

## Lecture 4 — January 16

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## 4.1 Tensors

A tensor is a multi-dimensional array, which are used in a variety of applications, such as weights and activations in deep neural networks. The order of a tensor (also known as the modes of a tensor) is the number of dimensions  $N$  of that tensor. An element  $(i, j, k)$  of a third-order tensor  $X$  is denoted by  $X_{i,j,k}$ . Fibers are defined by fixing every index but one; they are a higher-dimensional analogue of matrix rows and columns. Slices are defined by fixing all but two indices, i.e. two-dimensional sections of a tensor. Examples of fibers and slices are seen in figure 4.1. The (Frobenius) norm of a tensor is defined as

$$\|X\|_F = \sqrt{\sum_{i_1=1}^{I_1} \sum_{i_2=1}^{I_2} \dots \sum_{i_N=1}^{I_N} |X_{i_1 i_2 \dots i_N}|^2},$$

where the  $j$ -th dimension fiber is in  $\mathbb{R}^{I_j}$ .

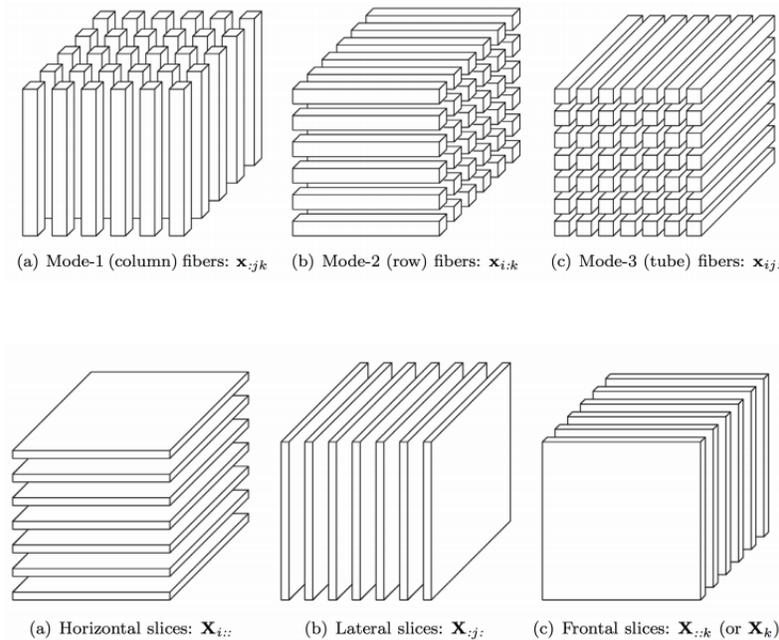


Figure 4.1: Sample fibers and slices of an order 3 tensor

## 4.2 Tensor Multiplication

### 4.2.1 Definition

The  $n$ -mode (matrix) product of a tensor  $A \in \mathbb{R}^{d_1 \times d_2 \times \dots \times d_N}$  with a matrix  $B \in \mathbb{R}^{p \times d_n}$  is done element-wise as below.

$$(A \times_n B)_{i_1, \dots, i_{n-1}, j, i_{n+1}, \dots, i_N} = \sum_{i_n=1}^{d_n} A_{i_1 i_2 \dots i_n \dots i_N} B_{j i_n}$$

In other words, each mode- $n$  fiber of  $A$  is multiplied by the matrix  $B$ .

### 4.2.2 Approximate Tensor Multiplication

The algorithm for approximate tensor multiplication is shown in Figure 4.2. The central idea is to reduce the dimensions of the tensor  $A$  and matrix  $B$  with sampling to get  $C$  and  $R$ , and perform an  $n$ -mode matrix product with  $C$  and  $R$  using the classical algorithm. The complexity of this algorithm is  $O(d_1 \dots d_{n-1} m d_n \dots d_N p)$ .

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#### Algorithm 1 Approximate Tensor $n$ -Mode Product via Sampling

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**Input:** An  $d_1 \times \dots \times d_n \times \dots \times d_N$  dimensional tensor  $A$  and an  $p \times d_n$  dimensional tensor  $B$ , an integer  $m$  and probabilities  $\{p_k\}_{k=1}^{d_n}$

**Output:** Tensors  $C, R$  such that  $CR \approx AB$

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- 1: **for**  $t = 1$  to  $m$  **do**
  - 2: Pick  $i_t \in \{1, \dots, d_n\}$  with probability  $\mathbb{P}[i_t = k] = p_k$  in i.i.d. with replacement
  - 3: Set  $C^{(t)} = \frac{1}{\sqrt{m p_{i_t}}} A_{:, i_t, :}$  and  $R_{(t)} = \frac{1}{\sqrt{m p_{i_t}}} B_{:, i_t, :}$
  - 4: **end for**
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Figure 4.2: Algorithm for approximate tensor multiplication

We now look at the mean and variance of the multiplication estimator. Define

$$M_{i \rightarrow j} \triangleq (A \times_n B)_{i_1, \dots, i_{n-1}, j, i_{n+1}, \dots, i_N} = \sum_{i_n=1}^{d_n} A_{i_1 i_2 \dots i_n \dots i_N} B_{j i_n}$$

and

$$\hat{M}_{i \rightarrow j} \triangleq \sum_{i_n=1}^m \frac{1}{p_{i_n}} A_{i_1 i_2 \dots i_n \dots i_N} B_{j i_n}.$$

This estimator is unbiased, i.e.  $\mathbb{E}[\hat{M}_{i \rightarrow j}] = M_{i \rightarrow j}$ . The variance is

$$\text{Var}[\hat{M}_{i \rightarrow j}] = \frac{1}{m} \sum_{i_n=1}^{d_n} \frac{1}{p_{i_n}} A_{i_1 i_2 \dots i_n \dots i_N}^2 B_{j i_n}^2 - \frac{1}{m} (M_{i \rightarrow j})^2.$$

To achieve the optimal multiplication estimator, we want to solve the following minimization problem.

$$\text{minimize}_p \mathbb{E} \|\hat{M} - M\|_F^2 = \text{minimize}_p \sum_{\vec{i}, \vec{j}} \mathbf{Var}[\hat{M}_{\vec{i}, \vec{j}}].$$

After some math, we find that the optimal  $p$  is defined by

$$p_k = \frac{\|A_{:\dots k \dots}\|_F \|B_{:k}\|_F}{\sum_k \|A_{:\dots k \dots}\|_F \|B_{:k}\|_F}.$$

### 4.3 Verifying Matrix Multiplication

We now consider a different problem. Suppose we are given three  $n \times n$  matrices  $A, B, M$ . We want to verify whether  $AB = M$ . The naive method is to multiply  $A$  and  $B$  with the classical method and compare each point in the product and  $M$  individually, which is  $O(n^3)$ . It turns out that a randomized algorithm can do this in  $O(n^2)$  and no faster.

The algorithm for this method is known as Freivald's Algorithm (1977). We first sample a random vector  $r = [r_1, \dots, r_n]^T$ . We compute  $Br$ , then  $A(Br)$ . We compute  $Mr$ . Finally, we compare our two products. If  $A(Br) \neq Mr$ , then  $AB \neq M$  with 100% probability. Otherwise, we return  $AB = M$ . Since there are three matrix-vector multiplications, we have a complexity of  $O(n^2)$ .

We would like to analyze the failure probability of this algorithm. Without knowing anything about the matrices  $A, B, M$ , we can't guarantee a high or low probability for this algorithm. However, if we pick each  $r_i$  in  $\mathbf{r} = [r_1, \dots, r_n]^T$  in an i.i.d. fashion to be  $+1$  or  $-1$  with probability  $\frac{1}{2}$ , we can claim  $\mathbb{P}[A\mathbf{B}\mathbf{r} = M\mathbf{r}] \leq \frac{1}{2}$ . Note that we can also choose  $r_i$  to be 0 or 1. To improve the error probability, we run the algorithm independently  $k$  times. If we ever find an  $\mathbf{r}^k$  such that  $A\mathbf{B}\mathbf{r}^k = M\mathbf{r}^k$ , then the algorithm correctly returns  $AB \neq M$ . If we always find  $A\mathbf{B}\mathbf{r} = M\mathbf{r}$ , then the error probability is at most  $\frac{1}{2^k}$ . For  $k = 25$ , we have an error probability  $\leq 10^{-9}$ .

### 4.4 Concentration Bounds

In order to achieve tighter success probabilities, we look at concentration bounds. Specifically for approximate matrix multiplication (AMM), the size of the sample is  $m = \frac{1}{\delta \epsilon^2}$ . We would like to have  $m$  not depend on the failure probability  $\delta$ .

#### 4.4.1 Specific Bounds

We provide a quick refresher on common bounds. Markov's Inequality states that for  $Z > 0$  and  $t > 0$ ,

$$\mathbb{P}[Z > a] \leq \frac{\mathbb{E}Z}{a}.$$

Chebyshev's Inequality is as follows. Let  $X$  be a random variable with expectation  $\mathbb{E}[X]$  and variance  $\mathbf{Var}[X]$ . Then,

$$\mathbb{P}[|X - \mathbb{E}[X]| \geq t] \leq \frac{\mathbf{Var}[X]}{t^2}.$$

Lastly, Chernoff's Bound has several versions with better constants, but we present this one. Let  $X_1, \dots, X_m$  be independent random variables  $\in [0, 1]$  and let  $\mu = \mathbb{E}X_1$ . Then

$$\mathbb{P}\left[\left|\frac{1}{m} \sum_{i=1}^m X_i - \mu\right| > t\mu\right] \leq 2e^{-m \frac{t^2}{3}}.$$

We will use this result in the following discussions.

### 4.4.2 Application 1: Monte Carlo Approximations

We look at applications in Monte Carlo Approximations. Suppose we want to estimate  $\pi$ . We uniformly sample  $z_1, \dots, z_m$  i.i.d. from  $[0, 1]^2$ . We define the random variable  $Z_i$  below.

$$Z_i = \begin{cases} 1 & \|z_i\|_2 \leq 1 \\ 0 & \text{otherwise} \end{cases}.$$

Thus,  $\mathbb{P}[Z_i = 1] = \frac{\pi}{4}$ . Applying the Chernoff Bound, we get

$$\left|\frac{1}{m} \sum_{i=1}^m Z_i - \frac{\pi}{4}\right| \leq \epsilon \frac{\pi}{4}$$

with probability at least  $1 - 2e^{-m\epsilon^2 \frac{\pi}{12}}$ . We can pick  $m \geq \frac{12}{\pi\epsilon^2} \log \frac{2}{\delta}$  and obtain an estimate of  $\hat{\pi}$  such that  $(1 - \epsilon)\pi \leq \hat{\pi} \leq (1 + \epsilon)\pi$  with probability at least  $1 - \delta$ . The range  $[(1 - \epsilon)\pi, (1 + \epsilon)\pi]$  is a confidence interval.

### 4.4.3 Application 2: Amplifying Probability of Success

Now we try to amplify the probability of success of a randomized algorithm. Suppose we have a randomized algorithm which produces an  $\epsilon$  approximation  $|\hat{x} - x^*| \leq \epsilon$  with probability at least 0.9. We repeat the algorithm  $m$  times independently, and take the median of the  $m$  outputs. Note that we take the median instead of the mean, because a failure case could result in very large/small values that shift the mean. Let the random variable  $X_i = 1$  if the  $i$ -th trial is good, i.e.  $|\hat{x}_i - x^*| \leq \epsilon$ . If at least half of the  $X_i$ 's are one, the median of the  $m$  outputs is also good, i.e.  $|\text{Median}(\hat{x}_i) - x^*| \leq \epsilon$ . The Chernoff Bound implies that  $|\frac{1}{m} \sum_{i=1}^m X_i - 0.9| \leq 0.9t$  with probability  $1 - e^{-t^2 0.9m/3}$ . Pick  $t = 0.4/0.9$ . Then, the median is an  $\epsilon$  approximation with probability at least  $1 - e^{-0.059m}$ , e.g., for  $m = 200$ , failure probability is  $\leq 7 \times 10^{-6}$ .

#### 4.4.4 Median for Approximate Matrix Multiplication

Since the Chernoff Bound implies that the majority of estimators are good, we would like to generalize the concept of a median to matrices. The median relies on the fact that  $\mathbb{R}^1$  is ordered; however, matrices aren't ordered. We could represent the median as the optimization problem,  $\operatorname{argmin}_y \sum |x_i - y|$ , but solving this for matrices is computationally expensive. The central idea is to have some concept of "centrality". We look at distances between estimates: the correct estimates will have many smaller distances, while the incorrect ones will have many larger distances.

We start with the AMM final probability bound. For any  $\delta > 0$ , set  $m = \frac{1}{\delta \epsilon^2}$  to obtain

$$\mathbb{P}[\|AB - CR\|_F > \epsilon \|A\|_F \|B\|_F] \leq \delta.$$

Suppose  $\|A\|_F = \|B\|_F = 1$  and let  $\epsilon = 0.1, \delta = 0.9$ . Repeat the algorithm independently and obtain  $C_1R_1, \dots, C_tR_t$  in  $t$  independent trials. Then,  $\|AB - C_iR_i\|_F < 0.1$  with probability 0.9 for each  $i$ . However, we don't know which ones are good, i.e.  $\|AB - C_iR_i\|_F < 0.1$ . Let  $X_i = 1$  if the  $i$ -th trial is good and  $X_i = 0$  otherwise. The Chernoff Bound implies that  $\frac{1}{m} \sum_{i=1}^m X_i \geq 0.5$  with probability  $1 - e^{-0.059m}$ , i.e. at least half of the matrices are good. Compute  $\rho_i \triangleq |\{j | j \neq i, \|C_iR_i - C_jR_j\|_F \leq 0.2\}|$ . Finally, output  $C_kR_k$  such that  $\rho_k \leq \frac{t}{2}$ . This results in a lemma that  $\|AB - C_kR_k\|_F \leq 0.3$  with probability at least  $1 - e^{-0.059m}$ .

We now prove this lemma. We use the triangle inequality which states that

$$\|X + Y\|_F \leq \|X\|_F + \|Y\|_F$$

and the reverse triangle inequality which states that

$$\|X + Y\|_F \geq \|X\|_F - \|Y\|_F.$$

Letting  $X = C_iR_i - AB, Y = AB - C_jR_j$ , we get

$$\|C_iR_i - C_jR_j\|_F \leq \|C_iR_i - AB\|_F + \|C_jR_j - AB\|_F$$

and

$$\|C_iR_i - C_jR_j\|_F \geq \|C_iR_i - AB\|_F - \|C_jR_j - AB\|_F.$$

If  $C_iR_i$  is good, i.e.  $\|AB - C_iR_i\|_F \leq 0.1$ , then it is close to at least half of the other  $C_jR_j$ 's. Thus,  $\rho_i \triangleq |\{j | j \neq i, \|C_iR_i - C_jR_j\|_F \leq 0.2\}| \geq \frac{t}{2}$  by the triangle inequality. If  $C_iR_i$  is bad, i.e.  $\|AB - C_iR_i\|_F > 0.3$ , then  $\|C_iR_i - C_jR_j\|_F \geq 0.2$  by the triangle inequality and  $\rho_i \leq \frac{t}{2}$ .