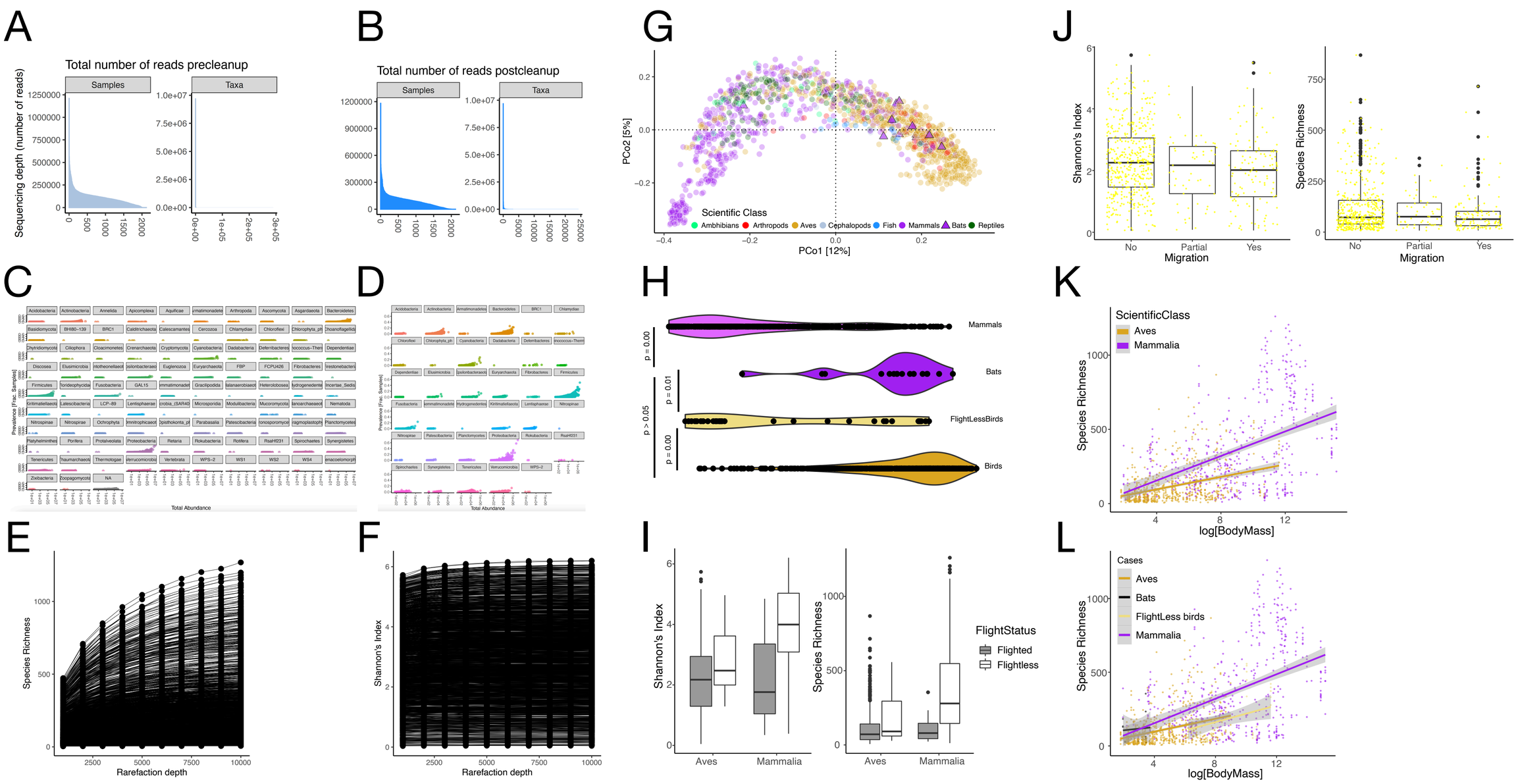
A
Total number of reads precleanup
Samples
Taxa
Sequencing depth (number of reads)
B
Total number of reads postcleanup
Samples
Taxa
G
PCo2 [5%]
PCo1 [12%]
Scientific Class
Ambhibians
Arthropods
Aves
Cephalopods
Fish
Mammals
Bats
Reptiles
J
Shannon's Index
Species Richness
No
Partial
Yes
Migration
C
Prevalence [Frac. Samples]
Total Abundance
D
Prevalence [Frac. Samples]
Total Abundance
H
p = 0.00
p = 0.01
p > 0.05
p = 0.00
Mammals
Bats
FlightLessBirds
Birds
K
ScientificClass
Aves
Mammalia
Species Richness
log[BodyMass]
E
Species Richness
Rarefaction depth
F
Shannon's Index
Rarefaction depth
I
Shannon's Index
Species Richness
Aves
Mammalia
FlightStatus
Flighted
Flightless
L
Cases
Aves
Bats
FlightLess birds
Mammalia
Species Richness
log[BodyMass]