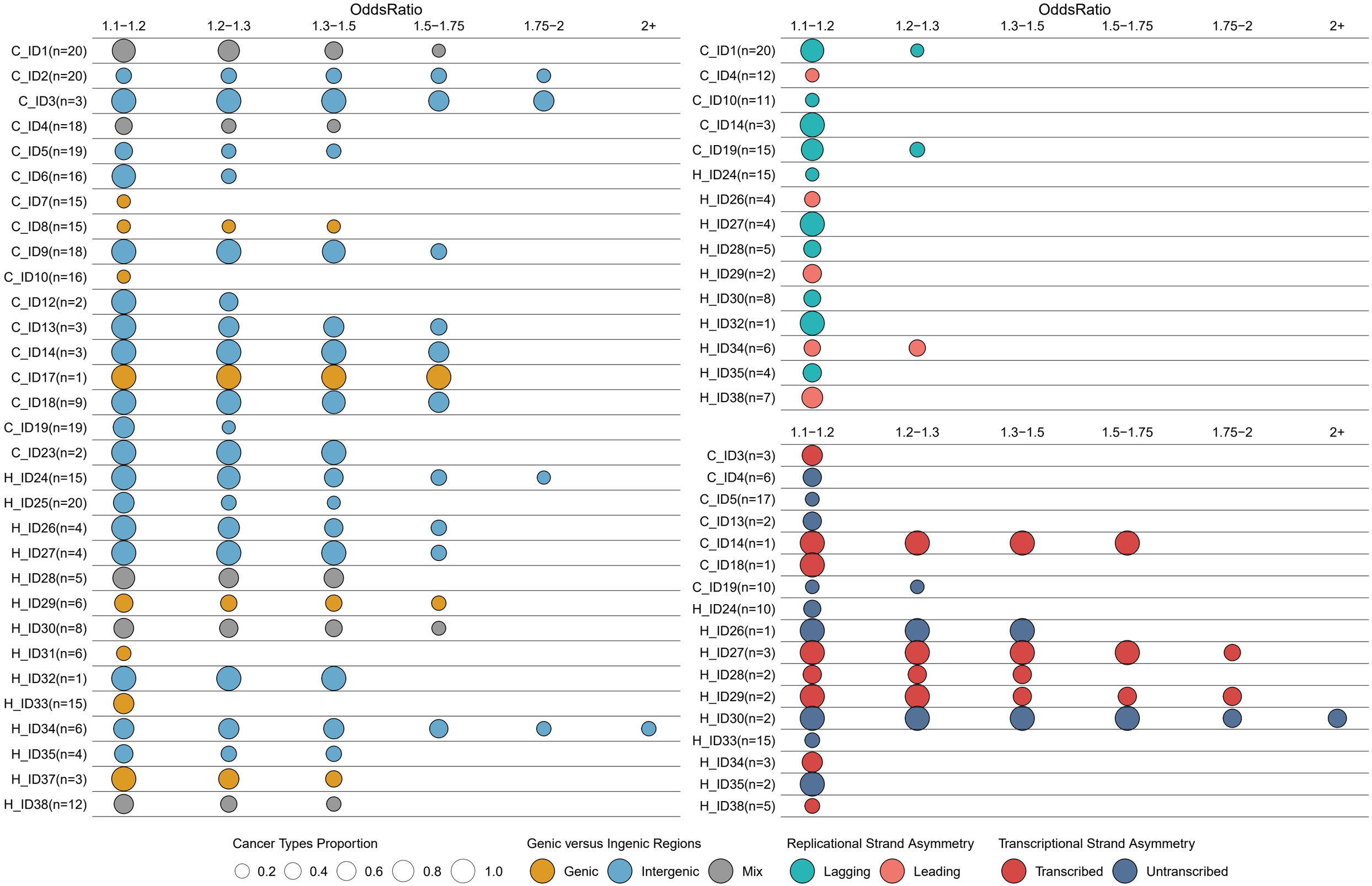
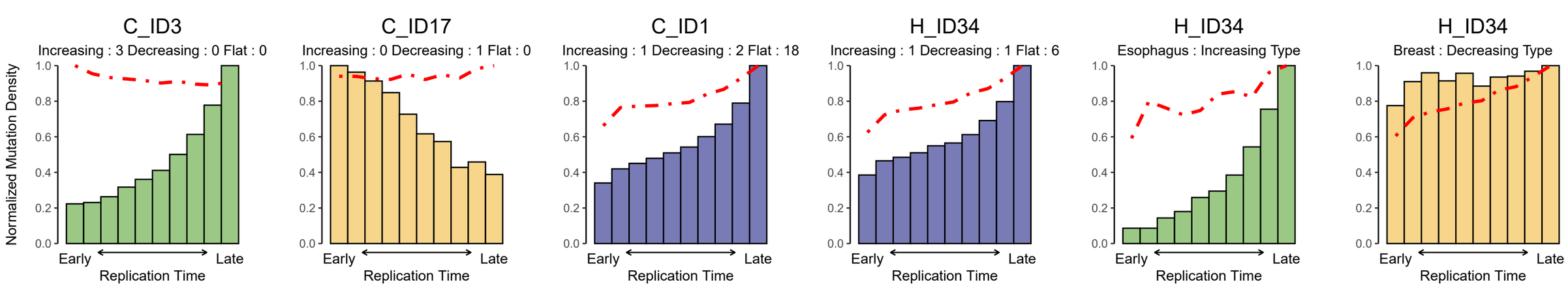

OddsRatio
1.1−1.2
1.2−1.3
1.3−1.5
1.5−1.75
1.75−2
2+
C_ID1(n=20)
C_ID2(n=20)
C_ID3(n=3)
C_ID4(n=18)
C_ID5(n=19)
C_ID6(n=16)
C_ID7(n=15)
C_ID8(n=15)
C_ID9(n=18)
C_ID10(n=16)
C_ID12(n=2)
C_ID13(n=3)
C_ID14(n=3)
C_ID17(n=1)
C_ID18(n=9)
C_ID19(n=19)
C_ID23(n=2)
H_ID24(n=15)
H_ID25(n=20)
H_ID26(n=4)
H_ID27(n=4)
H_ID28(n=5)
H_ID29(n=6)
H_ID30(n=8)
H_ID31(n=6)
H_ID32(n=1)
H_ID33(n=15)
H_ID34(n=6)
H_ID35(n=4)
H_ID37(n=3)
H_ID38(n=12)
OddsRatio
1.1−1.2
1.2−1.3
1.3−1.5
1.5−1.75
1.75−2
2+
C_ID1(n=20)
C_ID4(n=12)
C_ID10(n=11)
C_ID14(n=3)
C_ID19(n=15)
H_ID24(n=15)
H_ID26(n=4)
H_ID27(n=4)
H_ID28(n=5)
H_ID29(n=2)
H_ID30(n=8)
H_ID32(n=1)
H_ID34(n=6)
H_ID35(n=4)
H_ID38(n=7)
1.1−1.2
1.2−1.3
1.3−1.5
1.5−1.75
1.75−2
2+
C_ID3(n=3)
C_ID4(n=6)
C_ID5(n=17)
C_ID13(n=2)
C_ID14(n=1)
C_ID18(n=1)
C_ID19(n=10)
H_ID24(n=10)
H_ID26(n=1)
H_ID27(n=3)
H_ID28(n=2)
H_ID29(n=2)
H_ID30(n=2)
H_ID33(n=15)
H_ID34(n=3)
H_ID35(n=2)
H_ID38(n=5)
Cancer Types Proportion
0.2
0.4
0.6
0.8
1.0
Genic versus Ingenic Regions
Genic
Intergenic
Mix
Replicational Strand Asymmetry
Lagging
Leading
Transcriptional Strand Asymmetry
Transcribed
Untranscribed


C_ID3
Increasing : 3 Decreasing : 0 Flat : 0
C_ID17
Increasing : 0 Decreasing : 1 Flat : 0
C_ID1
Increasing : 1 Decreasing : 2 Flat : 18
H_ID34
Increasing : 1 Decreasing : 1 Flat : 6
H_ID34
Esophagus : Increasing Type
H_ID34
Breast : Decreasing Type
Normalized Mutation Density
0.0
0.2
0.4
0.6
0.8
1.0
Early
Late
Replication Time