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We thank the reviewers for their thoughtful feedbacks. We are encouraged they think M-HMC is addressing an important
problem (R1, R2), "is potentially going to be very influential" (R1) and is "a method with potential wide usage" (R2).
We are glad all reviewers find the paper to be well-written and clearly positioned w.r.t. prior work, the theory is correct
and "presented carefully" (R1), the experiments are "extensive" (R1), "well-designed" (R2), "very reasonable" (R4).
[R2] ...too many contents on the correctness of the proposed methods are in the appendix...I would like to
increase my score if a better sketch of the proof can be made in the main paper. Great point. The main idea is
to decompose the transition probability kernel R_T(s,B) into summation along countable number of deterministic
"probabilistic paths", then use similar proof techniques as in RRHMC. Starting from s \in \Omega \times \Sigma, for a given travel
time T, an M-HMC iteration (without MH correction) specifies a deterministic mapping for a fixed sequence of
random discrete proposals (Y in appendix). We introduce "probabilistic path" (\omega(s, T, Y) in appendix), containing all
information (sequence of proposals, indices/times&accept/reject decisions for discrete updates, and evolution of the
discrete/continuous states and auxiliary variables) of such deterministic trajectories for fixed Y's. There can only be a
countable number of possible probabilistic paths, since T and discrete state space are finite, so traveling from s for time T
gives a countable number of possible destinations s'. This implies the decomposition R_T(s,B) = \sum_{s'} \sum_{Y} r_{T,Y}(s,s'),
where the summation is done over possible destinations s' and all valid Y's that bring s to s' through \omega(s,T,Y).
r_{T,Y}(s,s') is the transition probability along the deterministic trajectory \omega(s,T,Y). Using similar proof techniques as
in RRHMC, we can prove detailed balance for these deterministic trajectories (Lemma 4 in appendix). This in turn
proves detailed balance of M-HMC. We will expand and include the above proof sketch in camera-ready version.
[R4] The distinction between this method and MH within HMC is not sufficiently strong for this work to be con-
sidered very novel...the implementation in Appendix L262...don't quite match the description in the paper...The
comparison of M-HMC to MH within HMC is questionable. This seems a factual misunderstanding. This compar-
ison is not meant to be novel. Rather, MH (with or without self-transition) within HMC isn't valid (Fig. 1c), while
M-HMC is (Fig. 1a). We can't imagine a stronger comparison. Appendix L262 is MH within HMC (L188, appendix
L178), so R4 is correct, it's invalid. M-HMC is in appendix L256, which is valid and matches Algo. 1. To us, the
distinction being simple in this 1 discrete variable case is a strength. This simple change naturally comes out of Algo. 1,
and corrects the inherent bias in MH within HMC (Fig. 1a/1c). In fact, we specifically included this function to highlight
this simple distinction, showing M-HMC is correct, more efficient, yet extremely easy to incorporate into existing HMC
implementations. For novelty: R1, R2 both think M-HMC solves an important problem and can potentially be very
influential and see wide usage. Even R4 mentioned "The work is clearly a novel extension of previous work"
[R4] M-HMC equally suffers from this issue (long trajectories for random-walk suppression vs frequent discrete
updates) because if the discrete variables are frequently updated...then the trajectory length between successive
discrete random variable updates is short. We respectively disagree. In HMC, using long trajectories gives distant
proposals/random-walk suppression (due to consistent momentum within an iteration), but means infrequent discrete
updates. Shorter trajectories enable more frequent discrete updates, but frequent momentum resampling between
iterations increases random-walk behavior. In contrast, M-HMC can always utilize consistent momentum/kinetic energy
in long trajectories, with potentially frequent discrete updates, to get distant proposals/random-walk suppression. The
trajectory lengths between successive discrete updates are irrelevant, as these updates involve no momentum resampling.
[R3] No theoretical insights about the speed-up... Intuitively speed-up is from more frequent discrete updates. More
detailed theoretical analysis is good future work. [R3] ...M-HMC will have to introduce much more continuous
variables than DHMC... Using Algo. 1, we only need 1 continuous variable (k_i^{\mathcal{D}}) in Algo. 1) for each discrete
variable. DHMC needs 2 (q_i^{\mathcal{D}} in L85, and p_i^{\mathcal{D}}). M-HMC also doesn't need to update all discrete variables every time,
making it more efficient than DHMC (see L265-268 for comparison on BLR). [R3] ...performance...very dependent
of...T...suggestions...? HMC is known to be sensitive to T (and step size). So is M-HMC. Automatically picking
these parameters is important future work (L325). [R3] It's not clear at all...the discrete part...needs...HMC... This
mechanism enables discrete updates within HMC, which, if done naively, is invalid. The proposals Q_i's are used instead
of gradients information. A random walk type implementation would be the invalid MH within HMC (Fig. 1c).
[R2, R4] HMC-within-Gibbs (HwG) should be used instead of NwG as baseline Additional experiments with HwG
show M-HMC is 2.5 (24D GMM)/3.6 (BLR)/3.3 (CTM) times more efficient than HwG in MRESS. HwG is indeed
better than NwG as R2 suggested. We will include HwG as an additional baseline in camera-ready version. [R2]
...MCMC component for discrete...should be emphasised... Will do. [R2] ...different MCMC components for
discrete variables should be experimented. We have started (and will include in camera-ready version) experiments
with Turing, if for particle Gibbs. But we share R2's expectation that M-HMC would still be better. [R2] How
adaptation for NUTS is set up? Dual averaging, 0.8 target acceptance. [R2] The outperforming of M-HMC over
NUTS may be due to NUTS stuck in some of the modes...using a larger step size manually...might be simply
better... This turns out to be exactly the case. NUTS MRESS increased to 1.37 \times 10^{-3} with step size 5 (v.s. 4 from
dual averaging), outperforming M-HMC as one expects. We will update the paper with these observations.
[R1] We will add a concise summary of experiments. [R2] We visualized the kernel for 1d GMM in 4.1, showing
discrete updates and distant proposals. We will include the visualization, along with suggested proof/algorithm
re-organizations. [R3] We will keep only Fig. 4a for 1 chain in main paper to improve readability of the traceplots.
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