

# PolytopeWalk: Sparse MCMC Sampling over Polytopes

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DOI: [10.xxxxxx/draft](https://doi.org/10.xxxxxx/draft)

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Submitted: 01 January 1970

Published: unpublished

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## Summary

High dimensional sampling is an important computational tool in statistics and other disciplines, with applications ranging from Bayesian statistical uncertainty quantification, metabolic modeling in systems biology, to volume computation. We present PolytopeWalk, a new scalable Python library designed for uniform sampling over polytopes. The library provides an end-to-end solution, that includes preprocessing algorithms such as facial reduction and initialization methods. Six state-of-the-art MCMC algorithms on polytopes are implemented, including the Dikin, Vaidya, and John Walk. Additionally, we introduce novel sparse constrained formulations of these algorithms, enabling efficient sampling from sparse polytopes of the form  $\mathcal{K}_2 = \{x \in \mathbb{R}^d \mid Ax = b, x \succeq_k 0\}$ . This implementation maintains sparsity in  $A$ , ensuring scalability to high dimensional settings ( $d > 10^4$ ). We also include novel implementation of several pre-processing steps, including the Facial Reduction algorithm. PolytopeWalk is available at [github.com/ethz-randomwalk/polytopewalk](https://github.com/ethz-randomwalk/polytopewalk) with documentation at [polytopewalk.readthedocs.io/](https://polytopewalk.readthedocs.io/).

## Statement of Need

High dimensional sampling is a fundamental problem in many computational disciplines such as statistics, probability, and operation research. For example, sampling is applied in portfolio optimization (Calès et al. (2018)), metabolic networks in systems biology (Heirendt et al. (2018)) and volume approximation over convex shapes (Simonovits (2003)). Markov chain Monte Carlo (MCMC) sampling algorithms offer a natural and scalable solution to this problem. These algorithms construct a Markov chain whose stationary distribution matches the target distribution. By running the chain for a large number of steps to ensure mixing, MCMC algorithms can efficiently generate approximately independent samples close to the target distribution, while not suffering from the curse of dimension issues.

This package focuses on sampling from a uniform distribution over a user-specified polytope. We define the polytope as the following. Let  $A \in \mathbb{R}^{n \times d}$ ,  $b \in \mathbb{R}^n$  and define  $x \succeq_k y$  to mean that the last  $k$ -coordinates of  $x$  are greater than or equal to the corresponding coordinates of  $y$ , i.e.,  $\{x_{d-k+1} - y_{d-k+1} \geq 0, \dots, x_d - y_d \geq 0\}$ . Depending on whether we allow equality constraints, the sampling problem can be formalized in two forms:

1. The full-dimensional form:

$$\mathcal{K}_1 = \{x \in \mathbb{R}^d \mid Ax \leq b\}, \quad (1)$$

where  $\mathcal{K}_1$  is specified via  $n$  inequality constraints.

2. The constrained form:

$$\mathcal{K}_2 = \{x \in \mathbb{R}^d \mid Ax = b, x \succeq_k 0\}, \quad (2)$$

37 where  $\mathcal{K}_2$  is specified via  $n$  equality constraints and  $k$  coordinate inequality constraints.

38 Large polytopes with sparse constraints are common in many applications. For instance,  
39 the largest human metabolic network RECON3D is modeled as a 13543-dimensional sparse  
40 polytope (King et al. (2015)). Additionally, linear programming datasets from NetLib are  
41 naturally in the constrained form, where  $A$  matrix is sparse. These applications motivate the  
42 need for MCMC algorithms that leverage the large and sparse  $\mathcal{K}_2$  formulation. We develop  
43 novel interior-point-method-based MCMC algorithms optimized for large and sparse constrained  
44 polytopes. By exploiting sparsity, our algorithms scale well in both per-iteration cost and  
45 sampling efficiency as a function of increasing dimension, enabling effective sampling from  
46 polytopes with dimensions exceeding  $10^4$ .

47 Interior-point-method-based MCMC sampling algorithms on a polytope are modifications of  
48 the Ball Walk (Vempala (2005)), incorporating key concepts from interior-point methods in  
49 optimization. These algorithms operate in two primary steps. First, the algorithm generates  
50 a proposal distribution whose covariance matrix is state-dependent and equal to the inverse  
51 of the Hessian matrix of a specified barrier function, capturing the local geometry of the  
52 polytope. Second, the algorithm employs the Metropolis-Hastings accept-reject step to ensure  
53 that its stationary distribution is uniform on the polytope. Using a state-dependent proposal  
54 distribution that adapts to the polytope's local geometry, these MCMC algorithms achieve  
55 an improved mixing rate. Specific algorithms in this class include the Dikin Walk (Sachdeva  
56 & Vishnoi (2015)), Vaidya Walk (Chen et al. (2018)), John Walk (Chen et al. (2018)), and  
57 Lee Sidford Walk (Laddha et al. (2019)), with theoretical guarantees therein. Each of these  
58 methods leverages different barrier functions and are specialized for sampling distributed  
59 truncated on a polytope.

60 In PolytopeWalk, we implement 4 interior-point-method-based MCMC sampling algorithms in  
61 both the sparse, constrained formulation and the full-dimensional form. PolytopeWalk makes  
62 meaningful strides in the open-source development of MCMC, speeding up calculations for sparse  
63 high-dimensional sampling. On our Github, we demonstrate the scalability of PolytopeWalk  
64 against packages like Volesti. Finally, we provide an an open-source implementation of the  
65 Facial Reduction algorithm, used to handle degeneracy in polytopes.

## 66 Package Overview

67 PolytopeWalk is an open-source library written in C++ with Python wrapper code. There are 3  
68 main components of the package: preprocessing, sampling, and post processing. PolytopeWalk  
69 provides accelerated MCMC sampling algorithms in both the  $\mathcal{K}_1$  formulation~(1) and  $\mathcal{K}_2$   
70 formulation~(2). The source code is written in low-level C++ with Eigen for optimized linear  
71 algebra operations, glpk for a fast sparse linear programming solver, and pybind to enable  
72 Python binding.

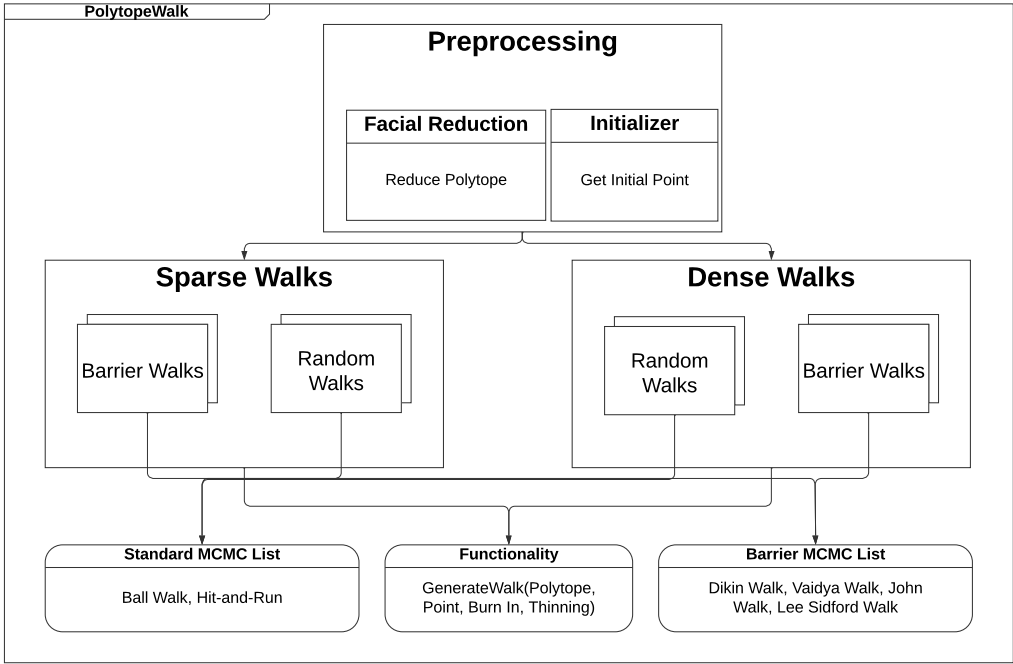


Figure 1: Code Structure of Package

Random Walk Algorithms

There are 2 main submodules: dense and sparse. The dense module includes implementations of the 6 random walks algorithms in the full-dimensional formulation. The sparse module includes implementations in the sparse, constrained formulation. Table I provides a broad overview of their respective mixing rates and cited paper.

Name	Mixing Time	Author
Ball Walk	$\tau(d^2 R^2 / r^2)$	Vempala (2005)
Hit and Run	$\tau(d^2 R^2 / r^2)$	Lovasz (1999)
Dikin Walk	$\tau(nd)$	Sachdeva et al. (2015)
Vaidya Walk	$\tau(n^{1/2} d^{3/2})$	Chen et al. (2018)
John Walk	$\tau(d^{2.5})$	Chen et al. (2018)
Lee Sidford Walk	$\tau(d^2)$	Laddha et al. (2019)

The mixing times refer to the required number of steps to converge to stationary distribution. In each,  $d$  refers to the dimension of the polytope and  $n$  refers to the number of boundaries ( $\mathcal{K}_1$  dimensions). In the first 2 walks,  $R^2/r^2$  means where the convex body contains a ball of radius  $r$  and is mostly contained in a ball of radius  $R$ .

Preprocessing Algorithms

PolytopeWalk comes with 2 preprocessing algorithms: initialization and facial reduction.

**Initialization:** If the user cannot specify a point inside of the polytope to start, PolytopeWalk provides a class to compute an initial point well within the polytope for both the full-dimensional formulation and constrained formulation. In  $\mathcal{K}_2$ , this is akin to solving the linear program

87 maximizing  $\delta \in \mathbb{R}$  s.t.  $Ax = b$  and  $x \succeq_k \vec{\delta}$ . In  $\mathcal{K}_1$ , this is equivalent to maximizing  $\delta \in \mathbb{R}$   
 88 such that  $Ax + \delta \cdot \mathbf{1} \leq b$  where  $\mathbf{1}$  is the all-ones vector  $\in \mathbb{R}^n$ .

89 **Facial Reduction:** We adopt the facial reduction algorithm implementation from Drusvyatskiy's  
 90 research (Drusvyatskiy et al. (2017)) (Im & Wolkowicz (2023)). The facial reduction  
 91 algorithm deals with cases of degeneracy in the polytope. In the constrained formulation  
 92  $\mathcal{K}_2 = \{x \in \mathbb{R}^d \mid Ax = b, x \succeq_k 0\}$ , degeneracy occurs when there is a lack of strict feasibility  
 93 in the polytope: there does not exist an  $x \in \mathbb{R}^d$  such that  $Ax = b$  and  $x \succ_k 0$ . The facial  
 94 reduction algorithm eliminates variables in the last  $k$  dimensions fixed at 0. It is far more  
 95 commonly used in linear programming than MCMC sampling. Degeneracy exists in polytopes  
 96 when the lower-dimensional polytope is embedded in a higher dimension. For example, if one  
 97 attempts to use MCMC algorithms to sample from a 2-dimensional hypercube in 5 dimensions,  
 98 the proposal distributions which generate an ellipsoid in 5 dimensions will certainly reject each  
 99 proposal. It is crucial that one projects the polytope back into the 2 dimensional form to run  
 100 the MCMC sampler efficiently. Facial Reduction accomplishes this task, ensuring numerical  
 101 stability for sampling.

## 102 Package Comparison

Feature	PolytopeWalk	Volesti	WalkR	Polyrun
Constrained Formulation	Y	N	Y	Y
Sparse Friendly	Y	N	N	N
C++ Implementation	Y	Y	N	N
Facial Reduction	Y	N	N	N
Dikin Walk	Y	Y	Y	N
Vaidya Walk	Y	Y	N	N
John Walk	Y	Y	N	N
Lee-Sidford Walk	Y	N	N	N

103 Table II contrasts the features of PolytopeWalk with Volesti (Chalkis & Fisikopoulos (2021)),  
 104 WalkR (Yao & Kane (2017)), and Polyrun (Ciomek & Kadziński (2021)). PolytopeWalk is  
 105 the first open-source package that enables users to leverage sparsity and use constrained  
 106 formulations of the polytopes to generate points uniformly with fast MCMC algorithms. We are  
 107 also one of the first to primarily focus on barrier walk MCMC samplers. PolytopeWalk includes  
 108 a C++ implementation with corresponding Python wrapper code. Conversely, Volesti is  
 109 implemented in C++ with some of its code represented in the Python library Dingo. Polyrun  
 110 only works on Java and WalkR on R. Notably, WalkR was removed from the CRAN repository,  
 111 motivating further open source MCMC sampling development.

## 112 Acknowledgements

113 Much of the work was done while Yuansi Chen was an assistant professor in the Department  
 114 of Statistical Science at Duke University. Both authors are partially supported by NSF  
 115 CAREER Award DMS-2237322, Sloan Research Fellowship and Ralph E. Powe Junior Faculty  
 116 Enhancement Awards.

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