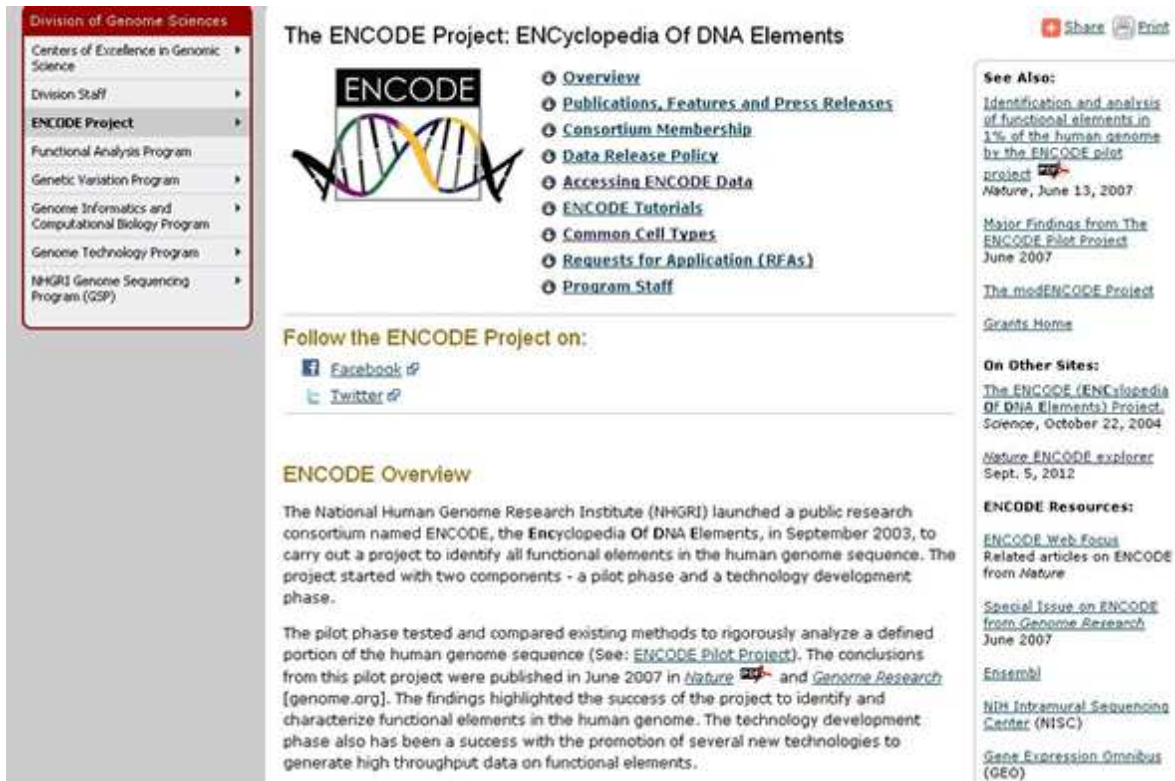


# Naturalment Webs

## Encode



The screenshot shows the ENCODE Project website. On the left is a navigation menu for the Division of Genome Sciences, with 'ENCODE Project' selected. The main content area is titled 'The ENCODE Project: ENcyclopedia Of DNA Elements' and features the ENCODE logo (a DNA double helix) and a list of links: Overview, Publications, Features and Press Releases, Consortium Membership, Data Release Policy, Accessing ENCODE Data, ENCODE Tutorials, Common Cell Types, Requests for Application (RFAs), and Program Staff. Below this is a 'Follow the ENCODE Project on:' section with Facebook and Twitter icons. The 'ENCODE Overview' section describes the project's launch in 2003 and its pilot phase. On the right, there are 'Share' and 'Print' buttons, a 'See Also:' section with links to Nature articles, 'On Other Sites:' with links to project pages and journals, and 'ENCODE Resources:' with links to web focus, special issues, Ensembl, NISC, and GEO.

<http://www.genome.gov/10005107>

Pàgina de l'Encode dins el National Human Genome Research Institute.





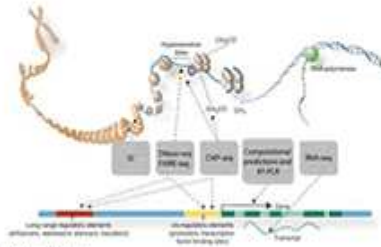
## Encyclopedia of DNA Elements

**General**

- Resources & FAQ
- Publications
- Software Tools
- Data Standards
- Human**
- Downloads
- Experiment Matrix
- Search
- Genome Browser (hg19)
- Integrative Analysis
- Session Gallery

**About ENCODE Data**

The Encyclopedia of DNA Elements (ENCODE) Consortium is an international collaboration of research groups funded by the National Human Genome Research Institute (NHGRI). The goal of ENCODE is to build a comprehensive parts list of functional elements in the human genome, including elements that act at the protein and RNA levels, and regulatory elements that control cells and circumstances in which a gene is active.



ENCODE data are now available for the entire human genome. **All ENCODE data are free and available for immediate use via:**

- [Search](#) for displayable tracks and downloadable files
- [Download](#) of data files
- [Visualization](#) in the UCSC Genome Browser (ENCODE data marked with the NHGRI logo)
- [Data mining](#) with the UCSC Table Browser and other [UCSC Genome Bioinformatics tools](#)

[Click to explore](#)

To search for ENCODE data related to your area of interest and set up a browser view, use the UCSC [Experiment Matrix](#) or [Track Search tool](#) (Advanced features). The [Experiment List \(Human\)](#) and [Experiment List \(Mouse\)](#) links provide comprehensive listings of ENCODE data that is released or in preparation.

All ENCODE data is freely available for download and analysis. However, before publishing research that uses ENCODE data, please read the [ENCODE Data Release Policy](#), which places some restrictions on publication use of data for nine months following data release. [Read more](#) about ENCODE data at UCSC.

<http://encodeproject.org/ENCODE/>

Pàgina oficial sobre el projecte ENCODE.



The screenshot shows the 'nature ENCODE explorer' website. At the top, there is a navigation bar with 'nature ENCODE' and a search box labeled 'Search ENCODE'. Below the navigation bar, there are two main sections: 'THREADS' and 'PAPERS'. The 'THREADS' section features a circular navigation menu with buttons labeled D1 through D11. A 'THREAD OVERVIEW' window is open, displaying the title 'RNA and chromatin modification patterns around promoters' and a brief description. The 'PAPERS' section shows a circular arrangement of paper icons, with one icon highlighted in red. A 'nature' paper detail window is open, showing the authors 'Sanyal, S. and Lajoie, B. et al.', the title 'The long-range interaction landscape of gene promoters', the journal 'Nature', and the DOI '10.1038/nature11279 (2012)'. There is also a link to 'View explorer & read full paper' and a section for 'Included in these threads' with buttons for D4, D8, D9, and D10. The website is produced with support from Illumina.

<http://www.nature.com/encode>

Pàgina de la revista Nature sobre ENCODE en la que hi podem trobar el conjunt d'articles publicats en el moment de la presentació dels primers resultats del projecte ENCODE, a les revistes *Nature*, *Genome Research* i *Genome Biology*.

