Introduction to Classical Error Correction

Wanqiang Liu, June 20, 2023. wanqiangl@uchicago.edu





Errors in Classical information



Fig. 1—Schematic diagram of a general communication system.

Warm up: Parity Check Code

- Rule: The coded message always has even number of 1 in a piece string 010...
- If I spot odd number of 1, there must be some error.
- Example: Use 9 bits to encode a piece of information
 - 000 000 000
 - 001 000 000
- Q1: How many different code words ("CORRECT MESSAGE") are allowed? (2^8)
- Q2: How many "flips" at least can make an undetectable error? (2)



[[n, k, d]] code

- n= # of bits used in "a piece of information"
- k = # of logical bits = log_2 (# of code words)
- d = code distance = minimum # of flips to make an undetectable error
- The parity check example is [[9, 8, 2]]

• Q3: If I detect an error in the parity check code, can I know how to correct it?



Make the code correctable! Rectangular parity check

Rule: If you receive a 9-bit message, stack it into a 3*3 square.

Each column and each row must have even number of 1.



Q1: [[n, k, d]]=? (Hint: how to go from one codeword to another?) [[9,4,4]] Q2: How many bit flips can make an UNCORRECTABLE error? 2

Another example: Majority Vote

- 111 111 111
- 000 000 000
- majority. Example: 111 000 111 -> 111 111 111

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Q: [[n, k, d]]=?
[[9,1,9]]
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Anything else is not a codeword, and should be corrected according to the

What makes a good code?

- Higher k/n: efficient use of bits
- Higher d: can detect more errors
- Efficient decoding
- Examples
 - Hamming code
 - Linear codes...

Tradeoff: if the message is "tightly packed" with codewords (high k/n), it's less likely to have room for errors (lower d)

What is the capacity limit? — Classical information theory Shannon theorem

- Length of the coded message to send: M_{C}
- Length of the coded stat message: M
- For any bit in M_C , the probability to flip is q

$$\frac{M}{M_C} \le 1 - q[\log_2(\frac{1}{q}) + (1 - q)\log_2(\frac{1}{1})]$$



Feynman Lectures on Computation, Chapter 4



Error correction in nature — DNA!



Redundancy — room to tolerate errors!

Q: Using Shannon's theorem, estimate what error rate q is tolerated in the DNA-to-protein process

Hint (you can disagree with this) • Shannon: $\frac{M}{M_C} \le 1 - q[\log_2(\frac{1}{q}) + (\frac{1}{q})]$ • LHS: $\frac{M}{M_C} \sim \frac{\log_4(21)}{3}$

• RHS: $log_2 \rightarrow log 4$, because there are 4 base pairs.

• Look up the true error rates in nature. How does the estimate compare?

$$(1-q)\log_2(\frac{1}{1-q})]$$