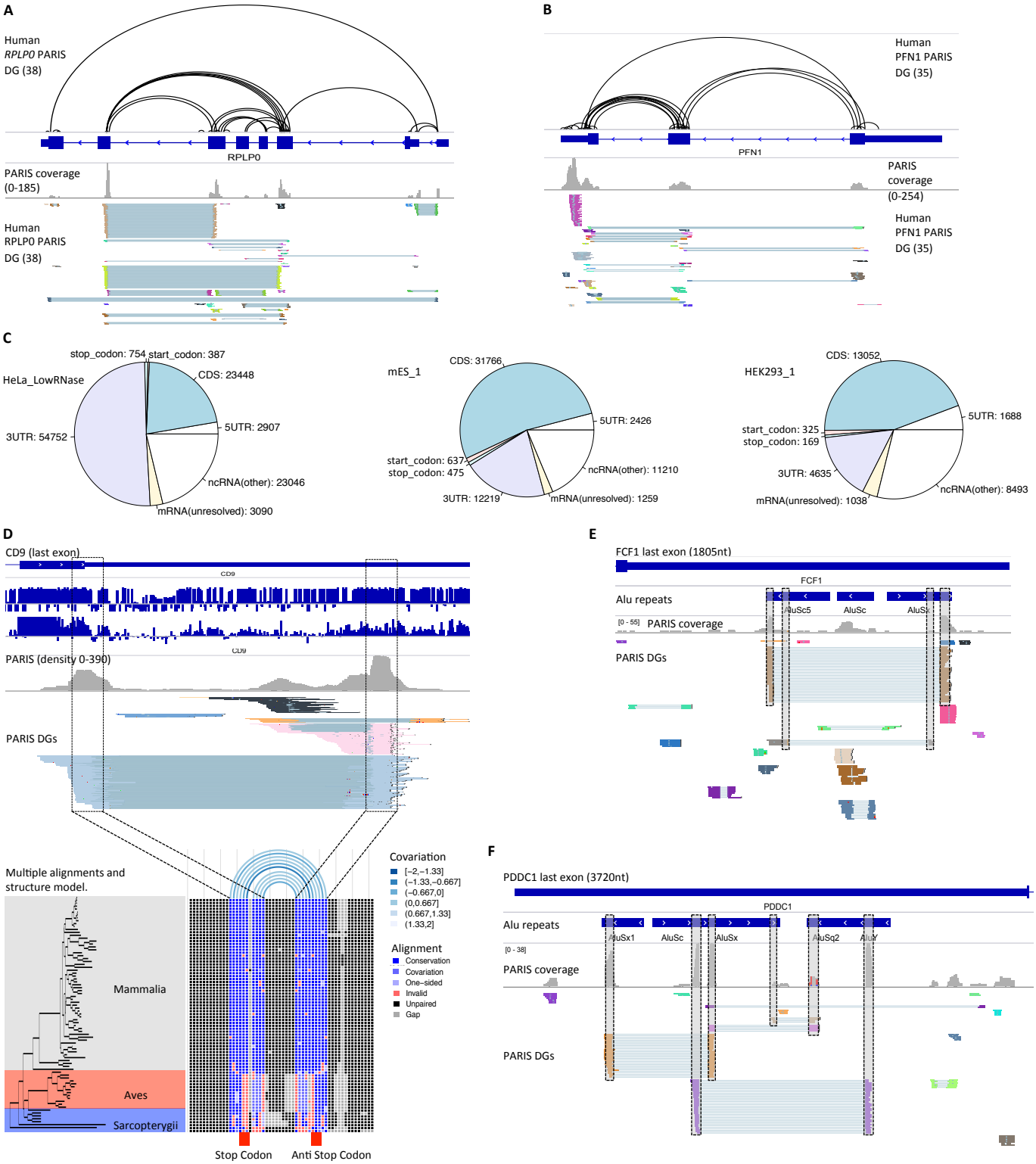
A
Human
RPLP0 PARIS
DG (38)
RPLP0
PARIS coverage
(0-185)
Human
RPLP0 PARIS
DG (38)
B
Human
PFN1 PARIS
DG (35)
PFN1
PARIS
coverage
(0-254)
Human
PFN1 PARIS
DG (35)
C
HeLa_LowRNase
stop_codon: 754
start_codon: 387
CDS: 23448
5UTR: 2907
3UTR: 54752
ncRNA(other): 23046
mRNA(unresolved): 3090
mES_1
CDS: 31766
5UTR: 2426
start_codon: 637
stop_codon: 475
ncRNA(other): 11210
3UTR: 12219
mRNA(unresolved): 1259
HEK293_1
CDS: 13052
5UTR: 1688
start_codon: 325
stop_codon: 169
3UTR: 4635
ncRNA(other): 8493
mRNA(unresolved): 1038
D
CD9 (last exon)
CD9
PARIS (density 0-390)
PARIS DGs
Multiple alignments and
structure model.
Mammalia
Aves
Sarcopterygii
Covariation
[−2,−1.33]
(−1.33,−0.667]
(−0.667,0]
(0,0.667]
(0.667,1.33]
(1.33,2]
Alignment
Conservation
Covariation
One−sided
Invalid
Unpaired
Gap
Stop Codon
Anti Stop Codon
E
FCF1 last exon (1805nt)
FCF1
Alu repeats
AluSc5
AluSc
AluSx
[0 - 55]
PARIS coverage
PARIS DGs
F
PDDC1 last exon (3720nt)
PDDC1
Alu repeats
AluSx1
AluSc
AluSx
AluSq2
AluY
[0 - 38]
PARIS coverage
PARIS DGs