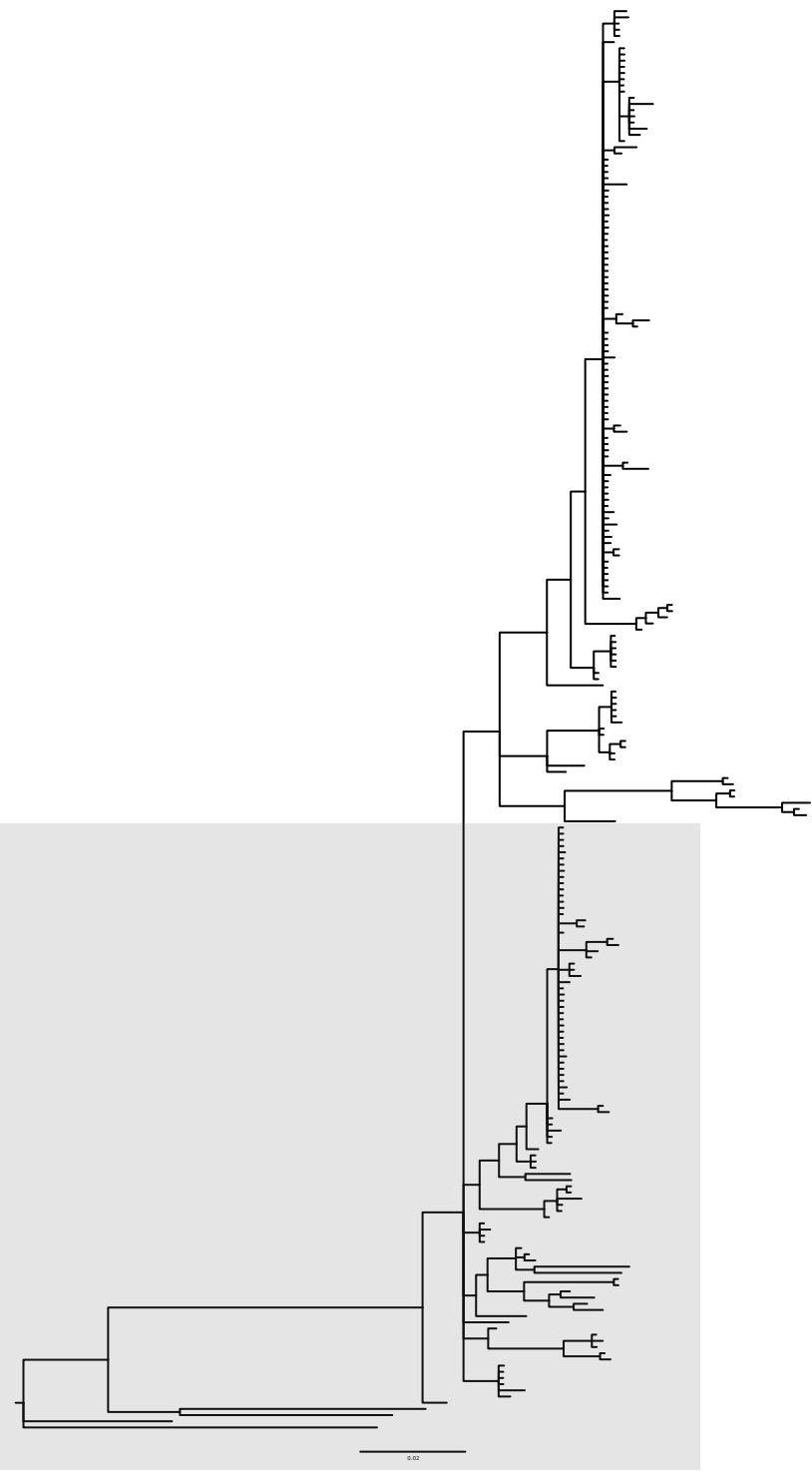


SPECIES DELIMITATION

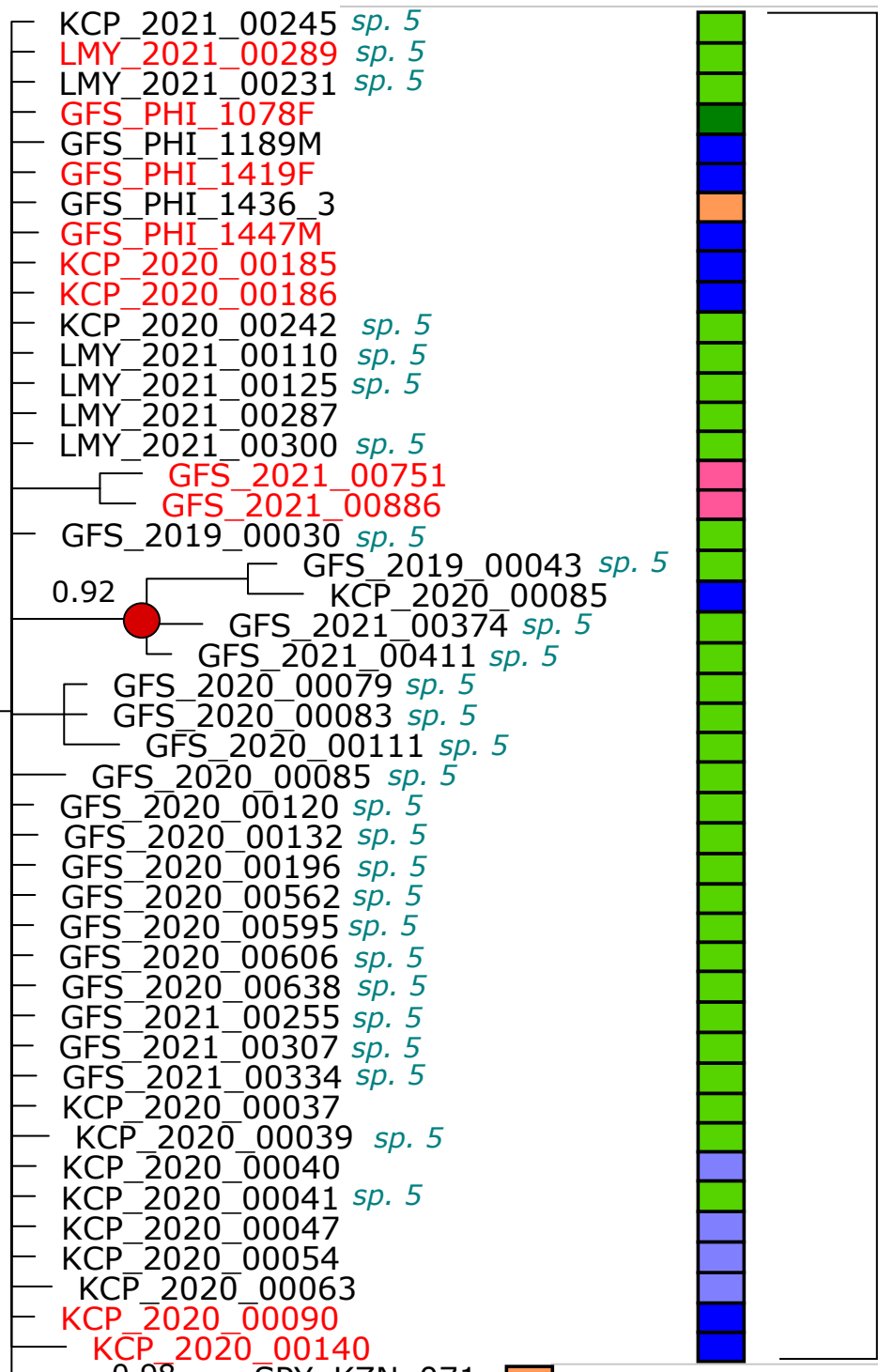


Scale
0.02 substitutions per site



2

NON-GOLDEN SHOULDER



NGS

NGS.SPY

HHIR.2

NGS.4

P1

P2

SYC

P3

P3.1

NGS.3

Clade number

ABGD (p = 0.0028, 1.3%)

ASAP

bPTP (BI)

PTP (ML)

GMYC (single)

HAPLOWEBS ($\epsilon = 0.2$)

P-DISTANCE = 0.3%

P-DISTANCE = 1%

P-DISTANCE = 1.5%

P-DISTANCE = 3%

CONSENSUS

OUTGROUPS

