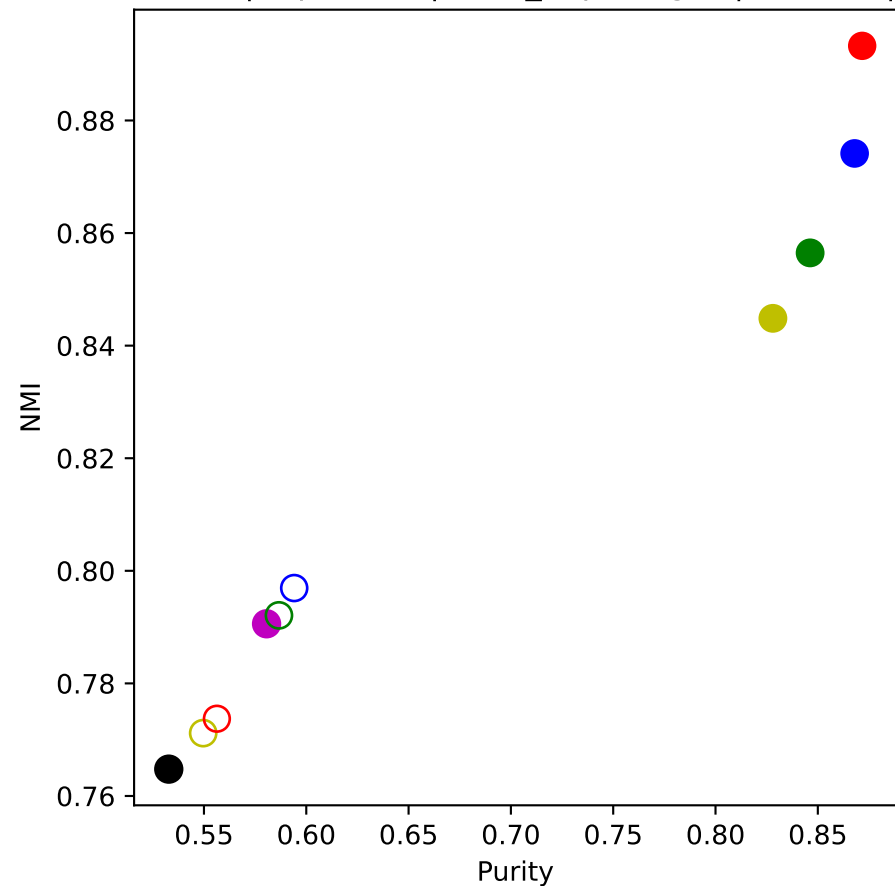
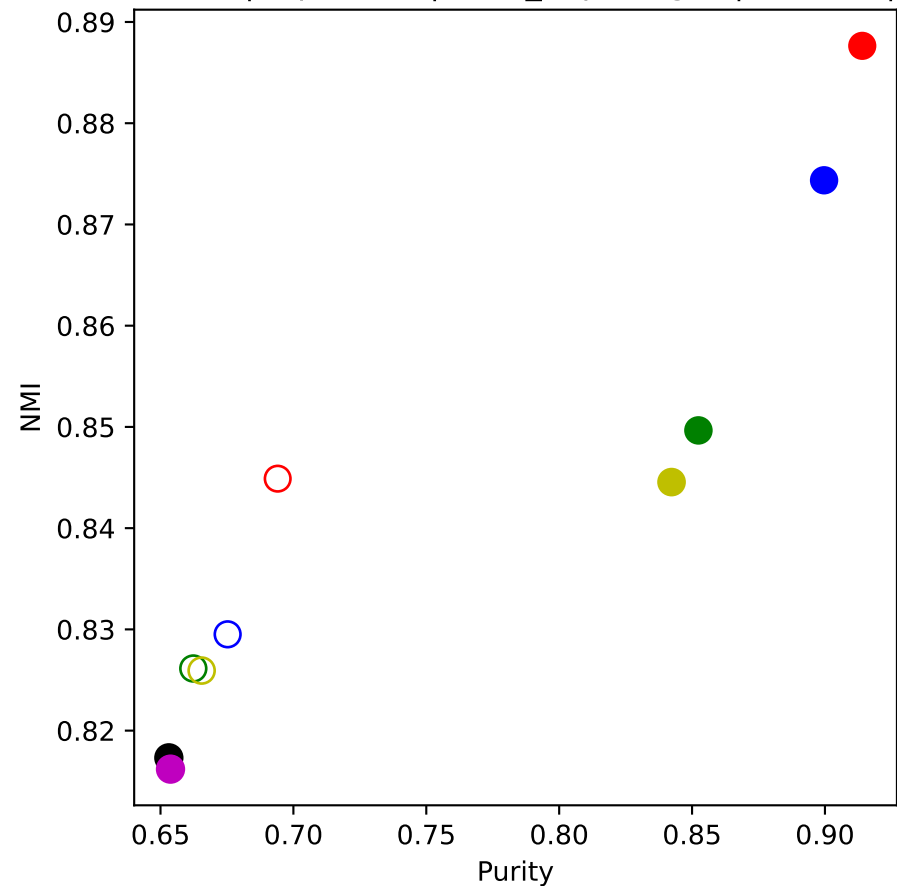


mRNA | top 20.0%| Num_hops: log(n)|res: 0.05|



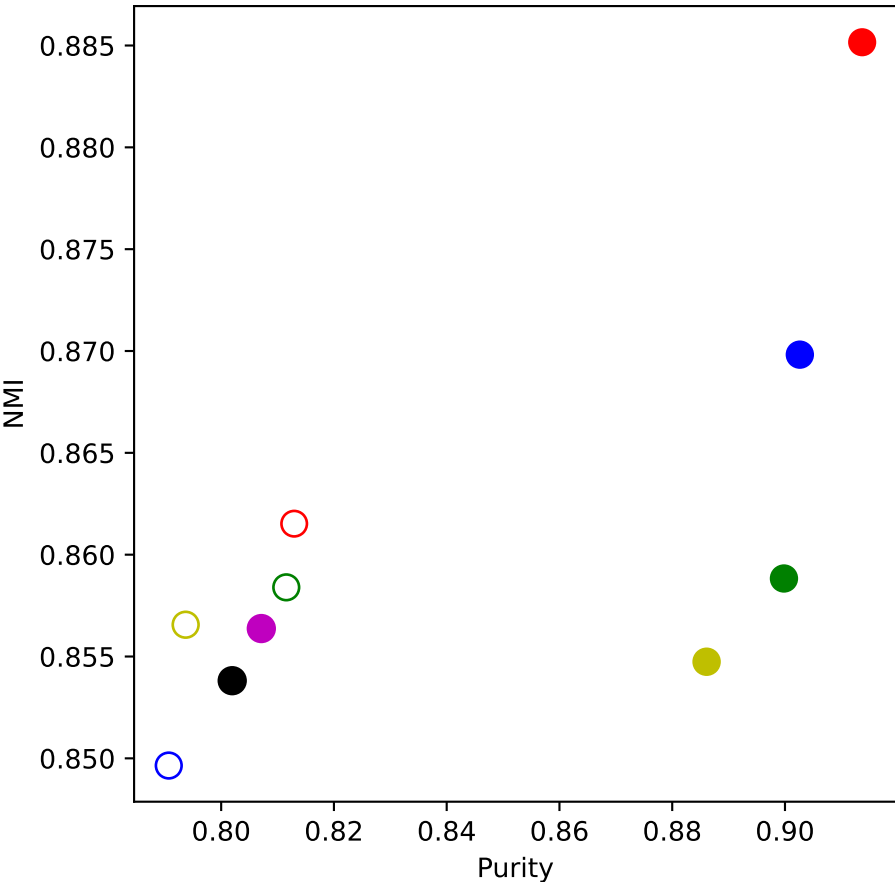
- FL|94.87% nodes
- Eff_checker|94.87% nodes
- Louv_baseline
- Leiden_baseline
- FL_ng|96.22% nodes
- Eff_checker|96.22% nodes
- FL_ng_prop|96.23% nodes
- Eff_checker|96.23% nodes
- PageRank|93.09% nodes
- Eff_checker|93.09% nodes

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



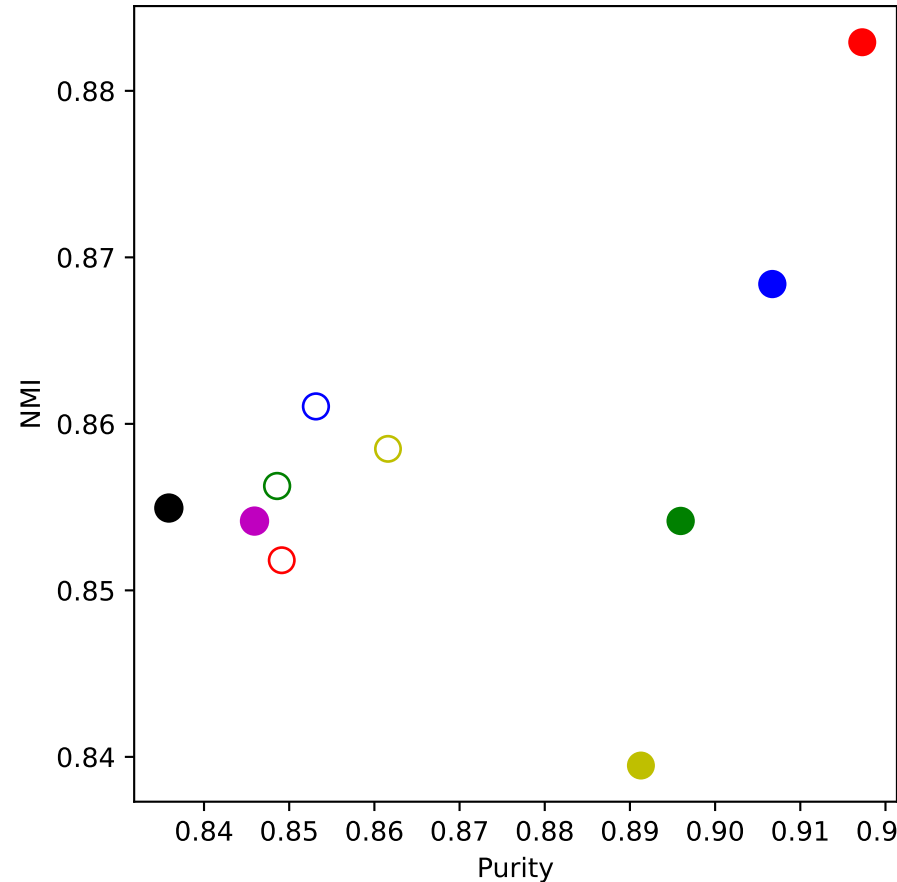
- FL|92.9% nodes
- Eff_checker|92.9% nodes
- Louv_baseline
- Leiden_baseline
- FL_ng|93.96% nodes
- Eff_checker|93.96% nodes
- FL_ng_prop|95.16% nodes
- Eff_checker|95.16% nodes
- PageRank|91.75% nodes
- Eff_checker|91.75% nodes

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



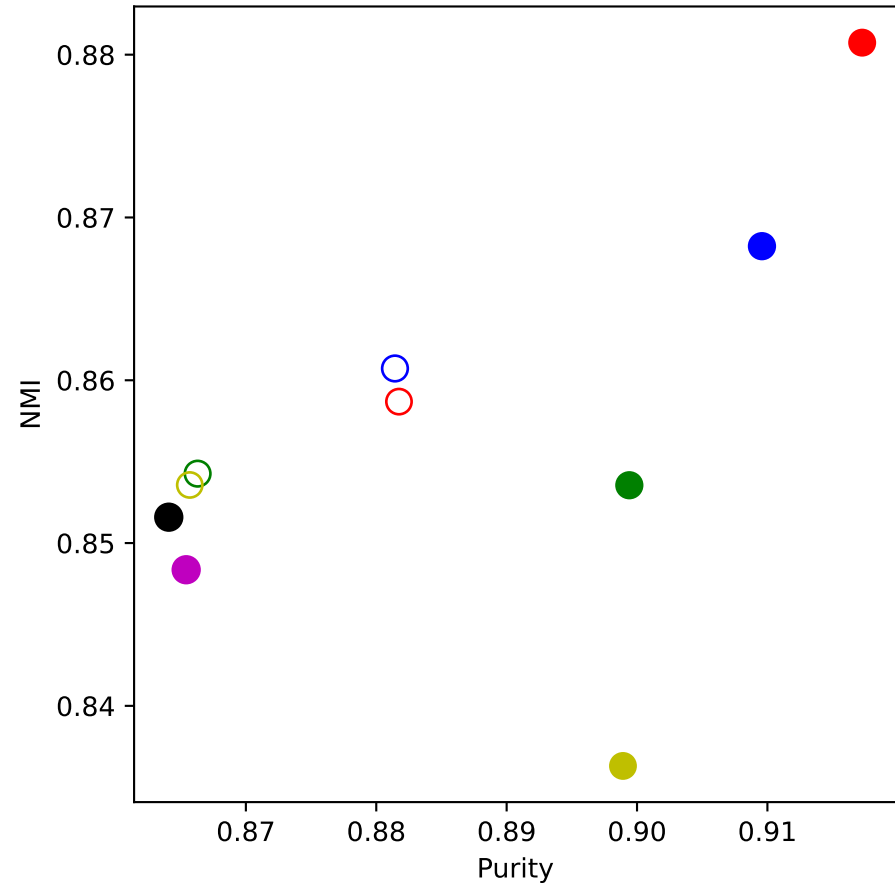
- FL|92.89% nodes
- Eff_checker|92.89% nodes
- Louv_baseline
- Leiden_baseline
- FL_ng|92.67% nodes
- Eff_checker|92.67% nodes
- FL_ng_prop|93.55% nodes
- Eff_checker|93.55% nodes
- PageRank|91.28% nodes
- Eff_checker|91.28% nodes

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



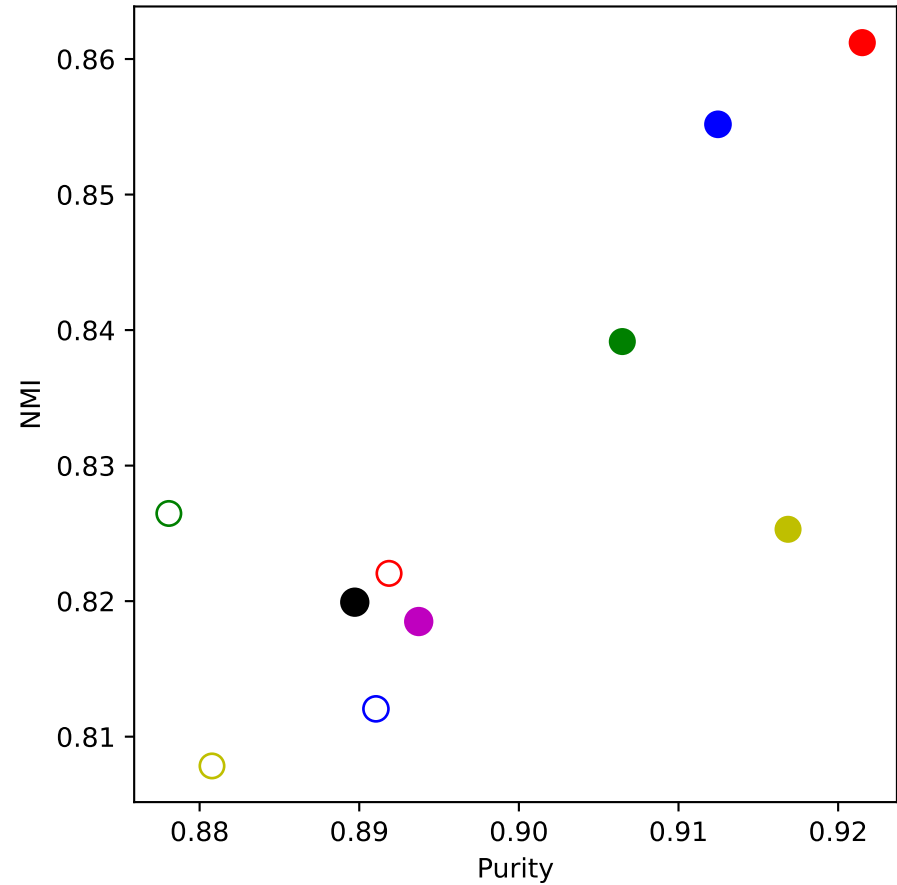
- FL|92.37% nodes
- Eff_checker|92.37% nodes
- Louv_baseline
- Leiden_baseline
- FL_ng|92.83% nodes
- Eff_checker|92.83% nodes
- FL_ng_prop|90.24% nodes
- Eff_checker|90.24% nodes
- PageRank|90.39% nodes
- Eff_checker|90.39% nodes

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



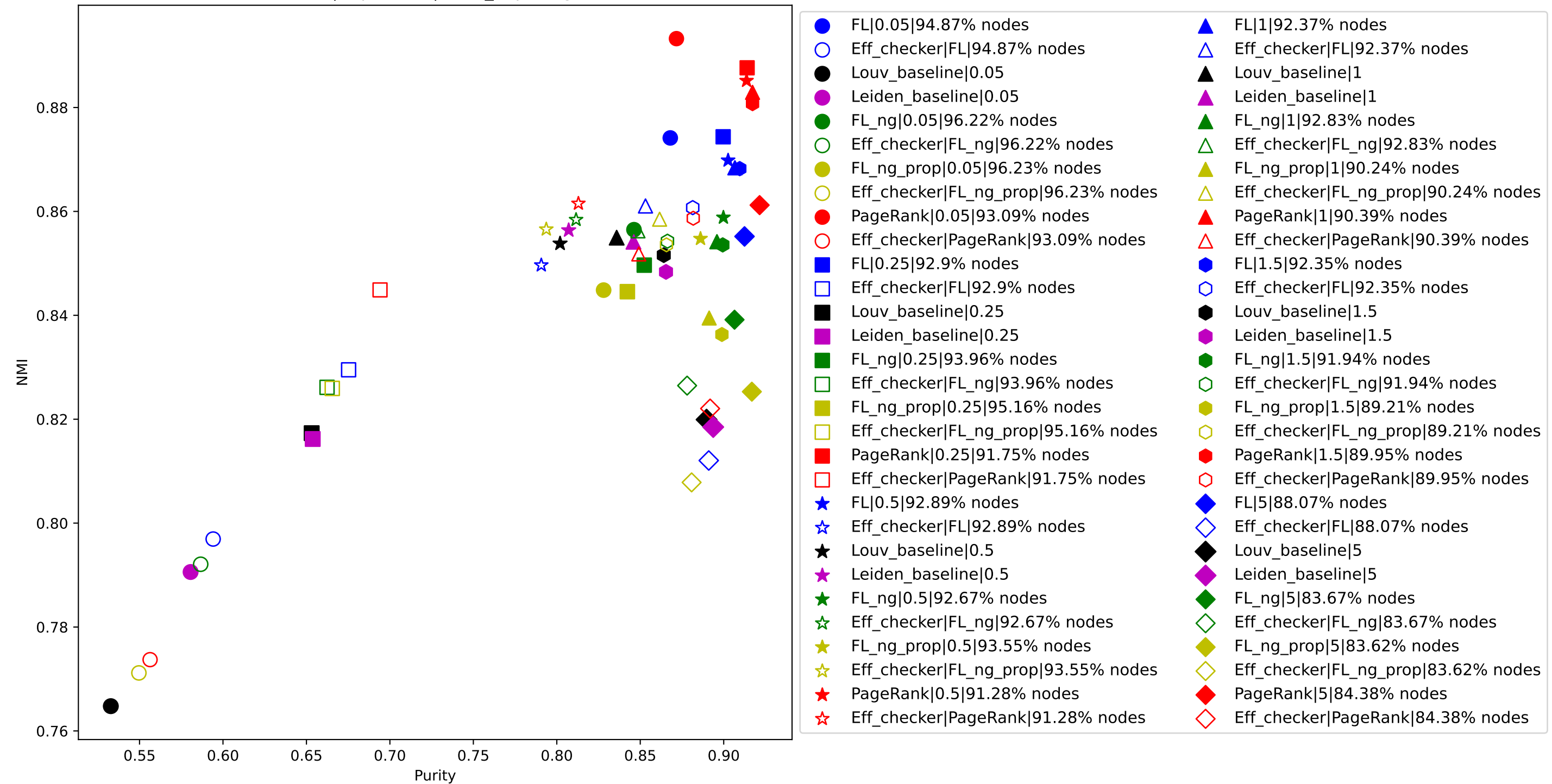
- FL|92.35% nodes
- Eff_checker|92.35% nodes
- Louv_baseline
- Leiden_baseline
- FL_ng|91.94% nodes
- Eff_checker|91.94% nodes
- FL_ng_prop|89.21% nodes
- Eff_checker|89.21% nodes
- PageRank|89.95% nodes
- Eff_checker|89.95% nodes

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

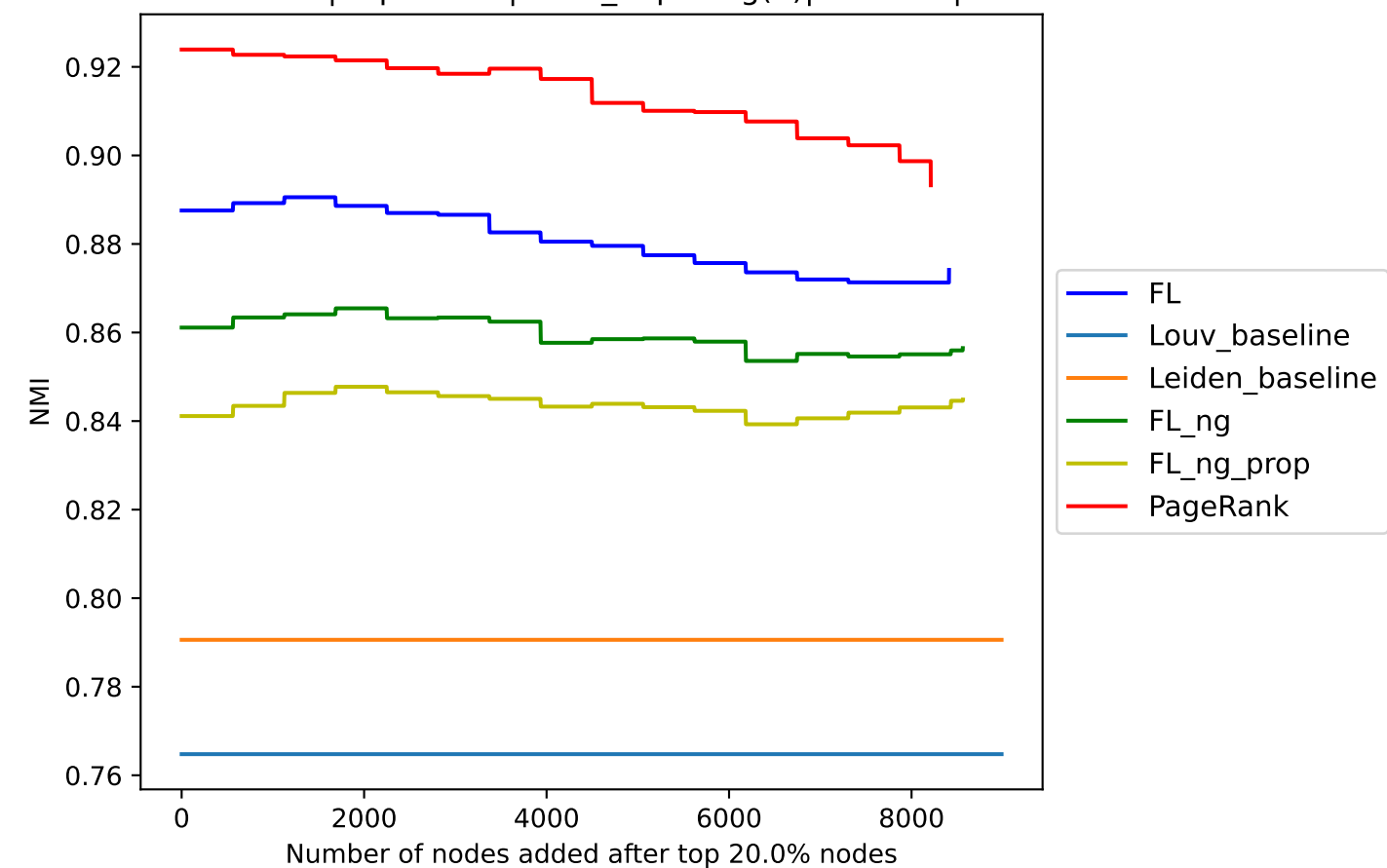


- FL|88.07% nodes
- Eff_checker|88.07% nodes
- Louv_baseline
- Leiden_baseline
- FL_ng|83.67% nodes
- Eff_checker|83.67% nodes
- FL_ng_prop|83.62% nodes
- Eff_checker|83.62% nodes
- PageRank|84.38% nodes
- Eff_checker|84.38% nodes

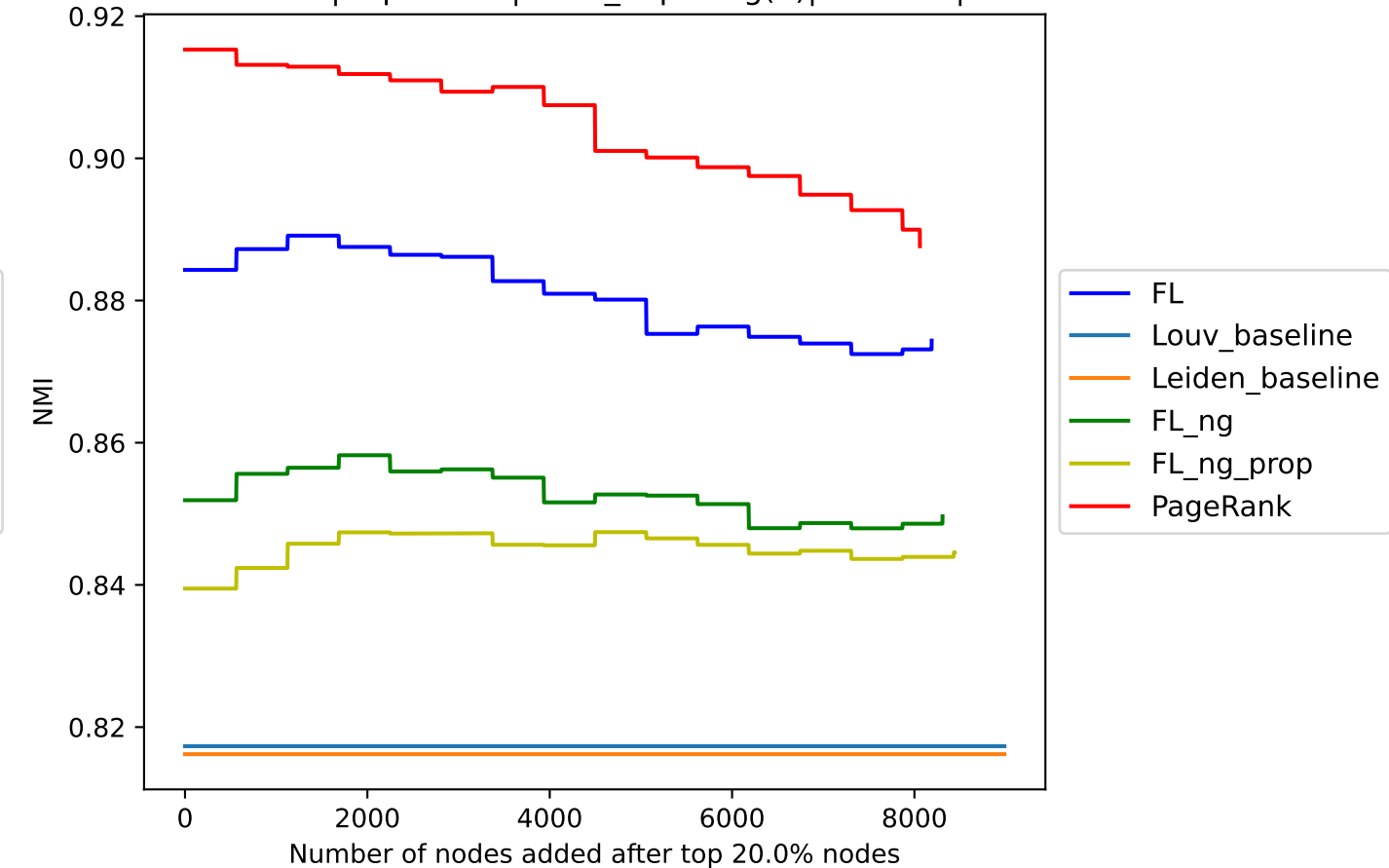
mRNA | top 20.0% | Num_hops: log(n)



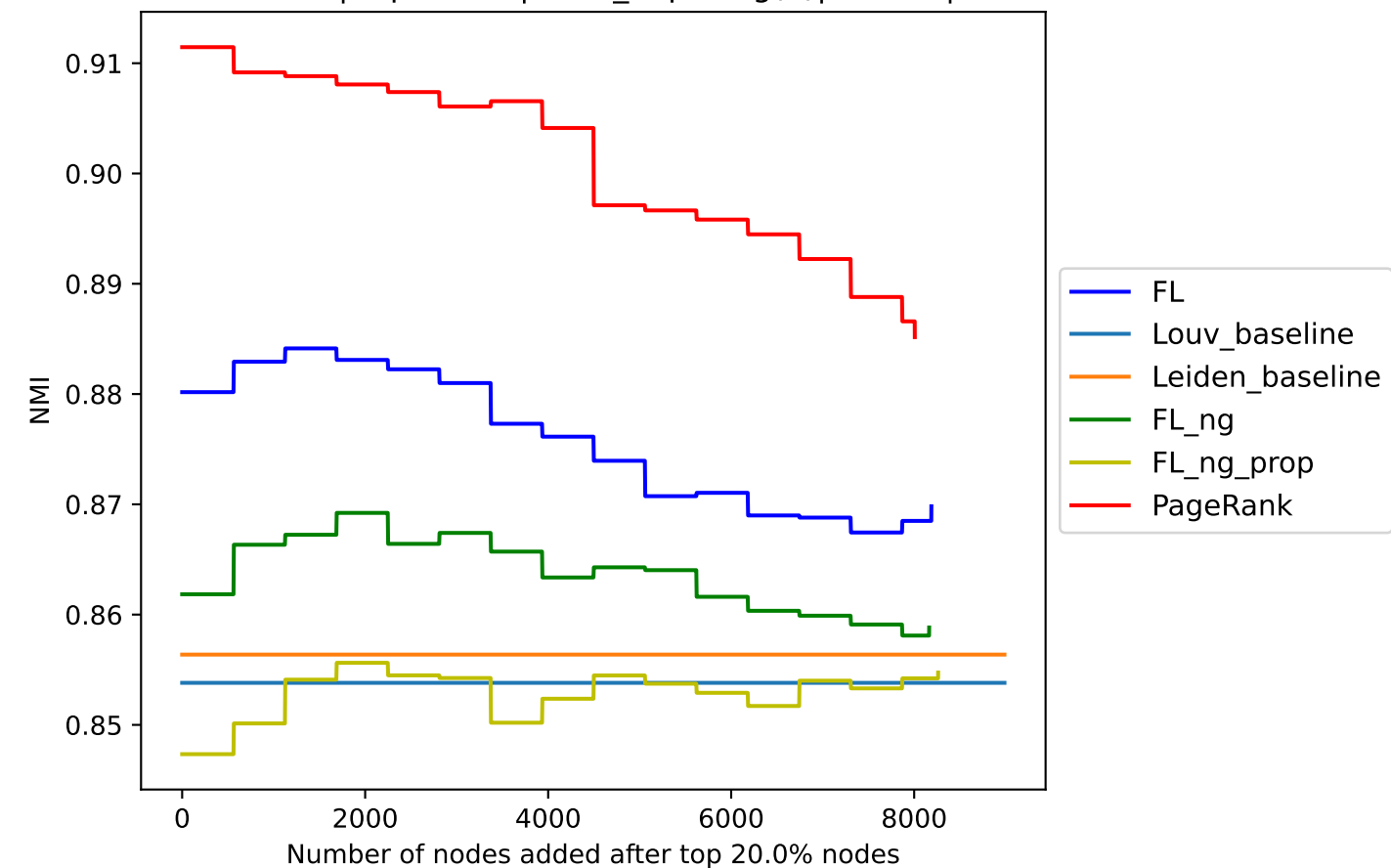
mRNA | top 20.0%| Num_hops: log(n)|res: 0.05|



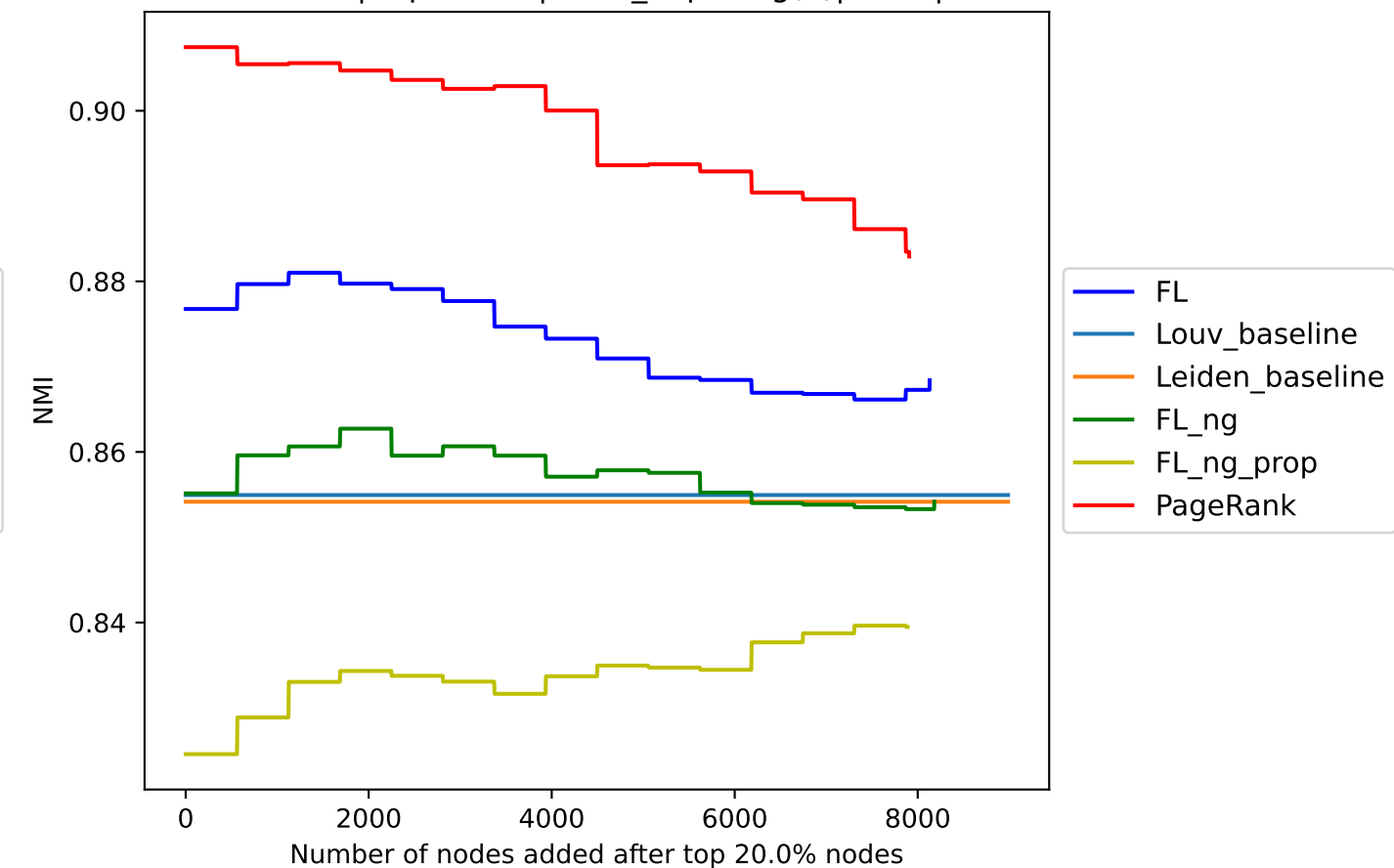
mRNA | top 20.0%| Num_hops: log(n)|res: 0.25|



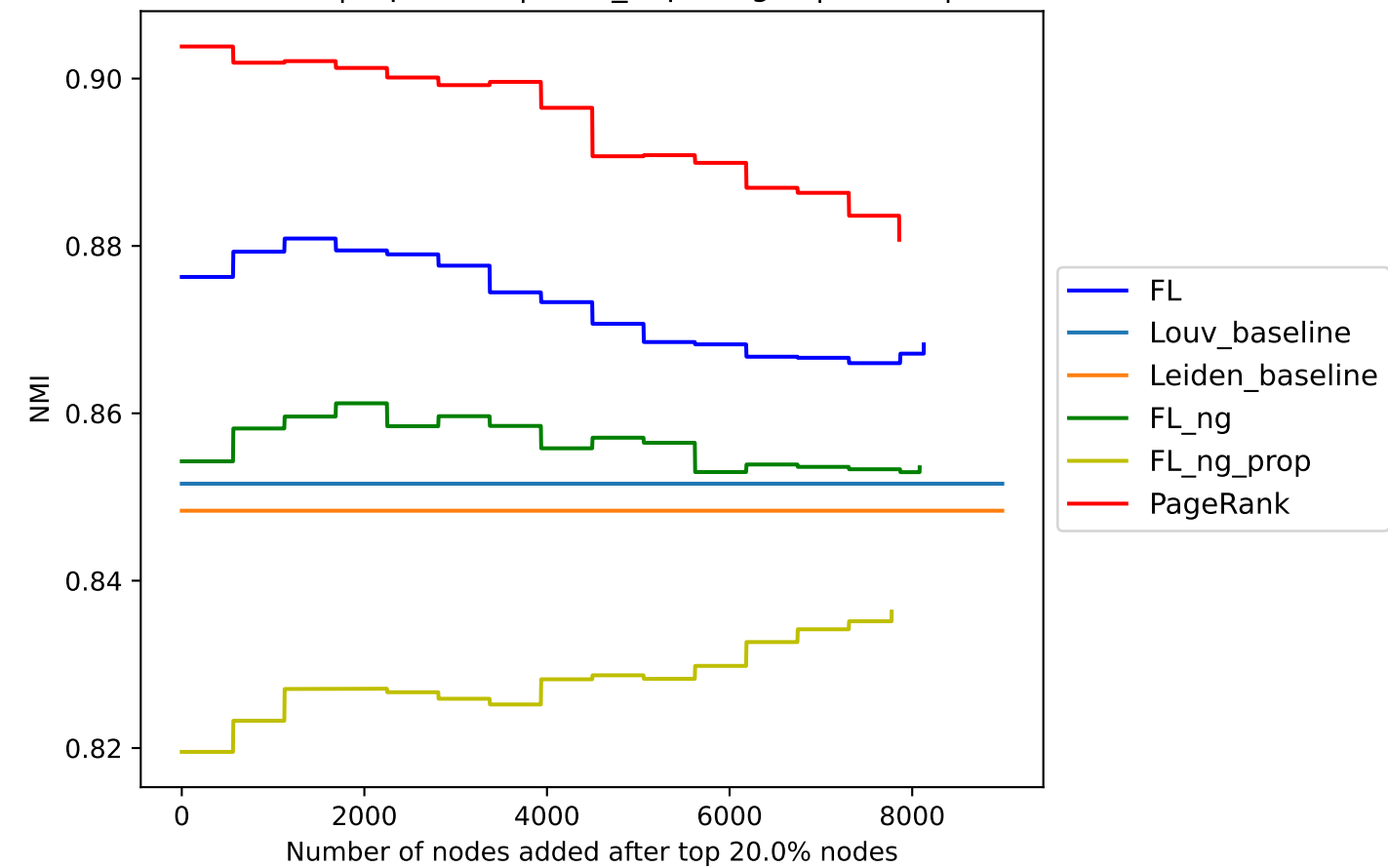
mRNA | top 20.0%| Num_hops: log(n)|res: 0.5|



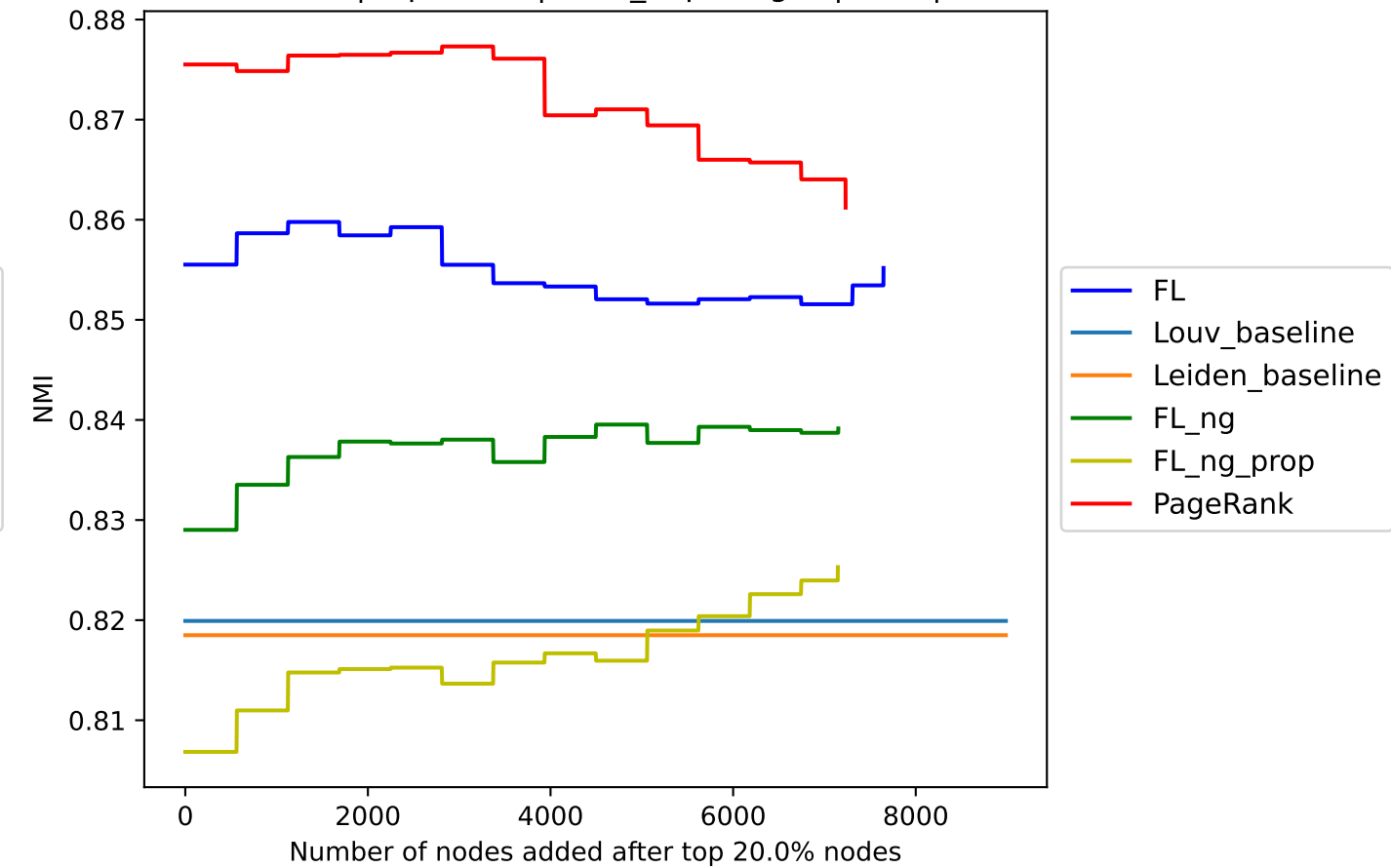
mRNA | top 20.0%| Num_hops: log(n)|res: 1|

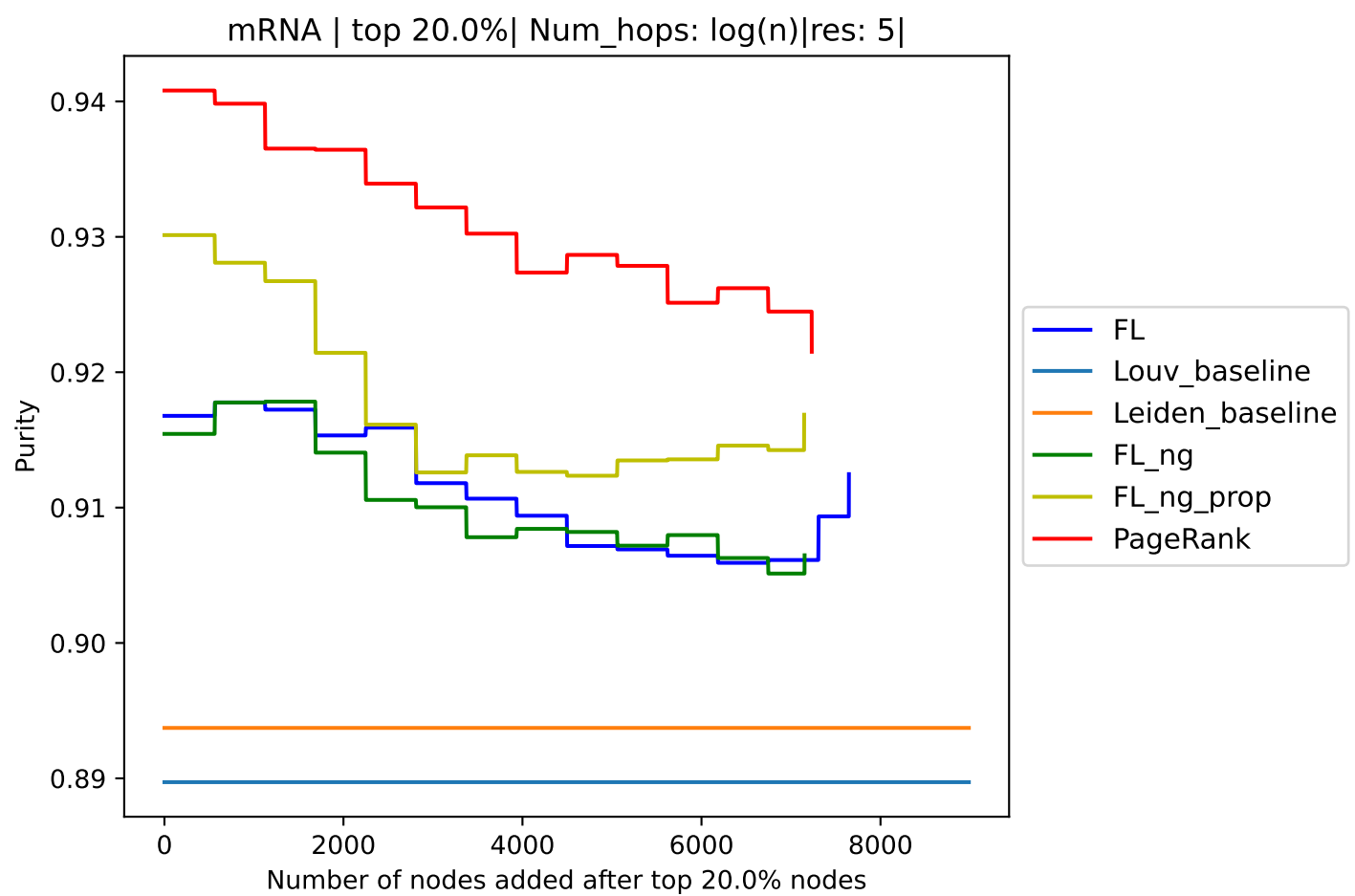
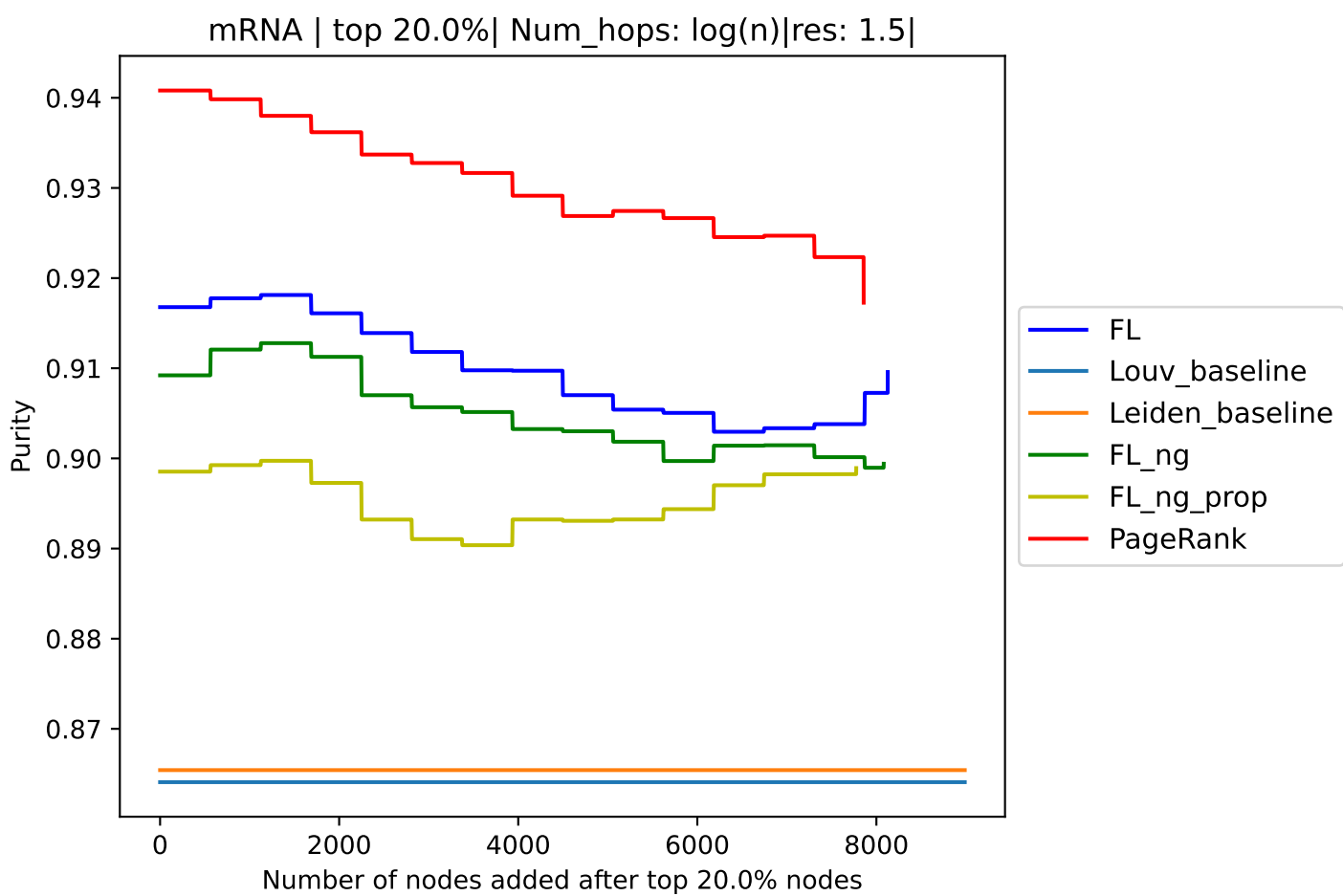
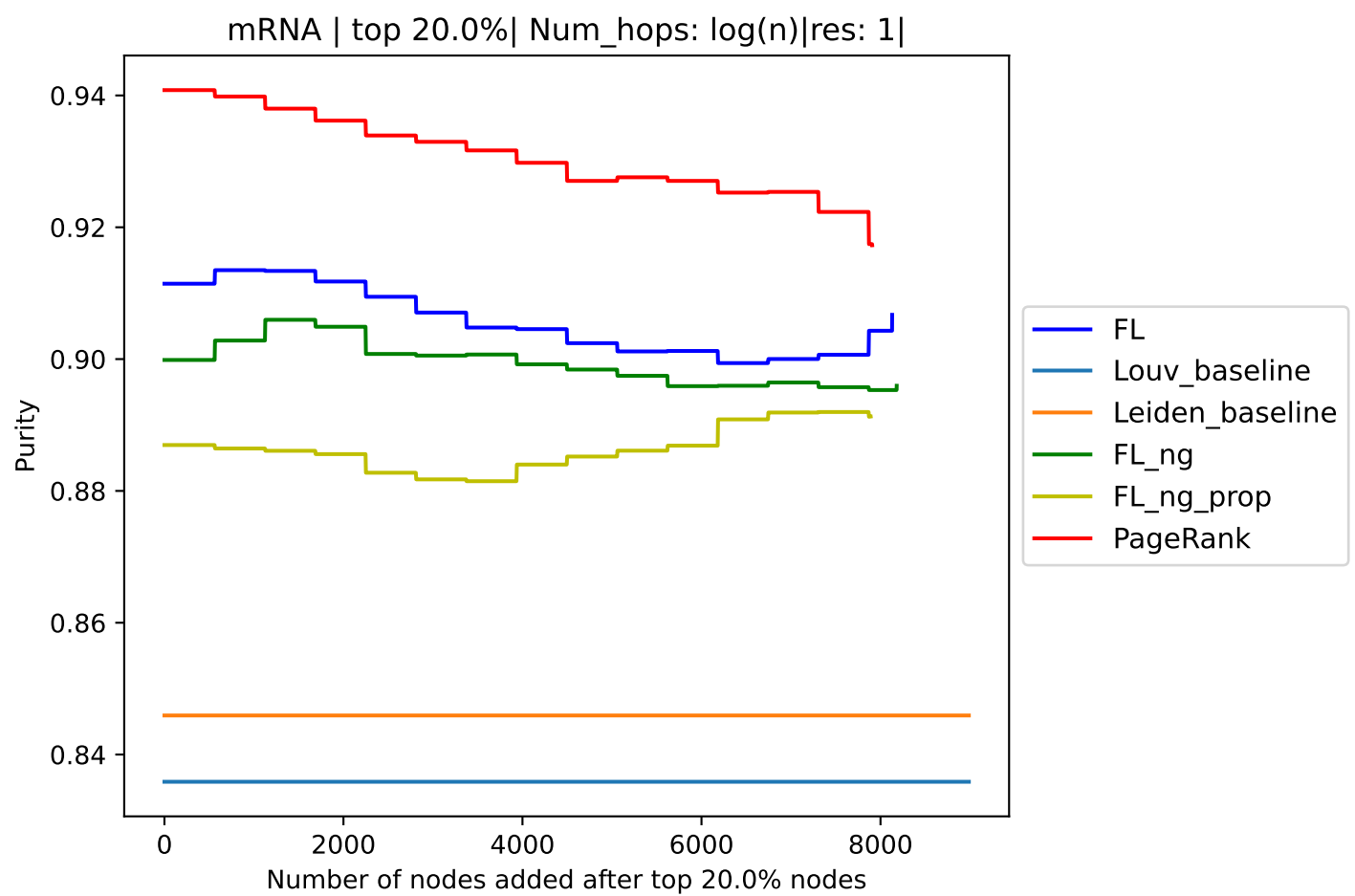
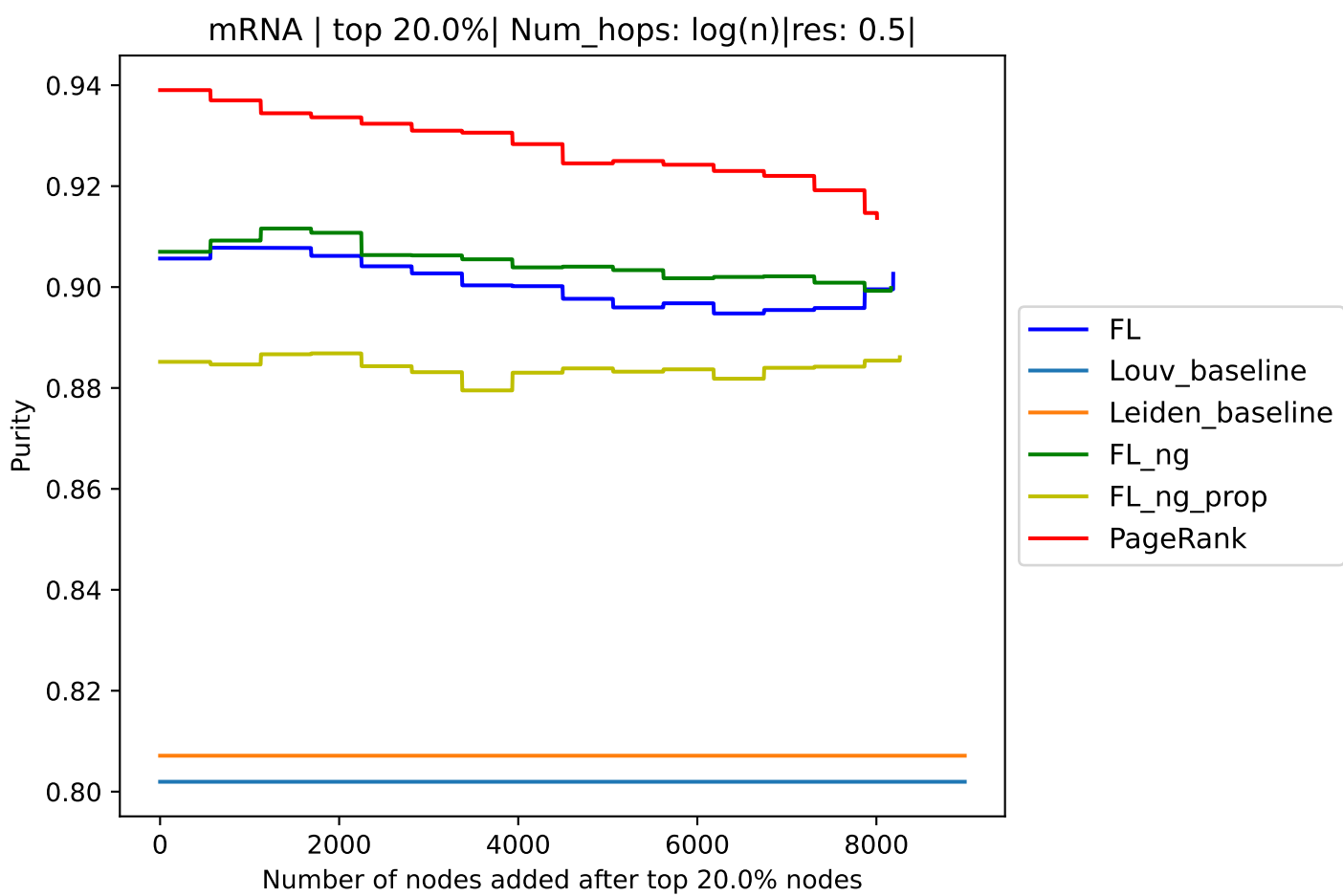
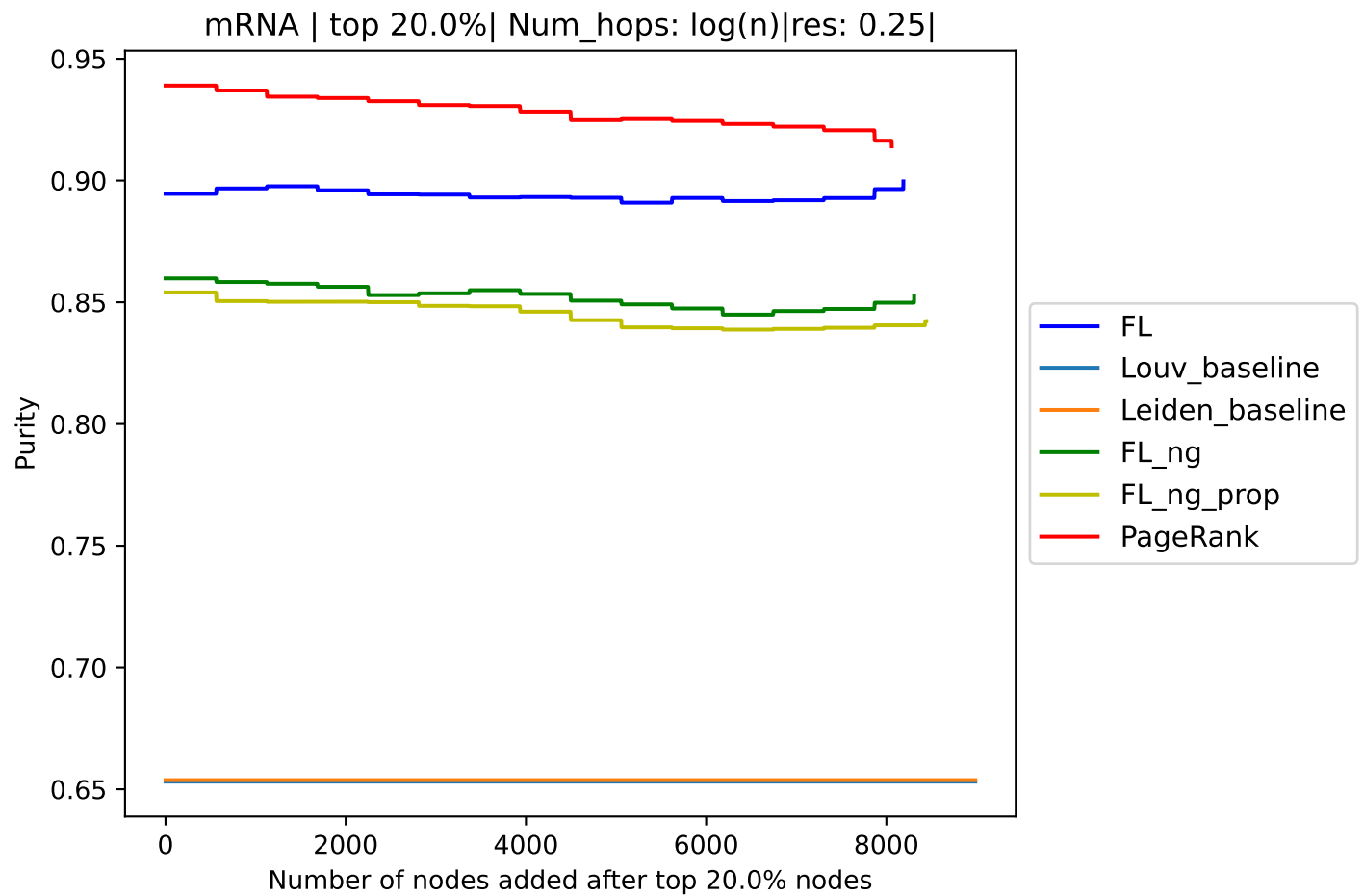
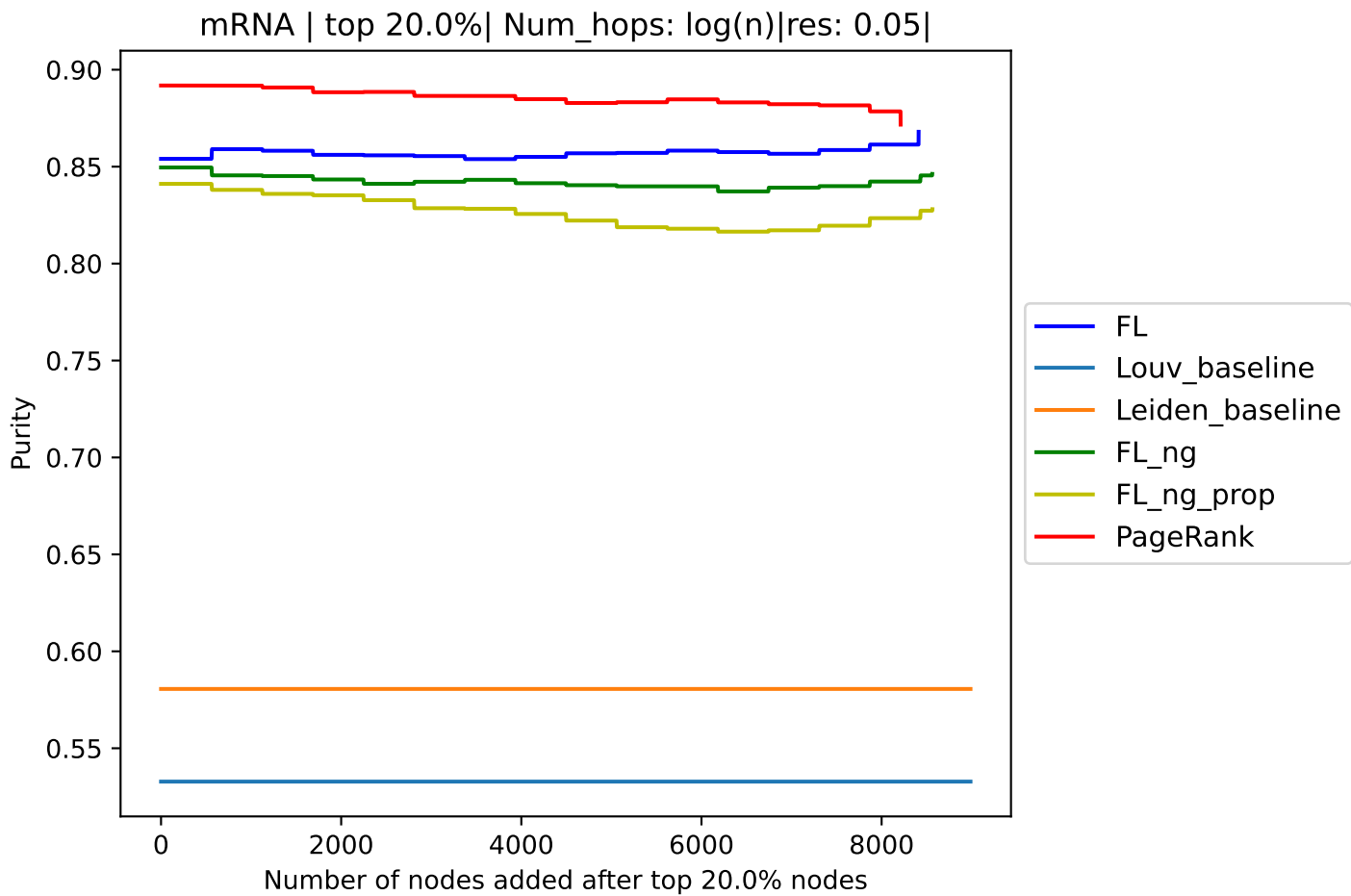


mRNA | top 20.0%| Num_hops: log(n)|res: 1.5|

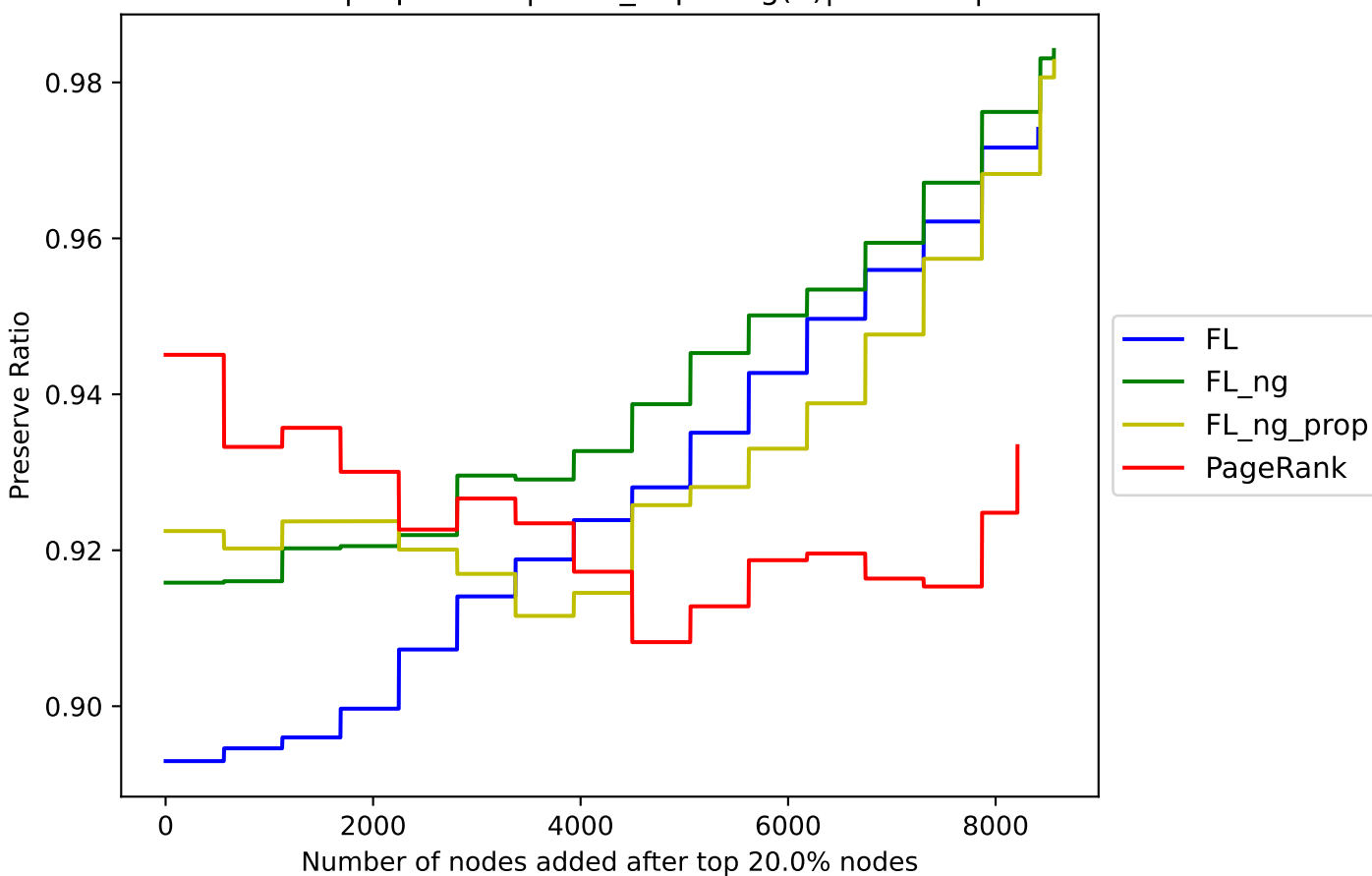


mRNA | top 20.0%| Num_hops: log(n)|res: 5|

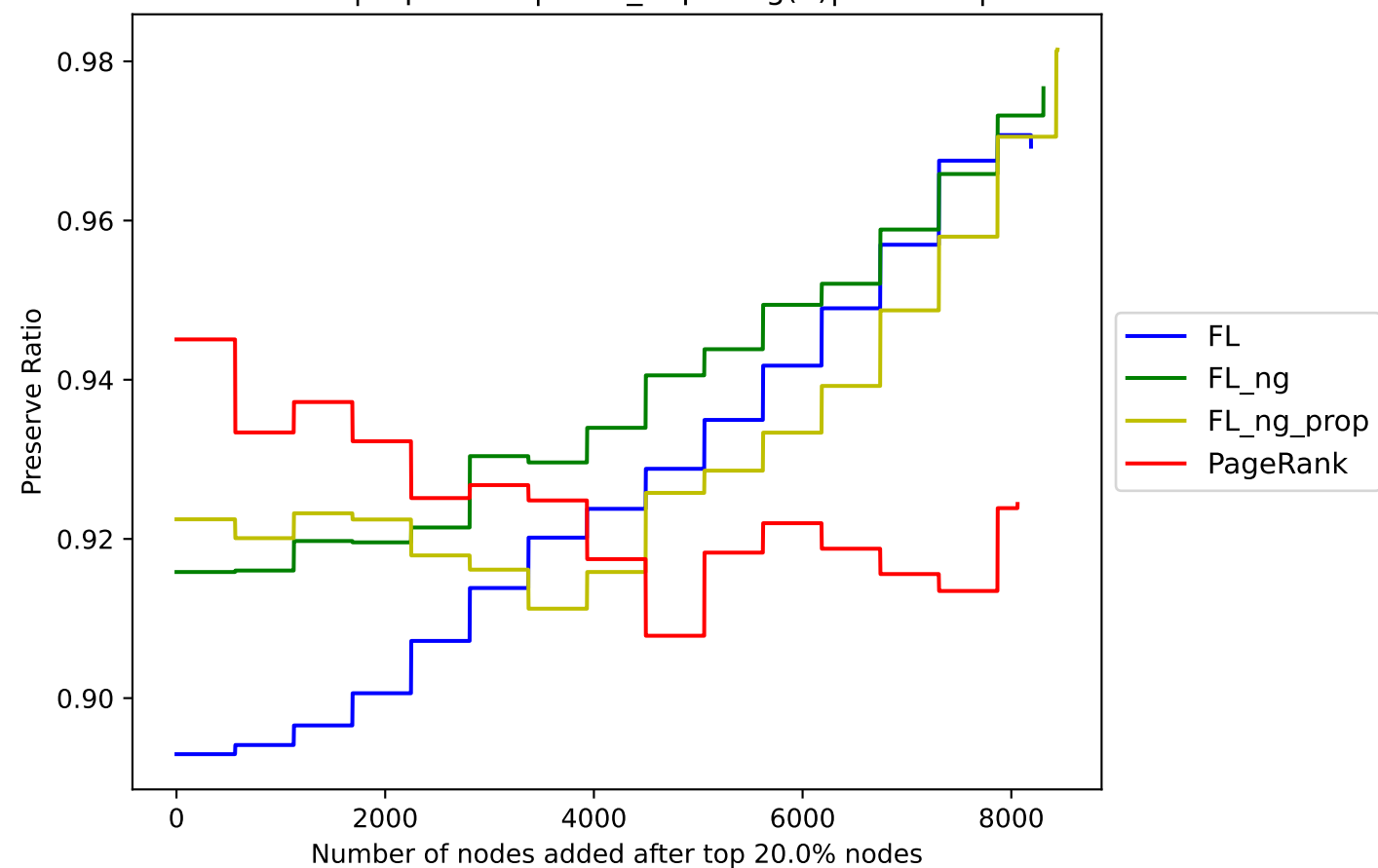




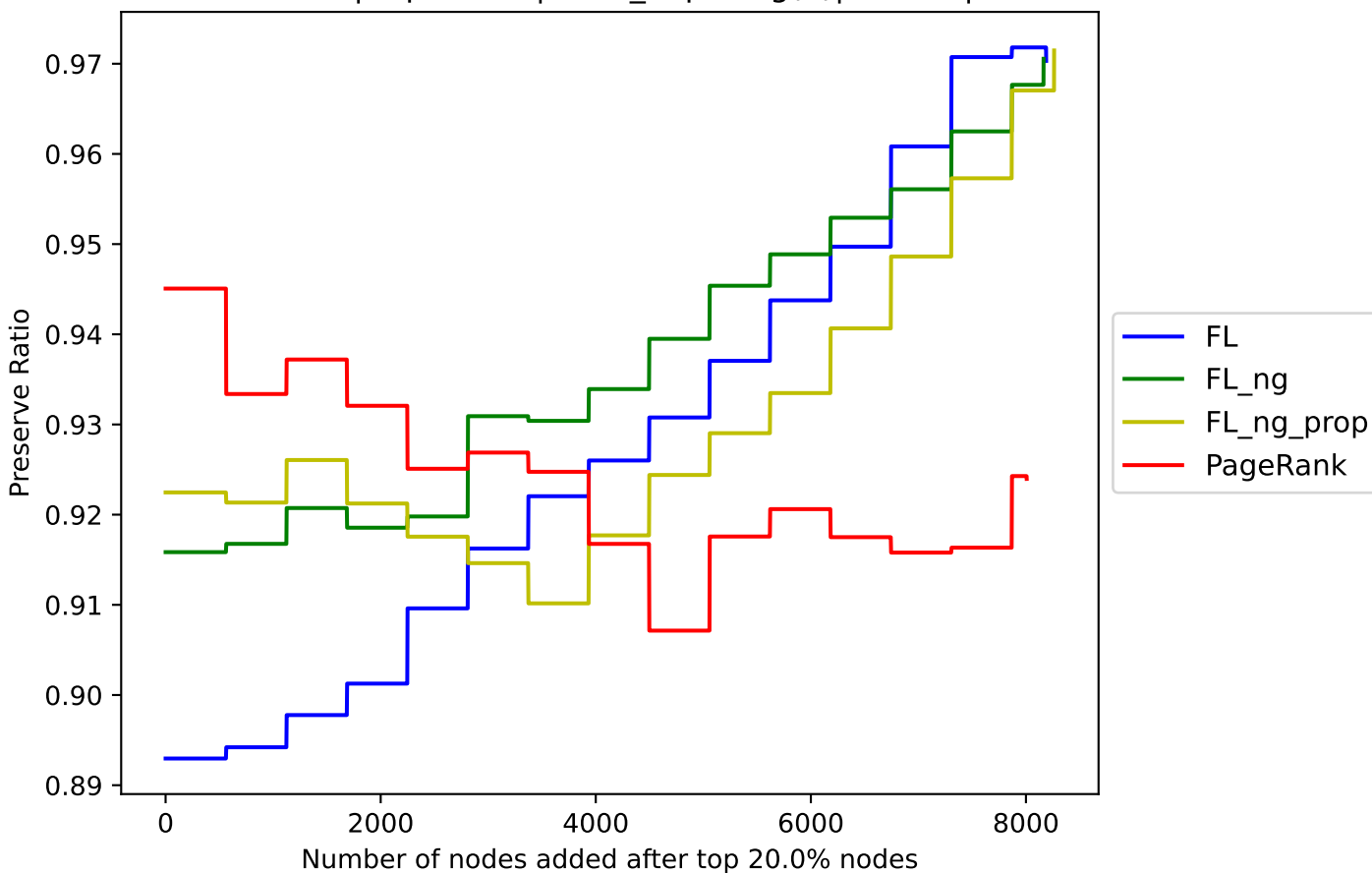
mRNA | top 20.0%| Num_hops: log(n)|res: 0.05|



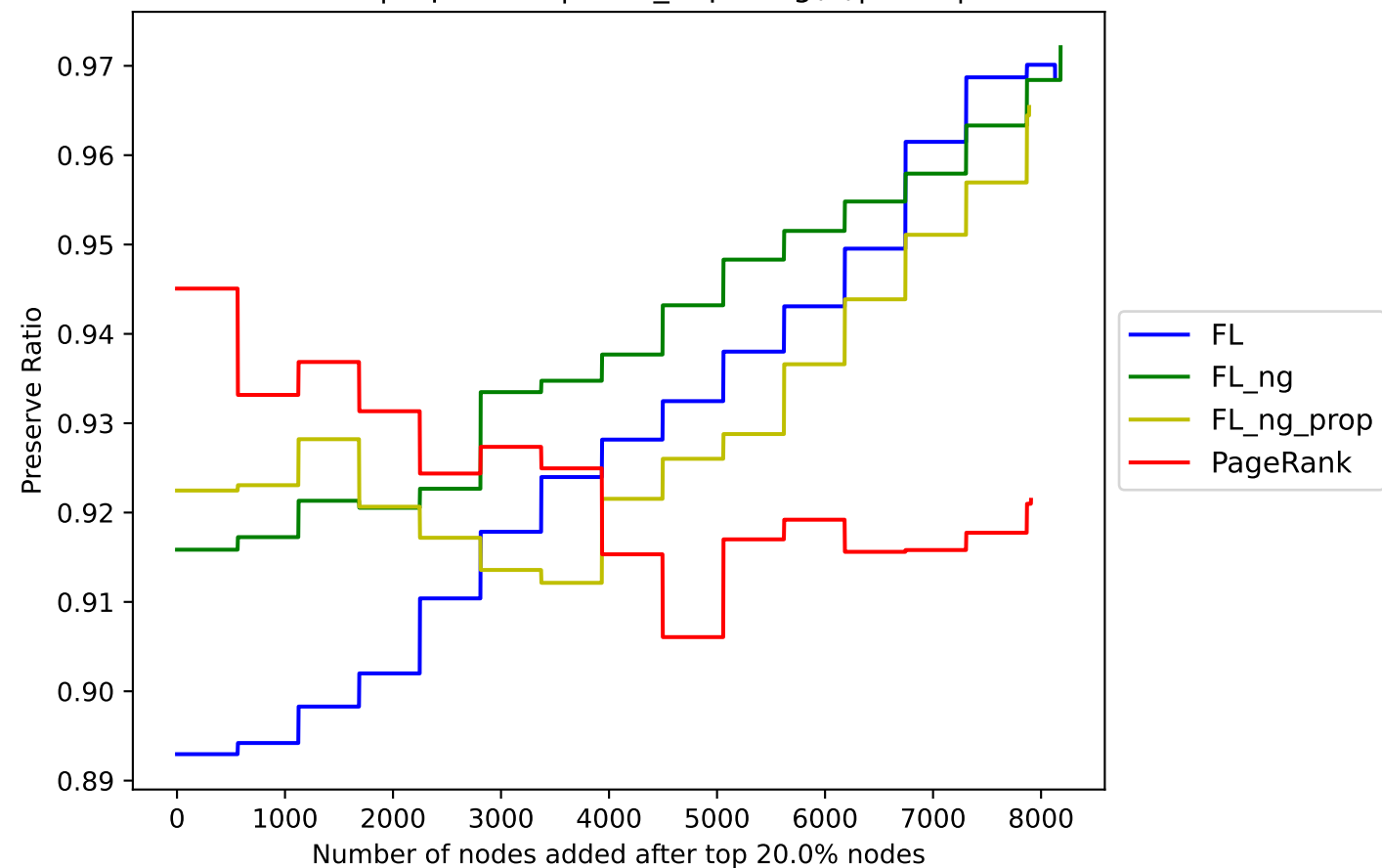
mRNA | top 20.0%| Num_hops: log(n)|res: 0.25|



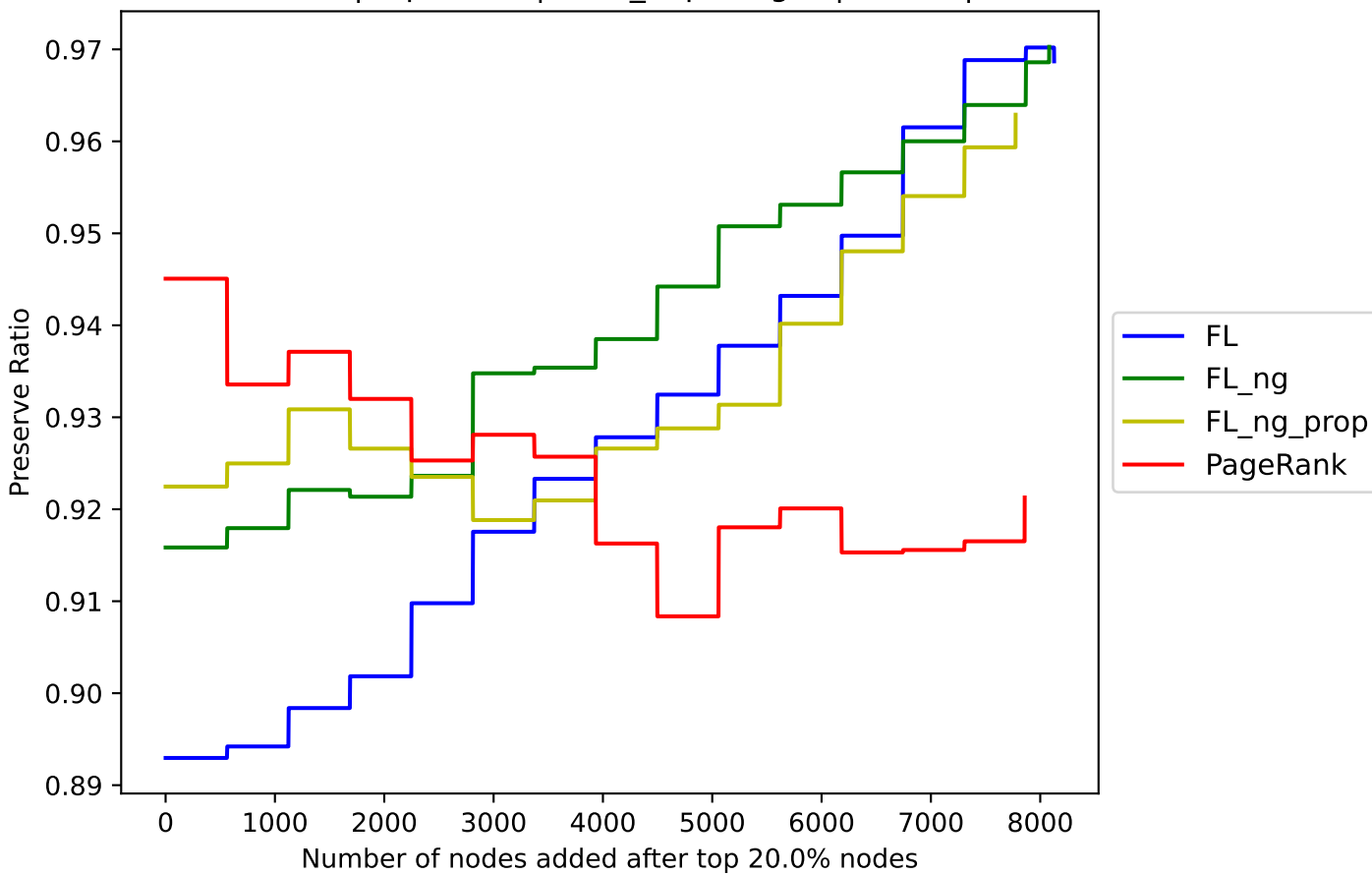
mRNA | top 20.0%| Num_hops: log(n)|res: 0.5|



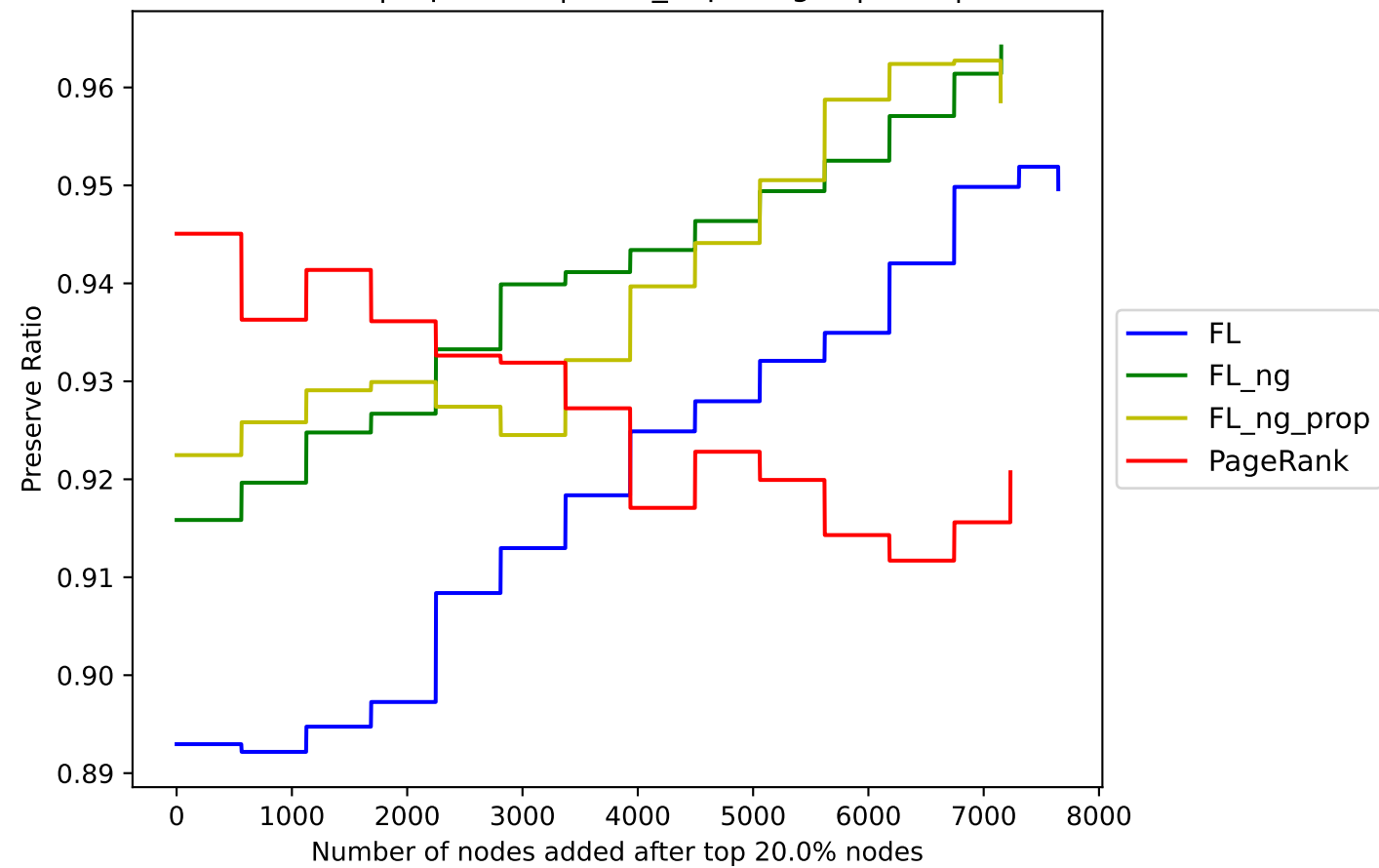
mRNA | top 20.0%| Num_hops: log(n)|res: 1|

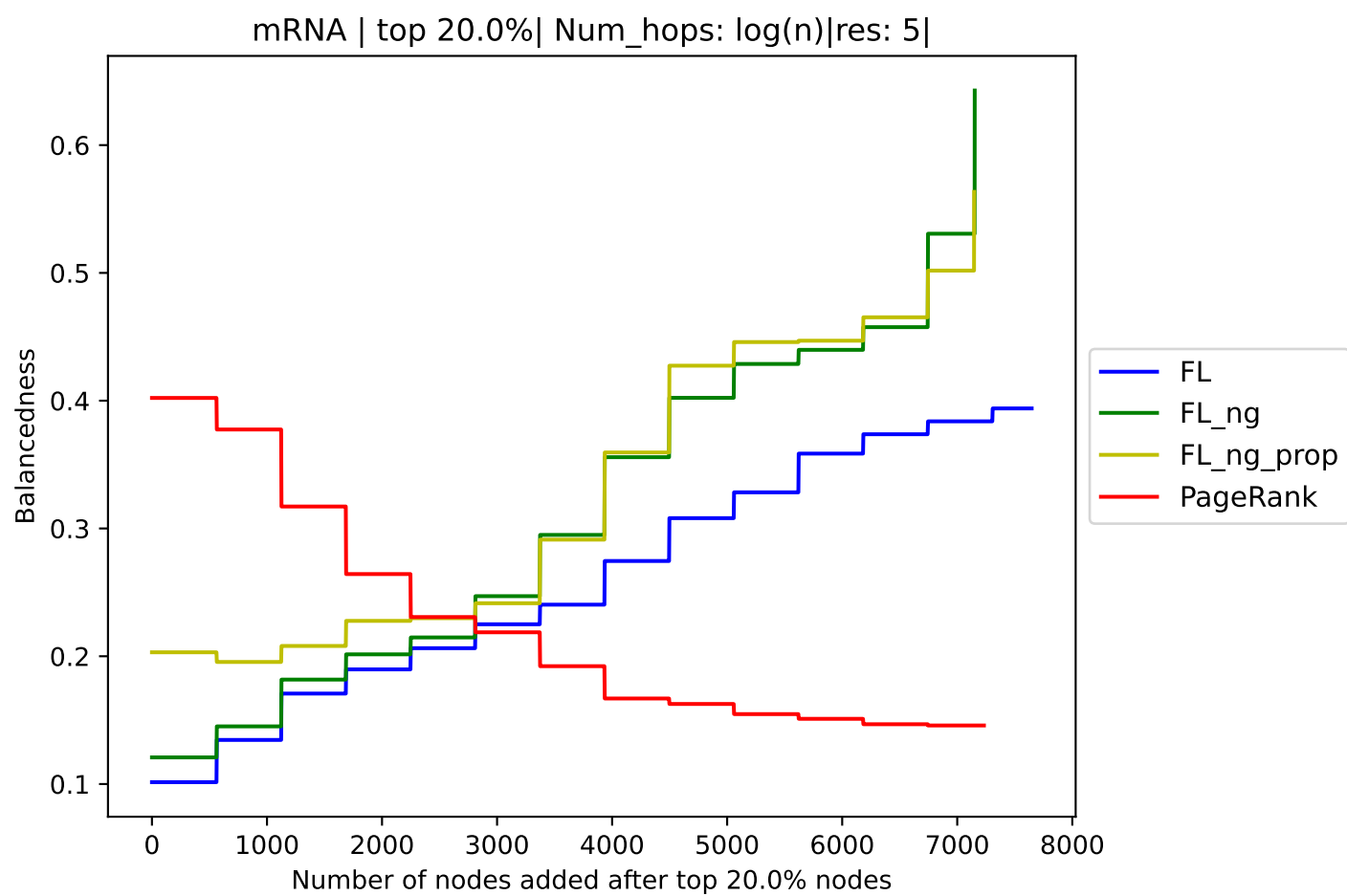
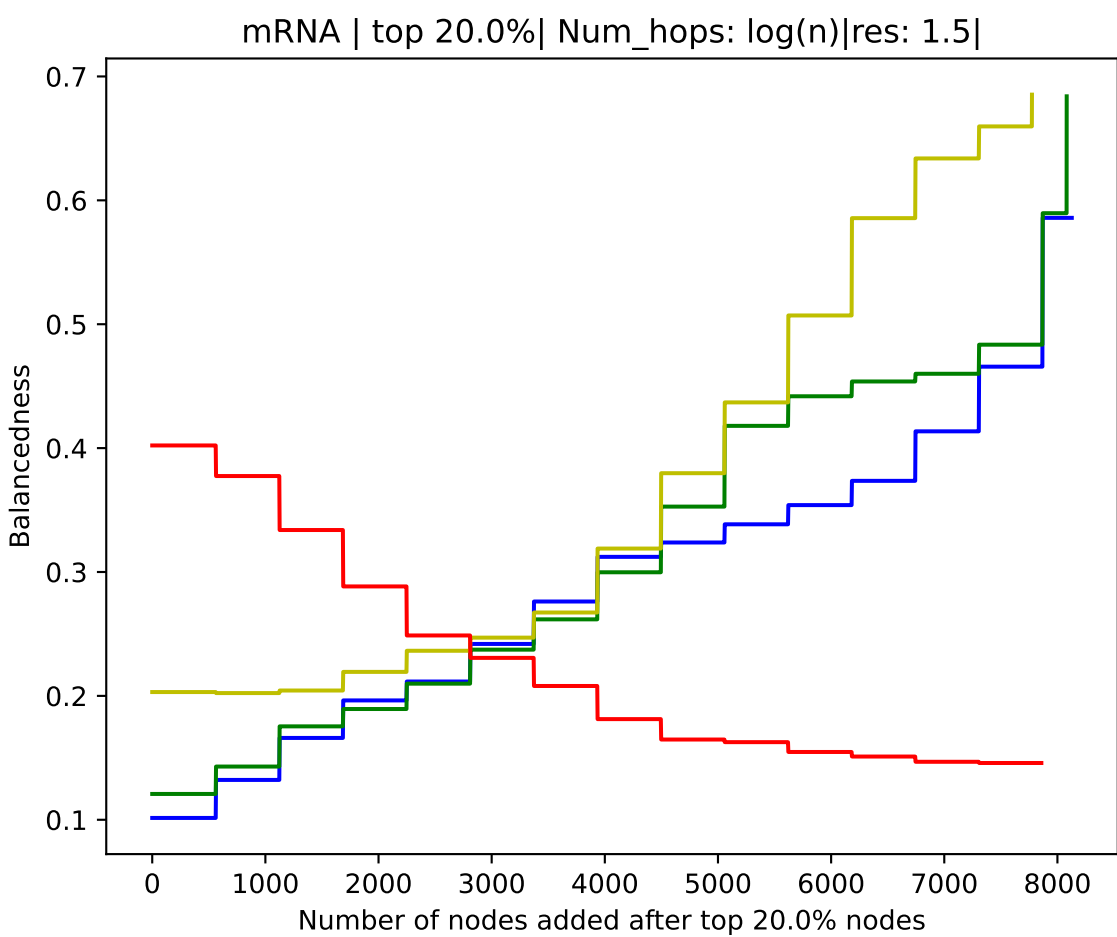
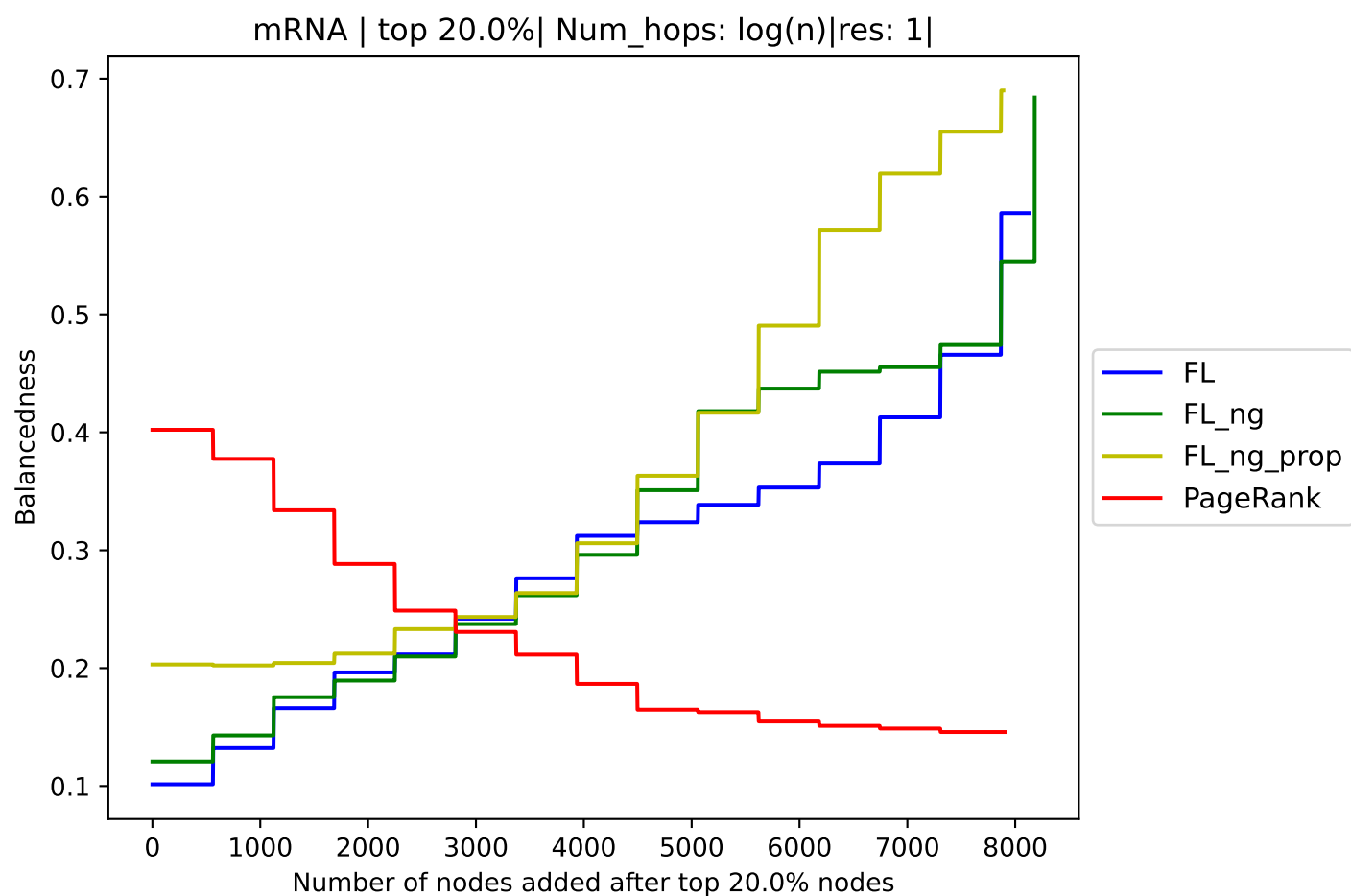
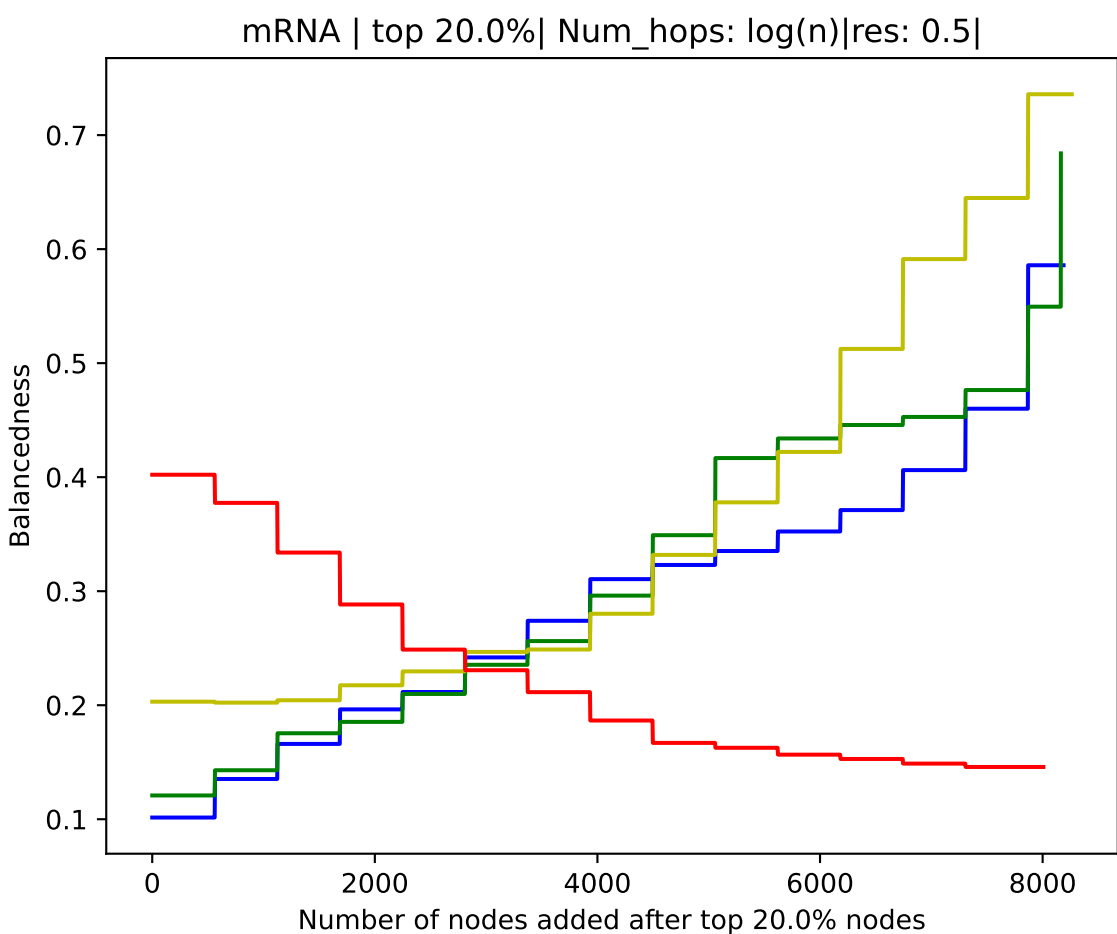
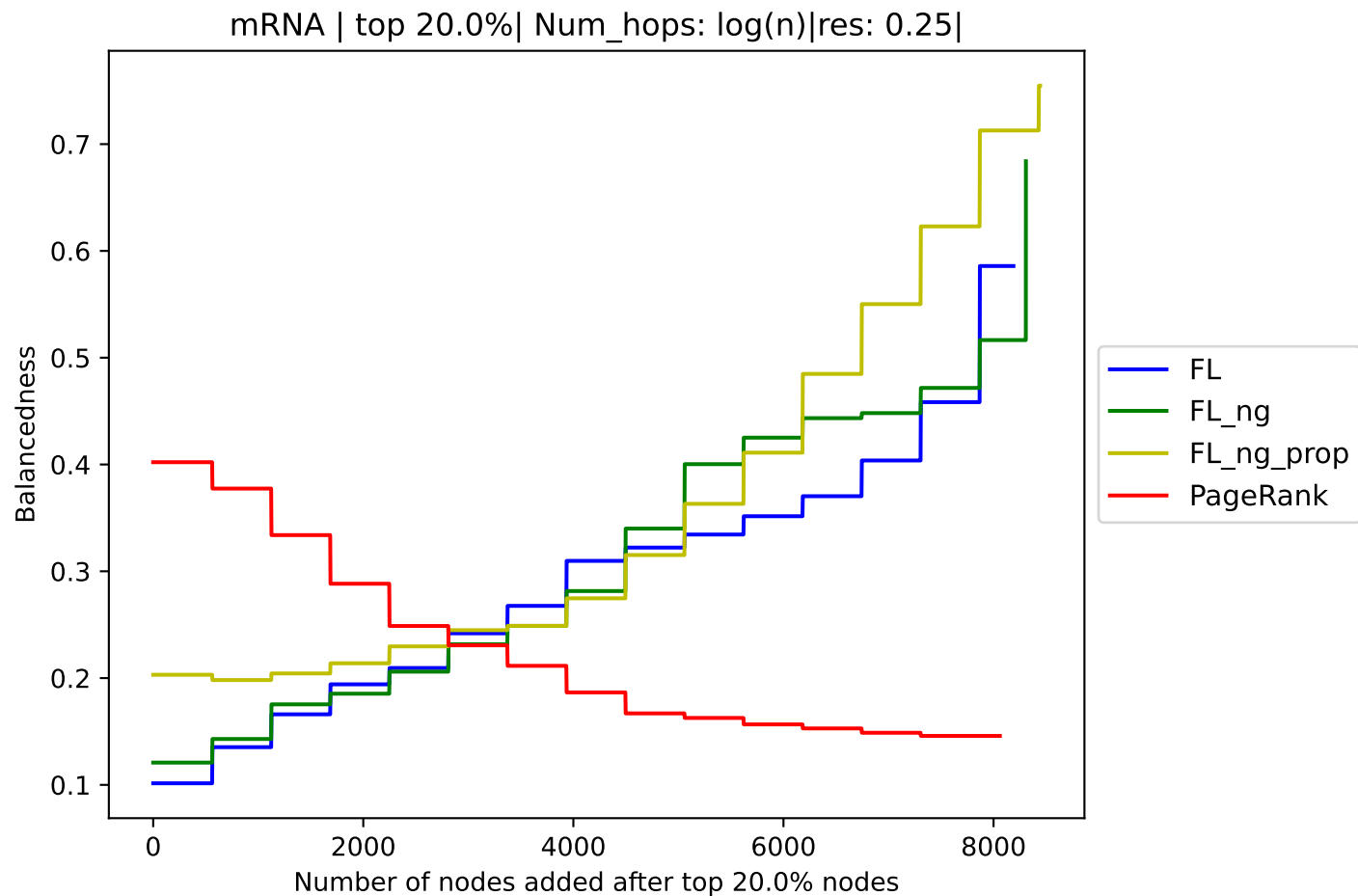
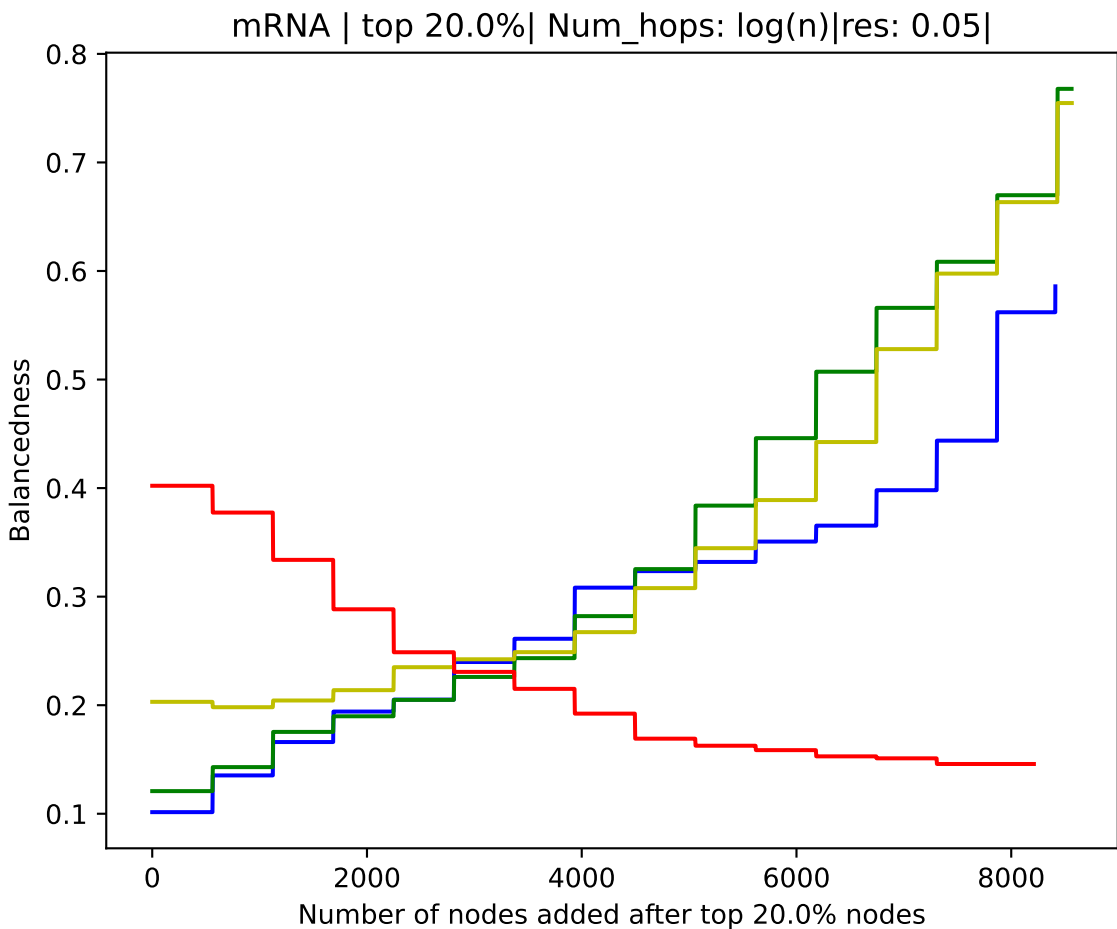


mRNA | top 20.0%| Num_hops: log(n)|res: 1.5|

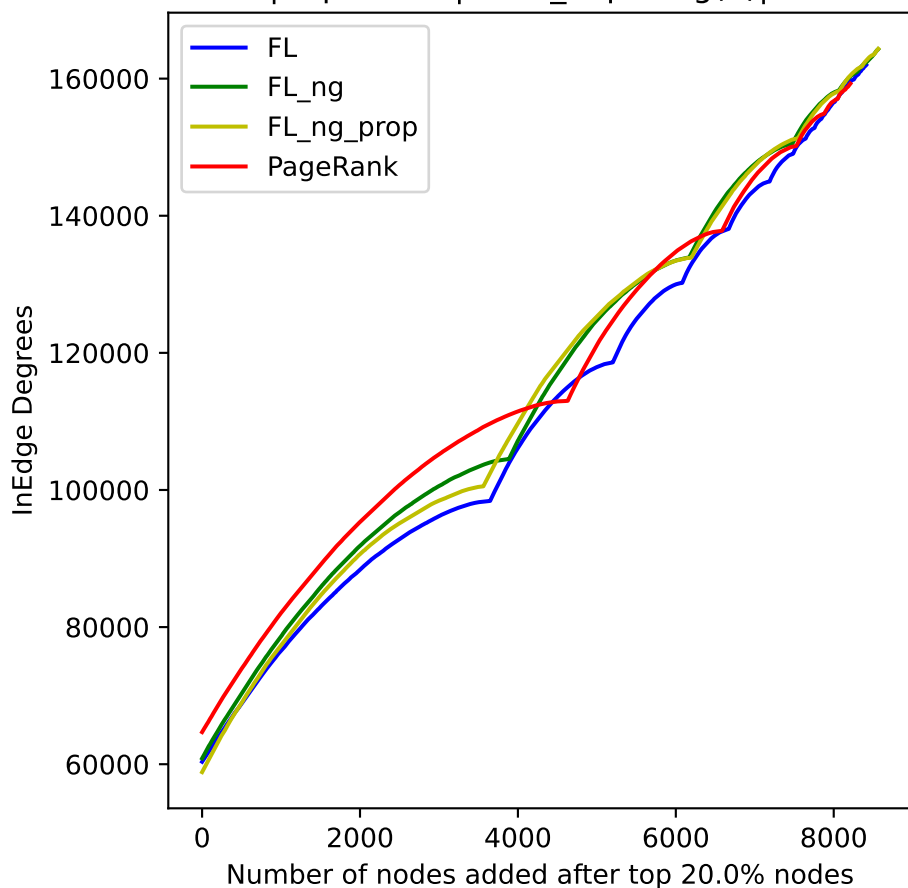


mRNA | top 20.0%| Num_hops: log(n)|res: 5|

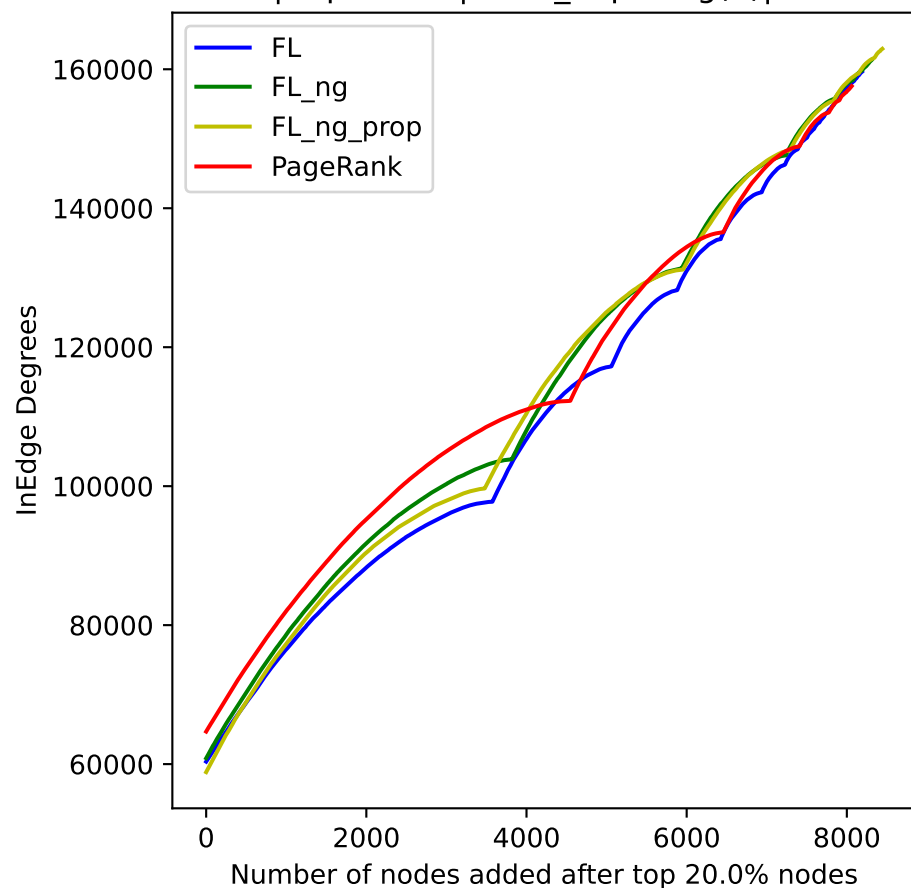




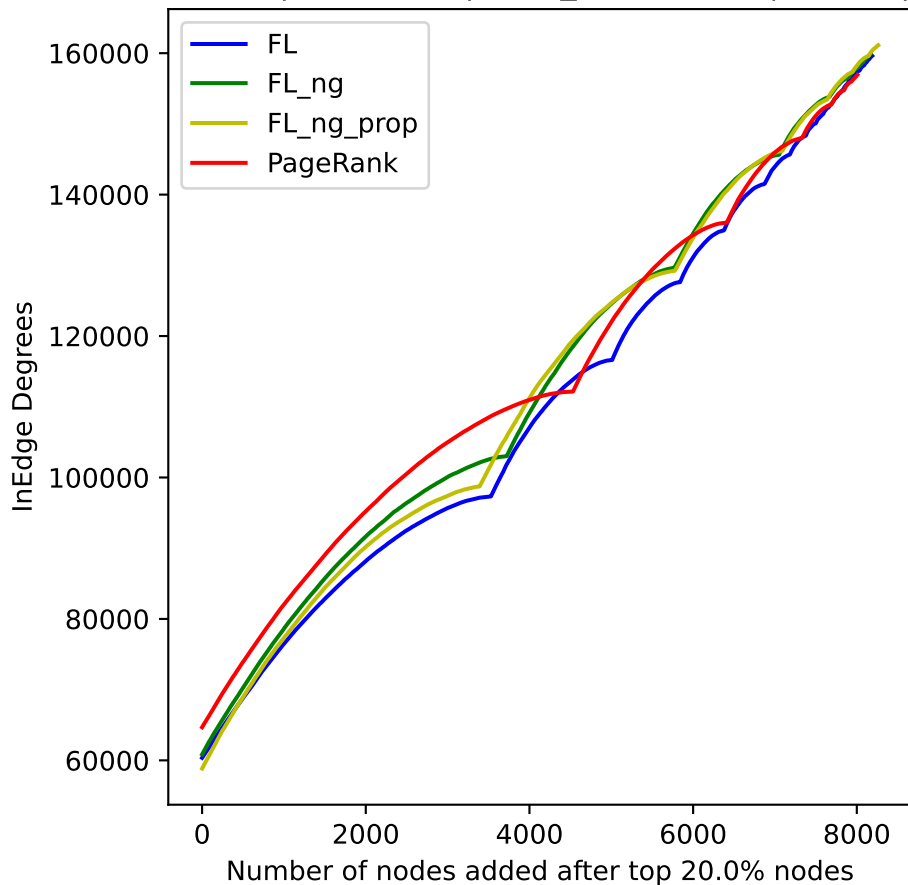
mRNA | top 20.0%| Num_hops: log(n)|res: 0.05|



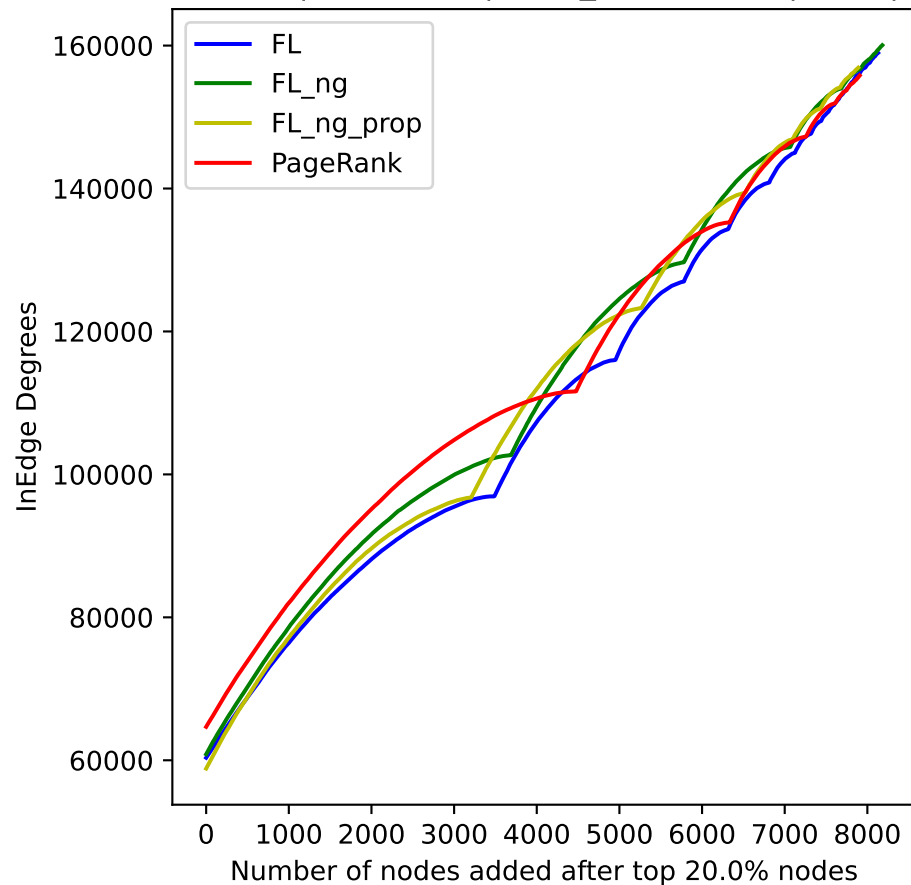
mRNA | top 20.0%| Num_hops: log(n)|res: 0.25|



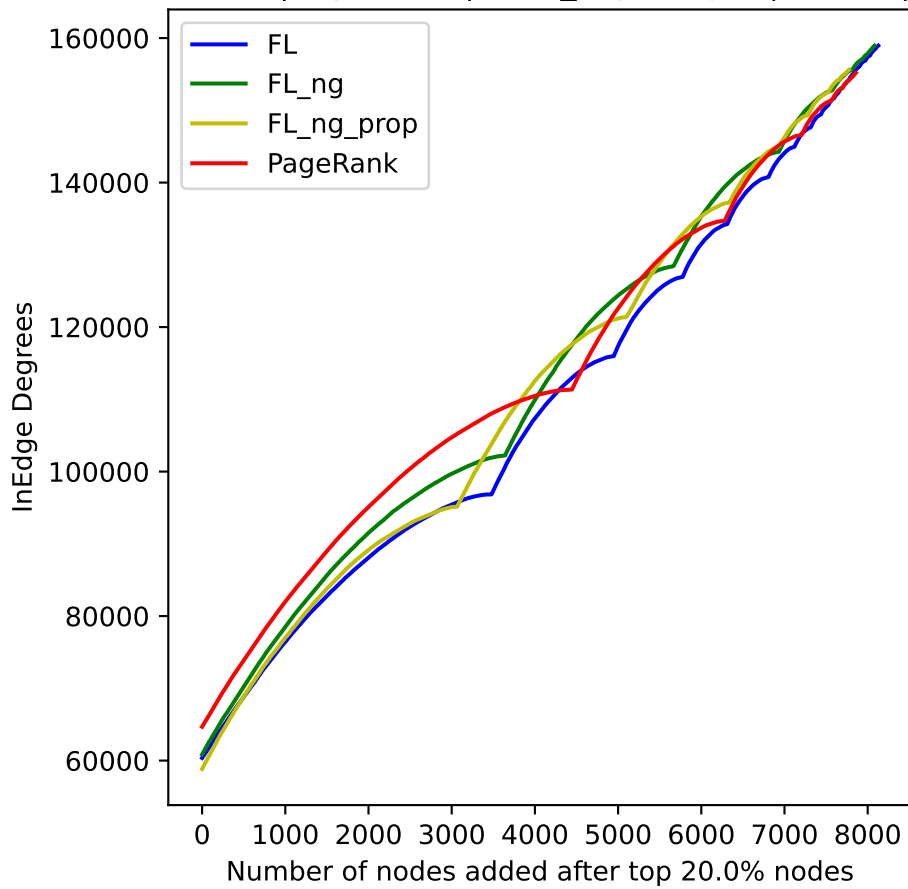
mRNA | top 20.0%| Num_hops: log(n)|res: 0.5|



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|

