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Generator and Parity Check Matrices Sumple in the interval of the interva







Proof that H is a Parity Check Matrix
Suppose that x is a message. Then
$H(xG)^{T} = H(G^{T}x^{T}) = (HG^{T})x^{T} = (A^{T}\mathbf{I}_{k}+\mathbf{I}_{n-k}A^{T})x^{T} = (A^{T}+A^{T})x^{T} = 0$
Now suppose that $Hy^T = 0$. Then $A^T_{i,*} \bullet y^T_{[1.k]} + y^T_{k+i} = 0$ (where $A^T_{i,*}$ is row i of A^T and $y^T_{[1.k]}$ are the first k elements of y^T]) for $1 \le i \le n-k$. Thus, $y_{[1.k]} \bullet A_{*,i} = y_{k+i}$ where $A_{*,i}$ is now column i of A, and $y_{[1.k]}$ are the first k elements of y, so $y_{[k+1n]} = y_{[1k]}A$.
Consider x = $y_{[1k]}$. Then xG = $[y_{[1k]} y_{[1k]}A] = y$.
Hence if $Hy^T = 0$, y is the codeword for x = $y_{[1k]}$.

The d of linear codes

<u>Theorem</u>: Linear codes have distance d if every set of (d-1) columns of H are linearly independent, but there is a set of d columns that are linearly dependent (i.,e., sum to 0).

<u>**Proof</u>**: if d-1 or fewer columns are linearly dependent, then for any codeword y, there is another codeword y', in which the bits in the positions corresponding to the columns are inverted, that both have the same syndrome (0).</u>

If every set of d-1 columns is linearly independent, then changing any d-1 bits in a codeword y must also change the syndrome (since the d-1 corresponding columns¹cannot sum to 0). Page25

Dual Codes

For every code with $G = I_k, A$ and $H = A^T, I_{n-k}$ we have a <u>dual code</u> with $G = I_{n-k}, A^T$ and $H = A, I_k$

- The dual of the Hamming codes are the binary simplex codes: $(2^{r}-1, r, 2^{r-1}-r)$
- The dual of the extended Hamming codes are the **first-order Reed-Muller** codes.
- Note that these codes are **highly redundant** and can fix many errors.

15-853

NASA Mariner:

Deep space probes from 1969-1977. Mariner 10 shown



Used (32,6,16) Reed Muller code (r = 5) Rate = 6/32 = .1875 (only 1 out of 5 bits are useful) Can fix up to 7 bit errors per 32-bit word



In **general** we can find the error location by creating a table that maps each syndrome to a set of error locations.

 $\label{eq:correspondence} \begin{array}{l} \hline \textbf{Theorem:}\\ \textbf{assuming s} \leq 2d\text{-1 every syndrome value}\\ \textbf{corresponds to a unique set of error locations.}\\ \hline \textbf{Proof: Exercise.} \end{array}$

Table has q^{n-k} entries, each of size at most n (i.e. keep a bit vector of locations).

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