We thank the reviewers for their time and valuable feedback. Overall, reviewers found our method G-META "novel". "scalability is particularly appealing", "theoretical analysis is great, substantial, valid, correct", experiments are "comprehensive", and the paper "well-written". Below, we clarify important points raised by reviewers: (1) relation to existing work, (2) local subgraphs in G-META vs. computation graphs in GNNs, (3) size of local subgraphs, (4) **ablation**, and (5) others. We believe these clarifications, together with our new analyses, resolve all key issues raised. (1) Relation to existing work. R1 questioned the difference between Meta-GNN and G-META, saying it "remains moderate." We respectfully disagree. The key difference are local subgraphs, and this is crucial because local subgraphs enable i) accurate and fast adaptation to new tasks, ii) learning in few-shot settings, and iii) theoretical justification. In contrast to our G-META, Meta-GNN and, similarly, Meta-Graph, use *entire graphs* for meta-learning. Because of that, they are unable to propagate label information across large graphs when only a few node labels are given. Theoretically, Meta-GNN/Meta-Graph works only on 1 out of 3 graph meta-learning problems (see Appendix B) and does not have any theoretical motivation whereas G-META works on all 3 problems and is theoretically justified. Empirically, G-META outperforms Meta-GNN, for example, by 65% on the ogbn-arxiv dataset (Meta-GNN cannot even be used on other datasets and meta-learning regimes). As suggested by reviewers, we will carefully discuss this in the final version.

(2) Local subgraphs vs. computation graphs. R3 raised a critical concern that "the idea of using local subgraphs to compute node representations is not novel." This points to a critical misunderstanding—computation graphs in GNNs are used to generate node embeddings vs. local subgraphs in our G-META are used to transfer knowledge for graph meta-learning. We are not claiming novelty in "the idea of using local subgraphs to compute node representations." Instead, as we write in the paper, we are claiming novelty in the idea of using local subgraphs for graph meta-learning, as recognized by R1 and R2. This innovation has important implications, which we show theoretically (i.e., proofs, solving classes of graph meta-learning problems not solved before) and empirically (i.e., considerable boost in accuracy over 9 baselines and 7 datasets). For example, a baseline method Meta-GNN, which uses a standard GNN on an entire graph together with MAML, performs 42.5% worse than G-META-MAML, a simplified variant of our G-META. Note that the only difference between Meta-GNN and this simplified G-META's variant is that it uses the entire graph vs. local subgraphs. We will clearly mention this contribution, which we agree is crucial for G-META.

(3) Size of local subgraphs and computational complexity. (3.1) R2, R3, and R4 nicely point that G-META's performance can vary with local subgraph size h. To address this, we evaluate G-META for h = 1, 2, 3 (Figure for Fold-PPI). We find that h = 2 gives the best performance, which nicely corroborates R2's hypothesis on the inverted U-shape relationship. We will include this analysis in our final version. (3.2) R4 also raises an important concern on computational complexity of subgraph extraction. We would like to

3

8 9

10

11

12

13

15

16

17

18

19

20

21

23

24

25

26

27

28

29

30

31

32

33

34

35

36

37

38

39

40

41

42

43

45

46

47

48

49

50

51

52

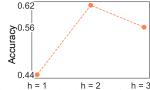
53

54

55

56

57



clarify that we don't need to compute "shortest paths ahead of the training". Instead, we simply do a lookup, retrieving neighbors of neighbors. Empirically, we find the subgraph construction takes 14.7% of training time, and this can be further reduced by implementing techniques for improving training efficiency, e.g., GraphSAINT [Zeng et al., ICLR 2020]. Regarding R4's comment of "evaluating each node label individually", we note that each mini-batch consists only of a few labels (e.g., 9 in 3-way 3-shot learning) and as such is a cheap operation.

(4) Further ablations. All reviewers raise an important point on ablation, which we agree is crucial especially for G-META. While we already included ablation in the form of baselines, we will make it explicit in the next version following R3 and R4. We thank R3 and R4 for nicely pointing out that baselines ProtoNet and MAML can be seen as G-META's ablations. In response to R2, new results in Table show that gradient-based meta-

Method	ogbn-arxiv	Tissue-PPI	Fold-PPI
G-META	0.451	0.768	0.561
- MAML	0.372	0.546	0.382
- Prototype	0.389	0.745	0.482

learning aids G-META more than metric-based meta-learning but both are indispensable for G-META's performance. (5) Baselines and further clarifications. (5.1) R1 raised an important point about performance under varying size of the training set, i.e., K in the K-shot problem. We conduct experiments and observe, as expected, a linear trend between K and performance (e.g., on arxiv-ogbn, accuracy goes from 0.373, 0.442 to 0.484 for K = 1, 3, and 10, respectively). We will include the full study in the final version. (5.2) R1 correctly pointed out G-META "covers node classification task and link prediction task where the target is discrete class label.". In contrast, existing Meta-GNN and Meta-Graph only work for one of these two tasks—G-META is the first to work on both tasks. Also, to avoid confusion, we will update reference for graph-level molecular prediction [Hu et al., ICLR 2019] to edge-level interaction prediction [Zitnik et al., Bioinf. 2018]. (5.3) We thank R4 for rightly pointing out that local subgraphs can alleviate over-smoothing because, in each iteration, different subgraphs are fed into GNN, which promotes inductive generalization. (5.4) R4 raised a question about hyper-parameters. We use random search on validation set to select hyper-parameters and find that model performance is stable for a broad range of values. We will include the recommended set of constant parameters in the next version. (5.5) R4 raised an important point on experimental setup. We follow standard episode training and semi-supervised setting in which most nodes are not labeled, i.e., few-shot learning. In K-shot N-way setup, for "Multiple Graphs and Shared Labels" problem, each task samples K nodes for each label N_i in the same label set of size N from one graph, and different tasks are associated with different graphs. For "Multiple Graph and Disjoint Labels" problem, each task defines an N-size label set, and samples K nodes for each label N_i .