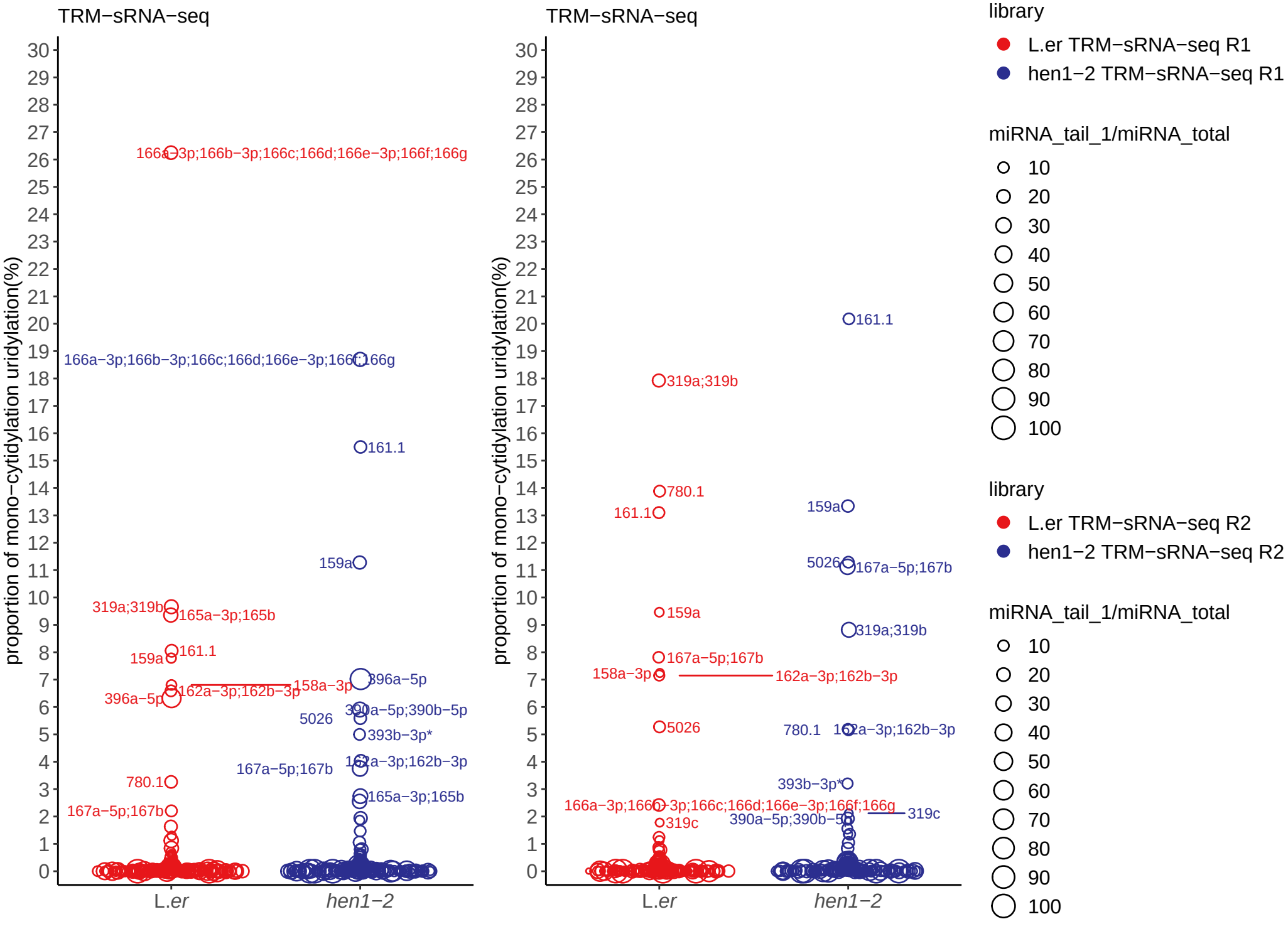
TRM−sRNA−seq
proportion of mono−cytidylation uridylation(%)
L.er
hen1−2
166a−3p;166b−3p;166c;166d;166e−3p;166f;166g
161.1
159a
319a;319b
165a−3p;165b
396a−5p
162a−3p;162b−3p
158a−3p
390a−5p;390b−5p
5026
393b−3p*
167a−5p;167b
780.1
319c
library
L.er TRM−sRNA−seq R1
hen1−2 TRM−sRNA−seq R1
miRNA_tail_1/miRNA_total
10
20
30
40
50
60
70
80
90
100
library
L.er TRM−sRNA−seq R2
hen1−2 TRM−sRNA−seq R2
miRNA_tail_1/miRNA_total
10
20
30
40
50
60
70
80
90
100