative model with 5 admixture events fitting not significantly worse than the published one

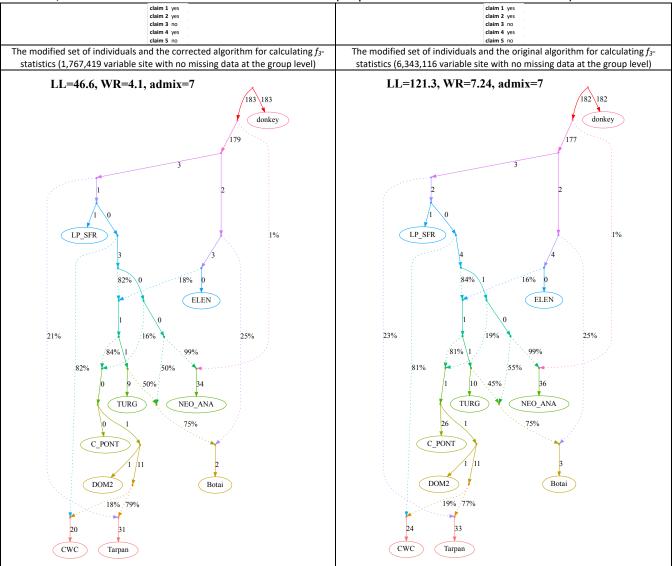


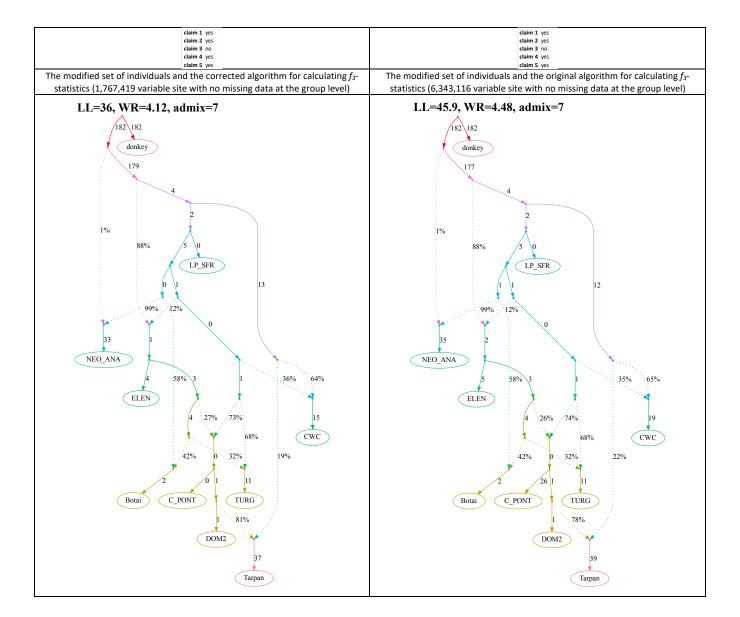
g, selected models with 6 admixture events



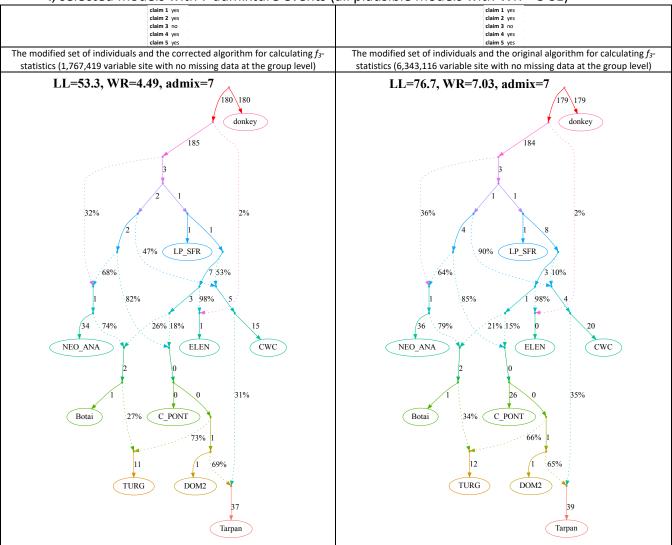


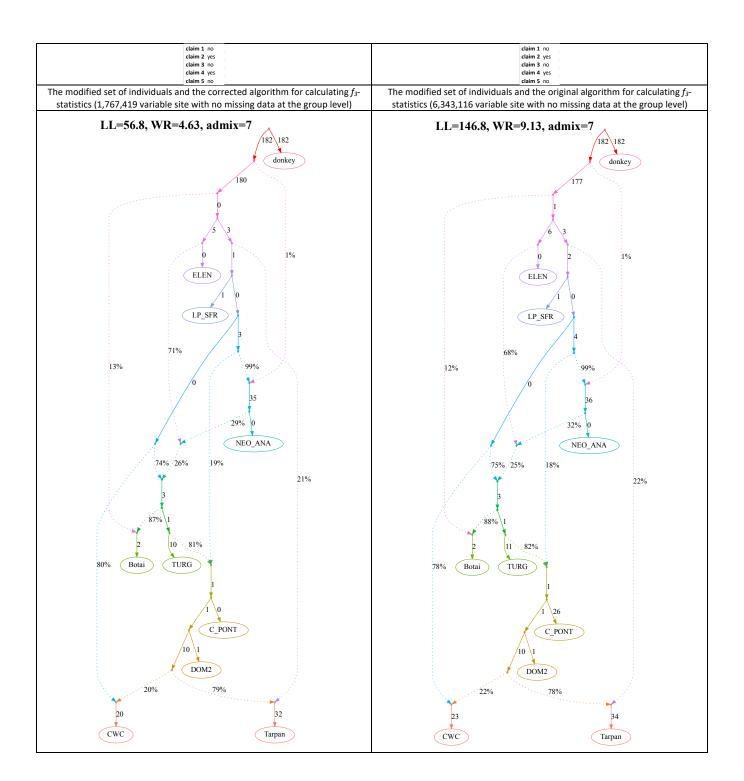
h, selected models with 7 admixture events (all plausible models with WR < 5 SE)





i, selected models with 7 admixture events (all plausible models with WR < 5 SE)





j, selected models with 8 admixture events (all plausible models with WR < 4 SE) claim 5 no The modified set of individuals and the corrected algorithm for calculating f_3 -The modified set of individuals and the original algorithm for calculating f_3 statistics (1,767,419 variable site with no missing data at the group level) statistics (6,343,116 variable site with no missing data at the group level) LL=37, WR=3.38, admix=8 LL=104.3, WR=7.81, admix=8 181 181 179 179 donkey donkey 30% 19% 1% LP SFR LP_SFR ELEN ELEN 0 42% 40% 60% 44% 56% 22% Botai 23% Botai 15% 99% 14% 99%

TURG

33

Tarpan

NEO_ANA

C PONT

DOM2

23

CWC

TURG

NEO_ANA

C_PONT

DOM2

21%

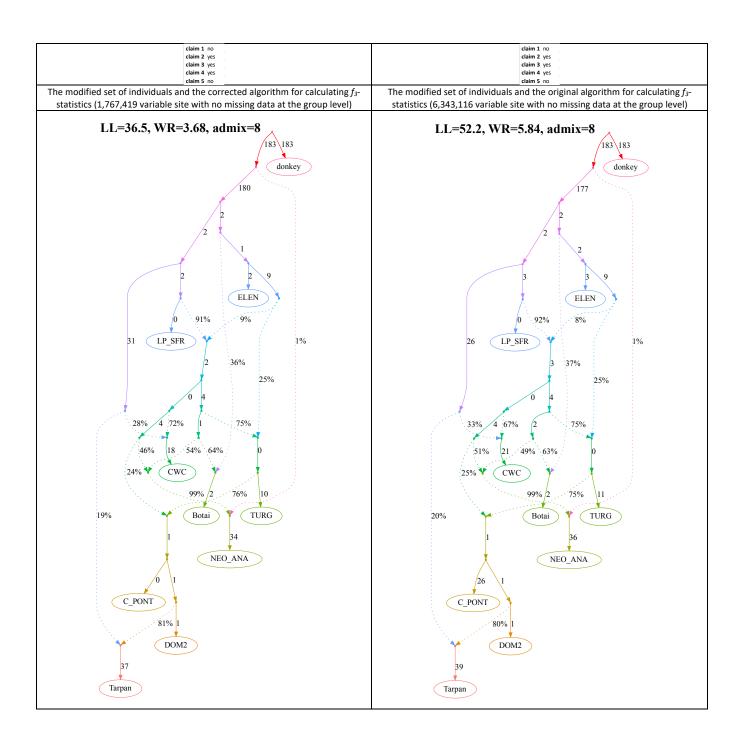
20

CWC

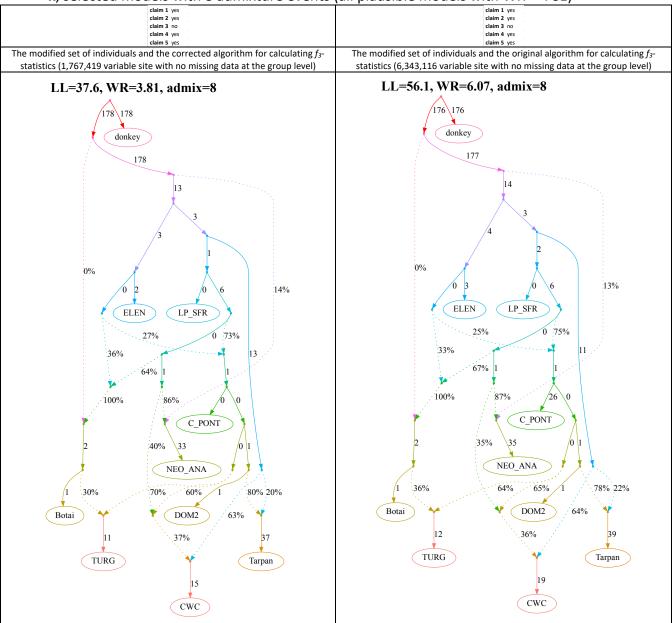
32

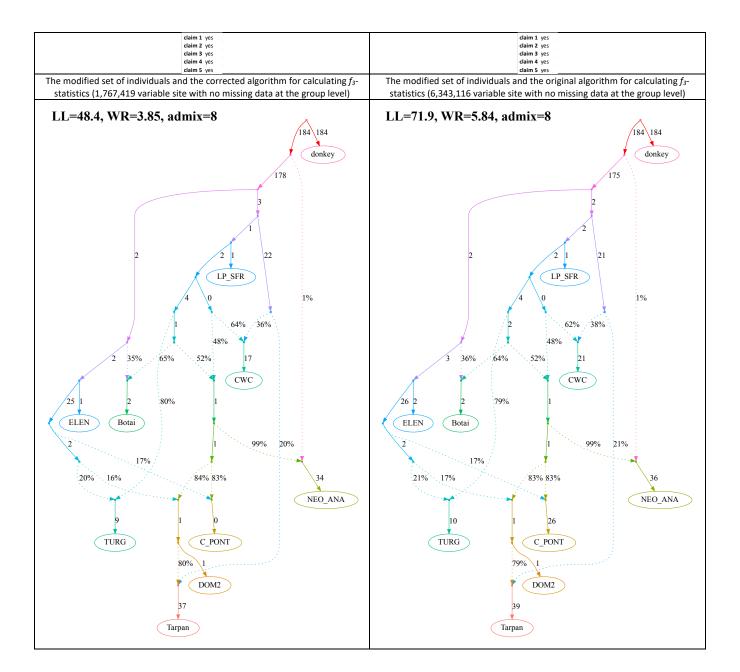
Tarpan

79%



k, selected models with 8 admixture events (all plausible models with WR < 4 SE)





I, selected models with 8 admixture events (all plausible models with WR < 4 SE) claim 5 yes claim 5 yes The modified set of individuals and the corrected algorithm for calculating f_{3} -The modified set of individuals and the original algorithm for calculating f_3 statistics (1,767,419 variable site with no missing data at the group level) statistics (6,343,116 variable site with no missing data at the group level) LL=43.2, WR=3.85, admix=8 LL=98.4, WR=7.86, admix=8 182 182 183 183 donkey donkey 1% 1% ELEN ELEN 21% 20% LP SFR LP_SFR 99% 18 99% 26% 74% 23% 77% 16% 86% 14% 36 88% 12% NEO_ANA NEO_ANA 71% 84% 70% 85% TURG Botai TURG Botai 30% 26 92% 29% 0 .93% DOM2 C_PONT DOM2 C_PONT

16 23%

Tarpan

CWC

78%

. 22%

Tarpan

CWC

m, selected models with 9 admixture events (all plausible models with WR < 4 SE) claim 5 yes claim 5 yes The modified set of individuals and the corrected algorithm for calculating The modified set of individuals and the original algorithm for calculating f_3 f_3 -statistics (1,767,419 variable site with no missing data at the group level) statistics (6,343,116 variable site with no missing data at the group level) LL=45.7, WR=5.66, admix=9 LL=27.2, WR=3.38, admix=9 136 136 134 134 donkey donkey 134 135 144 142 10 3% 3% LP_SFR LP_SFR /29 NEO_ANA NEO ANA 36% 0% 28% 97% 15 72% 72% 72% 32% 25% 11% 28% 28% 9%

ELEN

34%

TURG

Botai

75%

DOM2

25%

C_PONT

91%

16

CWC

ELEN

72%

TURG

Botai

C_PONT

12

CWC

71%

86%

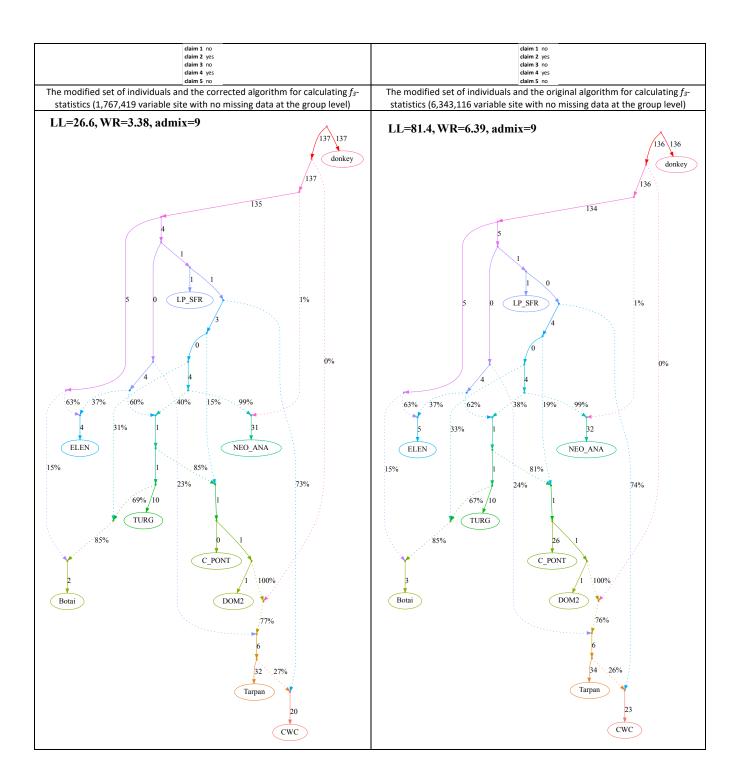
14%

100%

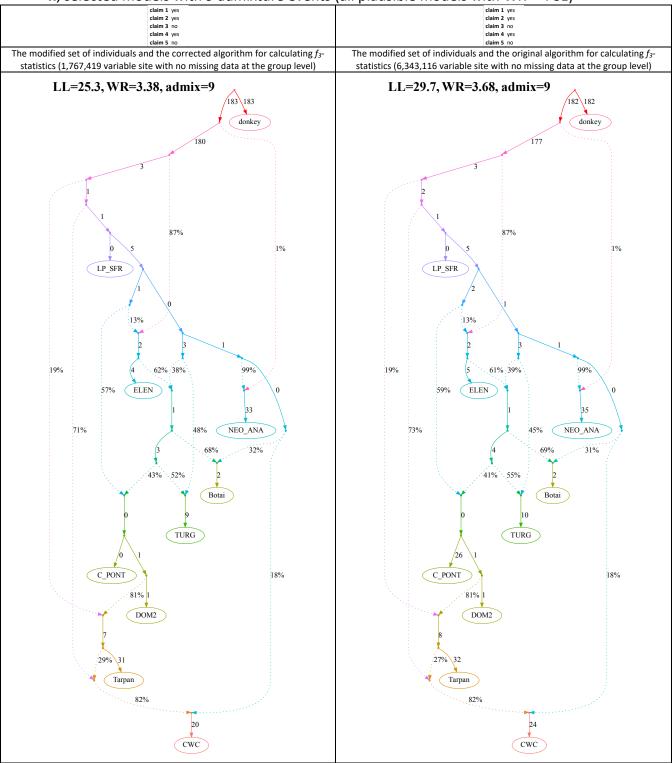
Tarpan

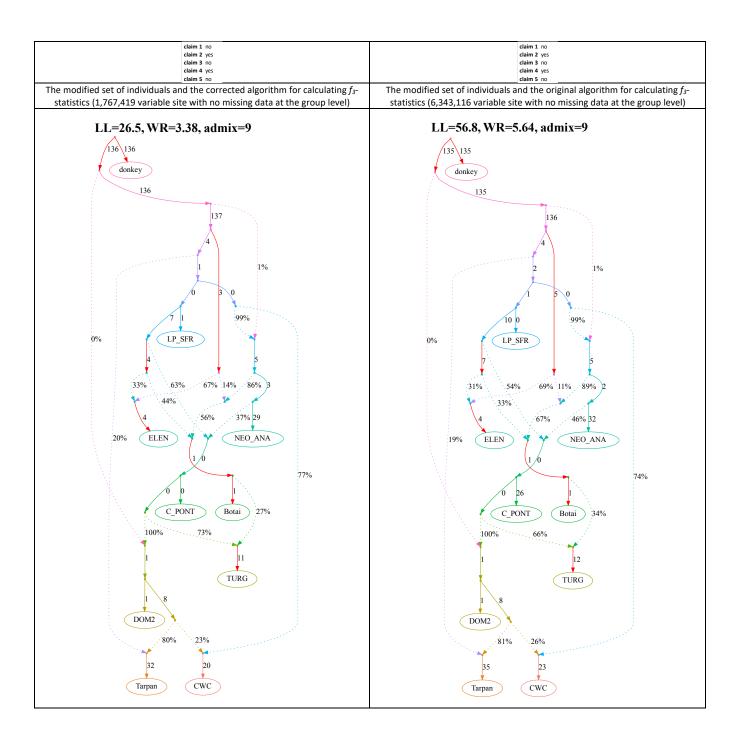
29%

DOM2

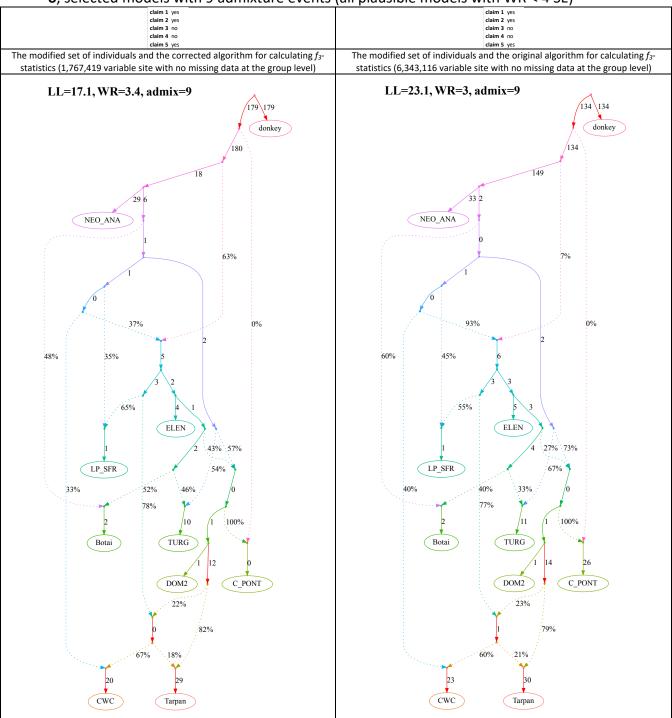


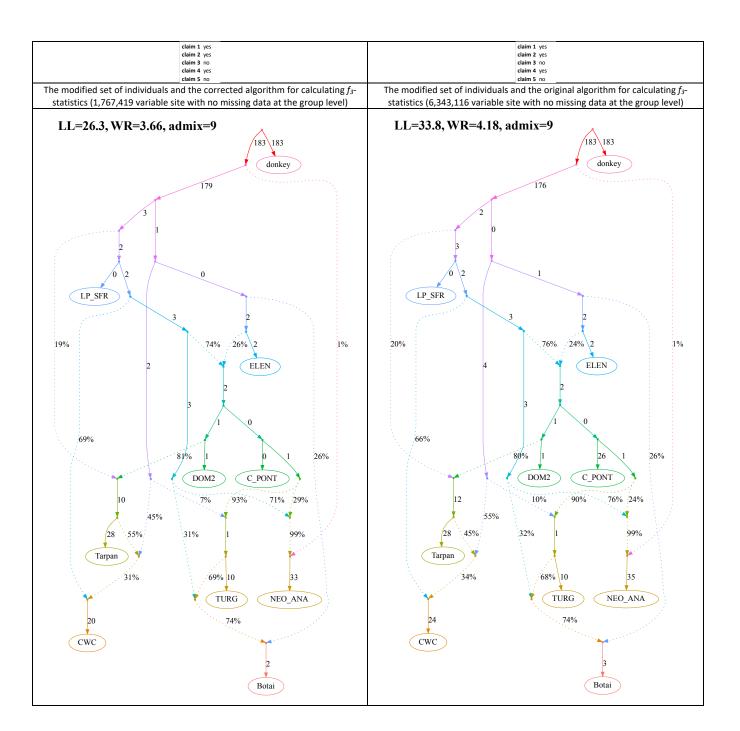
n, selected models with 9 admixture events (all plausible models with WR < 4 SE)



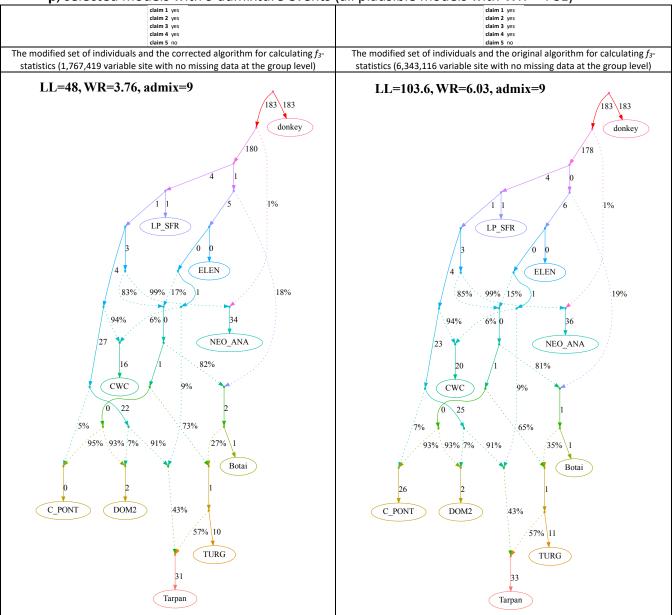


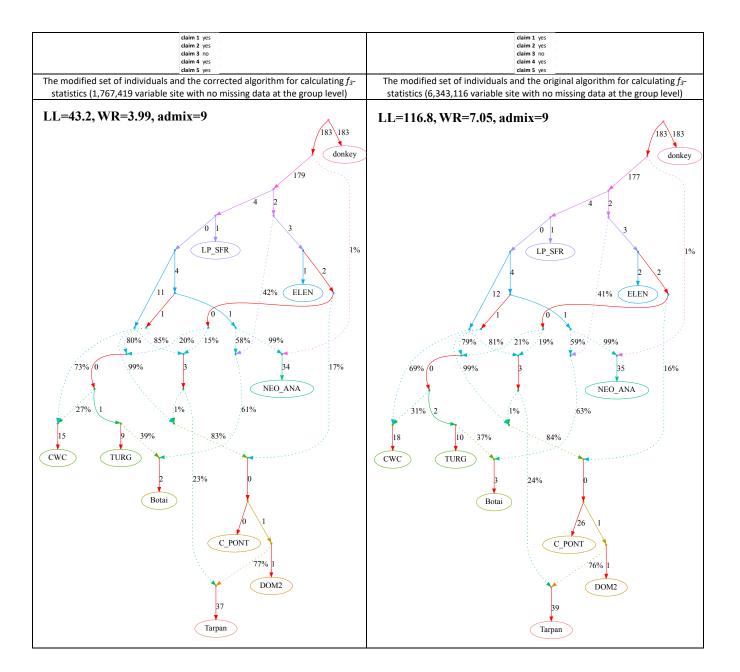
o, selected models with 9 admixture events (all plausible models with WR < 4 SE)



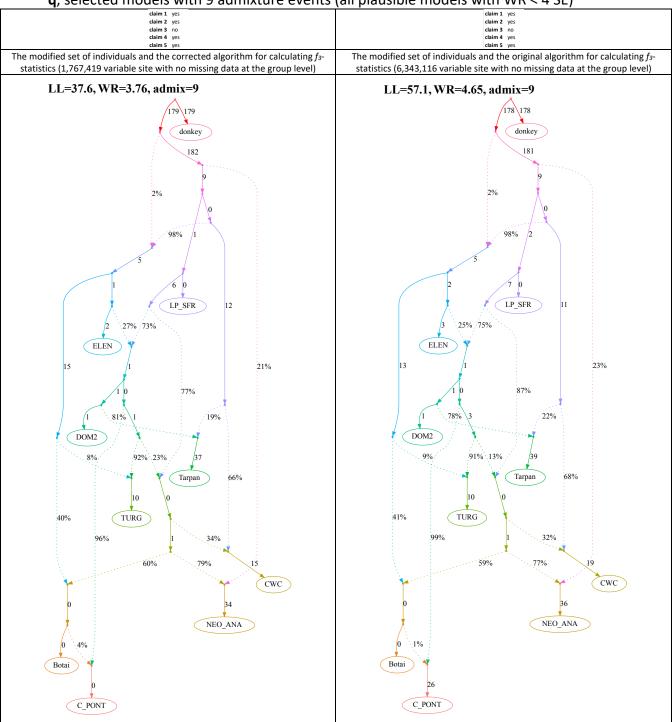


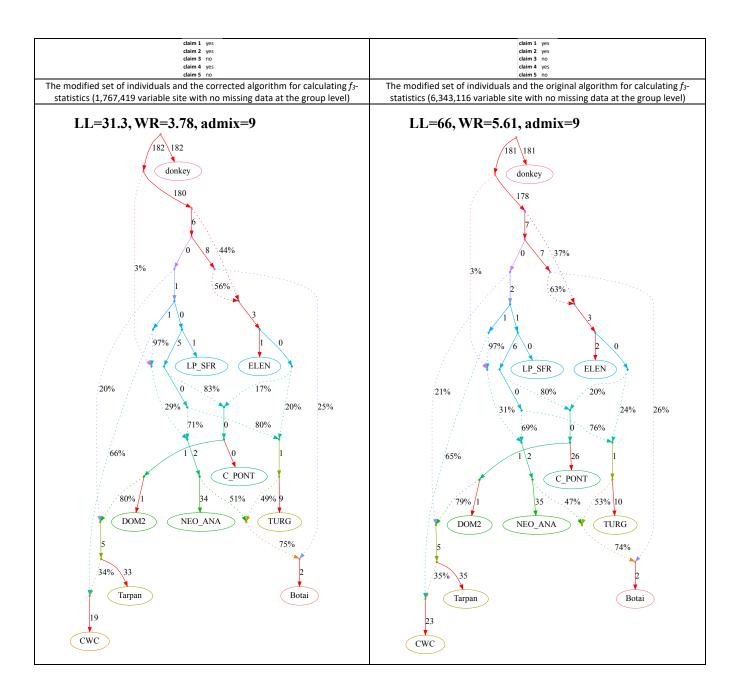






q, selected models with 9 admixture events (all plausible models with WR < 4 SE)





r, selected models with 9 admixture events (all plausible models with WR < 4 SE)

