

# **Introduction to Quantum Information Processing**

**QIC 710 / CS 678 / PH 767 / CO 681 / AM 871**

## **Lectures 17–18 (2013)**

**Richard Cleve**

DC 2117 / QNC 3129

[cleve@cs.uwaterloo.ca](mailto:cleve@cs.uwaterloo.ca)

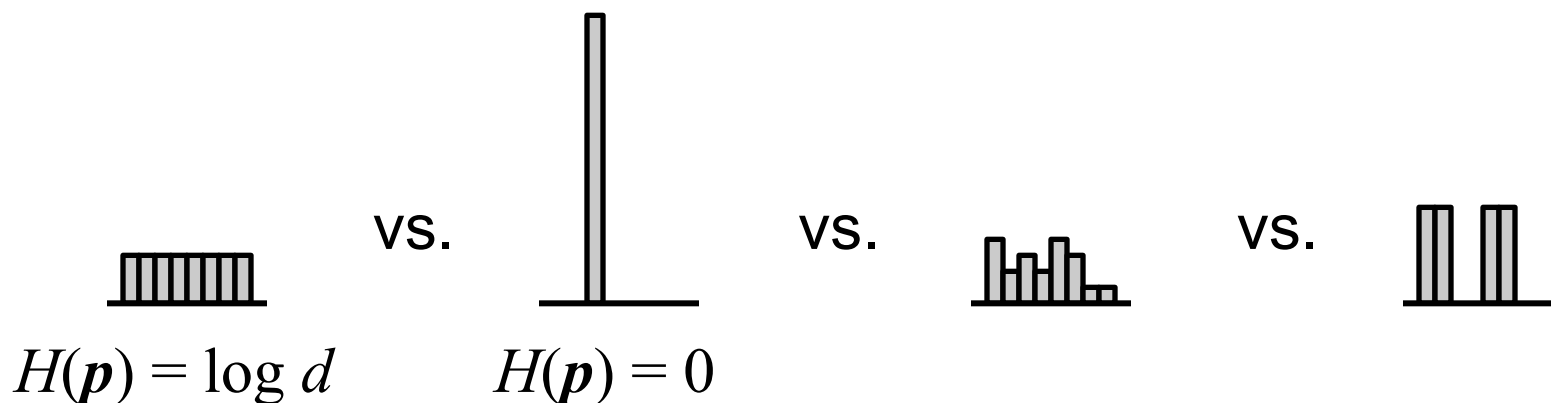
# Entropy and compression

# 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 random” the distribution  $\mathbf{p}$  is:



Operationally,  $H(\mathbf{p})$  is the number of bits needed to store the outcome (in a sense that will be made formal shortly)

# 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\*:** 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  $\varepsilon$

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

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

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

# Classical compression (2)

Define the random variable  $f: \{1, \dots, d\} \rightarrow \mathbf{R}$  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  $g: \{1, \dots, d\}^n \rightarrow \mathbf{R}$  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,  $g(j_1, \dots, j_n) = -\frac{1}{n} \log(p_{j_1} \cdots p_{j_n})$

# 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,  $H(p)$

More formally, call  $(j_1, \dots, j_n) \in \{1, \dots, d\}^n$   **$\varepsilon$ -typical** if

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

Then, the result is that, 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

# 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 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,  $(j, \dots, j)$  (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 atypical input 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:  $\left\{ \begin{array}{ll} |0\rangle & \text{prob. } \frac{1}{2} \\ |+\rangle & \text{prob. } \frac{1}{2} \end{array} \right.$

**Goal:** to “compress” this into as few qubits as possible so that the original state can be reconstructed with small error in the following sense ...

**$\epsilon$ -good:**

No procedure can distinguish between these two states

- (a) compressing and then uncompressing the data
- (b) the original data left as is

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

# Quantum compression (2)

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

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

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

**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  $\text{Tr}(\Pi_{\text{typ}} \rho^{\otimes n}) \geq 1 - \varepsilon$

This is because  $\rho$  is the density matrix of  $\left\{ \begin{array}{ll} |\psi_1\rangle & \text{prob. } q_1 \\ \vdots & \vdots \\ |\psi_d\rangle & \text{prob. } q_d \end{array} \right.$

# Quantum compression (4)

Calculation of the expected fidelity:

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

Abbreviations

$$I = i_1 i_2 \dots i_n$$

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

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

What does this mean?

If the generated state is  $| \phi_I \rangle$  and the compression process first applies the measurement  $\Pi_{\text{typ}}, \Pi_{\text{typ}}^\perp$  then the success probability is  $\langle \phi_I | \Pi_{\text{typ}} | \phi_I \rangle$  (call outcome  $\Pi_{\text{typ}}$  “success”)

Averaging over the possible choices of the index  $I$ , the success probability for the compression part is  $\geq 1 - \varepsilon$

# Quantum compression (5)

How good an approximation of the true data is the compressed state when the compression part succeeds?

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  $\left(I, \frac{1}{\gamma_I} \Pi_{\text{typ}} |\phi_I\rangle\right)$  where  $I$  is generated with probability  $p_I$

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

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

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

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

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