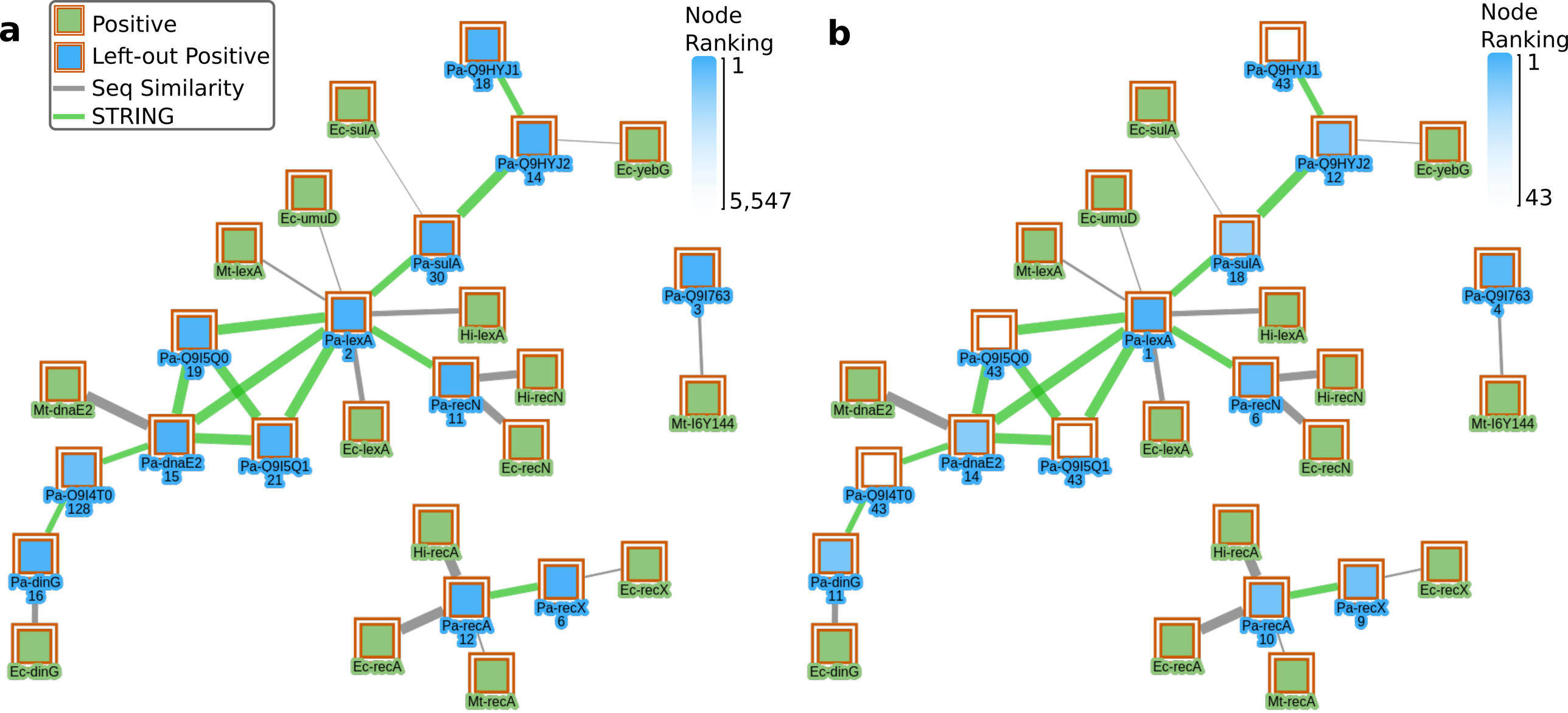
a
Positive
Left-out Positive
Seq Similarity
STRING
Node Ranking
1
5,547
Pa-Q9HYJ1 18
Ec-sulA
Pa-Q9HYJ2 14
Ec-yebG
Ec-umuD
Mt-lexA
Pa-sulA 30
Pa-Q9I763 3
Pa-lexA 2
Hi-lexA
Pa-Q9I5Q0 19
Mt-dnaE2
Pa-recN 11
Hi-recN
Mt-I6Y144
Ec-lexA
Pa-dnaE2 15
Pa-Q9I5Q1 21
Ec-recN
Pa-Q9I4T0 128
Hi-recA
Pa-dinG 16
Pa-recX 6
Ec-recX
Pa-recA 12
Ec-recA
Ec-dinG
Mt-recA
SinkSource
b
Node Ranking
1
43
Pa-Q9HYJ1 43
Ec-sulA
Pa-Q9HYJ2 12
Ec-yebG
Ec-umuD
Mt-lexA
Pa-sulA 18
Pa-Q9I763 4
Pa-lexA 1
Hi-lexA
Pa-Q9I5Q0 43
Mt-dnaE2
Pa-recN 6
Hi-recN
Mt-I6Y144
Ec-lexA
Pa-dnaE2 14
Pa-Q9I5Q1 43
Ec-recN
Pa-Q9I4T0 43
Hi-recA
Pa-dinG 11
Pa-recX 9
Ec-recX
Pa-recA 10
Ec-recA
Ec-dinG
Mt-recA
Local+