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Introduction

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- hierachical Bayesian model, sampling to TT approx

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Theoretical and Technical Background

In this chapter we give a brief introduction into the methods, that we use in this thesis. We try to keep it as general as possible, as we get more specific in the results, section **??**.
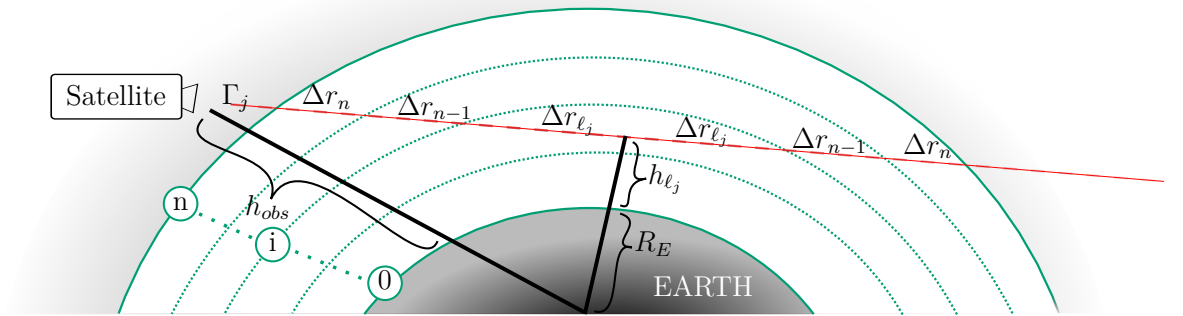
2.1 Forward Model

Figure 2.1: text

In this section we describe the forward model which we use to simulate data, as in section **??** and base the Bayesian inference, see section **??** and **??**.

As shown in Figure **??**, one measurement of a stationary satellite can be describes as the path integral along the line of sight Γ_j for $j = 1, 2, \ldots, m$. For

each measurement we can define a tangent height h_{ℓ_j} as the shortest distance along the line of sight to the earth.

The jth measurement, taken on line of sight Γ_j is modelled by the radiative transfer equation (RTE) [1]

$$y_j = \int_{\Gamma_j} B(\nu, T) k(\nu, T) \frac{p(T)}{k_{\text{B}} T(r)} x(r) \tau(r) \mathrm{d}r + \eta_j \tag{2.1}$$

$$\tau(r) = \exp\left\{ -\int_{r_{\text{obs}}}^{r} k(\nu, T) \frac{p(T)}{k_B T(r')} x(r') \mathrm{d}r' \right\} \tag{2.2}$$

where the path from the satellite along the line-of-sight of the jth pointing direction is Γ_j and the ozone concentration at distance r from the radiometer is $x(r)$ plus some noise η_j. Within the stratosphere the number density $p(T)/(k_{\text{B}} T(r))$ of molecules is dependent on the pressure $p(T)$, the temperature $T(r)$, and the Boltzmann constant k_{B}. The factor $\tau(r) \leq 1$ accounts for re-absorption of the radiation along the line-of-sight, which makes the RTE non linear. The absorption constant $k(\nu, T)$ for a single gas molecule at a specific wavenumber ν is given by the HITRAN database [2] and acts as a source function when multiplied with the black body radiation $B(\nu, T)$, given by Planck's law [3]. For fundamentals on the Radiative transfer equation we recommend Chapter one in [3].

To enable Matrix-Vector multiplication, we parametrise the ozone profile as a function of height, discretised into the n values in each of n layers of the discretised stratosphere where the ith layer is defined by two spheres of radii $h_{i-1} < h_i$, $i = 1, \ldots, n$, with h_0 and h_n. In between the heights h_{i-1} and h_i, each of the ozone concentration x_i, the pressure p_i, the temperature T_i, and thermal radiation is assumed to be constant. Above h_n and below h_0, the ozone concentration is set to zero, so no signal can be obtained. Then depending on the parameter of interest, which is either the ozone volume mixing ratio $\boldsymbol{x} = \{x_1, x_2, \ldots, x_n\} \in \mathbb{R}^n$ or the fraction of pressure and temperature $\boldsymbol{p/T} = \{p_1/T_1, p_2/T_2, \ldots, p_n/T_n\} \in \mathbb{R}^n$, we can rewrite the integral in Eq. (2.1) as e.g. as a vector multiplication $\boldsymbol{A_j}(\boldsymbol{x}, \boldsymbol{p}, \boldsymbol{T}) \boldsymbol{x}$, where the non-linear absorption $\tau(r)$ is included in $\boldsymbol{A_j}(\boldsymbol{x}, \boldsymbol{p}, \boldsymbol{T})$. Here, the row vector $\boldsymbol{A_j}(\boldsymbol{x}, \boldsymbol{p}, \boldsymbol{T}) \in \mathbb{R}^n$ defines a Kernel for each measurement so that the data vector

$$\boldsymbol{y} = \boldsymbol{A}(\boldsymbol{x}, \boldsymbol{p}, \boldsymbol{T}) \boldsymbol{x} + \boldsymbol{\eta} = \boldsymbol{A}(\boldsymbol{x}, \boldsymbol{p}, \boldsymbol{T}) \frac{\boldsymbol{p}}{\boldsymbol{T}} + \boldsymbol{\eta} \,. \tag{2.3}$$

can be written as a matrix vector multiplication, where the matrix $\boldsymbol{A}(\boldsymbol{x}, \boldsymbol{p}, \boldsymbol{T}) \in \mathbb{R}^{m \times n}$ and the noise vector $\boldsymbol{\eta} \in \mathbb{R}^m$.

Since the measurement process includes absorption $\tau(r)$ reducing measurements slighty and making the inverse problem only weakly non-linear. We use that to approximate the non-linear forward model $\boldsymbol{A}(\boldsymbol{x}, \boldsymbol{p}, \boldsymbol{T})$ with a map $\boldsymbol{M}$ so that $\boldsymbol{A}(\boldsymbol{x}, \boldsymbol{p}, \boldsymbol{T}) \approx \boldsymbol{M} \boldsymbol{A}_L$ Where each row $\boldsymbol{A}_{L,j}$ of matrix as $\boldsymbol{A}_L \in \mathbb{R}^{m \times n}$ is defined by the linear forward model, where absorption is neglected, e.g. $\tau = 1$. Then each entry in the row vector $\boldsymbol{A}_{L,j}$ is either defined by $B(\nu, T) S(\nu, T) \frac{\boldsymbol{p}(T)}{k_{\mathrm{B}} \boldsymbol{T}(r)} \mathrm{d}r$ or $B(\nu, T) S(\nu, T) \frac{\boldsymbol{x}}{k_{\mathrm{B}}} \mathrm{d}r$, as in Eq.. (2.1), depending on the parameter of interest. This poses a linear inverse problem with the forward map defined by the matrix $\boldsymbol{A} = \boldsymbol{M} \boldsymbol{A}_L$, where $\boldsymbol{M}$ is, more specifically, an affine map.

2.2 Affine Map

To approximate the non-linear forward model we use an affine map $M : \boldsymbol{A}_L \boldsymbol{x} \to \boldsymbol{A}(\boldsymbol{x}, \boldsymbol{p}, \boldsymbol{T}) \boldsymbol{x}$, which maps the linear forward model $\boldsymbol{A}_L \boldsymbol{x}$ onto the non-linear forward model $\boldsymbol{A}(\boldsymbol{x}, \boldsymbol{p}, \boldsymbol{T}) \boldsymbol{x}$.

An affine map is any linear map in between two vector spaces or affine spaces, where an affine space does not need to preserve a zero origin, see Def. 2.3.1. in [4]. In other words an affine map does not need to map to the origin of the associated vector space, or is a linear map on vector spaces including translation, or in the words of my supervisor, C. F., an affine map is a Taylor series of first order. For more information on affine spaces and maps we refer to the books [4, 5]

To find the affine map, we generate two affine subspaces spaces $V = \left\{ \boldsymbol{A}(\boldsymbol{x}^{(1)}, \boldsymbol{p}, \boldsymbol{T}), \dots, \boldsymbol{A}(\boldsymbol{x}^{(m)}, \boldsymbol{p}, \boldsymbol{T}) \right\}$ and $W = \left\{ \boldsymbol{A}_L \boldsymbol{x}^{(1)}, \dots, \boldsymbol{A}_L \boldsymbol{x}^{(m)} \right\}$ over the same field, with fixed $\boldsymbol{p}, \boldsymbol{T}$. Then we can use a linear solver to find the affine map $\boldsymbol{M}$ so that we can approximate the non linear forward model $\boldsymbol{A}(\boldsymbol{x}, \boldsymbol{p}, \boldsymbol{T}) \approx \boldsymbol{M} \boldsymbol{A}_L$ with a linear forward model, we go into further detail in section **??**. The parameter $\boldsymbol{x}$ is distributed as the so-called posterior distribution $\left\{ \boldsymbol{x}^{(1)}, \dots, \boldsymbol{x}^{(m)} \right\} \sim \pi(\boldsymbol{x}|\boldsymbol{\theta}, \boldsymbol{y})$ conditioned on the hyper-parameters $\boldsymbol{\theta}$, according to a Bayesian hierarchical model.

linear forward model
$$\boxed{V} \xrightarrow[\boldsymbol{M}]{\text{affine Map}} \boxed{W}$$
non-linear forward model

$$\boldsymbol{A}(\boldsymbol{x}, \boldsymbol{p}, \boldsymbol{T})\boldsymbol{x} \approx \boldsymbol{M}\boldsymbol{A}_L\boldsymbol{x} = \boldsymbol{A}\boldsymbol{x}$$
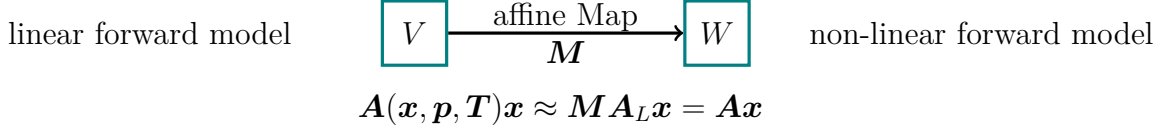
Figure 2.2: Schematics of Affine Map, which approximates the linear forward model to the non-linear forward model.

2.3   Bayesian Inference

In this this section we give a short introduction to Bayesian inference for a general parameter $\boldsymbol{x}$ given some data

$$\boldsymbol{y} = \boldsymbol{A}\boldsymbol{x} + \boldsymbol{\eta} \tag{2.4}$$

based on a linear forward model $\boldsymbol{A}$ and some noise $\boldsymbol{\eta}$. Later in section **??** we set up a more sophisticated Bayesian framework applied to the forward model in section **??**.

We can visualise the correlation structure of a measurement process through a hierarchiallly ordered directed acyclic graph (DAG), see Figure 2.3. As an observatory process naturally includes some random noise we include that in our DAG and classify the noise as a hyper-parameter in $\boldsymbol{\theta}$ [6]. Other hyper-parameters influence the parmeters $\boldsymbol{x}$ detemernistaclly, which are then mapped through the forward model onto the space of all measurables $\boldsymbol{u}$, from which we observe some data $\boldsymbol{y}$ including noise as previously mentioned. Drawing a DAG can help us to dependences within the measurement and modelling process. Given some data we inferer the distribution of the underling parameters and hyper-parameters by following the arrows in Figure 2.3 backwards and set up a Bayesian model, hierarchially ordered.

Within a linear Bayesian hierarchial model we need to define a likelihood function as well as a distribution over the unknown parameters $\boldsymbol{x}$ and hyper-parameters $\boldsymbol{\theta}$.

$$\boldsymbol{y}|\boldsymbol{x}, \boldsymbol{\theta} \sim \mathcal{N}(\boldsymbol{A}\boldsymbol{x}, \boldsymbol{\Sigma}(\boldsymbol{\theta})) \tag{2.5a}$$

$$\boldsymbol{x}|\boldsymbol{\theta} \sim \mathcal{N}(\boldsymbol{\mu}, \boldsymbol{Q}^{-1}(\boldsymbol{\theta})) \tag{2.5b}$$

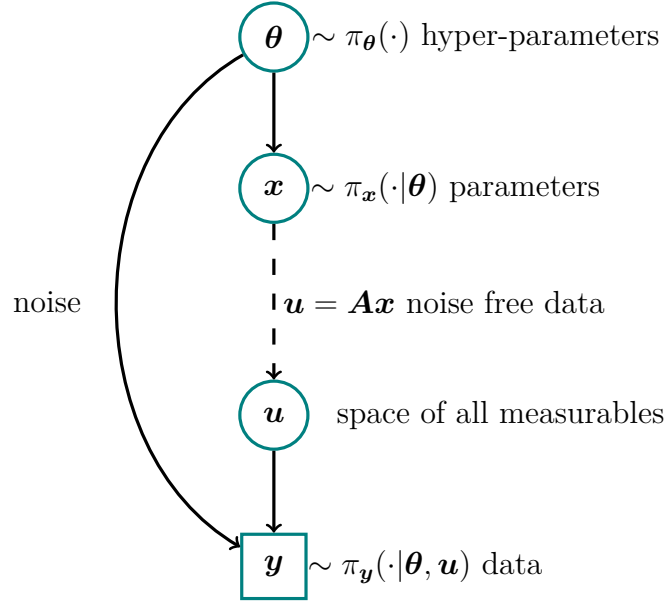$$\boldsymbol{\theta} \sim \pi(\boldsymbol{\theta})\,. \tag{2.5c}$$

Figure 2.3: The directed acyclic graph (DAG) for a linear inverse problem visualises statistical dependencies as solid line arrows and deterministic dependencies as dotted arrows. The parameters $\boldsymbol{x}$ have some statistical dependency of those hyper-parameters $\boldsymbol{\theta}$, which are distributed as $\pi(\boldsymbol{\theta})$. Then a parameter $\boldsymbol{x} \sim \pi_{\boldsymbol{x}}(\cdot|\boldsymbol{\theta})$ is mapped onto the space of all measurables $\boldsymbol{u} = \boldsymbol{A}\boldsymbol{x}$ deterministically through the linear forward model $\boldsymbol{A}$. From the space of all measurables we observe some data $\boldsymbol{y} = \boldsymbol{A}\boldsymbol{x} + \boldsymbol{\eta}$, statistically, so that $\boldsymbol{y} \sim \pi_{\boldsymbol{y}}(\cdot|\boldsymbol{\theta}, \boldsymbol{u})$, with naturally some random noise $\boldsymbol{\eta} \sim \pi_{\boldsymbol{\eta}}(\cdot|\boldsymbol{\theta})$.

Due to the assumption of Gaussian noise $\boldsymbol{\eta} \sim \mathcal{N}(0, \boldsymbol{\Sigma}(\boldsymbol{\theta}))$ the likelihood function $\pi(\boldsymbol{y}|\boldsymbol{x}, \boldsymbol{\theta})$, which includes information about the measurement process and gives a measure of how well parametersand hyperparemters fit given some data, is normally distributed with the noise covariance matrix $\boldsymbol{\Sigma}(\boldsymbol{\theta})$. More specifically because we choose a normally distributed prior $\pi(\boldsymbol{x}|\boldsymbol{\theta})$ with the prior precision matrix $\boldsymbol{Q}(\boldsymbol{\theta})$ and prior mean $\boldsymbol{\mu}$ this is a linear-Gaussian Bayesian hierarchical model as in [6], including some distribution over the hyper-parameters $\pi(\boldsymbol{\theta})$. One of the main strengths of a Bayesian framework is that through those prior and hyper-prior distributions we can incorporate functional dependencies as well model physical properties of the parameters $\boldsymbol{x}$.

Then Bayes Theorem gives the posterior distribution, the function of interest,

$$\pi(\boldsymbol{x}, \boldsymbol{\theta}|\boldsymbol{y}) = \frac{\pi(\boldsymbol{y}|\boldsymbol{x}, \boldsymbol{\theta})\pi(\boldsymbol{x}, \boldsymbol{\theta})}{\pi(\boldsymbol{y})} , \tag{2.6}$$

with the prior distribution $\pi(\boldsymbol{x}, \boldsymbol{\theta}) = \pi(\boldsymbol{x}|\boldsymbol{\theta})\pi(\boldsymbol{\theta})$ and the normalising constant $\pi(\boldsymbol{y})$. If the normalising constant is finite and non-zero we can approximate

the posterior distribution

$$\pi(\boldsymbol{x}, \boldsymbol{\theta}|\boldsymbol{y}) \propto \pi(\boldsymbol{y}|\boldsymbol{x}, \boldsymbol{\theta})\pi(\boldsymbol{x}, \boldsymbol{\theta}) \,. \tag{2.7}$$

and the expectation of any a function $h(\boldsymbol{x}(\boldsymbol{\theta}))$, where $\boldsymbol{x}$ may be depending on $\boldsymbol{\theta}$, can be described as

$$\mathrm{E}_{\boldsymbol{x}, \boldsymbol{\theta}|\boldsymbol{y}}[h(\boldsymbol{x}(\boldsymbol{\theta}))] = \int \int h(\boldsymbol{x}(\boldsymbol{\theta})) \, \pi(\boldsymbol{x}, \boldsymbol{\theta}|\boldsymbol{y}) \, \mathrm{d}\boldsymbol{x} \, \mathrm{d}\boldsymbol{\theta} \,, \tag{2.8}$$

which is usually a high dimensional integral and computationally not feasible to solve.

One way to work around the high dimensionality is to parameterise $\boldsymbol{x}$ using hyper-parameters $\boldsymbol{\theta}$ so that $\boldsymbol{x}(\boldsymbol{\theta})$. Another way is to seperate the posterior distribution over latent field $\boldsymbol{x}$ and the hyper-parameters $\boldsymbol{\theta}$. This is particular benefitial, when $\boldsymbol{x}$ is high dimensional, e.g. $\boldsymbol{x} \in \mathbb{R}^n$ with $n = 45$ and can not be parametrised, and $\boldsymbol{\theta}$ is low dimensional, e.g. two dimensional. Then by the law of total expectation [7] eq. 2.8 becomes

$$\mathrm{E}_{\boldsymbol{x}|\boldsymbol{y}}[h(\boldsymbol{x})] = \mathrm{E}_{\boldsymbol{\theta}|\boldsymbol{y}}\Big[\mathrm{E}_{\boldsymbol{x}|\boldsymbol{\theta}, \boldsymbol{y}}[h(\boldsymbol{x}(\boldsymbol{\theta}))]\Big] = \int \mathrm{E}_{\boldsymbol{x}|\boldsymbol{\theta}, \boldsymbol{y}}[h(\boldsymbol{x}(\boldsymbol{\theta}))] \, \pi(\boldsymbol{\theta}|\boldsymbol{y}) \, \mathrm{d}\boldsymbol{\theta} \,, \tag{2.9}$$

where in the case of a linear-Gaussian Bayesian hierarchical model $\mathrm{E}_{\boldsymbol{x}|\boldsymbol{\theta}, \boldsymbol{y}}[h(\boldsymbol{x}(\boldsymbol{\theta}))]$ is well defined. The marginal and then condtional method, does exactly that. Why MTC sperate hiughky correlated [**<empty citation>**]

Figure 4.1 in RUE and Held see appemdix Hight correlation bewten $\boldsymbol{\theta}$ and $\boldsymbol{x}$

2.3.1 Marginal and then Conditional

The marginal and then conditional (MTC) method factorises the full posterior distribution

$$\pi(\boldsymbol{x}, \boldsymbol{\theta}|\boldsymbol{y}) = \pi(\boldsymbol{x}|\boldsymbol{\theta}, \boldsymbol{y})\pi(\boldsymbol{\theta}|\boldsymbol{y}) \tag{2.10}$$

into the marginal posterior distribution $\pi(\boldsymbol{\theta}|\boldsymbol{y})$ and conditional posterior distribution $\pi(\boldsymbol{x}|\boldsymbol{\theta}, \boldsymbol{y})$.

For the in Eq. 2.5 specified linear-Gaussian Bayesian hierarchical model the marginal posterior distribution is given as

$$\pi(\boldsymbol{\theta}|\boldsymbol{y}) = \int \pi(\boldsymbol{x}, \boldsymbol{\theta}|\boldsymbol{y}) \, \mathrm{d}\boldsymbol{x} \tag{2.11}$$

$$\propto \sqrt{\frac{\det(\boldsymbol{\Sigma}^{-1}) \det(\boldsymbol{Q})}{\det(\boldsymbol{Q} + \boldsymbol{A}^T \boldsymbol{\Sigma}^{-1} \boldsymbol{A})}} \times \exp\left[-\frac{1}{2}(\boldsymbol{y} - \boldsymbol{A}\boldsymbol{\mu})^T \boldsymbol{Q}_{\boldsymbol{\theta}|\boldsymbol{y}}(\boldsymbol{y} - \boldsymbol{A}\boldsymbol{\mu})\right] \pi(\boldsymbol{\theta}), \tag{2.12}$$

with

$$\boldsymbol{Q}_{\boldsymbol{\theta}|\boldsymbol{y}} = \boldsymbol{\Sigma}^{-1} - \boldsymbol{\Sigma}^{-1} \boldsymbol{A}(\boldsymbol{A}^T \boldsymbol{\Sigma}^{-1} \boldsymbol{A} + \boldsymbol{Q})^{-1} \boldsymbol{A}^T \boldsymbol{\Sigma}^{-1}, \tag{2.13}$$

see Lemma 2 in [6]. Conditioned on the hyper-parameters $\boldsymbol{\theta}$ we can draw samples from the conditional posterior distribution

$$\boldsymbol{x}|\boldsymbol{\theta}, \boldsymbol{y} \sim \mathcal{N}\Big(\underbrace{\boldsymbol{\mu} + (\boldsymbol{A}^T \boldsymbol{\Sigma}^{-1} \boldsymbol{A} + \boldsymbol{Q})^{-1} \boldsymbol{A}^T \boldsymbol{\Sigma}^{-1}(\boldsymbol{y} - \boldsymbol{A}\boldsymbol{\mu})}_{\boldsymbol{\mu}_{\boldsymbol{x}|\boldsymbol{\theta}, \boldsymbol{y}}}, \underbrace{(\boldsymbol{A}^T \boldsymbol{\Sigma}^{-1} \boldsymbol{A} + \boldsymbol{Q})^{-1}}_{\boldsymbol{\Sigma}_{\boldsymbol{x}|\boldsymbol{y}, \boldsymbol{\theta}}} \Big) \tag{2.14}$$

using the Randomis then optimize (RTO) method, see section 2.5.2, or calculate weighted expectations e.g. $\mathrm{E}_{\boldsymbol{x}|\boldsymbol{\theta}, \boldsymbol{y}}[\boldsymbol{x}|\boldsymbol{\theta}, \boldsymbol{y}] = \boldsymbol{\mu}_{\boldsymbol{x}|\boldsymbol{\theta}, \boldsymbol{y}}$ or use Eq. 2.9 to calculate weighted expectations of $\mathrm{E}_{\boldsymbol{x}|\boldsymbol{\theta}, \boldsymbol{y}}[\mathrm{Var}(\boldsymbol{x}|\boldsymbol{\theta}, \boldsymbol{y})] = \boldsymbol{\Sigma}_{\boldsymbol{x}|\boldsymbol{y}, \boldsymbol{\theta}}$ with weights given by $\pi(\boldsymbol{\theta}|\boldsymbol{y})$. Note that the noise covariance $\boldsymbol{\Sigma} = \boldsymbol{\Sigma}(\boldsymbol{\theta})$ and the prior precision $\boldsymbol{Q} = \boldsymbol{Q}(\boldsymbol{\theta})$ are depending on hyper-parameters $\boldsymbol{\theta}$.

2.4   Regularisation

Another method to find a solution to a linear inverse problem as in Eq. 2.4 is to find a solution $\boldsymbol{x}_\lambda$ according to a data misfit norm and a regularisation semi-norm as in [6]. We will discuss the case of Tikhonov regularisation [8, 9] as this is the most similar to a linear-Gaussian hierarchial Bayesian model.

For a parameter $\boldsymbol{x}$ a linear forward model matrix $\boldsymbol{A}$ and some data $\boldsymbol{y}$ the data misfit norm

$$\|\boldsymbol{y} - \boldsymbol{A}\boldsymbol{x}\| \tag{2.15}$$

gives a measure of how good data fits to a mapped parameter $\boldsymbol{Ax}$ and the regularisation semi norm

$$\lambda \left\lVert \boldsymbol{Tx} \right\rVert \tag{2.16}$$

penalises $\boldsymbol{x}$ according to $\boldsymbol{T}$ and the regularisation parameter $\lambda > 0$. Given λ a regularised solution

$$\boldsymbol{x}_\lambda = \arg \min_{\boldsymbol{x}} \left\lVert \boldsymbol{y} - \boldsymbol{Ax} \right\rVert^2 + \lambda \left\lVert \boldsymbol{Tx} \right\rVert^2 \tag{2.17}$$

can be found by the derivativ

$$\nabla_{\boldsymbol{x}}\left\{(\boldsymbol{y} - \boldsymbol{Ax}_\lambda)^T(\boldsymbol{y} - \boldsymbol{Ax}_\lambda) + \lambda \boldsymbol{x}_\lambda^T \boldsymbol{T}^T \boldsymbol{T} \boldsymbol{x}_\lambda\right\} = 0 \tag{2.18}$$

$$\iff \quad \nabla_{\boldsymbol{x}}\left\{\boldsymbol{y}^T\boldsymbol{y} + \boldsymbol{x}_\lambda^T \boldsymbol{A}^T \boldsymbol{A} \boldsymbol{x}_\lambda - \boldsymbol{y}^T \boldsymbol{A} \boldsymbol{x}_\lambda - \boldsymbol{x}_\lambda^T \boldsymbol{A}^T \boldsymbol{y} + \lambda \boldsymbol{x}_\lambda^T \boldsymbol{T}^T \boldsymbol{T} \boldsymbol{x}_\lambda\right\} = 0 \tag{2.19}$$

$$\iff \qquad\qquad\qquad 2\boldsymbol{A}^T \boldsymbol{A} \boldsymbol{x}_\lambda - 2\boldsymbol{A}^T \boldsymbol{y} + 2\lambda \boldsymbol{T}^T \boldsymbol{T} \boldsymbol{x}_\lambda = 0. \tag{2.20}$$

Then a regularised solution is given as:

$$(\boldsymbol{A}^T \boldsymbol{A} + \lambda \boldsymbol{L})^{-1} \boldsymbol{A}^T \boldsymbol{y} = \boldsymbol{x}_\lambda, \tag{2.21}$$

where we can set $\boldsymbol{T}^T \boldsymbol{T} = \boldsymbol{L}$, which is typically matrix approximation of the nth derivative [9]. In practise $\boldsymbol{x}_\lambda$ is calculated for a range of λ, and is evaluated by the data-misfit norm with respect to the regularised semi-norm so that the best $\boldsymbol{x}_\lambda$ lays on the point of maximum curvature of a so-called L-Curve [10], which we will show in section **??**.

2.5 Sampling Methods

In this chapter we present the sampling methods used in this thesis and also argue why we can use sampling methods to calculate the expectation

$$\mathrm{E}_{\boldsymbol{x},\boldsymbol{\theta}|\boldsymbol{y}}[h(\boldsymbol{x}(\boldsymbol{\theta}))] = \underbrace{\int \int h(\boldsymbol{x}(\boldsymbol{\theta}))\pi(\boldsymbol{x},\boldsymbol{\theta}|\boldsymbol{y})\,\mathrm{d}\boldsymbol{\theta}\,\mathrm{d}\boldsymbol{x}}_{\boldsymbol{\mu}_{\mathrm{int}}} \tag{2.22}$$

of a function $h(\boldsymbol{x}(\boldsymbol{\theta}))$ with respect to a probability density $\pi(\boldsymbol{x}, \boldsymbol{\theta}|\boldsymbol{y})$. In the case where the calculation of the integral is not feasible we can approximate Eq. 2.22 with a sample based estimate

$$\mathrm{E}_{\boldsymbol{x}, \boldsymbol{\theta}|\boldsymbol{y}}[h(\boldsymbol{x}(\boldsymbol{\theta}))] \approx \underbrace{\frac{1}{N} \sum_{k=1}^{N} h(\boldsymbol{x}^{(k)}(\boldsymbol{\theta}^{(k)}))}_{\boldsymbol{\mu}_{\mathrm{samp}}}, \tag{2.23}$$

for large enough samples size N of a sample set $\mathcal{M} = \{(\boldsymbol{x}, \boldsymbol{\theta})^{(1)}, \dots, (\boldsymbol{x}, \boldsymbol{\theta})^{(k)}, \dots, (\boldsymbol{x}, \boldsymbol{\theta})^{(N)}\} \sim \pi(\boldsymbol{x}, \boldsymbol{\theta}|\boldsymbol{y})$. The variance of this unbiased estimator is $\mathcal{O}(1/N)$ so large samples sizes can reduce the uncertainty of $\boldsymbol{\mu}_{samp}$ [11]. We can do this as the central limit theorem states that the samples means $\boldsymbol{\mu}_{samp}^{(i)}$, of samples sets $\mathcal{M}_i$ for $i = 1, \dots, n$ of any distribution , converge in distribution to a normal distribution so that

$$\sqrt{n}(\boldsymbol{\mu}_{\mathrm{samp}}^{(i)} - \boldsymbol{\mu}_{\mathrm{int}}) \xrightarrow{\mathcal{D}} \mathcal{N}(0, \sigma^2)[12], \tag{2.24}$$

and if $\sigma^2 < \infty$ the error $\boldsymbol{\mu}_{\mathrm{samp}}^{(i)} - \boldsymbol{\mu}_{\mathrm{int}}$ is bound.

Now, we have to show that the methods we use actually draw samples $\mathcal{M} = \{(\boldsymbol{x}, \boldsymbol{\theta})^{(1)}, \dots, (\boldsymbol{x}, \boldsymbol{\theta})^{(k)}, \dots, (\boldsymbol{x}, \boldsymbol{\theta})^{(N)}\} \sim \pi(\boldsymbol{x}, \boldsymbol{\theta}|\boldsymbol{y})$ from the desired target distribution so that we can use sample based estimates as in Eq. 2.23. Here $\mathcal{M}$ is a Markov-Chain, where each sample $(\boldsymbol{x}, \boldsymbol{\theta})^{(k)}$ is only depending on the previous sample $(\boldsymbol{x}, \boldsymbol{\theta})^{(k-1)}$ [**<empty citation>**]. Markov-chain Monte Carlo methods generates such a chain $\mathcal{M}$ with random (Monte Carlo) proposals $(\boldsymbol{x}, \boldsymbol{\theta})^{(k)} \sim q(\cdot|(\boldsymbol{x}, \boldsymbol{\theta})^{(k-1)})$ according to a proposal distribution, where ergodicity of $\mathcal{M}$ is a sufficient criterium to use samples based estimates [9, 11]. The ergodicy theorem in [9] states that, if an aperiodic and irreducible Markov chain $\mathcal{M}$ is reversible then it converges towards a stationary unique equilibirum distribution $\pi(\boldsymbol{x}, \boldsymbol{\theta}|\boldsymbol{y})$. In other words if from any state in the chain we can reach any other state in the sampling space and the previous state, and we do not get stuck in periodic loop, then the chain converges towards a stationary distribution. In practise one can look at the trace $\pi(\boldsymbol{x}^{(k)}, \boldsymbol{\theta}^{(k)}|\boldsymbol{y})$ for $k = 1, \dots, N$ of the samples and eyeball ergodicity.

The sampling methods used in this thesis have proven ergodic properties, so we will cite and refer the reader to the respective documents.

2.5.1 Metropolis- within Gibbs sampling

As introduced in section 2.3.1 when using the MTC method will sample seperatly from $\pi(\boldsymbol{\theta}|\boldsymbol{y})$ and $\pi(\boldsymbol{x}|\boldsymbol{\theta},\boldsymbol{y})$. To sample from $\pi(\boldsymbol{\theta}|\boldsymbol{y})$ we use a Metropolis-within-Gibbs sampler as in [6] and discuss the 2 dimesnional case, as used in this thesis, with $\boldsymbol{\theta} = (\theta_1, \theta_2)$, where we do a metroplis step in θ_1 deirection and a gibbs stepp in θ_2 direction. Ergodicity is proven here [13].

The Metropolis-within-Gibbs algorithm starts with a initial guess $\boldsymbol{\theta}^{(t)}$ at $t = 0$. Then, we propose a new sample $\theta_1 \sim q(\theta_1|\theta_1^{(t-1)})$ conditioned on the previous state according to a symmetric proposal distribution $q(\theta_1|\theta_1^{(t-1)}) = q(\theta_1^{(t-1)}|\theta_1)$, which is a special case of the Metropolis-Hastings algorithm [13] and cancels when computing the acceptance probability α. We accept and set $\theta_1^{(t)} = \theta_1$ with

$$\alpha(\theta_1|\theta_1^{(t-1)}) = \min\left\{1, \frac{\pi(\theta_1|\theta_2^{(t-1)}, \boldsymbol{y})q(\theta_1^{(t-1)}|\theta_1)}{\pi(\theta_1^{(t-1)}|\theta_2^{(t-1)}, \boldsymbol{y})q(\theta_1|\theta_1^{(t-1)})}\right\} \tag{2.25}$$

or reject and keep $\theta_1^{(t)} = \theta_1^{(t-1)}$, which we do by comparing α to a uniform random number $u \sim \mathcal{U}(0, 1)$. Next, we do a Gibbs step in θ_2 direction, where Gibbs sampling is a special case of the Metropolis-Hastings algorithm with acceptance probaility of 1, and draw the next sample $\theta_2^{(t)} \sim \pi(\cdot|\theta_1^{(t)}, \boldsymbol{y})$ conditioned on $\theta_1^{(t)}$ at step t. We reapet this N times and assure convergence independent of the intial sample (irreducibility) as we discard samples after the so-called burn-in period so that we produce a Markov-Chain of length $N - N_{\text{burn-in}}$.

2.5.2 Draw a sample from a multivariate normal distribution

after sampling from $\pi(\boldsymbol{\theta}|\boldsymbol{y})$ we draw samples from $\pi(\boldsymbol{x}|\boldsymbol{\theta},\boldsymbol{y})$ within the MTC scheme. For Linear Gaussian Bayesian hierarchical model we can draw a sample $\boldsymbol{x}$ from the multivariate normal distribution$\pi(\boldsymbol{x}|\boldsymbol{\theta},\boldsymbol{y})$ using the radomize then optimise (RTO) method [14].

In doing so we can rewrite the full conditional normal distribution $\pi(\boldsymbol{x}|\boldsymbol{y},\boldsymbol{\theta})$ to:

$$\pi(\boldsymbol{x}|\boldsymbol{y},\boldsymbol{\theta}) \propto \pi(\boldsymbol{y}|\boldsymbol{x},\boldsymbol{\theta})\pi(\boldsymbol{x}|\boldsymbol{\theta}) \tag{2.26}$$

$$= \exp\|\hat{\boldsymbol{A}}\boldsymbol{x} - \hat{\boldsymbol{y}}\|^2 , \tag{2.27}$$

Algorithm 1: Metropolis within Gibbs

1: Initialize and suppose two dimensional vector $\boldsymbol{\theta}^{(0)} = (\theta_1^{(0)}, \theta_2^{(0)})$
2: **for** $k = 1, \ldots, N$ **do**
3:    Propose $\theta_1 \sim q(\cdot|\theta_1^{(t-1)}) = q(\theta_1^{(t-1)}|\cdot)$
4:    Compute

$$\alpha(\theta_1|\theta_1^{(t-1)}) = \min\left\{1, \frac{\pi(\theta_1|\theta_2^{(t-1)}, \boldsymbol{y})q(\theta_1^{(t-1)}|\theta_1)}{\pi(\theta_1^{(t-1)}|\theta_2^{(t-1)}, \boldsymbol{y})q(\theta_1|\theta_1^{(t-1)})}\right\}$$

5:    Draw $u \sim \mathcal{U}(0, 1)$
6:    **if** $\alpha \geq u$ **then**
7:       Accept and set $\theta_1^{(t)} = \theta_1$
8:    **else**
9:       Reject and keep $\theta_1^{(t)} = \theta_1^{(t-1)}$
10:    **end if**
11:    Draw $\theta_2^{(t)} \sim \pi(\cdot|\theta_1^{(t)}, \boldsymbol{y})$
12: **end for**
13: Output: $\boldsymbol{\theta}^{(0)}, \ldots, \boldsymbol{\theta}^{(k)}, \ldots, \boldsymbol{\theta}^{(N)} \sim \pi(\boldsymbol{\theta}|\boldsymbol{y})$

---

where

$$\hat{\boldsymbol{A}} = \begin{bmatrix} \boldsymbol{\Sigma}^{-1/2}(\boldsymbol{\theta})\boldsymbol{A} \\ \boldsymbol{Q}^{1/2}(\boldsymbol{\theta}) \end{bmatrix}, \quad \hat{\boldsymbol{y}} = \begin{bmatrix} \boldsymbol{\Sigma}^{-1/2}(\boldsymbol{\theta})\boldsymbol{y} \\ \boldsymbol{Q}^{1/2}(\boldsymbol{\theta})\boldsymbol{\mu} \end{bmatrix} \text{[15]}. \tag{2.28}$$

Then one sample can be computed by minimising the following equation with respect to $\hat{\boldsymbol{x}}$:

$$\boldsymbol{x}_i = \arg\min_{\hat{\boldsymbol{x}}} \|\hat{\boldsymbol{A}}\hat{\boldsymbol{x}} - (\hat{\boldsymbol{y}} + \boldsymbol{b})\|^2, \quad \boldsymbol{b} \sim \mathcal{N}(\boldsymbol{0}, \boldsymbol{I}), \tag{2.29}$$

where we add a randomised perturbation $\boldsymbol{b}$. Similarly as in section 2.4 we can rewrite the argument of Eq. 2.28 to

$$(\boldsymbol{A}^T\boldsymbol{\Sigma}^{-1}(\boldsymbol{\theta})\boldsymbol{A} + \boldsymbol{Q}(\boldsymbol{\theta}))\boldsymbol{x}_i = \boldsymbol{A}^T\boldsymbol{\Sigma}^{-1}(\boldsymbol{\theta})\boldsymbol{y} + \boldsymbol{Q}(\boldsymbol{\theta})\boldsymbol{\mu} + \boldsymbol{v}_1 + \boldsymbol{v}_2, \tag{2.30}$$

where we substitute $-\hat{\boldsymbol{A}}^T\boldsymbol{b} = \boldsymbol{v}_1 + \boldsymbol{v}_2$ so that $\boldsymbol{v}_1 \sim \mathcal{N}(\boldsymbol{0}, \boldsymbol{A}^T\boldsymbol{\Sigma}^{-1}(\boldsymbol{\theta})\boldsymbol{A})$ and $\boldsymbol{v}_2 \sim \mathcal{N}(\boldsymbol{0}, \boldsymbol{Q}(\boldsymbol{\theta}))$ are independent random variables [6, 14].

If within the MTC scheme the Markov chain from the marginal posterior is ergodic then with independent samples $\boldsymbol{x}^{(k)}$ from the full conditional $\pi(\boldsymbol{x}|\boldsymbol{\theta}, \boldsymbol{y})$ the combined chain $\{(\boldsymbol{x}, \boldsymbol{\theta})^{(1)}, \ldots, (\boldsymbol{x}, \boldsymbol{\theta})^{(k)}, \ldots, (\boldsymbol{x}, \boldsymbol{\theta})^{(N)}\} \sim \pi(\boldsymbol{x}, \boldsymbol{\theta}|\boldsymbol{y})$ is ergodic as well [16].

2.5.3   t-walk

If there is a functional dependency of the parameters $\boldsymbol{x}$ and the hyper-parameters $\boldsymbol{\theta}$ so that $\boldsymbol{x}(\boldsymbol{\theta})$ we can use the t-walk algorithm by Christens and Fox on $\pi(\boldsymbol{\theta}|\boldsymbol{y})$. We use the t-walk as a black box sampler, where convergence is guaranteed by construction [17].

2.6   Numerical Approxiamtion Methods - Tensor Train

Using the tensor train (TT) format to approximate a d-dimensional function $\pi(x)$, with $x \in \mathbb{R}^d$ on a discrete grid with n^d points, enables us to compute marginal posterior probability distributions cheaply. As the name suggest the tensor train format is a train of tensors, more specifically two and three dimensional tensors which we call cores $\pi_k \in \mathbb{R}^{r_{k-1} \times n \times r_k}$ and are connected through ranks r_k and r_{k-1} for each $k = 1, \ldots, d$, with the number of grid points in each dimension n, as in Figures 2.4 and **??** displayed. For the first and last dimensional the core outer ranks are $r_0 = r_d = 1$, so that for $x = x_1, \ldots, x_d$ the function value $\pi(x) = \left(\pi_1(x_1) \ldots \pi_d(x_d)\right) \in \mathbb{R}$ is a vector-matrix-vector multiplication and each core $\pi_k(x_k)$ at a fixed x_k on the approximated grid has dimensions $r_{k-1} \times 1 \times r_k = r_{k-1} \times r_k$.
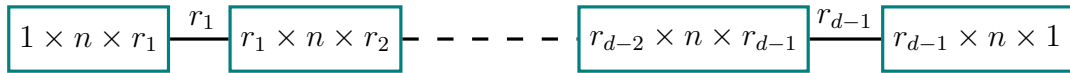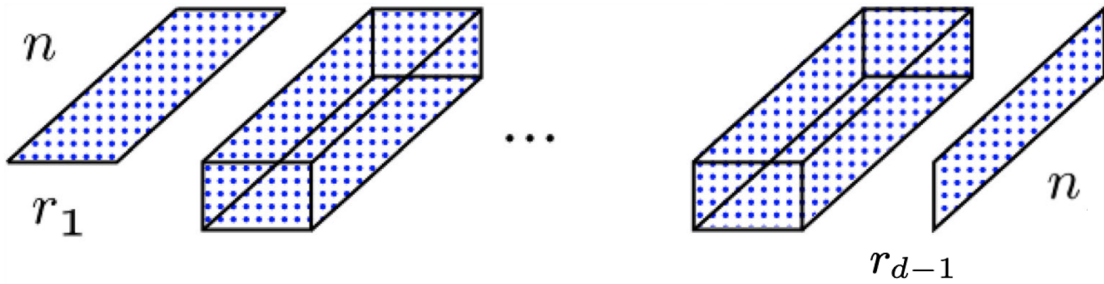

Figure 2.4: text

Figure 2.5: nice matries picture [18]

Asumme that the triple $(\Omega, \mathcal{F}, \mathbb{P})$ is called a probability space over the whole sample space Ω with the collection $\mathcal{F}$ of very countable subset $\{A_n\}_{n \in \mathbb{N}}$, where a Event A_n is a subset of Ω, $A_n \subseteq \Omega$, and a map $\mathbb{P} : \mathcal{F} \longrightarrow \mathbb{R}$. We assume $\mathbb{P}$ fullfils probality measure and $\mathcal{F}$ is a σ-algebra see Appendix **??**. We call $\mathbb{P}(A)$ the probaility of an event $A \subseteq \mathcal{F}$

$$\mathbb{P}(A) = \int_A \mathrm{d}\mathbb{P}[\textbf{<empty citation>}]. \tag{2.31}$$

using the Radon-Nikodym theroem [**<empty citation>**]

$$\mathbb{P}(A) = \int_A \frac{\mathrm{d}\mathbb{P}}{\mathrm{d}x} \, \mathrm{d}x \tag{2.32}$$

where $\mathrm{d}x$ is a measure on the same probality space and we call this the lebesgue measure and $\frac{\mathrm{d}\mathbb{P}}{\mathrm{d}x}$ is often called Radon-Nikodym derivativ, which can be set to the probaility density $\pi(x)$ of a random variable $X : \Omega \longrightarrow \mathbb{R}$. Next we intrdouce a weight function $\lambda(x)$, which can be helpful for qraudrature rules, so that

$$\mathbb{P}(A) = \int_A \lambda(x)\pi(x) \, \mathrm{d}x \, , \tag{2.33}$$

where Cui et al. [19] call $\lambda(x)$ the ' product-from Lebsegue measurable weighting functions" and define it as $\lambda(\mathcal{X}) = \prod_{i=1}^{d} \lambda_i(\mathcal{X}_i)$ and $\lambda_i(\mathcal{X}_i) = \int_{\mathcal{X}_i} \lambda_i(x_i)\mathrm{d}x_i$.

For the caclultion of marginal PDF, asuume the random variabes $(X_1, \ldots, X_d) : \Omega \longrightarrow \mathbb{R}^d$ define a joint distributiont trought the probailty density function $\pi(x)$ then the marginal distribution is

$$f_{X_k}(x_k) = \int_{\mathbb{R}} \cdots \int_{\mathbb{R}} \lambda(x)\pi(x) \, \mathrm{d}x_1 \cdots \mathrm{d}x_{k-1} \, \mathrm{d}x_{k+1} \cdots \mathrm{d}x_d \tag{2.34}$$

Consequently, using a tensor train approxiamtion, the marginal target function

$$\begin{aligned} f_{X_k}(x_k) = \frac{1}{z} \bigg| & \left(\int_{\mathbb{R}} \pi_1(x_1)\lambda_1(x_1)\mathrm{d}x_1 \right) \cdots \left(\int_{\mathbb{R}} \pi_{k-1}(x_{k-1})\lambda_{k-1}(x_{k-1})\mathrm{d}x_{k-1} \right) \\ & \pi_k(x_k)\lambda_k(x_k) \\ & \left(\int_{\mathbb{R}} \pi_{k+1}(x_{k+1})\lambda_{k+1}(x_{k+1})\mathrm{d}x_{k+1} \right) \cdots \left(\int_{\mathbb{R}} \pi_d(x_d)\lambda_d(x_d)\mathrm{d}x_d \right) \bigg| \end{aligned} \tag{2.35}$$

is given by integration over each core [20] including some normalising constant z [19].

From here we follow the notation and procedure mostly from Cui et al. [19] and approximate the square root

$$\sqrt{\pi(x)} \approx g(x) = \boldsymbol{G}_1(x_1), \dots, \boldsymbol{G}_k(x_k), \dots, \boldsymbol{G}_d(x_d) \qquad (2.36)$$

for numerical stability, where each TT-core

$$G_k^{(\alpha_{k-1}, \alpha_k)}(x_k) = \sum_{i=1}^{n_k} \phi_k^{(i)}(x_k) \boldsymbol{A}_k[\alpha_{k-1}, i, \alpha_k], \quad k = 1, \dots, d, \qquad (2.37)$$

is associated kth coefficient tensor $\boldsymbol{A}_k \in \mathbb{R}^{r_{k-1} \times n_k \times r_k}$ and the k-th basis functions $\phi_k^{(i)}(x_k)$. We assume the function

$$\pi(x) \approx \gamma' + g^2(x) , \qquad (2.38)$$

is approximated through the TT decomposition $g(x)$, where the error γ' assures positivity and is chosen according to the L_2 norm

$$\gamma' \leq \frac{1}{\lambda(\mathcal{X})} \|g - \sqrt{\pi}\|_2^2 . \qquad (2.39)$$

Then the normalised target function is

$$f_X(x) = \frac{1}{z}\pi(x)\lambda(x) = \frac{1}{z}(\gamma'\lambda(x) + g^2(x)\lambda(x)) . \qquad (2.40)$$

Given the tensor train approximation of the sqared rooted function $\sqrt{\pi}$ can be expressed as

$$
\begin{aligned}
f_{X_k}(x_k) = \frac{1}{z}\Bigg(&\gamma' \prod_{i=1}^{k-1} \lambda_i(\mathcal{X}_i) \prod_{i=k+1}^{d} \lambda_i(\mathcal{X}_i) \\
&+ \left(\int_{\mathbb{R}} \boldsymbol{G}_1^2(x_1)\lambda_1(x_1)\mathrm{d}x_1\right) \cdots \left(\int_{\mathbb{R}} \boldsymbol{G}_{k-1}^2(x_{k-1})\lambda_{k-1}(x_{k-1})\mathrm{d}x_{k-1}\right) \\
&\boldsymbol{G}_k^2(x_k)\lambda_k(x_k) \\
&\left(\int_{\mathbb{R}} \boldsymbol{G}_{k+1}^2(x_{k+1})\lambda_{k+1}(x_{k+1})\mathrm{d}x_{k+1}\right) \cdots \left(\int_{\mathbb{R}} \boldsymbol{G}_d^2(x_d)\lambda_d(x_d)\mathrm{d}x_d\right)\Bigg) .
\end{aligned}
$$
$$(2.41)$$

To efficiently calculate these marginals on can use a procedure similar to something that is called left and right orthogonalisation of cores [21]. To do so we define the mass matrix $\boldsymbol{M}_k \in \mathbb{R}^{n_k \times n_k}$ by

$$\boldsymbol{M}_k[i, j] = \int_{X_k} \phi_k^{(i)}(x_k)\phi_k^{(j)}(x_k)\lambda(x_k) \, dx_k, \quad i = 1, \dots, n_k, \quad j = 1, \dots, n_k, \qquad (2.42)$$

where $\{\phi_k^{(i)}(x_k)\}_{i=1}^{n_k}$ is the set of basis functions for the k-th coordinate.

2.6.1 Marginal Functions

We calculate the marginal functions through procedures, which we call backward

marginalisation [19] and forward marginalisation. We gain the coefficient matrices

$\boldsymbol{B}_k$ through backward marginalisation and the coefficient matrices $\boldsymbol{B}_{pre,n}$ through

forward marginalisation, which enables us to calculate marginal function similar

to [19]. The proposition 1 to caculte $\boldsymbol{B}_k$ is taken from [19].

Proposition 1 (Backward Marginalisation)**:** Starting with the last coordinate $k = d$, we set $\boldsymbol{B}_d = \boldsymbol{A}_d$. The following procedure can be used to obtain the coefficient tensor $\boldsymbol{B}_{k-1} \in \mathbb{R}^{r_{k-2} \times n_{k-1} \times r_{k-1}}$, which we need for defining the marginal function $f_{X_k}(x_k)$:

1. Use the Cholesky decomposition of the mass matrix, $\boldsymbol{L}_k \boldsymbol{L}_k^\top = \boldsymbol{M}_k \in \mathbb{R}^{n_k \times n_k}$, to construct a tensor $\boldsymbol{C}_k \in \mathbb{R}^{r_{k-1} \times n_k \times r_k}$:

$$\boldsymbol{C}_k[\alpha_{k-1}, \tau, l_k] = \sum_{i=1}^{n_k} \boldsymbol{B}_k[\alpha_{k-1}, i, l_k] \boldsymbol{L}_k[i, \tau]. \qquad (2.43)$$

2. Unfold $\boldsymbol{C}_k$ along the first coordinate and compute the thin QR decomposition, so that $\boldsymbol{C}_k^{(R)} \in \mathbb{R}^{r_{k-1} \times (n_k r_k)}$:

$$\boldsymbol{Q}_k \boldsymbol{R}_k = \left(\boldsymbol{C}_k^{(R)}\right)^\top. \qquad (2.44)$$

3. Compute the new coefficient tensor:

$$\boldsymbol{B}_{k-1}[\alpha_{k-2}, i, l_{k-1}] = \sum_{\alpha_{k-1}=1}^{r_{k-1}} \boldsymbol{A}_{k-1}[\alpha_{k-2}, i, \alpha_{k-1}] \boldsymbol{R}_k[l_{k-1}, \alpha_{k-1}]. \quad (2.45)$$

We start the forward marginalisation with the first dimension as in Proposition 2.

Proposition 2 (Forward Marginalistaion)**:** Starting with the first coordinate $k = 1$, we set $\boldsymbol{B}_{pre,1} = \boldsymbol{A}_1$. The following procedure can be used to obtain the coefficient tensor $\boldsymbol{B}_{pre,k+1} \in \mathbb{R}^{r_k \times n_{k+1} \times r_{k+1}}$ for defining the marginal function $f_{X_k}(x_k)$:

1. Use the Cholesky decomposition of the mass matrix, $\boldsymbol{L}_k \boldsymbol{L}_k^\top = \boldsymbol{M}_k \in \mathbb{R}^{n_k \times n_k}$, to construct a tensor $\boldsymbol{C}_k \in \mathbb{R}^{r_{k-1} \times n_k \times r_k}$:

$$\boldsymbol{C}_{pre,k}[\alpha_{k-1}, \tau, l_k] = \sum_{i=1}^{n_k} \boldsymbol{L}_k[i, \tau] \boldsymbol{B}_{pre,k}[\alpha_{k-1}, i, l_k]. \qquad (2.46)$$

2. Unfold $\boldsymbol{C}_{pre,k}$ along the first coordinate and compute the thin QR decomposition, so that $\boldsymbol{C}_{pre,k}^{(R)} \in \mathbb{R}^{(r_{k-1} n_k) \times r_k}$:

$$\boldsymbol{Q}_{pre,k} \boldsymbol{R}_{pre,k} = (\boldsymbol{C}_{pre,k}^{(R)}). \qquad (2.47)$$

3. Compute the new coefficient tensor $\boldsymbol{B}_{pre,k+1} \in \mathbb{R}^{r_{k-1} \times n_k \times r_k}$:

$$\boldsymbol{B}_{pre,k+1}[l_{k+1}, i, \alpha_{k+1}] = \sum_{\alpha_k=1}^{r_k} \boldsymbol{R}_{pre,k}[l_{k+1}, \alpha_k] \boldsymbol{A}_{k+1}[\alpha_k, i, \alpha_{k+1}]. \qquad (2.48)$$

The marginal PDF of X_k can be expressed as

$$f_{X_k}(x_k) = \frac{1}{z}\left(\gamma' \prod_{i=1}^{k-1} \lambda_i(X_i) \prod_{i=k+1}^{d} \lambda_i(X_i) + \sum_{l_{k-1}=1}^{r_{k-1}} \sum_{l_k=1}^{r_k} \left(\sum_{i=1}^{n_k} \phi_k^{(i)}(x_k) \boldsymbol{D}_k[l_{k-1}, i, l_k]\right)^2\right) \lambda_k(x_k),$$
$$(2.49)$$

where $\boldsymbol{D}_k \in \mathbb{R}^{r_{k-1} \times n_k \times r_k}$ and $\boldsymbol{R}_{pre,k-1} \in \mathbb{R}^{r_{k-1} \times r_{k-1}}$ and $\boldsymbol{B}_k \in \mathbb{R}^{r_{k-1} \times n_k \times r_k}$

$$\boldsymbol{D}_k[l_{k-1}, i, l_k] = \sum_{\alpha_{k-1}=1}^{r_{k-1}} \boldsymbol{R}_{pre,k-1}[l_{k-1}, \alpha_{k-1}] \boldsymbol{B}_k[\alpha_{k-1}, i, l_k]. \qquad (2.50)$$

In the special case for the first dimension, $f_{X_1}(x_1)$ can be expressed as

$$f_{X_1}(x_1) = \frac{1}{z}\left(\gamma' \prod_{i=2}^{d} \lambda_i(\mathcal{X}_i) + \sum_{l_1=1}^{r_1} \left(\sum_{i=1}^{n_1} \phi_1^{(i)}(x_1) \boldsymbol{D}_1[i, l_1]\right)^2\right) \lambda_1(x_1), \qquad (2.51)$$

where $\boldsymbol{D}_1[i, l_1] = \boldsymbol{B}_1[\alpha_0, i, l_1]$ and $\alpha_0 = 1$, and similiarly in the last dimesnion

$$f_{X_n}(x_n) = \frac{1}{z}\left(\gamma' \prod_{i=1}^{d-1} \lambda_i(\mathcal{X}_i) + \sum_{l_{n-1}=1}^{r_{n-1}} \left(\sum_{i=1}^{n_1} \phi_1^{(i)}(x_1) \boldsymbol{D}_n[l_{n-1}, i]\right)^2\right) \lambda_n(x_n), \qquad (2.52)$$

where $\boldsymbol{D}_n[l_{n-1}, i] = \boldsymbol{B}_{pre,n}[l_{n-1}, i, \alpha_n]$ and $\alpha_n = 1$.

Appendices

A
Correlation Structure

In the book Gaussian Markov Random Fields, Rue and Held show the correlation structure between the hyper-parameter μ and the latent field $\boldsymbol{x}$, which slows down convergence especially when using Gibbs samplers. They consider the hierarchichal sturcture

$$\mu \sim \mathcal{N}(0,1) \tag{A.1}$$

$$\boldsymbol{x}|\mu \sim \mathcal{N}(\mu\mathbf{1}, \boldsymbol{Q}^{-1}), \tag{A.2}$$

and use a Gibbs sampler over the full conditionals

$$\mu^{(k)}|\boldsymbol{x}^{(k)} \sim \mathcal{N}\left(\frac{\mathbf{1}^T\boldsymbol{Q}\boldsymbol{x}^{(k-1)}}{1+\mathbf{1}^T\boldsymbol{Q}\mathbf{1}}, (1+\mathbf{1}^T\boldsymbol{Q}\mathbf{1})^{-1}\right) \tag{A.3}$$

$$\boldsymbol{x}^{(k)}|\mu^{(k)} \sim \mathcal{N}(\mu^{(k)}\mathbf{1}, \boldsymbol{Q}^{-1}). \tag{A.4}$$

In Figure A.1 one can clearly see that when steps only in the μ-direction, x-axis, or in the $\boldsymbol{x}$-direction, vertical-(y)-axis, are allowed it takes a lot of steps to explore the parameter space due to high correlation in between μ and $\boldsymbol{x}$. A solution is to update (μ, x) jointly, where, since μ is one dimensional, effectively only marginal density of μ, by integrating out $\boldsymbol{x}$ of the joint density $\pi(\mu, x)$, is needed.

$$\mu^\star \sim q(\mu^\star|\mu^{(k-1)}) \tag{A.5}$$

$$\boldsymbol{x}^{(k)}|\mu^\star \sim \mathcal{N}(\mu^\star\mathbf{1}, \boldsymbol{Q}^{-1}) \tag{A.6}$$
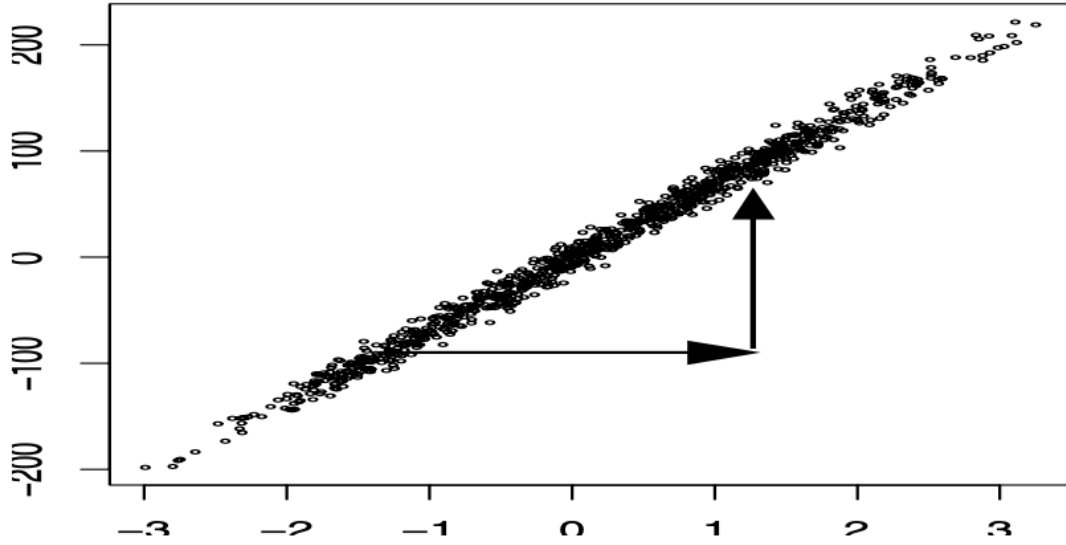
Figure A.1: Figure 4.1 Figure (a) shows the marginal chain for µ over 1000 iterations of the marginal chain for the hyperparmeter with a specuifuc autoregressive proces defined in Q The algorithm updates successively μ and x from their full conditionals. Figure (b) displays the pairs $(\mu(k)$, $1^T Q x(k)$), with$\mu(k)$ on the horizontal axis. The slow mixing (and convergence) of $\mu$is due to the strong dependence with $1^T Q x(k)$ as only horizontal and vertical moves are allowed. The arrows illustrate how a joint update can improve the mixing (and convergence)

With a simple MCMC algorithm on μ one can explore the sample space of μ and only draw a sample $\boldsymbol{x}$ from its full conditional after e.g. $\mu^\star$ is accepted.

Mesure theroy

Recall that the probability space $(\Omega, \mathcal{F}, \mathbb{P})$, where we call Ω the samples spae with a collection $\mathcal{F}$, which is a σ-algebra, of very countable subset $\{A_n\}_{n \in \mathbb{N}}$. We call A_n an Event in Ω, $A_n \subseteq \Omega$, and a map $\mathbb{P} : \mathcal{F} \longrightarrow \mathbb{R}$ a probavlity measure. Now we like to define a σ algebra and a prpbailty measure.

B.1   sigma alrgbea

A collections of subsets $\mathcal{F}$ is called sigma algebra if

- $\emptyset, \Omega \in \mathcal{F}$

- if $A \in \mathcal{F}$ then $A^C := A/\Omega \in A$

- if $A_1, A_2, \cdots \in \mathcal{F}$ then $\bigcup_{j \in \mathbb{N}} A_j \in \mathcal{F}$

In other words the empty set $\emptyset$ and the whole sample space Ω should always lay in $\mathcal{F}$ If we take any subset A in $\mathcal{F}$ the complement A^C, which is the sample space without A, A/Ω, has to lay in $\mathcal{F}$ as well. So if we able to calculate the probability $\mathbb{P}(A)$ we have to calculate the probability of not A,$\mathbb{P}(A^C)$. Finally, if the collection of countable subsets $A_1, A_2, \ldots$ lays in $\mathcal{F}$ then the union $\bigcup_{j \in \mathbb{N}} A_j$ also has to lay in $\mathcal{F}$.

B.2 probailty measure

For a probability measure we require

- $\mathbb{P}(\Omega) = 1$ and $\mathbb{P}(\emptyset) = 0$

- $\mathbb{P}(A) \in [0,1]$

- $\mathbb{P}(A \cup B) = \mathbb{P}(A) + \mathbb{P}(B)$ if A, B are disjoint or $A \cap B = \emptyset$

- $\mathbb{P}(\bigcup_{j \in \mathbb{N}} A_j) = \sum_{j \in \mathbb{N}} \mathbb{P}(A_j)$ if we have pairwise disjoint sets or $A_i \cap A_j = \emptyset$ for $i \neq j$

In other words the probability over the whole sample space should be equal to one and the probability over the empty set is zero. So for every subset A of the sample space Ω the probability $\mathbb{P}(A)$ lays in between zero and one. If we have two subsets A and B with no overlap then the probability of the union of those two subset , $\mathbb{P}(A \cup B)$,is equal to the sum of the probability of each of those subsets, $\mathbb{P}(A) + \mathbb{P}(B)$. This has to hold for the more general case of all countable unions of subsets $\bigcup_{j \in \mathbb{N}} A_j$.

References

[1]   C. Readings. *Envisat MIPAS An Instrument for Atmospheric Chemistry and Climate Research*. Noordwijk: ESA Publications Division, 2000.

[2]   Iouli E Gordon et al. "The HITRAN2020 molecular spectroscopic database". In: *Journal of Quantitative Spectroscopy and Radiative Transfer* 277 (2022), p. 107949.

[3]   George B. Rybicki and Alan P. Lightman. *Radiative processes in Astrophysics*. Weinheim: Wiley-VCH, 2004.

[4]   Marcel Berger. *Geometry I. 4th Edition*. Berlin Heidelberg: Springer-Verlag, 2009.

[5]   Katsumi Nomizu and Takeshi Sasaki. *Affine differential geometry*. Cambridge: Cambridge University Press, 1994.

[6]   Colin Fox and Richard A Norton. "Fast sampling in a linear-Gaussian inverse problem". In: *SIAM/ASA Journal on Uncertainty Quantification* 4.1 (2016), pp. 1191–1218.

[7]   Charles W. Champ and Andrew V. Sills. "The Generalized Law of Total Covariance". In: *preprint* (2022). URL: https://arxiv.org/abs/2205.14525.

[8]   Jari P. Kaipio and Erkki Somersalo. *Statistical and Computational Inverse Problems*. New York: Springer-Verlag New York, 2005.

[9]   Sze M Tan, Colin Fox, and Geoff K. Nicholls. *Course notes for ELEC 445 – Inverse Problems and Imaging*. https://coursesupport.physics.otago.ac.nz/wiki/pmwiki.php/ELEC445/HomePage. [Online; accessed 10/12/23]. 2016.

[10] Per Christian Hansen and Dianne Prost O'Leary. "The use of the L-curve in the regularization of discrete ill-posed problems". In: *SIAM Journal on Scientific Computing* 14.6 (1993), pp. 1487–1503.

[11] Gareth O. Roberts and Jeffrey S Rosenthal. "General state space Markov chains and MCMC algorithms". In: *Probability Surveys* 1 (2004), pp. 20–71.

[12] Charles J Geyer. "Practical markov chain monte carlo". In: *Statistical science* (1992), pp. 473–483.

[13] Gareth O. Roberts and Jeffrey S Rosenthal. "Harris recurrence of Metropolis-within-Gibbs and trans-dimensional Markov chains". In: *The Annals of Applied Probability* 16.4 (2006), 2123–2139.

[14] Johnathan M Bardsley. "MCMC-based image reconstruction with uncertainty quantification". In: *SIAM Journal on Scientific Computing* 34.3 (2012), A1316–A1332.

[15] Johnathan M Bardsley et al. "Randomize-then-optimize: A method for sampling from posterior distributions in nonlinear inverse problems". In: *SIAM Journal on Scientific Computing* 36.4 (2014), A1895–A1910.

[16]   Felipe Acosta, Mark L Huber, and Galin L Jones. "Markov chain Monte Carlo with linchpin variables". In: *preprint* (2014). URL: https://arxiv.org/abs/2205.14525.

[17]   J. Andrés Christen and Colin Fox. "A general purpose sampling algorithm for continuous distributions (the t-walk)". In: *Bayesian Analysis* 5.2 (2010), pp. 263 –281. URL: https://doi.org/10.1214/10-BA603.

[18]   Colin Fox et al. "Grid methods for Bayes-optimal continuous-discrete filtering and utilizing a functional tensor train representation". In: *Inverse Problems in Science and Engineering* 29.8 (2021), pp. 1199–1217.

[19]   Tiangang Cui and Sergey Dolgov. "Deep composition of tensor-trains using squared inverse rosenblatt transports". In: *Foundations of Computational Mathematics* 22.6 (2022), pp. 1863–1922.

[20]   Sergey Dolgov et al. "Approximation and sampling of multivariate probability distributions in the tensor train decomposition". In: *Statistics and Computing* 30 (2020), pp. 603–625.

[21]   Ivan V Oseledets. "Tensor-train decomposition". In: *SIAM Journal on Scientific Computing* 33.5 (2011), pp. 2295–2317.