

# Introduction to Quantum Information Processing

QIC 710 / CS 768 / PH 767 / CO 681 / AM 871

## Lecture 16 (2019)

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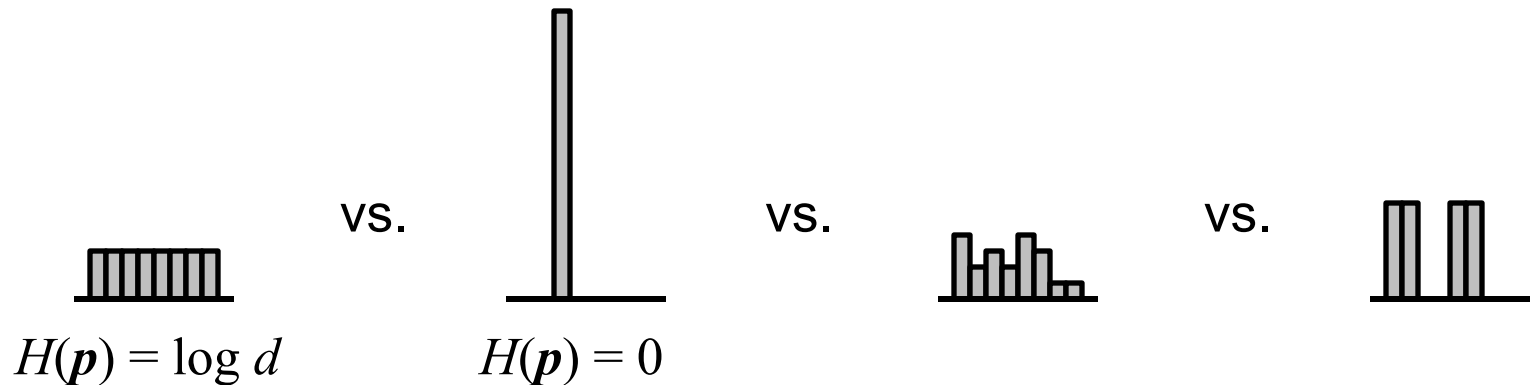
# Entropy and compression

# Shannon entropy

Let  $\mathbf{p} = (p_1, \dots, p_d)$  be a probability distribution on a set  $\{1, \dots, d\}$

Then the (Shannon) **entropy** of  $\mathbf{p}$  is  $H(p_1, \dots, p_d) = -\sum_{j=1}^d p_j \log p_j$

Intuitively, this turns out to be a good measure of how much “randomness” (or “uncertainty”) is there is in  $\mathbf{p}$ :



We'll see that, operationally,  $H(\mathbf{p})$  is the number of bits needed to store the outcome (in a sense that will be made formal)

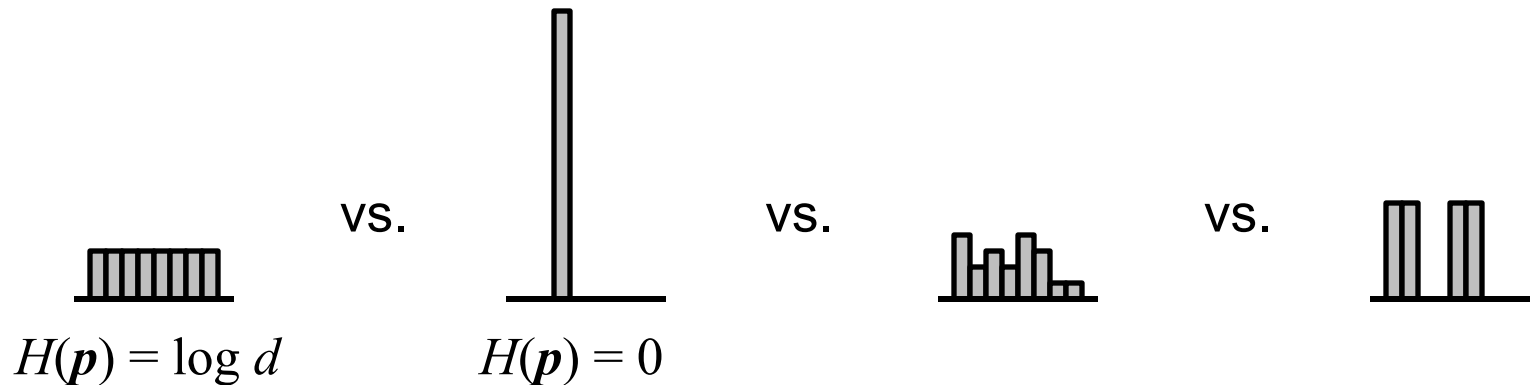
# Shannon entropy

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Then the (Shannon) **entropy** of  $\mathbf{p}$  is  $H(p_1, \dots, p_d) = \sum_{j=1}^d p_j \log p_j$

Intuitively, this turns out to be a good measure of how much “randomness” (or “uncertainty”) is there is in  $\mathbf{p}$ :



We'll see that, operationally,  $H(\mathbf{p})$  is the number of bits needed to store the outcome (in a certain sense)

# Von Neumann entropy

For a density matrix  $\rho$ , it turns out that  $S(\rho) = -\text{Tr} \rho \log \rho$  is a good quantum analogue of entropy

**Note:**  $S(\rho) = H(p_1, \dots, p_d)$ , where  $p_1, \dots, p_d$  are the eigenvalues of  $\rho$  (with multiplicity)

Operationally,  $S(\rho)$  is the number of **qubits** needed to store  $\rho$  (in a sense that will be made formal later on)

Both the classical and quantum compression results pertain to the case of large blocks of  $n$  independent instances of data:

- probability distribution  $\mathbf{p}^{\otimes n}$  in the classical case, and
- quantum state  $\rho^{\otimes n}$  in the quantum case

# Classical compression (1)

Let  $\mathbf{p} = (p_1, \dots, p_d)$  be a probability distribution on a set  $\{1, \dots, d\}$  where  $n$  independent instances are sampled:

$(j_1, \dots, j_n) \in \{1, \dots, d\}^n$  ( $d^n$  possibilities,  $n \log d$  bits to specify one)

**Theorem\* (Shannon data compression):** for all  $\varepsilon > 0$ , for sufficiently large  $n$ , there is a scheme that compresses the specification to  $n(H(\mathbf{p}) + \varepsilon)$  bits while introducing an error with probability at most  $\varepsilon$

**Example:** an  $n$ -bit binary string with each bit distributed as  $\Pr[0] = 0.9$  and  $\Pr[1] = 0.1$  can be compressed to  $\approx 0.47n$  bits

Intuitively, there is a subset  $T \subseteq \{1, \dots, d\}^n$ , called the “typical sequences”, that has size  $2^{n(H(\mathbf{p}) + \varepsilon)}$  and probability  $1 - \varepsilon$  of occurring

Note that, in the above example,  $|T| \ll 2^n$  even though  $\Pr[T] \geq 1 - \varepsilon$

\* “Plain vanilla” version that ignores, for example, the tradeoffs between  $n$  and  $\varepsilon$

# Classical compression (2)

A nice way to prove the theorem, is based on two cleverly defined random variables ...

Define the random variable  $f: \{1, \dots, d\} \rightarrow \mathbb{R}$  as  $f(j) = -\log p_j$

Note that  $E[f] = \sum_{j=1}^d p_j f(j) = -\sum_{j=1}^d p_j \log p_j = H(p_1, \dots, p_d)$

Define  $g: \{1, \dots, d\}^n \rightarrow \mathbb{R}$  as  $g(j_1, \dots, j_n) = \frac{f(j_1) + \dots + f(j_n)}{n}$

Thus  $E[g] = H(p_1, \dots, p_d)$

Also,  $g(j_1, \dots, j_n) = -\frac{1}{n} \log(p_{j_1} \dots p_{j_n}) = -\frac{1}{n} \log(\Pr[(j_1, \dots, j_n)])$

which implies  $\Pr[(j_1, \dots, j_n)] = 2^{-ng(j_1, \dots, j_n)}$

# Classical compression (3)

By standard results in statistics\*, as  $n \rightarrow \infty$ , the observed value of  $g(j_1, \dots, j_n)$  approaches its expected value,  $H(p)$ , in this sense:

$\Pr[|g(j_1, \dots, j_n) - H(p)| \leq \varepsilon] \geq 1 - \varepsilon$  for all  $\varepsilon > 0$ , for sufficiently large  $n$

[recall that  $g(j_1, \dots, j_n)$  is an average of independent  $f(j)$  ]

Define  $(j_1, \dots, j_n) \in \{1, \dots, d\}^n$  to be  **$\varepsilon$ -typical** if

$$|g(j_1, \dots, j_n) - H(p)| \leq \varepsilon$$

Then, the above implies, for all  $\varepsilon > 0$ , for sufficiently large  $n$ ,

$$\Pr[(j_1, \dots, j_n) \text{ is } \varepsilon\text{-typical}] \geq 1 - \varepsilon$$

We can also bound the **number of** these  $\varepsilon$ -typical sequences:

- By definition, each such sequence has probability  $\geq 2^{-n(H(p) + \varepsilon)}$
- Therefore, there can be at most  $2^{n(H(p) + \varepsilon)}$  such sequences (otherwise, the sum of probabilities would exceed 1)

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\* The weak law of large numbers



# Classical compression (4)

In summary, the compression procedure is as follows:

The input data is  $(j_1, \dots, j_n) \in \{1, \dots, d\}^n$ , each independently sampled according to the probability distribution  $\mathbf{p} = (p_1, \dots, p_d)$

The compression procedure is to leave  $(j_1, \dots, j_n)$  intact if it is  $\varepsilon$ -typical and otherwise change it to some fixed  $\varepsilon$ -typical sequence, say, some  $(j_k, \dots, j_k)$  (which will result in an error)

Since there are at most  $2^{n(H(\mathbf{p}) + \varepsilon)}$   $\varepsilon$ -typical sequences, the data can then be converted into  $n(H(\mathbf{p}) + \varepsilon)$  bits

The error probability is at most  $\varepsilon$ , the probability of an input that is not typical arising

# Quantum compression (1)

**The scenario:**  $n$  independent instances of a  $d$ -dimensional state are randomly generated according some distribution:

$$\left\{ \begin{array}{ll} |\varphi_1\rangle & \text{prob. } p_1 \\ \vdots & \vdots \\ |\varphi_r\rangle & \text{prob. } p_r \end{array} \right.$$

Example:	$\left\{ \begin{array}{ll}  0\rangle & \text{prob. } \frac{1}{2} \\  +\rangle & \text{prob. } \frac{1}{2} \end{array} \right.$
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**Goal:** to “compress” this into as few qubits as possible so that the original state can be reconstructed “with small error”

What’s a good formal definition of error in a quantum compression scheme?

Define a quantum compression scheme to be  **$\varepsilon$ -good** if no procedure can distinguish between these two states

- a) the state resulting from compressing and then uncompressing the data
- b) the original state

with probability more than  $\frac{1}{2} + \frac{1}{4} \varepsilon$

# Quantum compression (2)

Define  $\rho = \sum_{i=1}^r p_i |\phi_i\rangle\langle\phi_i|$

**Theorem (Schumacher data compression):** for all  $\varepsilon > 0$ , for sufficiently large  $n$ , there is a scheme that compresses the data to  $n(S(\rho) + \varepsilon)$  qubits, that is  $\sqrt{2\varepsilon}$ -good

For the aforementioned example,  $\approx 0.6n$  qubits suffices

$ 0\rangle$	prob. $\frac{1}{2}$
$ +\rangle$	prob. $\frac{1}{2}$

**The compression method:**

Express  $\rho$  in its eigenbasis as  $\rho = \sum_{j=1}^d q_j |\psi_j\rangle\langle\psi_j|$

With respect to this basis, we will define an  $\varepsilon$ -typical subspace of dimension  $2^{n(S(\rho) + \varepsilon)} = 2^{n(H(q) + \varepsilon)}$

# Quantum compression (3)

The  **$\varepsilon$ -typical subspace** is that spanned by  $|\psi_{j_1}, \dots, \psi_{j_n}\rangle$   
 where  $(j_1, \dots, j_n)$  is  $\varepsilon$ -typical with respect to  $(q_1, \dots, q_d)$

**Define:**  $\Pi_{\text{typ}}$  as the projector into the  $\varepsilon$ -typical subspace

By the same argument as in the classical case, the subspace has  
 dimension  $\leq 2^{n(S(\rho) + \varepsilon)}$  and  $\text{Tr}(\Pi_{\text{typ}} \rho^{\otimes n}) \geq 1 - \varepsilon$

Why? Because  $\rho$  is the density matrix of  $\begin{cases} |\psi_1\rangle & \text{prob. } q_1 \\ \vdots & \vdots \\ |\psi_d\rangle & \text{prob. } q_d \end{cases}$  ← “eigenstate” mixture

$$\begin{aligned} \text{and } \text{Tr}(\Pi_{\text{typ}} \rho^{\otimes n}) &= \text{Tr}\left(\Pi_{\text{typ}} \sum_{j_1 \dots j_n} q_{j_1} \dots q_{j_n} |\psi_{j_1} \dots \psi_{j_n}\rangle \langle \psi_{j_1} \dots \psi_{j_n}|\right) \\ &= \sum_{j_1 \dots j_n} q_{j_1} \dots q_{j_n} \langle \psi_{j_1} \dots \psi_{j_n} | \Pi_{\text{typ}} | \psi_{j_1} \dots \psi_{j_n} \rangle \\ &= \sum_{j_1 \dots j_n} q_{j_1} \dots q_{j_n} \chi_{[j_1 \dots j_n \text{ is typical}]} \geq 1 - \varepsilon \end{aligned}$$

# Quantum compression (4)

We would now be done if our *actual mixture* was an *eigenstate mixture*

**actual mixture:**

$$\left\{ \begin{array}{ll} |\phi_1\rangle & \text{prob. } p_1 \\ \vdots & \vdots \\ |\phi_r\rangle & \text{prob. } p_r \end{array} \right.$$

**eigenstate mixture:**

$$\left\{ \begin{array}{ll} |\psi_1\rangle & \text{prob. } q_1 \\ \vdots & \vdots \\ |\psi_r\rangle & \text{prob. } q_r \end{array} \right.$$

Calculation of the “expected fidelity” for our actual mixture:

$$\begin{aligned} \sum_I p_I \langle \phi_I | \Pi_{\text{typ}} | \phi_I \rangle &= \sum_I p_I \text{Tr}(\Pi_{\text{typ}} | \phi_I \rangle \langle \phi_I |) \\ &= \text{Tr} \left( \sum_I p_I \Pi_{\text{typ}} | \phi_I \rangle \langle \phi_I | \right) \\ &= \text{Tr}(\Pi_{\text{typ}} \rho^{\otimes n}) \\ &\geq 1 - \varepsilon \end{aligned}$$

Abbreviations used

$$I = i_1 i_2 \dots i_n$$

$$p_I = p_{i_1} p_{i_2} \dots p_{i_n}$$

$$|\phi_I\rangle = |\phi_{i_1} \phi_{i_2} \dots \phi_{i_n}\rangle$$

**Does this mean that the scheme is  $\varepsilon'$ -good for some  $\varepsilon'$ ?**

# Quantum compression (5)

The **true data** is of the form  $(I, |\phi_I\rangle)$  where the  $I$  is generated with probability  $p_I$

The **approximate data** is of the form  $(I, \frac{1}{\gamma_I} \Pi_{\text{typ}} |\phi_I\rangle)$  where  $I$  is generated with probability  $p_I$

$\gamma_I = \sqrt{\langle \phi_I | \Pi_{\text{typ}} | \phi_I \rangle}$  is a normalization factor

Above two states **at least** as hard to distinguish as these two purifications:

$$|\Phi\rangle = \sum_I \sqrt{p_I} |I\rangle \otimes |\phi_I\rangle \quad |\Phi'\rangle = \sum_I \sqrt{p_I} |I\rangle \otimes \frac{1}{\gamma_I} \Pi_{\text{typ}} |\phi_I\rangle$$

$$\text{Fidelity: } \langle \Phi | \Phi' \rangle = \sum_I p_I \frac{1}{\gamma_I} \langle \phi_I | \Pi_{\text{typ}} | \phi_I \rangle \geq \sum_I p_I \langle \phi_I | \Pi_{\text{typ}} | \phi_I \rangle \geq 1 - \varepsilon$$

$$\text{Trace distance: } \| |\Phi\rangle - |\Phi'\rangle \|_{\text{tr}} \leq \sqrt{2\varepsilon}$$

Therefore the scheme is  $\approx \sqrt{2\varepsilon}$ -good