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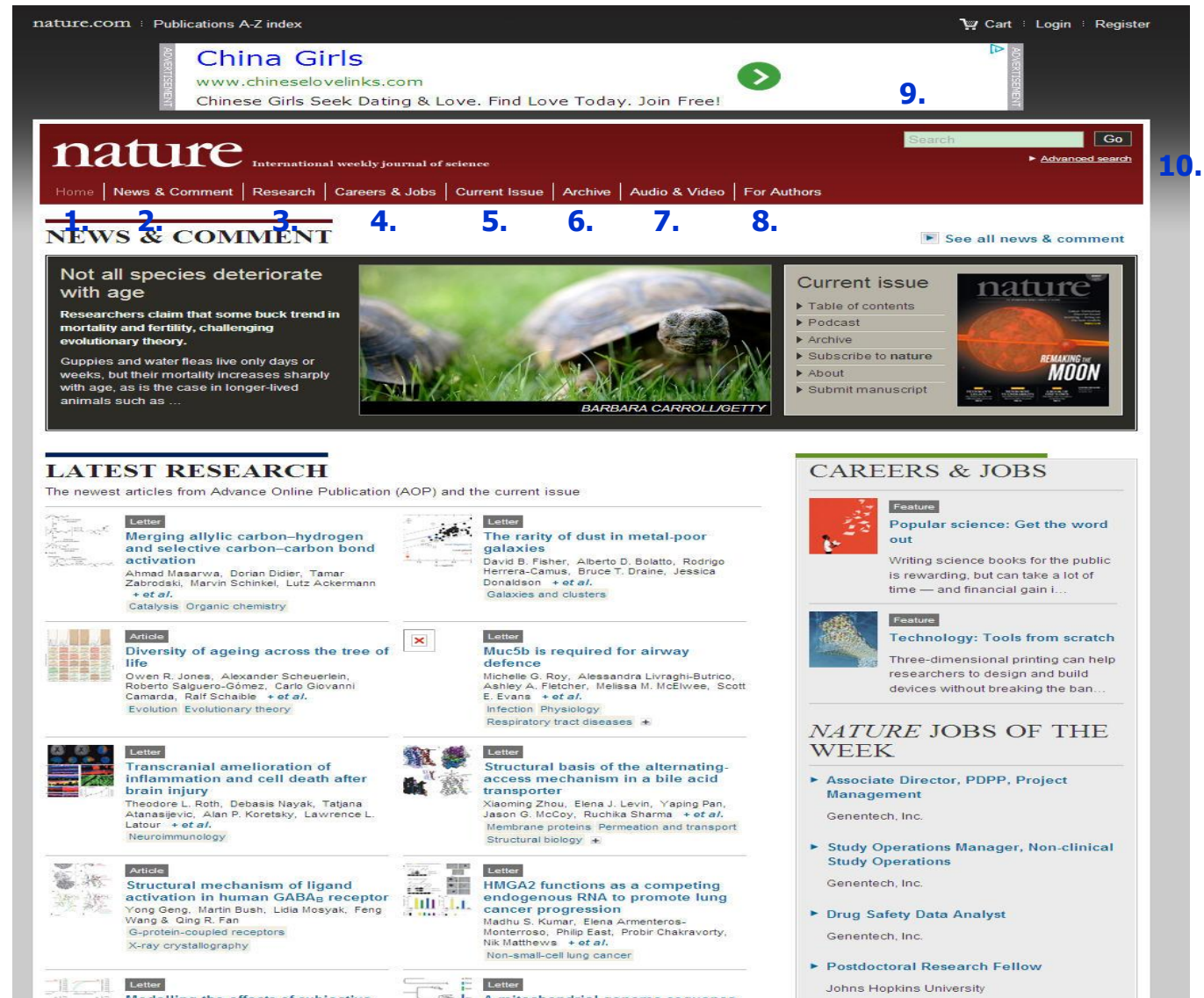
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- 5.** Points to the "LATEST RESEARCH" section header.
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- 7.** Points to the "Current issue" section.
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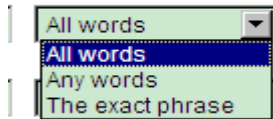
The screenshot shows the Nature journal website's 'CURRENT ISSUE' page for Volume 504, Number 7478, dated 5 December 2013. The page is annotated with four numbered points:

- 1.** Points to the top navigation menu (Home, News & Comment, Research, Careers & Jobs, Current Issue, Archive, Audio & Video, For Authors).
- 2.** Points to the 'CURRENT ISSUE' section header.
- 3.** Points to the 'THIS WEEK' section, which includes sub-sections like Editorials, World View, Research Highlights, Seven Days, News in Focus, and Careers.
- 4.** Points to the search bar at the top right of the page.

The page content includes a cover image of the journal, a 'THIS WEEK' section with articles like 'Call the cops' and 'The FDA and me', a 'WORLD VIEW' section with 'How sexual harassment changed the way I work', and a 'RESEARCH HIGHLIGHTS' section with various scientific topics. There are also sidebars for 'Journal home', 'Subscribe', 'Nature Publishing Index - China', and 'nature events directory'.

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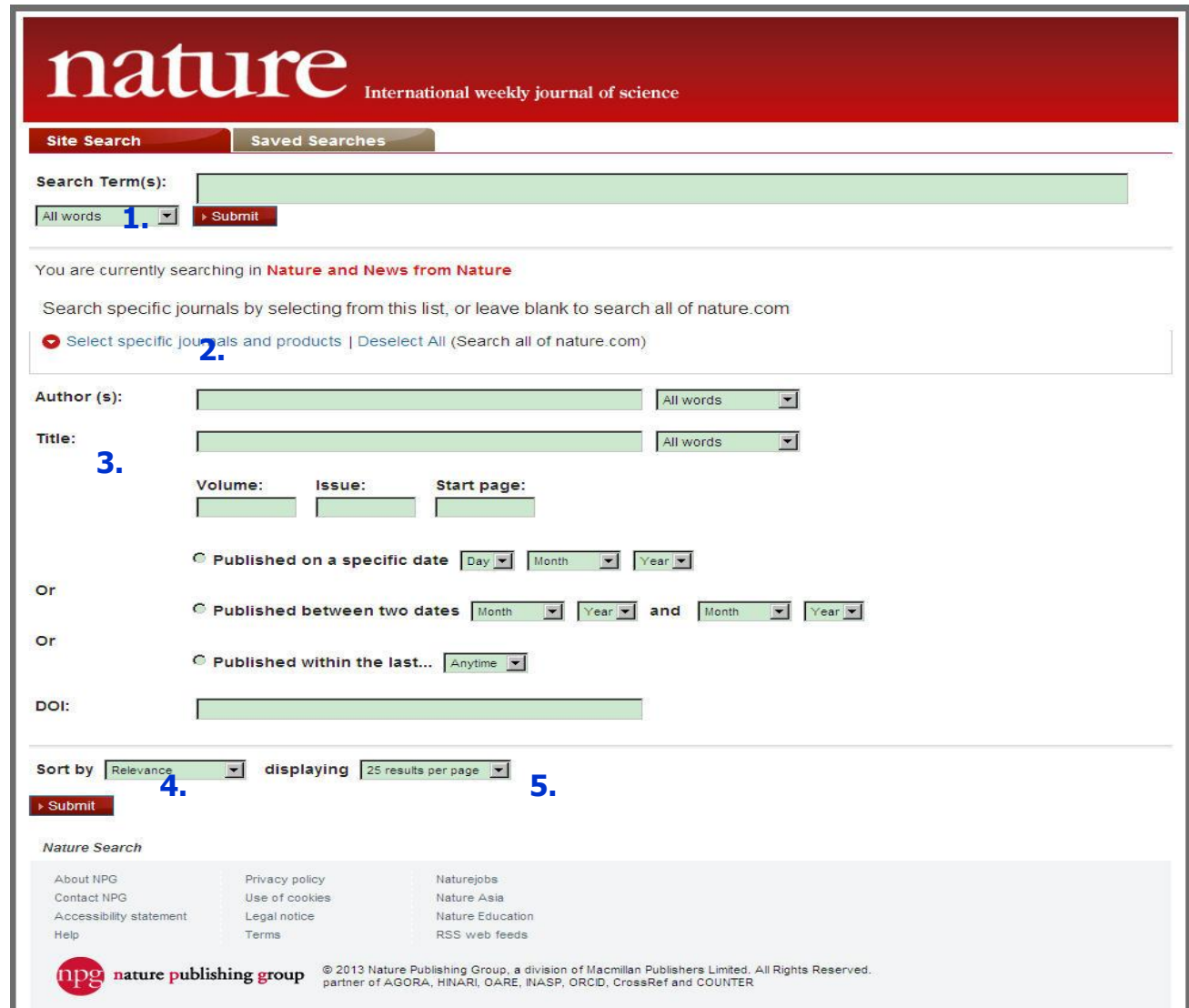
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Erratum: Sickle cell detection: 5.
Nature 329, 678-678 doi:10.1038/329678d0
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2. 22 December 1956
Genetic Studies of the Relationship of Tumour-Host Cells: Detection of an Allelic Difference at a Single Gene Locus in a Small Fraction of a Large Tumour-Cell Population
GEORGE KLEIN & EVA KLEIN
Nature 178, 1389-1391 doi:10.1038/1781389a0
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3. 4 September 2013
Immunology: Lipopolysaccharide sensing on the inside
Vijay A. K. Rathinam & Katherine A. Fitzgerald

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1. Letters to Nature

Nature **401**, 79–82 (2 September 1999) | doi:10.1038/43459 | Received 21 May 1999; Accepted 16 July 1999

2. **3.** **Cell transformation by the superoxide-generating oxidase Mox1**

Young-Ah Suh^{1,2}, Rebecca S. Arnold^{1,2}, Bernard Lassegue², Jing Shi¹, Xiangxi Xu¹, Dan Sorescu², Andrew B. Chung², Kathy K. Griendling² & J. David Lambeth²

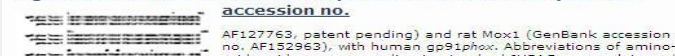
4. ¹ Departments of Biochemistry and ² Medicine, Emory University Medical School, Atlanta, Georgia 30322, USA
³ These authors contributed equally to this work.

Correspondence to: J. David Lambeth² Correspondence and requests for materials should be addressed to J.D.L. (e-mail: dlamba@bimcore.emory.edu).

5. **Reactive oxygen species (ROS) generated in some non-phagocytic cells are implicated in mitogenic signalling and cancer^{1,2,3,4,5,6,7}. Many cancer cells show increased production of ROS⁷, and normal cells exposed to hydrogen peroxide or superoxide show increased proliferation⁸ and express growth-related genes^{9,10,11}. ROS are generated in response to growth factors, and may affect cell growth^{2,3,12,13}, for example in vascular smooth-muscle cells^{6,13,14,15}. Increased ROS in Ras-transformed fibroblasts correlates with increased mitogenic rate¹⁶. Here we describe the cloning of *mox1*, which encodes a homologue of the catalytic subunit of the superoxide-generating NADPH oxidase of phagocytes^{17,18}, *gp91phox*. *mox1* messenger RNA is expressed in colon, prostate, uterus and vascular smooth muscle, but not in peripheral blood leukocytes. In smooth-muscle cells, platelet-derived growth factor induces *mox1* mRNA production, while antisense *mox1* mRNA decreases superoxide generation and serum-stimulated growth. Overexpression of *mox1* in NIH3T3 cells increases superoxide generation and cell growth. Cells expressing *mox1* have a transformed appearance, show anchorage-independent growth and produce tumours in athymic mice. These data link ROS production by Mox1 to growth control in non-phagocytic cells.**

A human expressed sequence tag (EST) sequence which showed homology to human *gp91phox* was identified, and complete sequencing revealed a predicted amino-acid sequence homologous to the carboxy-terminal half of *gp91phox*. Sequencing was completed with 5'-rapid amplification of complementary DNA ends (5'-RACE) using human colon cDNA, and the predicted amino-acid sequence is shown in Fig. 1. The predicted protein is 564 amino acids long, compared with 569 residues for *gp91phox*, and the two show 56% identity. The gene is located at Xq22 (Genebank accession no. Z83819), but the locus is not informative with regard to known diseases. Rat *mox1*, cloned by using degenerate primers based on sequences that are highly conserved between *gp91phox* and human *mox1*, is 82% identical to human *mox1* (Fig. 1). Human and rat Mox1 lack asparagine-linked candidate glycosylation sites, which are seen in the highly glycosylated human and mouse *gp91phox*^{13,21} (the positions of which are indicated by ♣ and †, respectively, in Fig. 1). Regions in *gp91phox* (underlined), which were previously proposed^{21,22} to be binding sites for flavin (regions 1a and 1b) and pyridine nucleotide (regions 2a–2d), are identical or nearly identical between *gp91phox* and Mox1. Also shown (filled circles) are conserved histidines, which are candidates for haem ligation. The hydropathy profiles of human *gp91phox* and Mox1 are nearly identical (not shown), and include five very hydrophobic stretches in the amino-terminal half of the molecules which are predicted to be membrane-spanning regions.

6. **Figure 1: Predicted amino-acid sequence of human Mox1 (GenBank accession no.**



AF127763, patent pending) and rat Mox1 (GenBank accession no. AF152963); with human *gp91phox*. Abbreviations of amino-acid residues are indicated by standard three-letter codes.

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