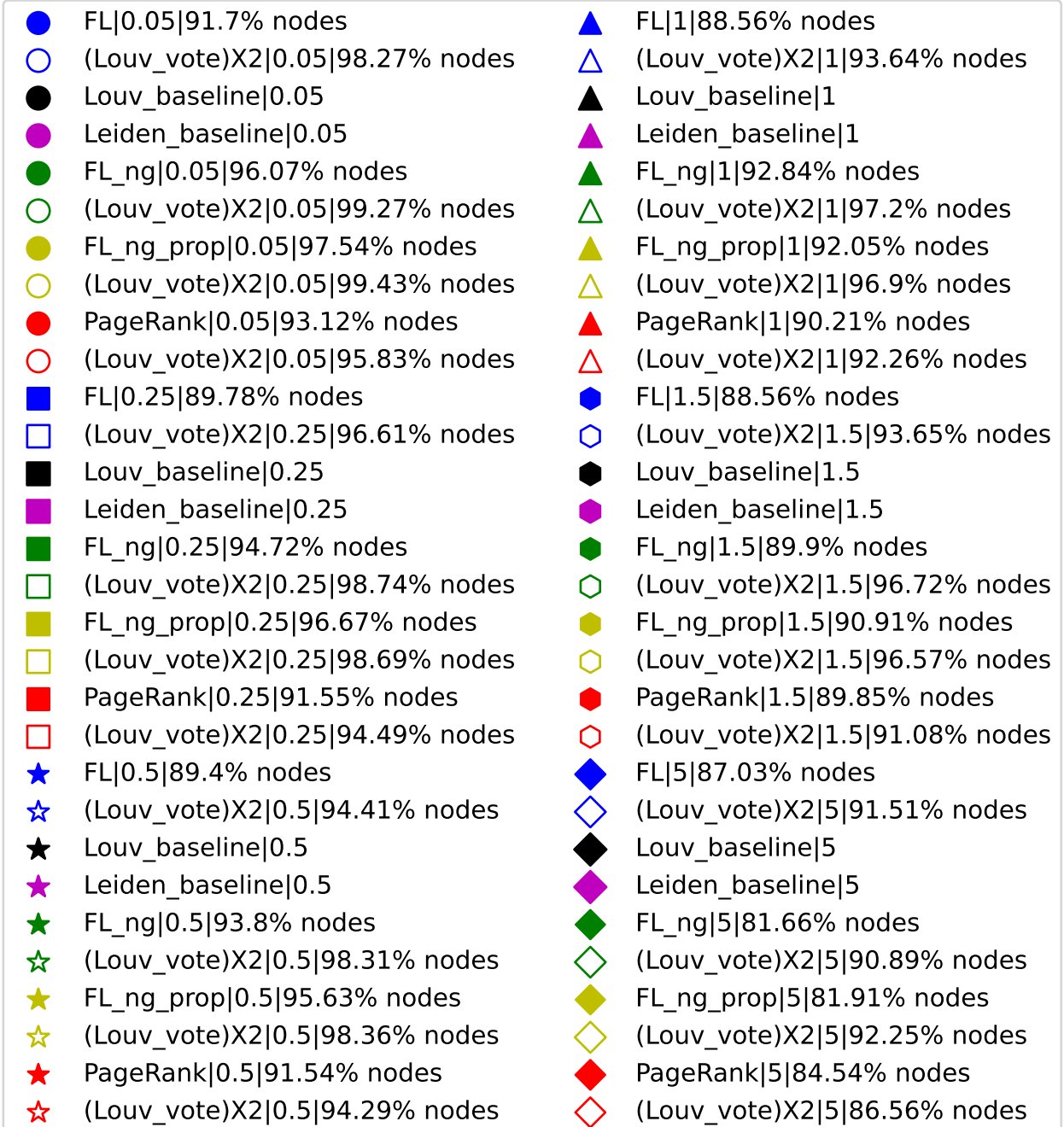
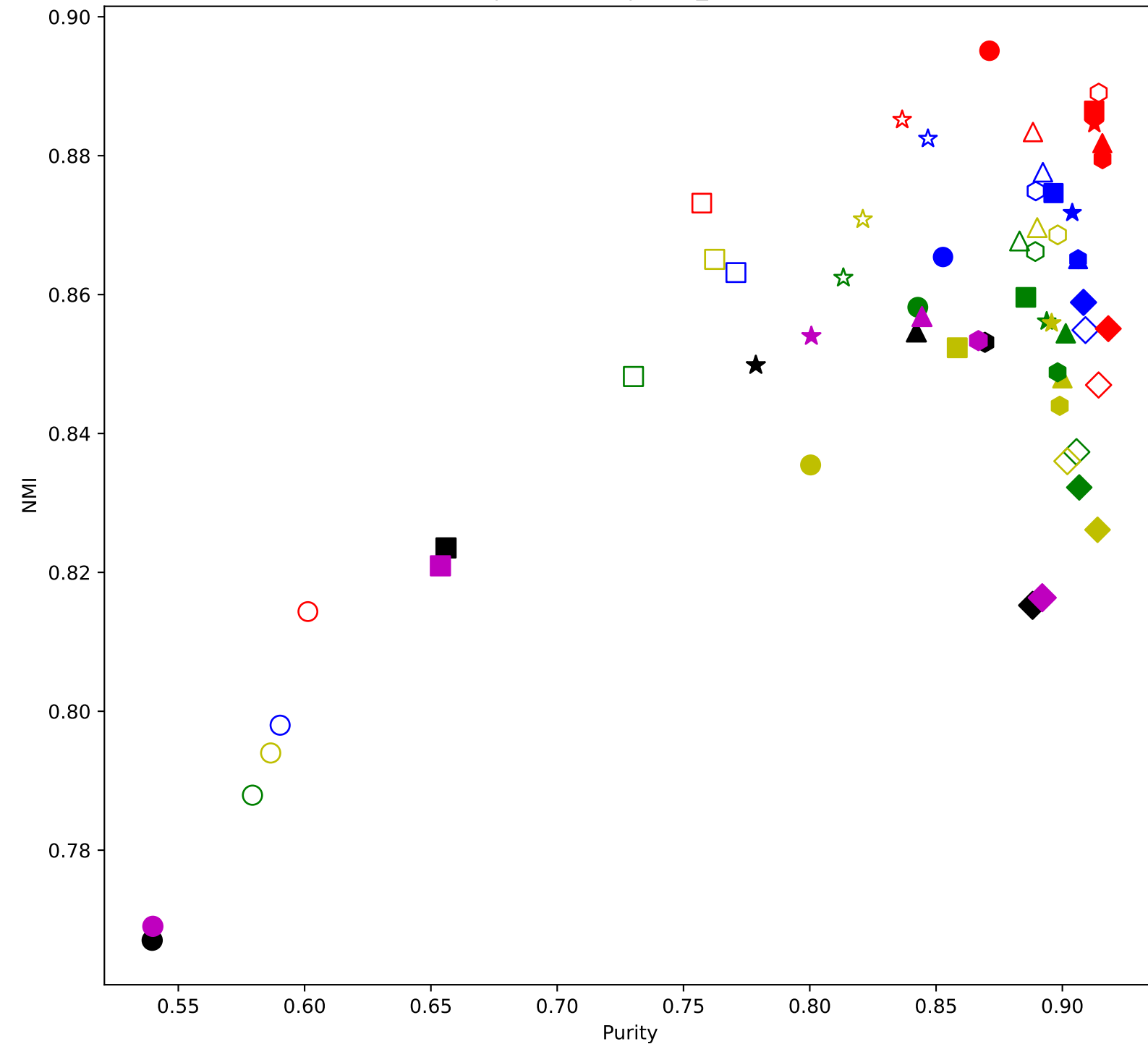
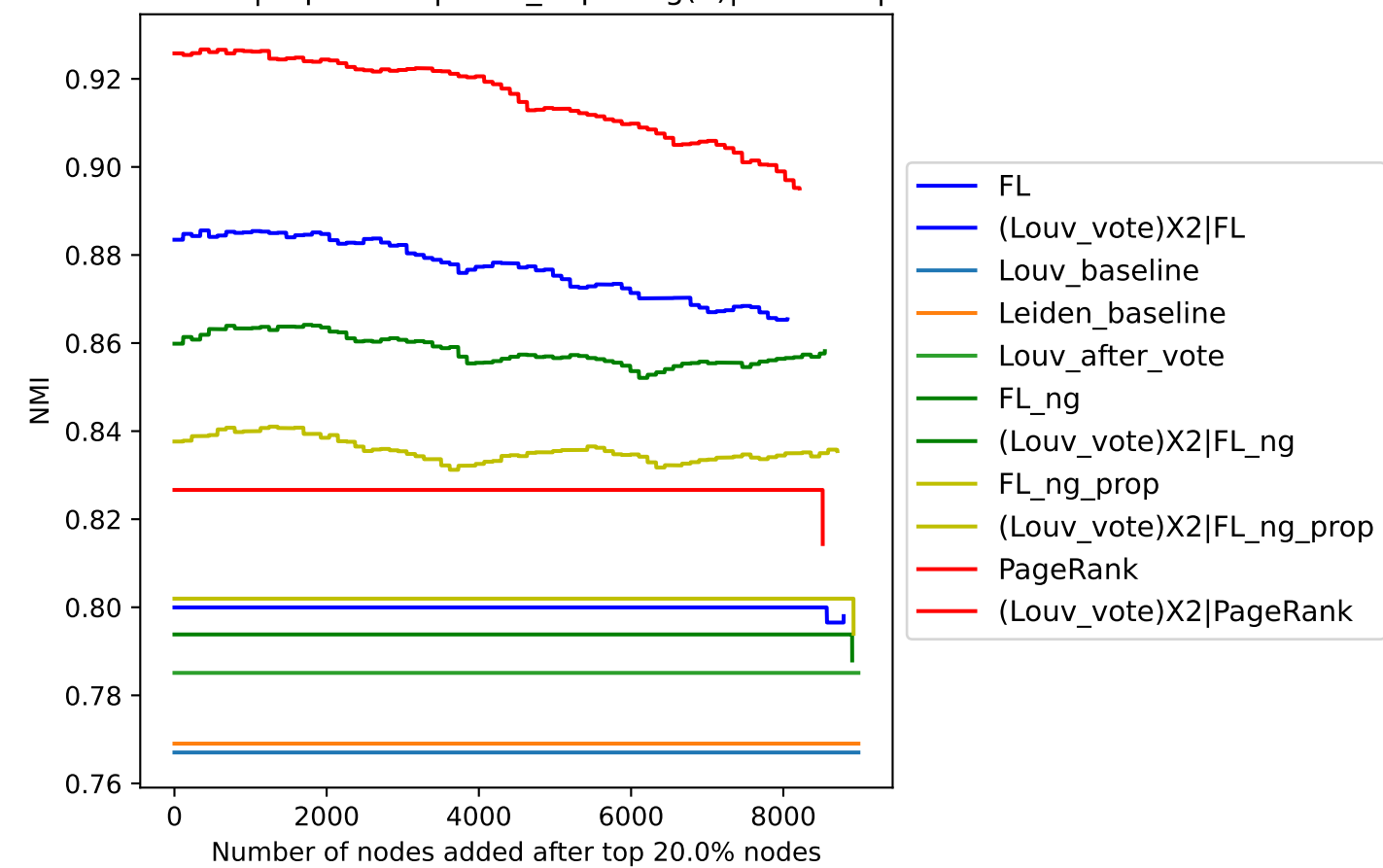


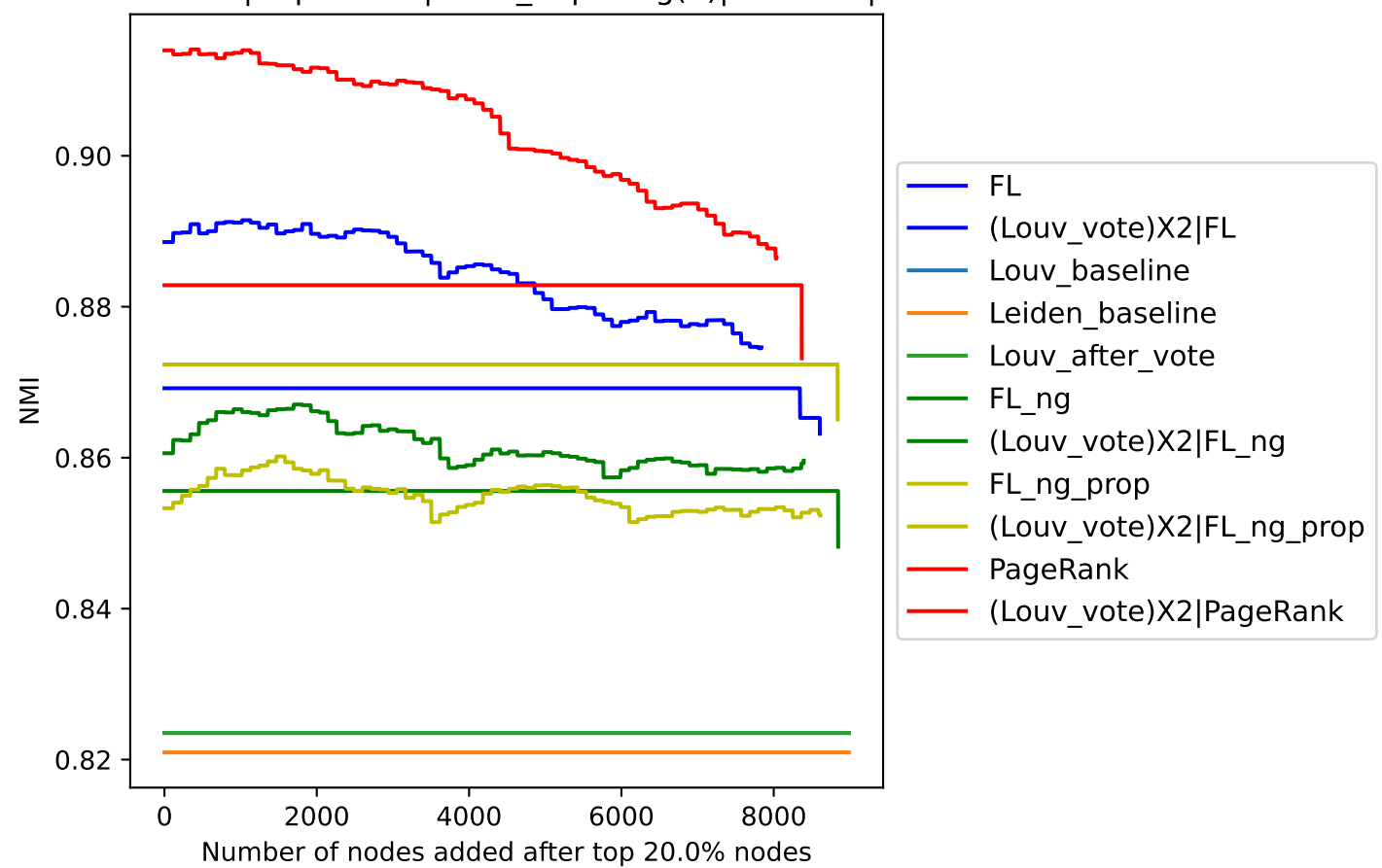
mRNA | top 20.0%| Num_hops: log(n)



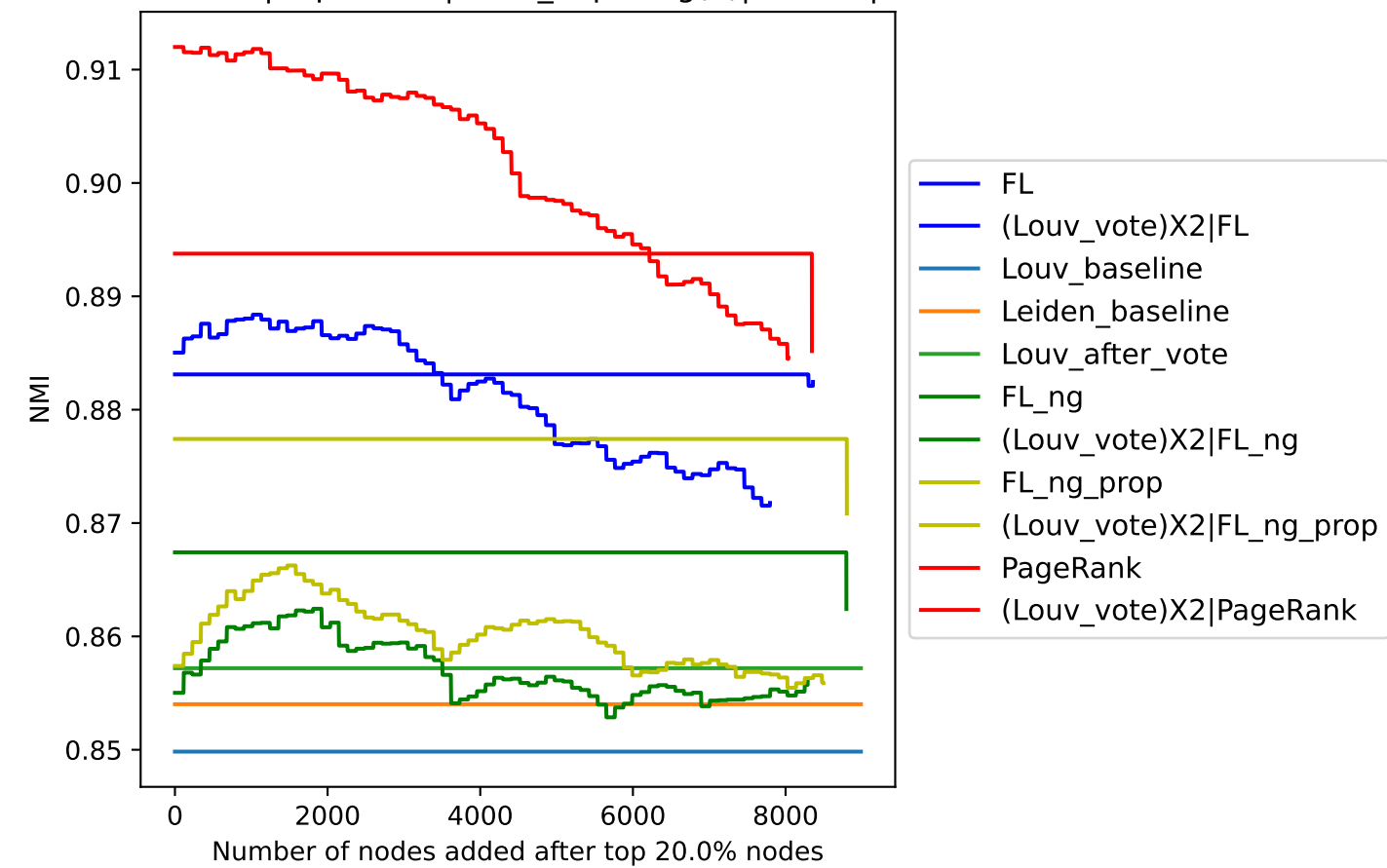
mRNA | top 20.0%| Num_hops: log(n)|res: 0.05|



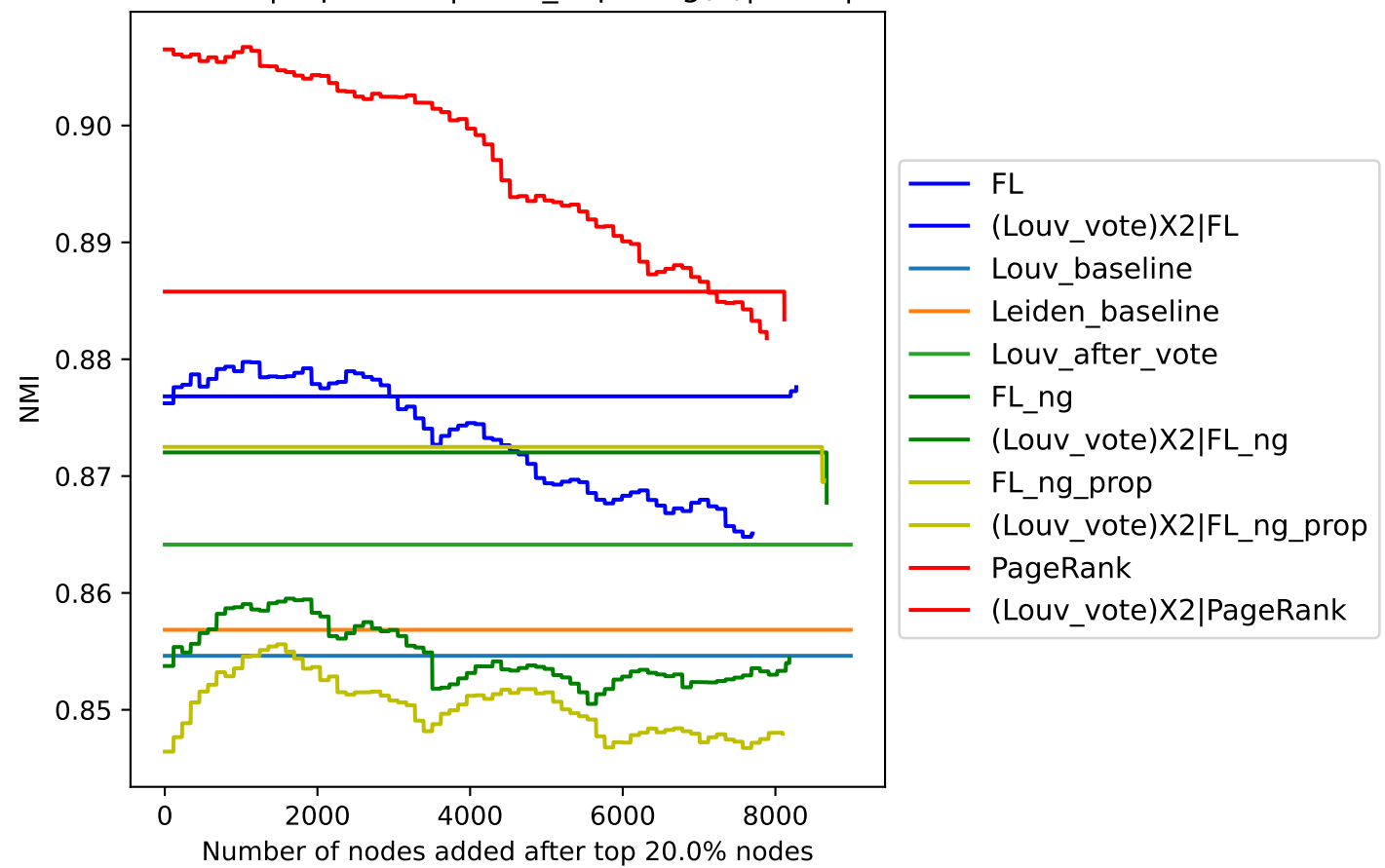
mRNA | top 20.0%| Num_hops: log(n)|res: 0.25|



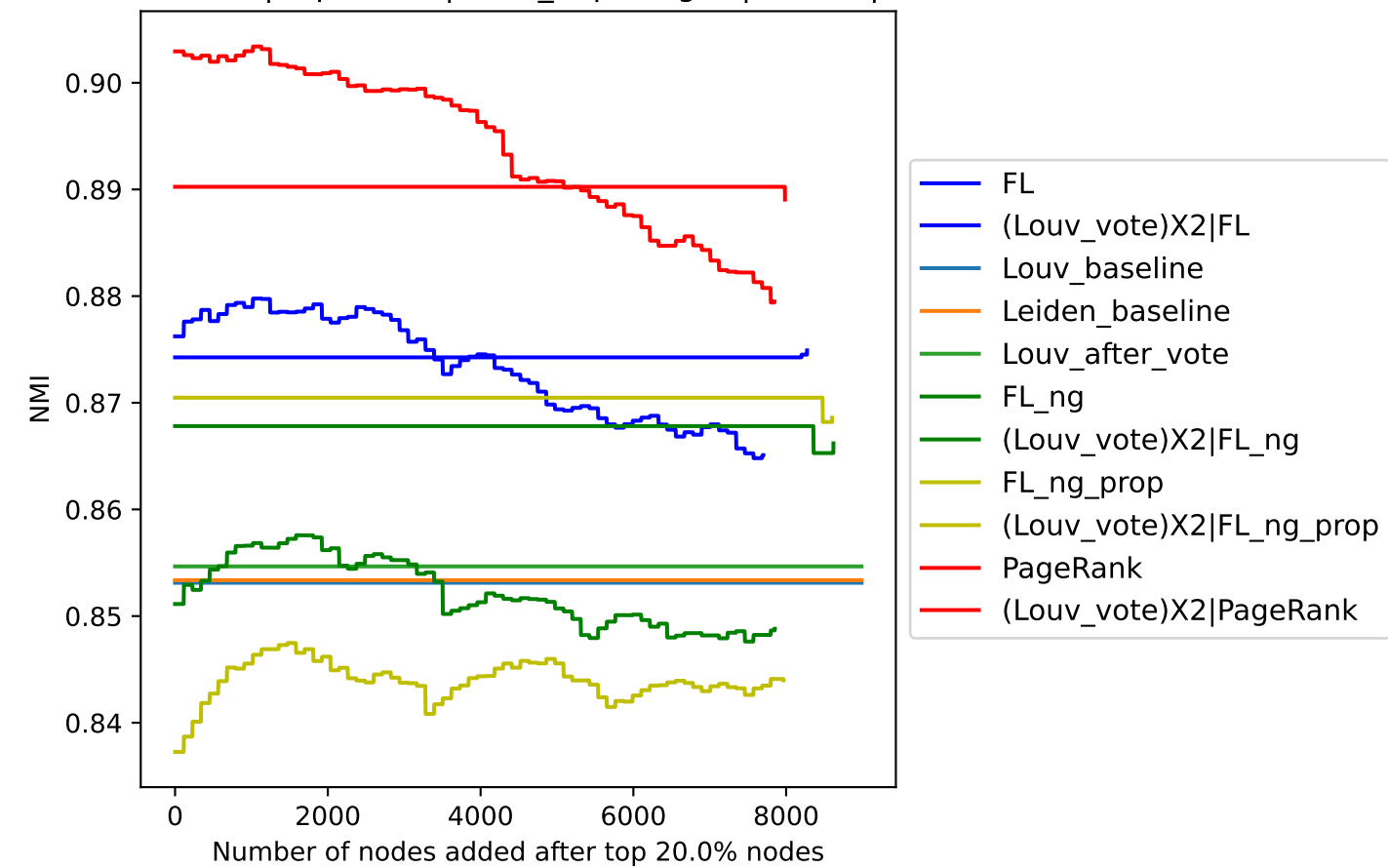
mRNA | top 20.0%| Num_hops: log(n)|res: 0.5|



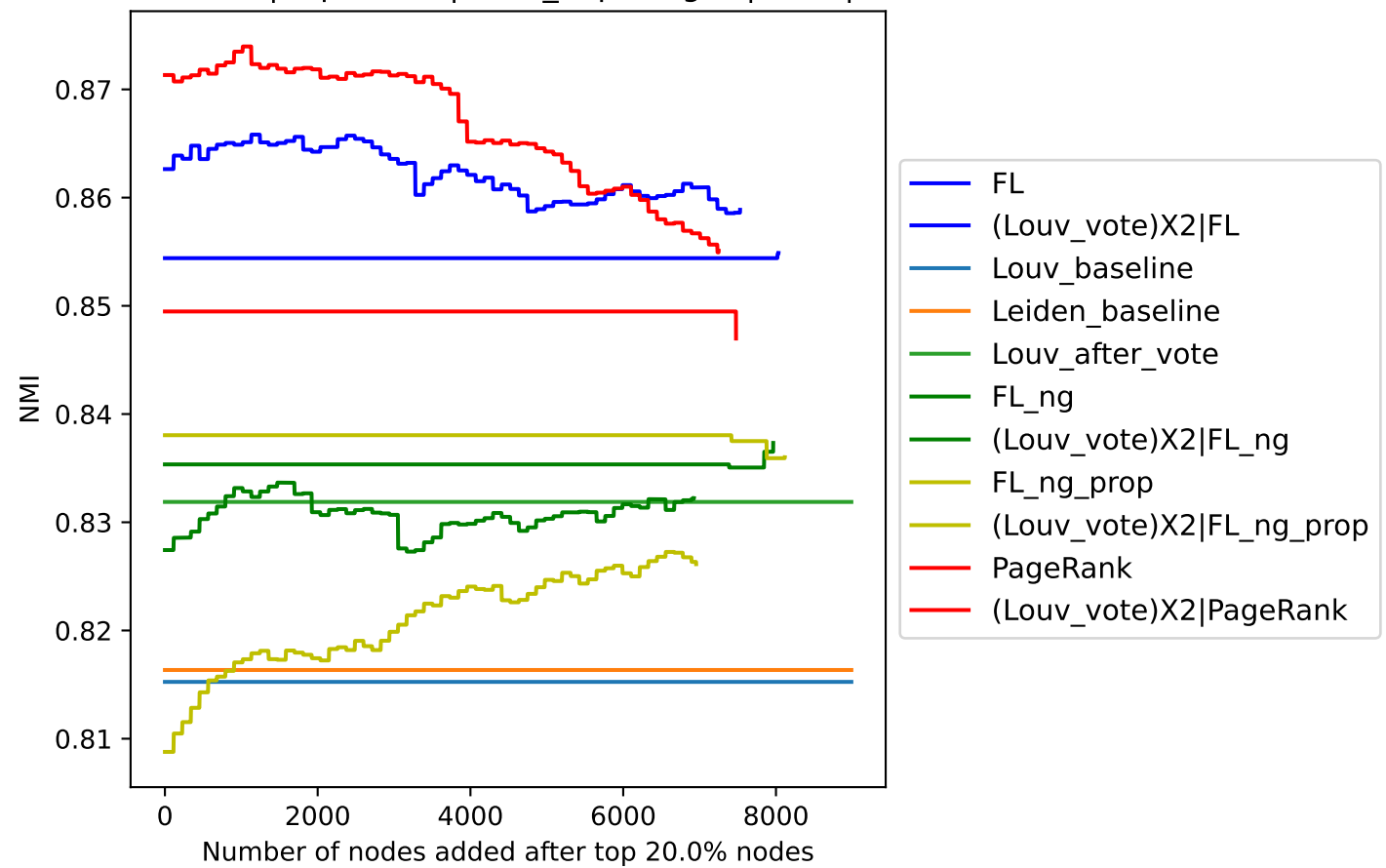
mRNA | top 20.0%| Num_hops: log(n)|res: 1|

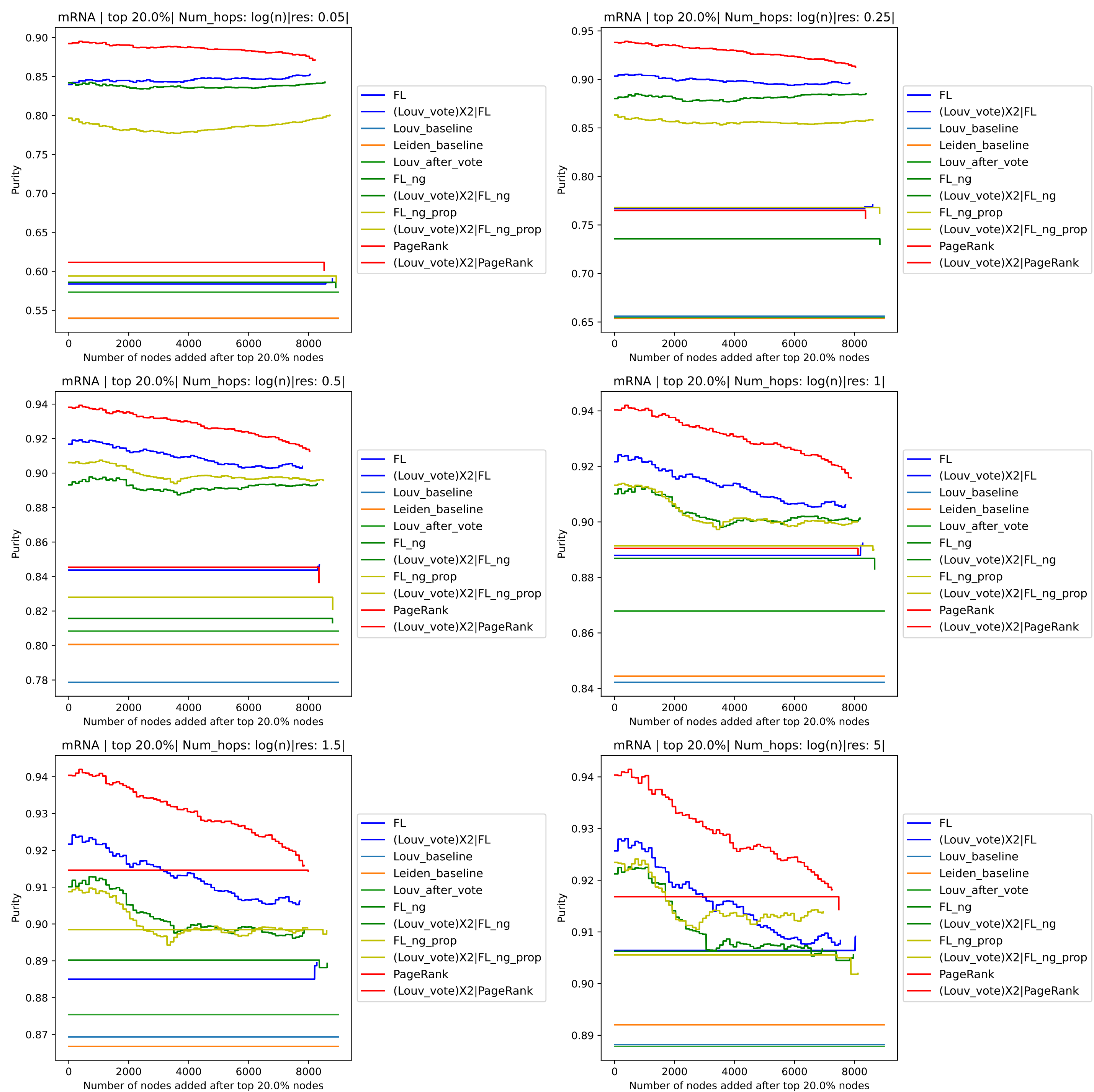


mRNA | top 20.0%| Num_hops: log(n)|res: 1.5|

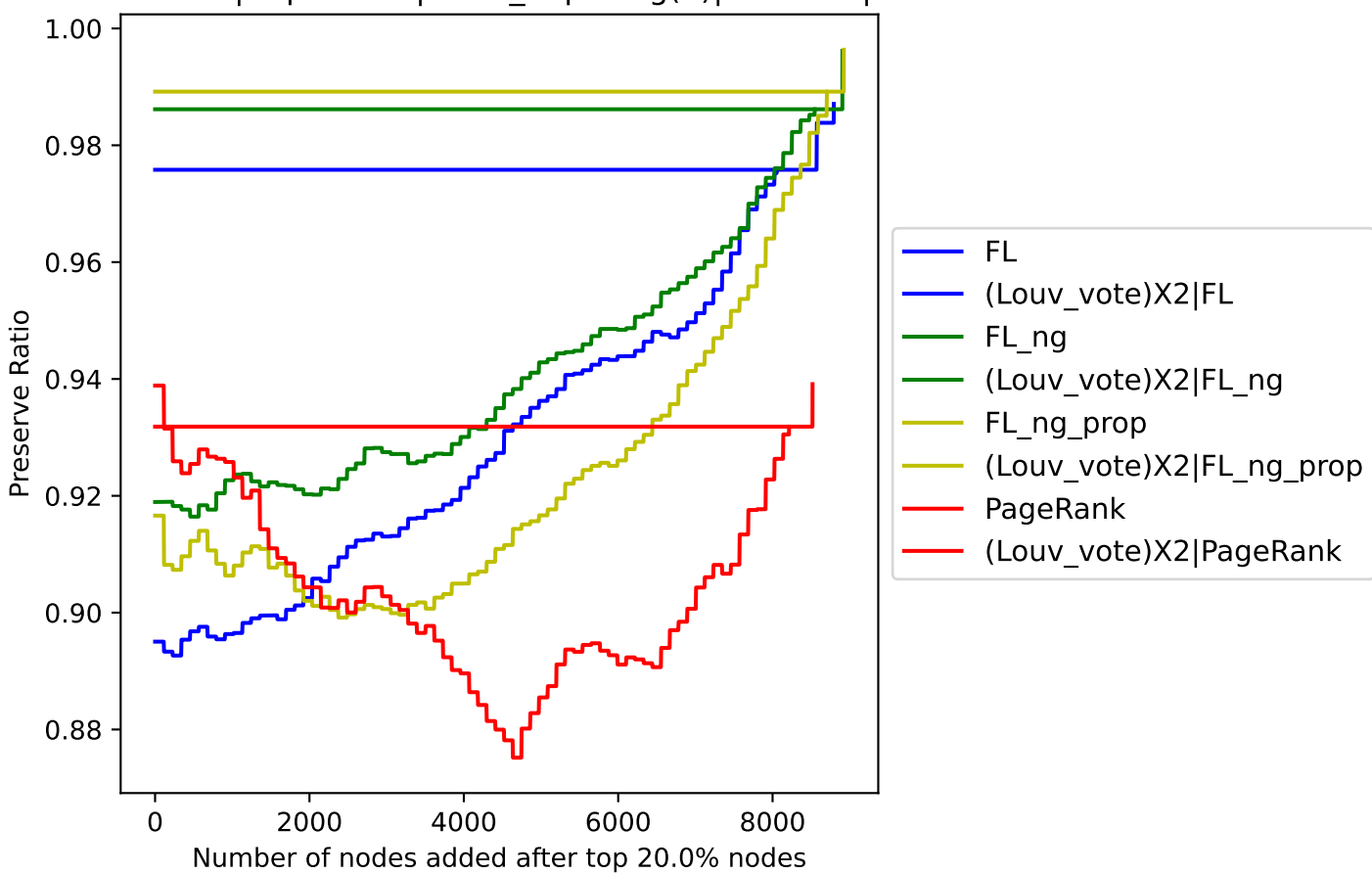


mRNA | top 20.0%| Num_hops: log(n)|res: 5|

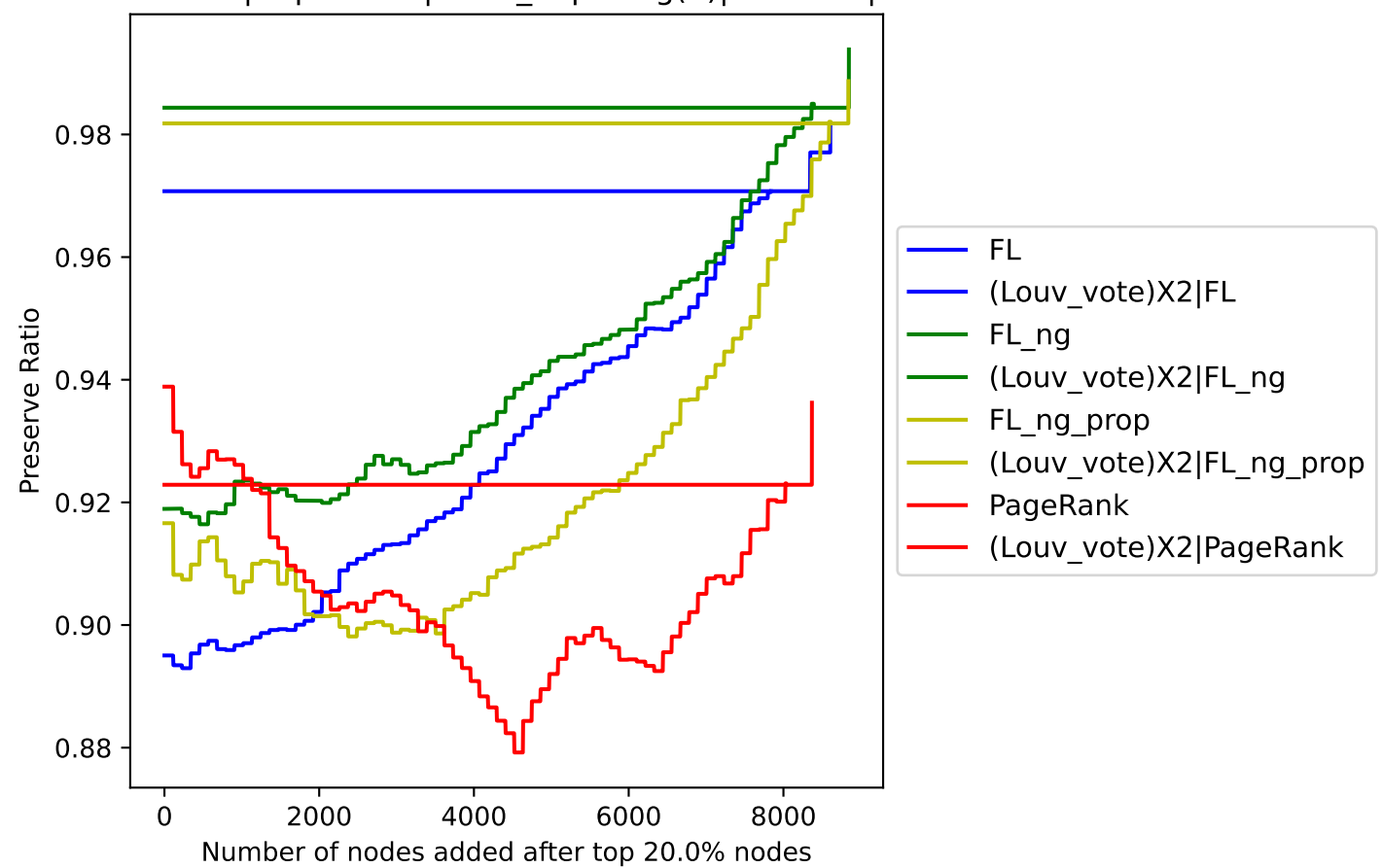




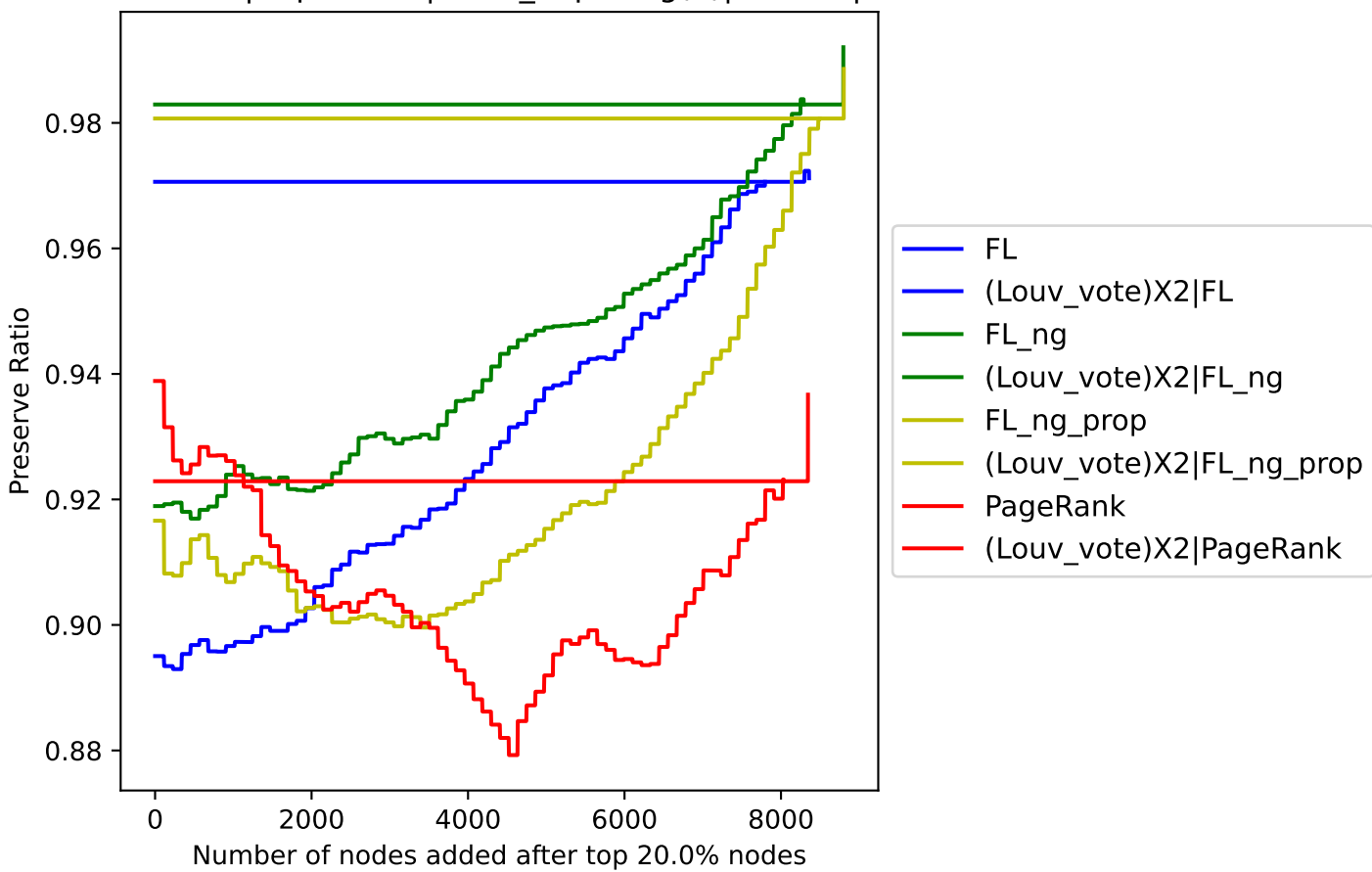
mRNA | top 20.0%| Num_hops: log(n)|res: 0.05|



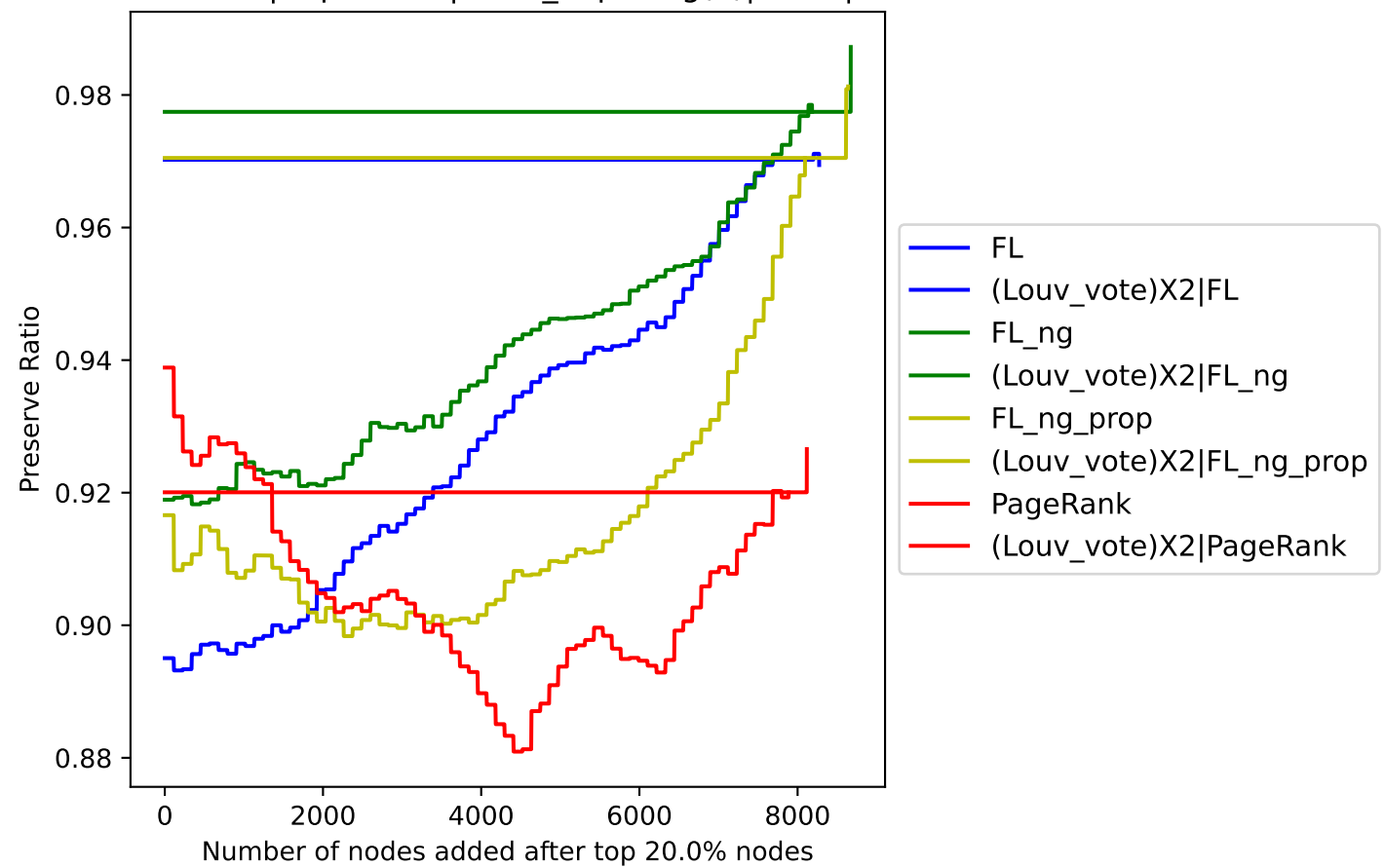
mRNA | top 20.0%| Num_hops: log(n)|res: 0.25|



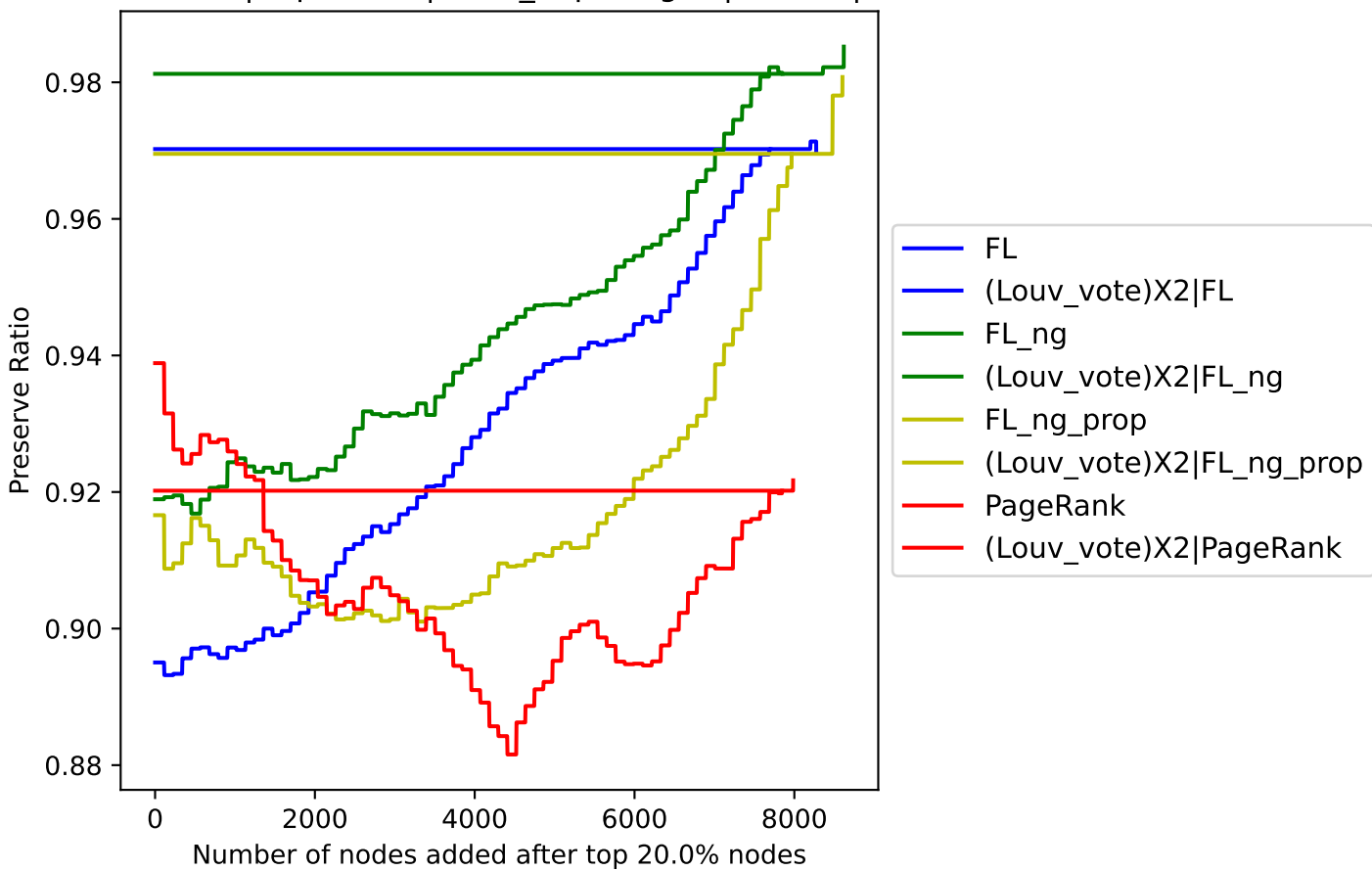
mRNA | top 20.0%| Num_hops: log(n)|res: 0.5|



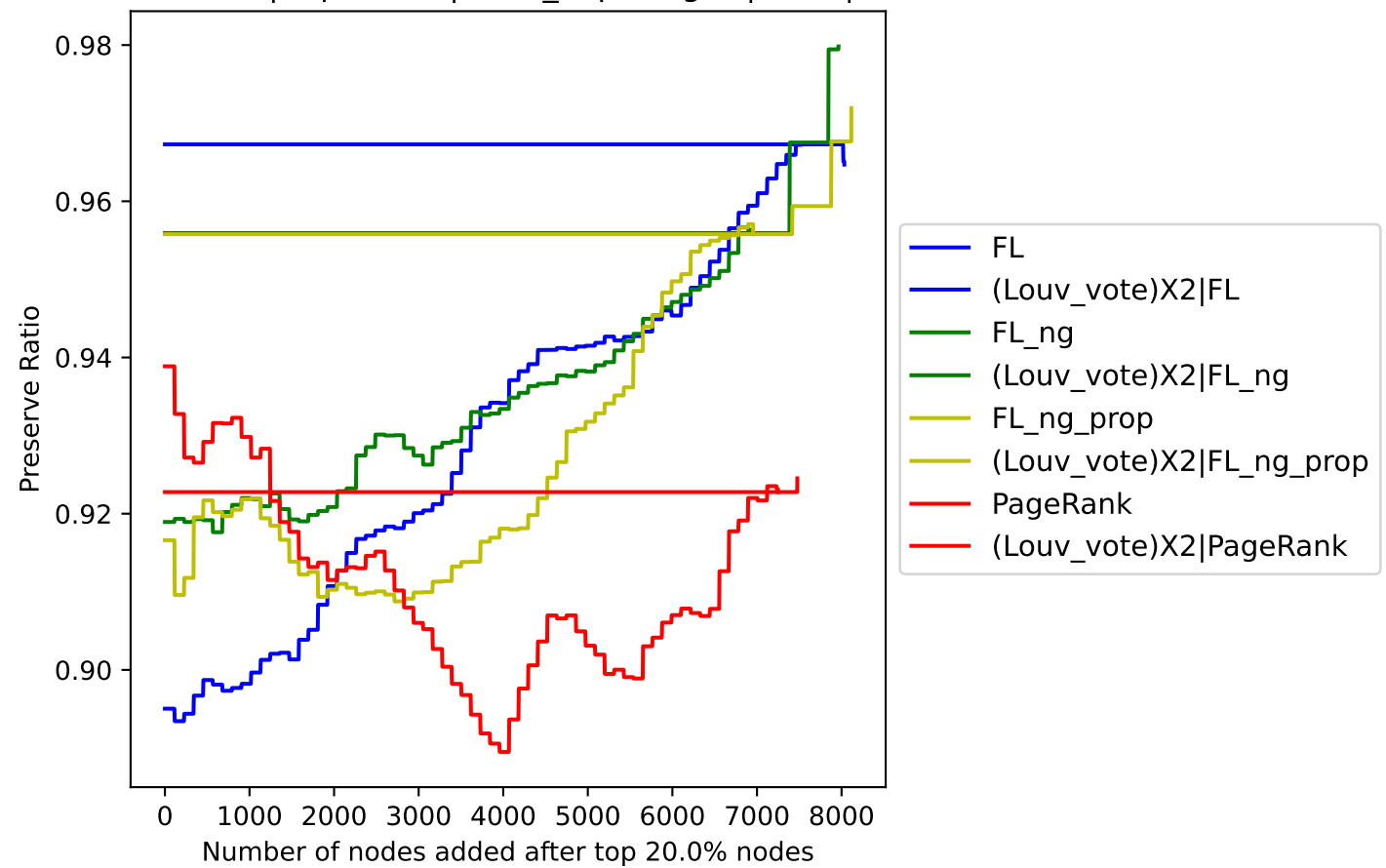
mRNA | top 20.0%| Num_hops: log(n)|res: 1|



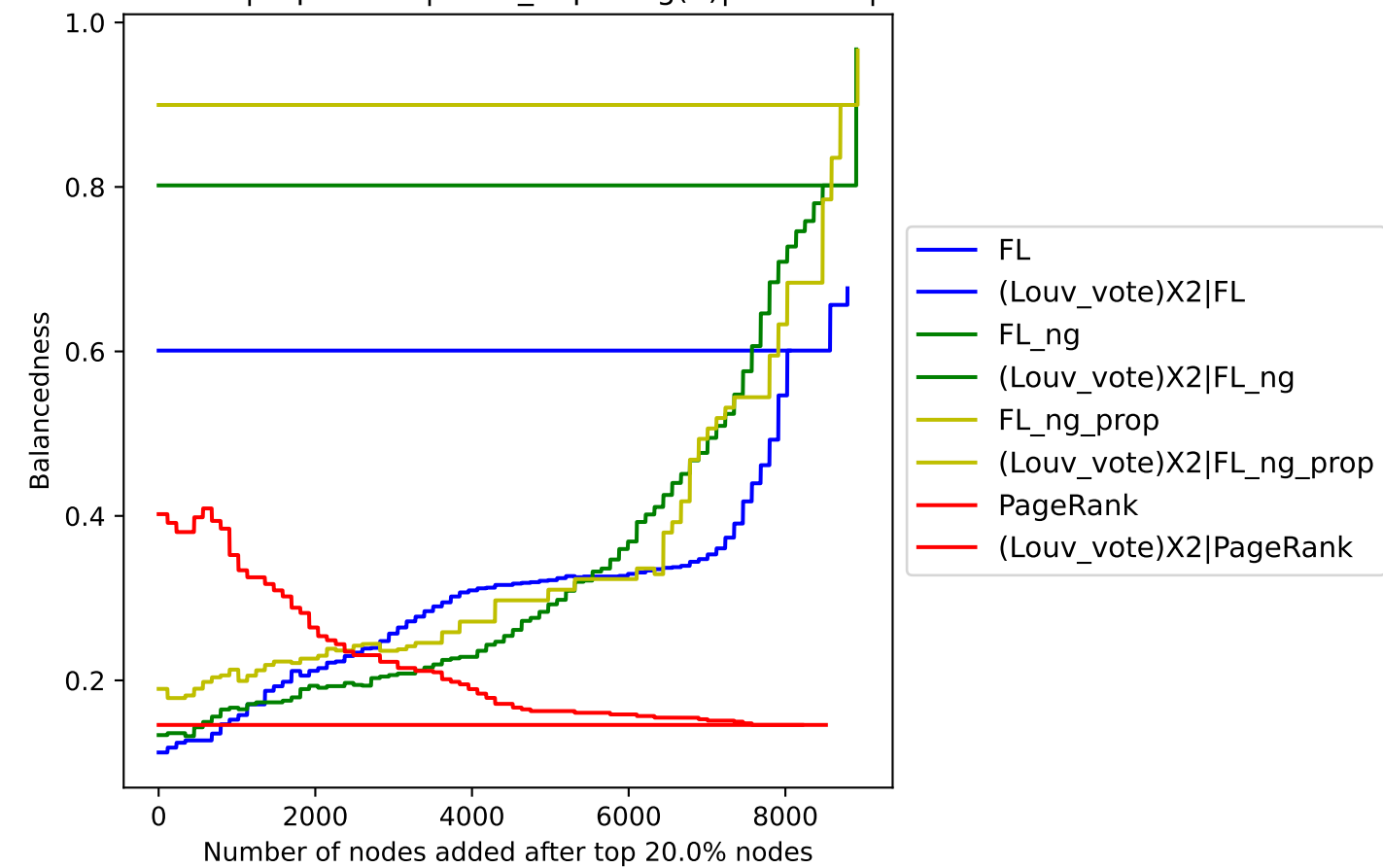
mRNA | top 20.0%| Num_hops: log(n)|res: 1.5|



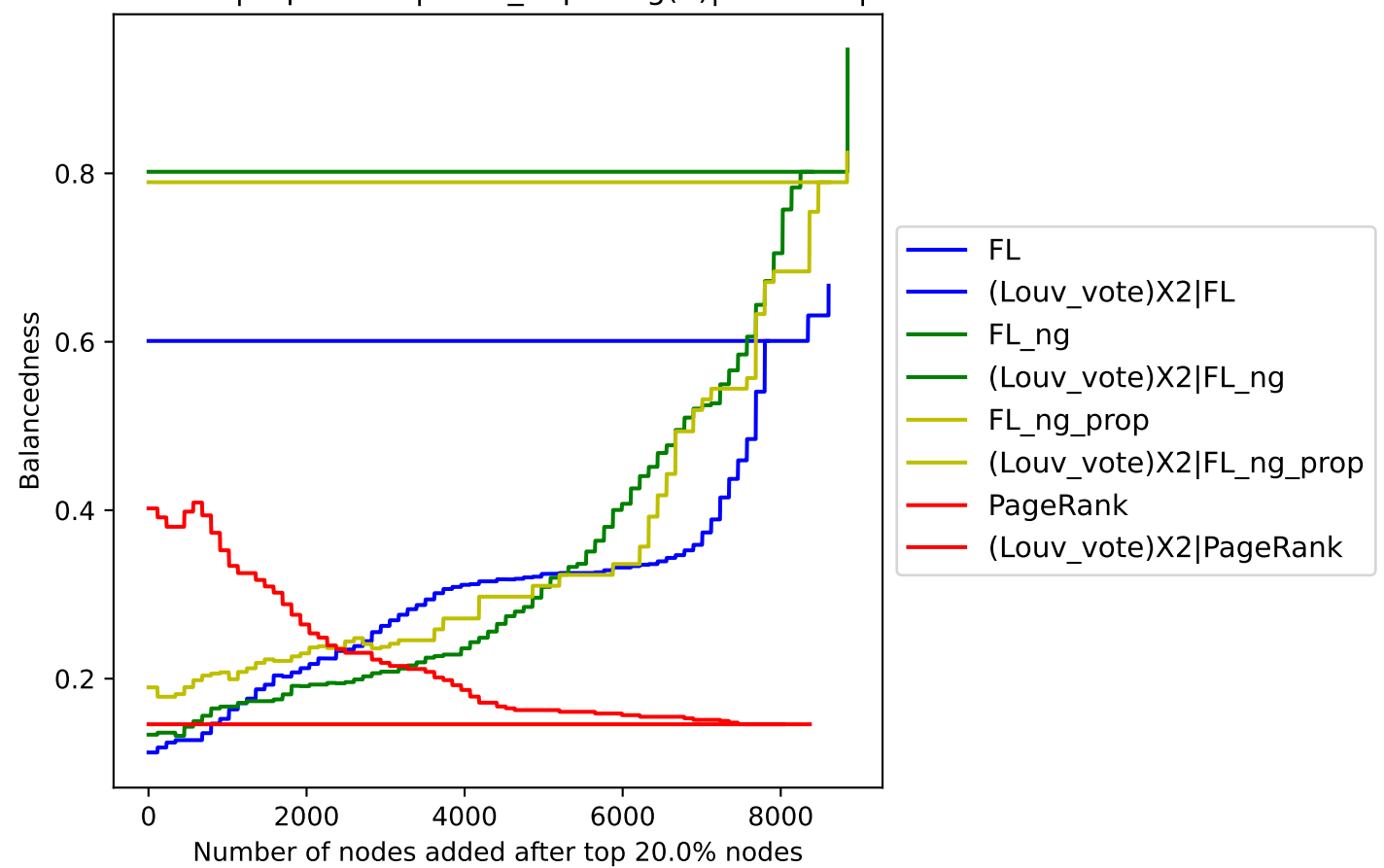
mRNA | top 20.0%| Num_hops: log(n)|res: 5|



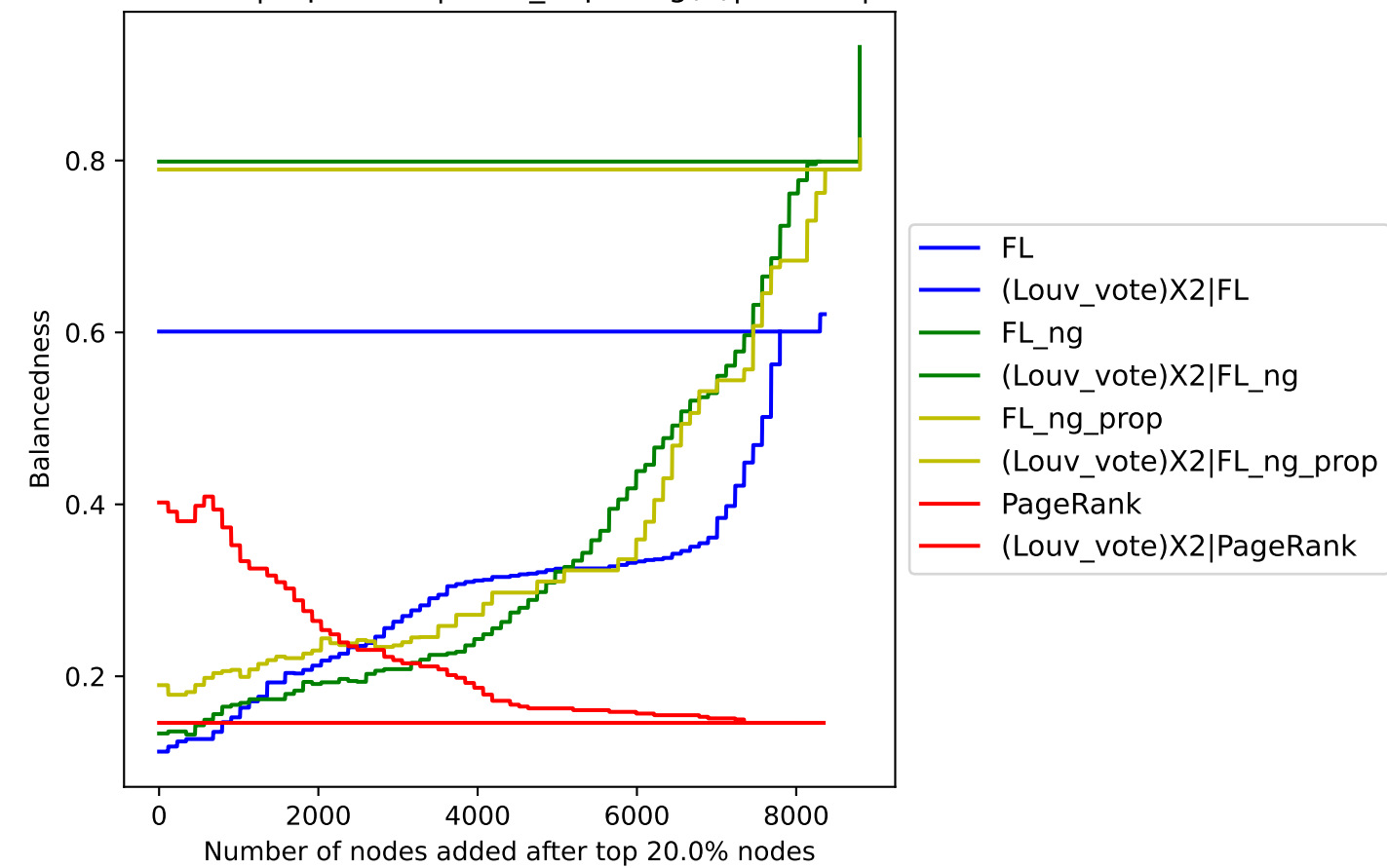
mRNA | top 20.0%| Num_hops: log(n)|res: 0.05|



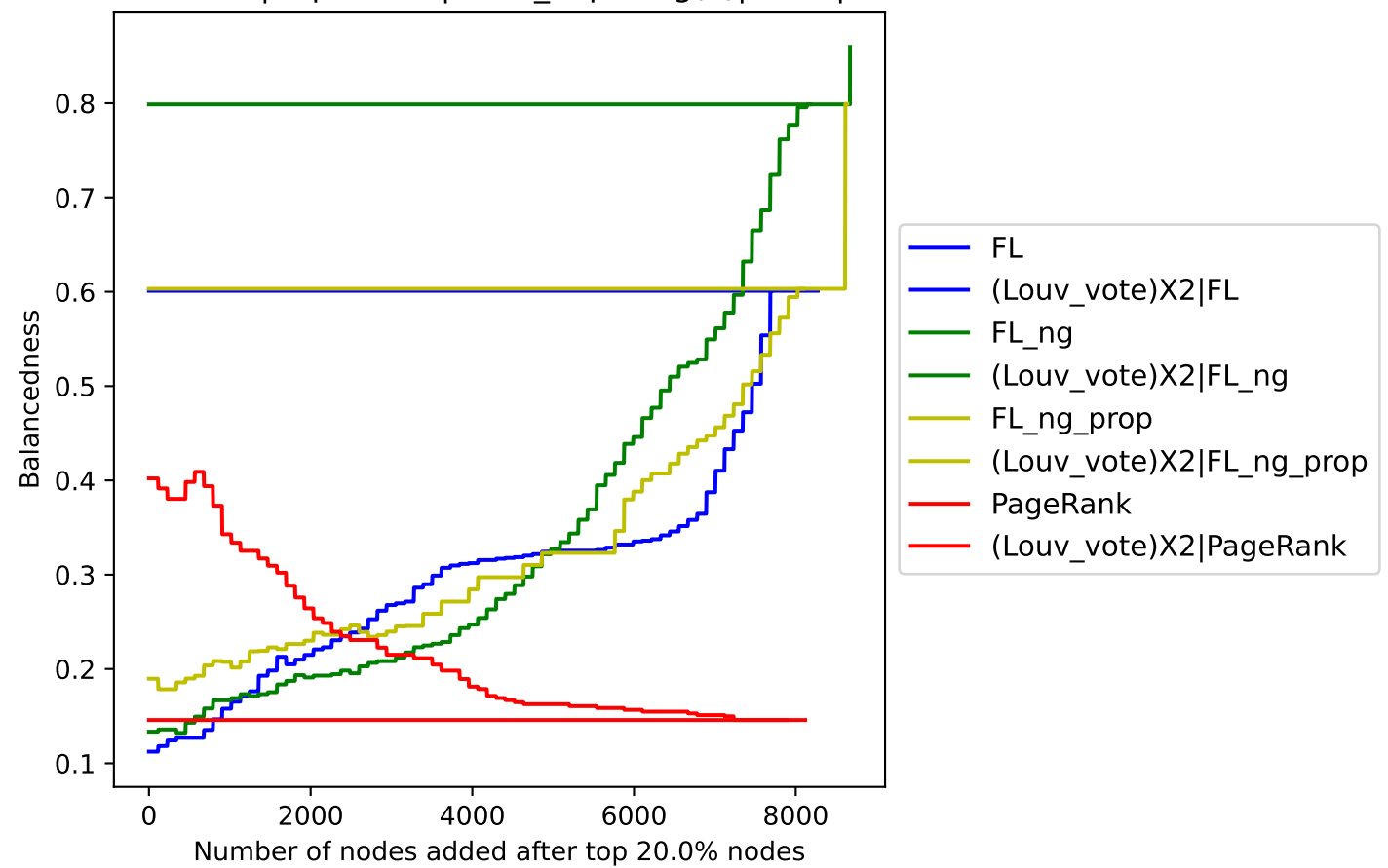
mRNA | top 20.0%| Num_hops: log(n)|res: 0.25|



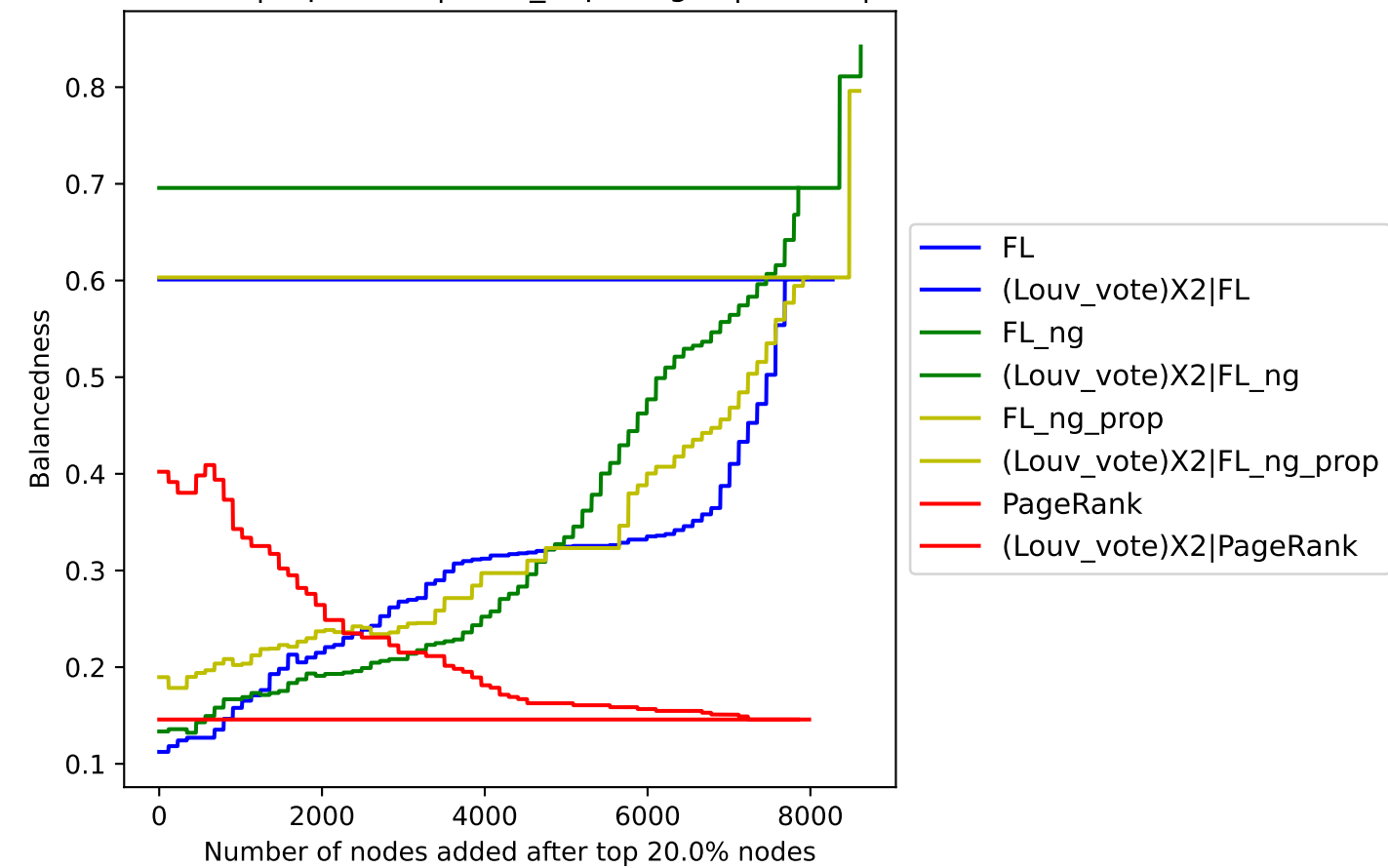
mRNA | top 20.0%| Num_hops: log(n)|res: 0.5|



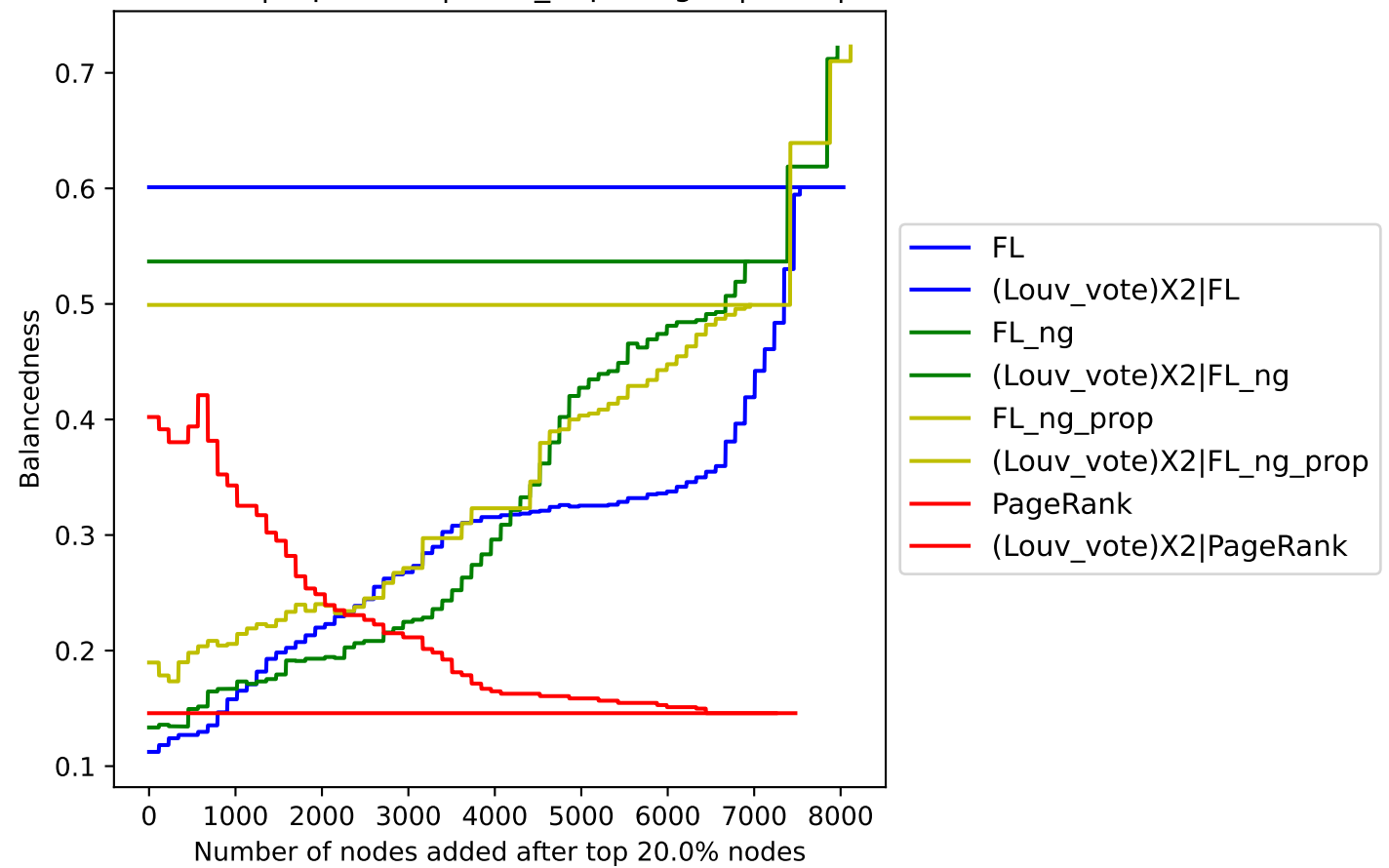
mRNA | top 20.0%| Num_hops: log(n)|res: 1|



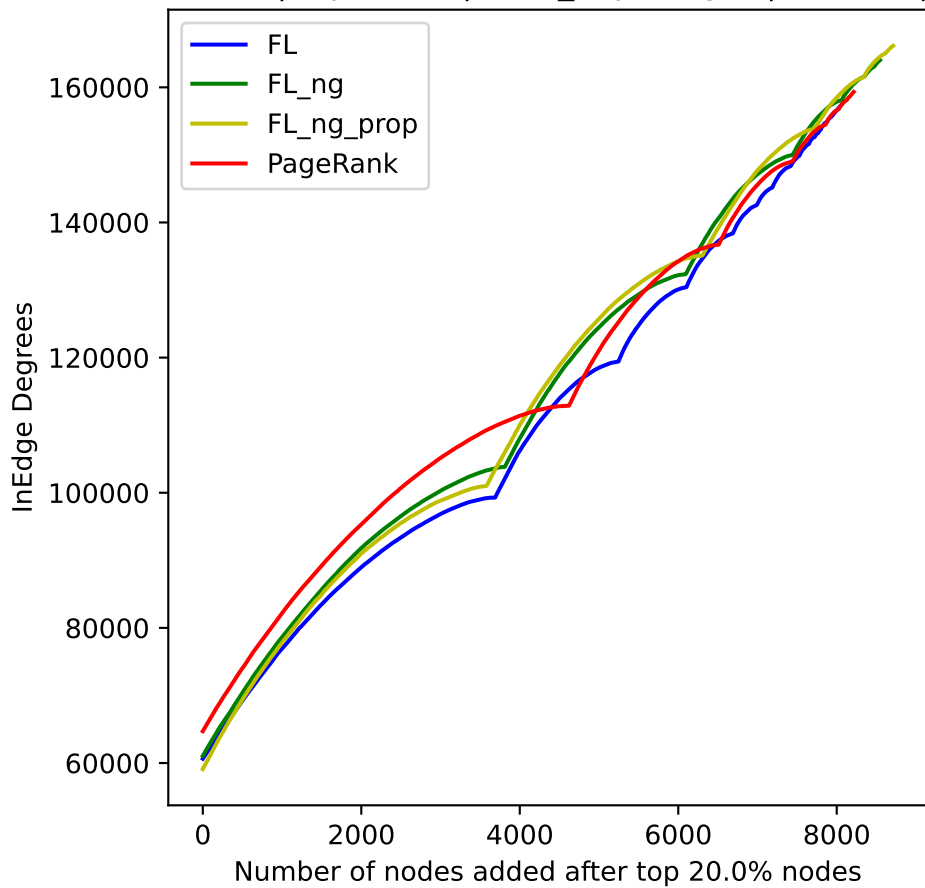
mRNA | top 20.0%| Num_hops: log(n)|res: 1.5|



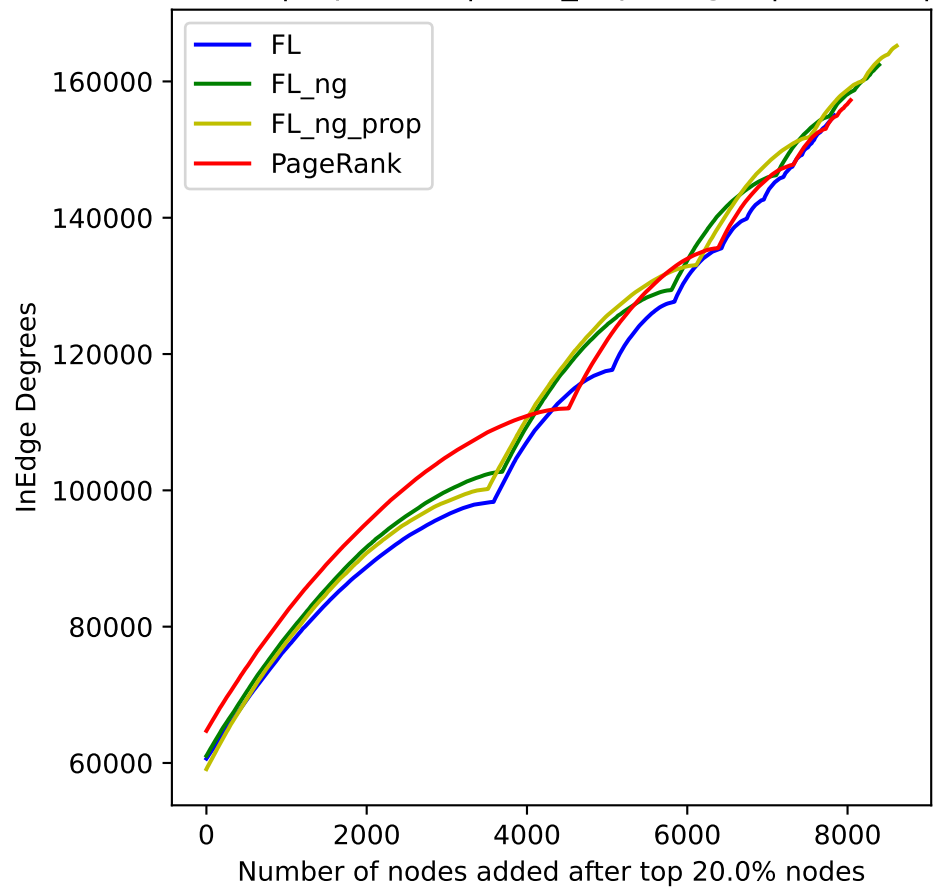
mRNA | top 20.0%| Num_hops: log(n)|res: 5|



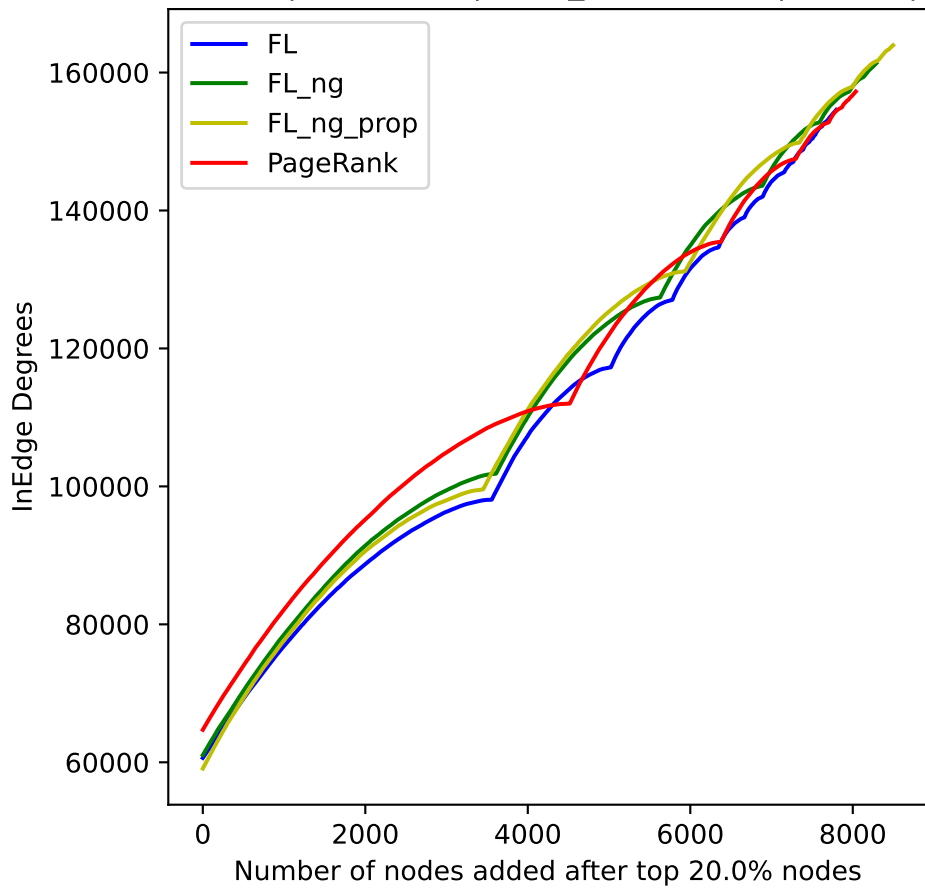
mRNA | top 20.0%| Num_hops: log(n)|res: 0.05|



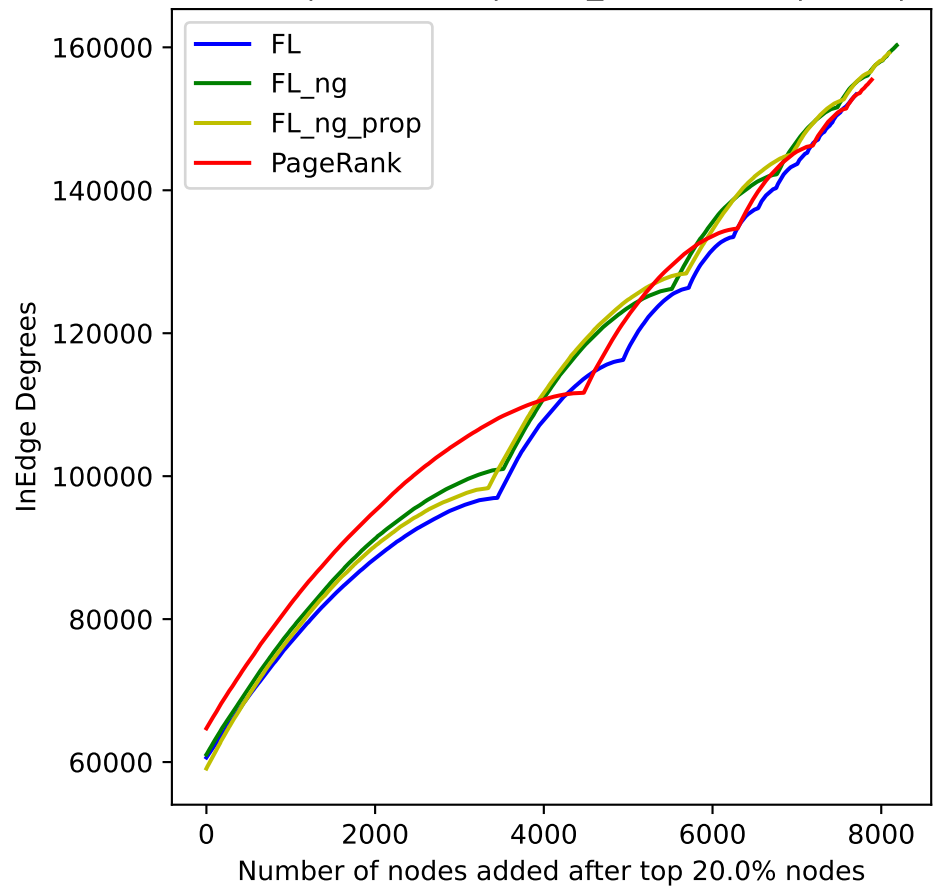
mRNA | top 20.0%| Num_hops: log(n)|res: 0.25|



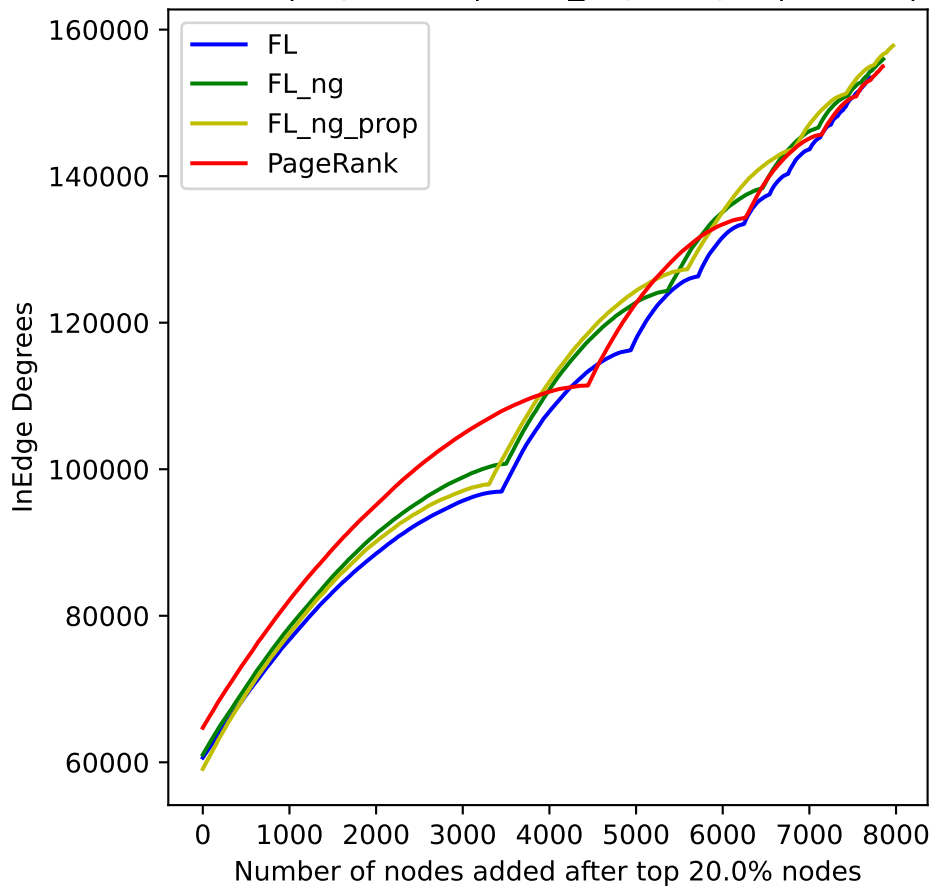
mRNA | top 20.0%| Num_hops: log(n)|res: 0.5|



mRNA | top 20.0%| Num_hops: log(n)|res: 1|



mRNA | top 20.0%| Num_hops: log(n)|res: 1.5|



mRNA | top 20.0%| Num_hops: log(n)|res: 5|

