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# Australopithecus sediba: A New Species of Homo-Like Australopith from South Africa 

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#### Abstract

Despite a rich African Plio-Pleistocene hominin fossil record, the ancestry of Homo and its relation to earlier australopithecines remain unresolved. Here we report on two partial skeletons with an age of 1.95 to 1.78 million years. The fossils were encased in cave deposits at the Malapa site in South Africa. The skeletons were found close together and are directly associated with craniodental remains. Together they represent a new species of Australopithecus that is probably descended from Australopithecus africanus. Combined craniodental and postcranial evidence demonstrates that this new species shares more derived features with early Homo than any other australopith species and thus might help reveal the ancestor of that genus.


The origin of the genus Homo is widely debated, with several candidate ancestors being proposed in the genus Australopithecus (1-3) or perhaps Kenyanthropus (4). The earliest occurrence of fossils attributed to Homo (H. aff. H. habilis) at 2.33 million years ago (Ma) in Ethiopia (5) makes it temporally antecedent to all other known species of the genus Homo. Within early Homo, the hypodigms and phylogenetic relationships between $H$. habilis and another early species, $H$. rudolfensis, remain unresolved ( $6-8$ ), and the placement of these species within Homo has been challenged (9). H. habilis is generally thought to be the ancestor of H. erectus (10-13), although this might be questioned on the basis of the considerable temporal overlap that existed between them (14). The identity of the direct ancestor of the genus Homo, and thus its link to earlier Australopithecus, remains controversial. Here we describe two recently discovered, directly associated, partially articulated Australopithecus skeletons from the Malapa site in South Africa, which allow us to investigate several competing hypotheses regarding the ancestry of Homo. These skeletons cannot be accommodated within any existing fossil taxon; thus, we establish a new species, Australopithecus sediba, on the basis of a com-

[^0]bination of primitive and derived characters of the cranium and postcranium.

The following is a description of Au. sediba: Order Primates Linnaeus 1758; suborder Anthropoidea Mivart 1864; superfamily Hominoidea Gray 1825; family Hominidae Gray 1825; genus Australopithecus DART 1925; species Australopithecus sediba sp. nov.

Etymology. The word sediba means "fountain" or "wellspring" in the seSotho language.

Holotype and paratype. Malapa Hominin 1 (MH1) is a juvenile individual represented by a partial cranium, fragmented mandible, and partial postcranial skeleton that we designate as the species holotype [Figs. 1 and 2, supporting online material (SOM) text S1, figs. S1 and S2, and table S1]. The first hominin specimen recovered from Malapa was the right clavicle of MH1 (UW88-1), discovered by Matthew Berger on 15 August 2008. MH2 is an adult individual represented by isolated maxillary teeth, a partial mandible, and partial postcranial skeleton that we designate as the species paratype. Although MH1 is a juvenile, the second molars are already erupted and in occlusion. Using either a human or an ape model, this indicates that MH1 had probably attained at least $95 \%$ of adult brain size (15). Although additional growth would have occurred in the skull and skeleton of this individual, we judge that it would not have appreciably altered the morphology on which this diagnosis is based.

Locality. The two Au. sediba type skeletons were recovered from the Malapa site (meaning "homestead" in seSotho), situated roughly 15 km NNE of the well-known sites of Sterkfontein, Swartkrans, and Kromdraai in Gauteng Province, South Africa. Detailed information regarding geology and dating of the site is in (16).


Fig. 1. Craniodental elements of Au. sediba. UW88-50 (MH1) juvenile cranium in (A) superior, (B) frontal, and (C) left lateral views. (D) UW88-8 (MH1) juvenile mandible in right lateral view, (E) UW88-54 (MH2) adult mandible in right lateral view, (F) UW88-8 mandible in occlusal view, (G) UW 88-54 mandible in occlusal view, and (H) UW 88-50 right maxilla in occlusal view (scale bars are in centimeters).

Diagnosis. Au. sediba can be distinguished from other species of Australopithecus by a combination of characters presented in Table 1; comparative cranial measures are presented in Table 2. A number of derived characters separate $A u$. sediba from the older chronospecies $A u$. anamensis and Au. afarensis. Au. sediba exhibits neither the extreme megadontia, extensive cranial cresting, nor facial prognathism of Au. garhi. The suite of derived features characterizing Au. aethiopicus, Au. boisei, and Au. robustus, in particular the pronounced cranial muscle markings, derived facial morphology, mandibular corpus robusticity, and postcanine megadontia, are absent in Au. sediba. The closest morphological comparison for $A u$. sediba is Au. africanus, as these taxa share numerous similarities in the cranial vault, facial skeleton, mandible, and teeth (Table 1). Nevertheless, Au. sediba can be readily differentiated from $A u$. africanus on both craniodental and postcranial evidence. Among the more notable differences, we observe that although the cranium is small, the vault is relatively transversely expanded with vertically oriented parietal walls and widely spaced temporal lines; the face lacks the pro-
nounced, flaring zygomatics of $A u$. africanus; the arrangement of the supraorbital torus, nasoalveolar region, infraorbital region, and zygomatics result in a derived facial mask; the mandibular symphysis is vertically oriented with a slight bony chin and a weak post-incisive planum; and the teeth are differentiated by the weakly defined buccal grooves of the maxillary premolars, the weakly developed median lingual ridge of the mandibular canine, and the small absolute size of the postcanine dentition. These exact differences also align Au. sediba with the genus Homo (see SOM text S2 for hypodigms used in this study). However, we consider Au. sediba to be more appropriately positioned within Australopithecus, based on the following craniodental features: small cranial capacity, pronounced glabelar region, patent premaxillary suture, moderate canine jugum with canine fossa, small anterior nasal spine, steeply inclined zygomaticoalveolar crest, high masseter origin, moderate development of the mesial marginal ridge of the maxillary central incisor, and relatively closely spaced premolar and molar cusps.

Postcranially, Au. sediba is similar to other australopiths in its small body size, its relatively


Fig. 2. Associated skeletal elements of MH1 (left) and MH2 (right), in approximate anatomical position, superimposed over an illustration of an idealized $A u$. africanus skeleton (with some adjustment for differences in body proportions). The proximal right tibia of MH1 has been reconstructed from a natural cast of the proximal metaphysis.
long upper limbs with large joint surfaces, and the retention of apparently primitive characteristics in the upper and lower limbs (table S2). Au. sediba differs from other australopiths, but shares with Homo a number of derived features of the os coxa, including increased buttressing of the ilium and expansion of its posterior portion, relative reduction in the distance between the sacroiliac and hip joints, and reduction of distance from the acetabulum to the ischial tuberosity. These synapomorphies with Homo anticipate the reorganization of the pelvis and lower limb in H. erectus and possibly the emergence of more energetically efficient walking and running in that taxon (17). As with the associated cranial remains, the postcranium of $A u$. sediba is defined not by the presence of autapomorphic features but by a unique combination of primitive and derived traits.

Cranium. The cranium is fragmented and slightly distorted. The minimum cranial capacity of MH1 is estimated at $420 \mathrm{~cm}^{3}$ (SOM text S4). The vault is ovoid, with transversely expanded, vertically oriented parietal walls. The widely spaced temporal lines do not approach the midline. Postorbital constriction is slight. The weakly arched supraorbital torus is moderately developed and laterally extended, with sharply angled lateral corners and a weakly defined supratoral sulcus. A robust glabelar region is evident, with only a faint depression of the supraorbital torus at the midline. The frontal process of the zygomatic faces primarily laterally and is expanded medially but not laterally. The zygomatic prominence does not show anterolateral expansion. The zygomatics are weakly flared laterally, resulting in an uninterrupted frontal profile of the facial mask that is squared superiorly and tapered inferiorly. The zygomaticoalveolar crests are long, straight, and steeply inclined, resulting in a high masseter origin. The root of the zygomatic begins at the anterior margin of $\mathrm{M}^{1}$. The nasal bones are widened superiorly, become narrowest about one-third of the way down, and flare to their widest extent at their inferior margin. The nasal bones are elevated as a prominent ridge at the internasal suture, with an increasingly anterior projection inferiorly. The bone surface of the maxilla retreats gently away from the nasal aperture laterally, resulting in an everted margin of the superolateral portion of the aperture relative to the infraorbital region. The inferolateral portion of the nasal aperture becomes bluntly rounded. The infraorbital region is slightly convex (18) and is oriented at an approximately right angle to the alveolar plane. There is a trace of a premaxillary suture near the superolateral margin of the nasal aperture. Prominent canine juga delineate moderately developed canine fossae. Anterior pillars are absent. The inferior margin of the nasal aperture is marked by a stepped nasal sill and a small but distinct anterior nasal spine. The subnasal region is straight in the coronal plane and only weakly projecting relative

Table 1. List of characters used to diagnose Au. sediba. These characters are commonly used Numbers in parentheses in the first column refer to measures presented in Table 2 ; in hominin phylogenetic studies (11, 38-40) or have been recorded as diagnostic for various descriptions of these character states are provided in SOM text S3. Abbreviations are as hominin taxa in the past $(3,10,36)$. Recognizing the potential pitfalls of performing a follows: A-M, anteromedial; costa supr., costa supraorbitalis; intermed., intermediate; lat., cladistic analysis on possibly interdependent characters of uncertain valence, we produced a lateral; med., medial; mesognath., mesognathic; mod., moderately; MMR, mesial marginal cladogram from the data in this table as a test of the phylogenetic position of Au. sediba (fig. ridge; orthogn., orthognathic; procumb., procumbent; proj., projecting; TMJ, temperomanS3). Our most parsimonious cladogram places Au. sediba at the stem of the Homo clade. dibular joint.

| Characters | Au. afarensis | $A u$. garhi | $A u$. africanus | $A u$. sediba | H. habilis | H. rudolfensis | H. erectus | $A u$. aethiopicus | $A u$. boisei | $A u$. robustus |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| Vault |  |  |  |  |  |  |  |  |  |  |
| Cranial capacity (1) | Small | Small | Small | Small | Intermed. | Large | Large | Small | Small | Small |
| A-M incursion of temporal lines on frontal bone (9) | Strong | Moderate | Moderate | Weak | Weak | Weak | Weak | Strong | Strong | Strong |
| Position of temporal lines on parietal bones | Crest | Crest | Variable | Wide | Variable | Wide | Wide | Crest | Crest | Crest |
| Compound temporal nuchal crest (males) | Extensive | ? | Absent | Absent | Variable | Absent | Absent | Extensive | Variable | Absent |
| Postorbital constriction (5) | Marked | Moderate | Moderate | Slight | Moderate | Moderate | Slight | Marked | Marked | Marked |
| Pneumatization of temporal squama | Extensive | ? | Extensive | Reduced | Reduced | Reduced | Reduced | Extensive | Variable | Reduced |
| Facial hafting | Low | Low | Low | Low | Low | Low | Low | High | High | High |
| Frontal trigon | Present | Present | Absent | Absent | Absent | Absent | Absent | Present | Present | Present |
| Supraglenoid gutter width | Narrow | ? | Narrow | Narrow | Narrow | Narrow | Narrow | Wide | Wide | Wide |
| Horizontal distance between TMJ and M2/M3 (6) | Long | ? | Long | Short | Short | Long | Short | Long | Long | Long |
| Parietal transverse expansion/tuber | Absent | Absent | Absent | Present | Present | Present | Present | Absent | Absent | Absent |
| Facial skeleton |  |  |  |  |  |  |  |  |  |  |
| Supraorbital expression | Costa supr. | Costa supr. | Intermed. | Torus | Torus | Intermed. | Torus | Costa supr. | Costa supr. | Costa supr. |
| Supraorbital contour | Less arched | Less arched | Variable | Arched | Arched | Arched | Arched | Less arched | Variable | Arched |
| Glabellar region forms as prominent block | No | No | Variable | Yes | No | Variable | No | No | Yes | Yes |
| Lat. half of infraorbital margin blunt and protruding | No | ? | No | No | No | No | No | Yes | No | Yes |
| Zygomatic arch relative to inferior orbital margin | Above | ? | Level | Level | Level | ? | Level | Above | Above | Above |
| Convexity/concavity of infraorbital region | ? | ? | Convex | Convex | Concave | Concave | Convex | Concave | Concave | Concave |
| Nasal bone projection above frontomaxillary suture | Expanded | ? | Variable | No | No | No | No | Tapered | Expanded | Expanded |
| Inferior width of projecting nasal bone (25) | Wide | ? | Variable | Wide | Variable | Narrow | Wide | Not proj. | Not proj. | Not proj. |
| Infraorbital foramen height (32) | High | ? | Variable | High | High | ? | High | Low | Low | Low |


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| Characters | Au. afarensis | Au. garhi | Au. africanus | Au. sediba | H. habilis | H. rudolfensis | H. erectus | Au. aethiopicus | Au. boisei | Au. robustus |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| Palate |  |  |  |  |  |  |  |  |  |  |
| Protrustion of incisors beyond bi-canine line | Yes | Yes | Yes | Yes | Yes | No | Yes | No | No | No |
| Anterior palatal depth | Shallow | Shallow | Deep | Deep | Variable | Deep | Variable | Shallow | Deep | Shallow |
| Dental arcade shape | Rectangle | Rectangle | Variable | Parabolic | Parabolic | Parabolic | Parabolic | Rectangle | Parabolic | Parabolic |
| Maxillary I2/C diastema | Present | Present | Absent | Absent | Variable | Absent | Absent | Absent | Absent | Absent |
| Mandible |  |  |  |  |  |  |  |  |  |  |
| Orientation of mandibular symphysis | Receding | ? | Receding | Vertical | Vertical | Vertical | Vertical | Vertical | Vertical | Vertical |
| Bony chin (mentum osseum) | Absent | ? | Slight | Slight | Slight | Slight | Slight | Slight | Slight | Slight |
| Direction of mental foramen opening | Variable | ? | Variable | Lateral | Lateral | Lateral | Lateral | Lateral | Lateral | Lateral |
| Post-incisive planum | Prominent | ? | Prominent | Weak | Prominent | Weak | Weak | Prominent | Prominent | Prominent |
| Torus marginalis and marginal tubercles | Prominent | ? | Moderate | Moderate | Moderate | Prominent | Prominent | ? | Prominent | Prominent |
| Mandibular corpus cross-sectional area at $M_{1}(50)$ | Small | ? | Small | Small | Small | Variable | Small | Large | Large | Large |
| Teeth |  |  |  |  |  |  |  |  |  |  |
| Incisor-to-postcanine ratio (maxillary) (60) | Large | Moderate | Moderate | Moderate | Moderate | Moderate | Large | ? | Small | Small |
| Canine-to-postcanine ratio (maxillary/mandibular) $(61,62)$ | Large | Large | Large | Large | Large | Large | Large | ? | Small | Small |
| Postcanine crown area (maxillary/mandibular) $(57,59)$ | Moderate | Large | Large | Moderate | Moderate | Large | Small | Large | Large | Large |
| Maxillary $\mathrm{I}^{1}$ : MMR development, lingual face | Moderate | ? | Moderate | Moderate | Weak | Weak | Weak | ? | Moderate | Moderate |
| Maxillary C: development of lingual ridges | Marked | Marked | Marked | Weak | Weak | Marked | Marked | ? | Marked | Weak |
| Maxillary premolar molarization | None | Minor | Minor | None | Minor | Minor | None | Marked | Marked | Marked |
| Maxillary premolars: buccal grooves | Marked | Marked | Marked | Weak | Weak | Marked | Weak | ? | Weak | Weak |
| Median lingual ridge of mandibular canine | Prom. | ? | Prom. | Weak | Weak | Weak | Weak | ? | Weak | Weak |
| Mandibular $\mathrm{P}_{3}$ root number | 2 | ? | 2 | 2 | 1 | 2 | 1 | ? | 2 | 2 |
| Protoconid/metaconid more mesial cusp (molars) | Equal | ? | Equal | Protoconid | Protoconid | Protoconid | Protoconid | ? | Equal | Equal |
| Peak of enamel forms between roots of molars | No | ? | Yes | Yes | No | No | No | ? | No | Yes |
| Relative enamel thickness | Thick | Thick | Thick | Thick | Thick | Thick | Thick | Hyper | Hyper | Hyper |
| Positions of apices of lingual (LC) and buccal (BC) cusps of premolars and molars relative to occlusal margin | LC at margin, BC slightly lingual | LC at margin, $B C$ slightly lingual | LC slightly buccal, BC moderately lingual | LC slightly buccal, BC moderately lingual | LC at margin, $B C$ slightly lingual | LC at margin, $B C$ slightly lingual | LC at margin, BC slightly lingual | LC mod. buccal, BC strongly lingual | LC mod. buccal, BC strongly lingual | LC mod. buccal, BC strongly lingual |

## RESEARCH ARTICLES

otherwise indicated. Descriptions of character states presented in Table 1 that are based on measurements from this table are provided in SOM text S3. Abbreviations are as follows: br, bregma; ek, ectoconchion; ekm, ectomolare; fmt, frontomolare temporale; ft , frontotemporale;
g , glabella; mf, maxillofrontale; n , nasion; ns, nasospinale; or, orbitale; po, porion; pr,
prosthion; rhi, rhinion; zm , zygomaxillare; zy, zygion; zyo, zygoorbitale.
$A u$. $\underset{\text { boisei }}{\text { Au }}$
-n

ェ
H.
habilis

Au.
sediba

## $A u$.

$A u$. Table 1 were estimated. Several character states in Table 1 are recorded as variable, although only species average values are presented here. Measurements are in millimeters unless
Item $\begin{gathered}\text { Measurement } \\ \text { description in (6) }\end{gathered}$




 끙






| Item | Measurement description in (6) | Measurement | $A u$. afarensis | $A u$. africanus | $A u$. sediba | H. habilis | H. rudolfensis | H. erectus | $A u$. aethiopicus | Au. boisei | $A u$. robustus |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 41 | 98 | Intercanine distance | 26 | 30 | 30 | 30 | 33 | 31 | - | 29 | 27 |
| 42 | 88 | Palate breadth (ekm-ekm) | 68 | 64 | 63 | 70 | 80 | 66 | 83 | 82 | 67 |
| 43 | 141 | Mandibular symphysis height | 39 | 38 | 32 | 27 | 36 | 34 | - | 47 | 42 |
| 44 | 142 | Mandibular symphysis depth | 60 | 20 | 19 | 19 | 24 | 19 | - | 28 | 25 |
| 45 | 147 | Mandibular corpus height at $\mathrm{P}_{4}$ | 34 | 33 | 28 | 30 | 38 | 30 | - | 42 | 38 |
| 46 | 148 | Mandibular corpus depth at $\mathrm{P}_{4}$ | 19 | 21 | 18 | 20 | 22 | 19 | - | 28 | 24 |
| 47 | 149 | Cross-sectional area at $\mathrm{P}_{4}$ (calculated as an ellipse) | 511 | 558 | 382 | 427 | 653 | 458 | - | 910 | 709 |
| 48 | 150 | Mandibular corpus height at $\mathrm{M}_{1}$ | 33 | 32 | 28 | 29 | 36 | 30 | 35 | 41 | 37 |
| 49 | 151 | Mandibular corpus depth at $M_{1}$ | 19 | 21 | 18 | 20 | 23 | 20 | 26 | 28 | 26 |
| 50 | 152 | Cross-sectional area at $M_{1}$ (calculated as an ellipse) | 488 | 532 | 396 | 421 | 667 | 469 | 715 | 913 | 759 |
| 51 | 154 | Mandibular corpus height at $\mathrm{M}_{2}$ | 31 | 31 | 25 | 31 | 36 | 30 | - | 41 | 35 |
| 52 | 155 | Mandibular corpus depth at $M_{2}$ | 22 | 25 | 22 | 23 | 26 | 21 | - | 31 | 28 |
| 53 | 156 | Cross-sectional area at $\mathrm{M}_{2}$ (calculated as an ellipse) | 536 | 612 | 436 | 537 | 745 | 504 | - | 980 | 770 |
| 54 | 162 | Height of mental foramen relative to alveolar margin | 20 | 19 | 13 | 13 | 17 | 13 | - | 20 | 20 |
| 55 |  | Maxillary incisor crown area $\left(1^{1}+1^{2}\right)$ | 143 | 135 | 109 | 132 | 137 | 136 | - | 117 | 109 |
| 56 |  | Maxillary canine crown area | 107 | 104 | 79 | 95 | 118 | 96 | - | 76 | 79 |
| 57 |  | Maxillary postcanine crown area | 713 | 868 | 731 | 755 | 829 | 617 | - | 1012 | 941 |
| 58 |  | Mandibular canine crown area | 87 | 95 | 68 | 83 | - | 79 | - | 72 | 61 |
| 59 |  | Mandibular molar crown area | 550 | 651 | 536 | 565 | 668 | 466 | - | 781 | 678 |
| 60 |  | Maxillary incisor to postcanine ratio | 20.0 | 15.6 | 14.9 | 17.4 | 16.6 | 22.1 | - | 11.5 | 11.6 |
| 61 |  | Maxillary canine to postcanine ratio | 15.0 | 11.9 | 10.8 | 12.6 | 14.2 | 15.5 | - | 7.5 | 8.4 |
| 62 |  | Mandibular canine to molar ratio | 15.8 | 14.6 | 12.7 | 14.6 | - | 16.7 | - | 9.2 | 9.0 |

to the facial plane. The face is mesognathic. The palate is consistently deep along its entire extent, with a parabolic dental arcade.

Mandible. Descriptions apply to the more complete juvenile (MH1) mandible unless otherwise stated. The nearly vertical mandibular symphysis presents a weak lateral tubercle, resulting in a slight mental trigone, and a weak mandibular incurvation results in a slight mentum osseum. The post-incisive planum is weakly developed and almost vertical. Both mandibular corpora are relatively gracile, with a low height along the alveolar margin. The extramolar sulcus is relatively narrow in both mandibles. In MH1, a moderate lateral prominence displays its greatest protrusion at the mesial extent of $\mathrm{M}_{2}$, with a marked decrease in robusticity to $\mathrm{P}_{4}$; in MH2 the moderate lateral prominence shows its greatest protrusion at $M_{3}$, with a marked decrease in robusticity to $\mathrm{M}_{2}$. The alveolar prominence is moderately deep with a notable medial projection posteriorly. The anterior and posterior subalveolar fossae are continuous. The ramus of MH1 is tall and narrow, with nearly parallel, vertically oriented anterior and posterior borders; the ramus of MH2 is relatively broader, with nonparallel anterior and posterior borders (fig. S2). The mandibular notch is relatively deep and narrow in MH1 and more open in MH2. The coronoid extends farther superiorly than the condyle. The condyle is mediolaterally broad and anteroposteriorly narrow. The endocondyloid buttress is absent in MH1, whereas in MH2 a weak endocondyloid buttress approaches the condyle without reaching it.

Dental size and proportions. The dentition of the juvenile (MH1) is relatively small, whereas preserved molars of the adult (MH2) are even smaller (Fig. 3 and fig. S4). For MH1, the maxillary central incisor is distinguishable only from the reduced incisors of $A u$. robustus. The maxillary canine is narrower than all canines of $A u$. africanus except TM 1512, whereas the mandibular canine falls well below the range of $A u$. africanus. Premolars and molars are at the lower end of the $A u$. africanus range and within that of $H$. habilis-H. rudolfensis and $H$. erectus. Molar dimensions of the adult individual (MH2) are smaller than those of Au . africanus, are at or below the range of those of $H$. habilisH. rudolfensis, and are within the range of those of $H$. erectus. Au. sediba mirrors the Au. africanus pattern of maxillary molars that increase slightly in size posteriorly, though it differs in that the molars tend to be considerably larger in the latter taxon. Conversely, the Au. sediba pattern varies slightly from that seen in specimens KNM-ER 1813, OH 13, and OH 65 and H. erectus, wherein the molars increase from $M^{1}$ to $M^{2}$ but then decrease to $\mathrm{M}^{3}$. In broad terms, the teeth of $A u$. sediba are similar in size to teeth of specimens assigned to Homo but share the closely spaced cusp apices seen in Australopithecus.

Postcranium. Preserved postcranial remains of $A u$. sediba (table S1) denote small-bodied
hominins that retain an australopith pattern of long upper limbs, a high brachial index, and relatively large upper limb joint surfaces (table S2). In addition to these aspects of limb and joint proportions, numerous other features in the upper limb are shared with sibling species of Australopithecus (to the exclusion of later Homo), including a scapula with a cranially oriented glenoid fossa and a strongly developed axillary border, a prominent conoid tubercle on the clavicle, with a pronounced angular margin; low proximal-to-distal humeral articular proportions; a distal humerus with a marked crest for the brachioradialis muscle, a large and deep olecranon fossa with a septal aperture, and a marked trochlear/capitular keel (19); an ulna with a pronounced flexor carpi ulnaris tubercle; and long, robust, and curved manual phalanges that preserve strong attachment sites for the flexor digitorum superficialis muscle.

Numerous features of the hip, knee, and ankle indicate that Au. sediba was a habitual biped. In terms of size and morphology, the proximal and distal articular ends of the femur and tibia fall within the range of variation of specimens attributed to Au. africanus. However, several derived features in the pelvis link the Malapa specimens with later Homo. In the os coxa (Fig. 4), Au. sediba shares with Homo a pronounced acetabulocristal buttress; a more posterior position of the cristal tubercle; a superoinferiorly extended posterior iliac blade, with an expanded retroauricular area; a sigmoid-shaped anterior inferior iliac spine; a reduced lever arm for weight transfer between the auricular surface and the acetabulum; an enlarged and rugose iliofemoral ligament attachment area; a tall and thin pubic symphyseal face; and a relatively short ischium with a deep and narrow tuberoacetabular sulcus. These features are present in taxonomically un-
assigned postcranial remains from Koobi Fora (KNM-ER 3228) and Olduvai Gorge (OH 28), which have been argued to represent early Homo (20), as well as in early Homo erectus (21). An os coxa from Swartkrans (SK 3155) has been considered by some to also represent early Homo (22) but can be seen to possess the australopith pattern in most of these features. In addition, Au. sediba shares with later Homo the humanlike pattern of low humeral-to-femoral diaphyseal strength ratios, in contrast to the ape-like pattern seen in the H. habilis specimen OH 62 (table S2).

Although aspects of the pelvis are derived, the foot skeleton is more primitive overall, sharing with other australopiths a flat talar trochlea articular surface with medial and lateral margins with equal radii of curvature, and a short, stout, and medially twisted talar neck with a high horizontal angle and a low neck torsion angle


Fig. 3. Dental size of a selection of Au. sediba teeth compared to other early hominin taxa; see fig. S 4 for additional teeth. Dental measurements were taken as described by Wood (6). Owing to small sample sizes, H. habilis and H. rudolfensis were combined. (A) Upper central incisor mesiodistal (MD) length. (B) Upper canine MD length. (C) Lower canine MD length. (D) Square root of calculated [MD $\times$ BL (BL, buccolingual)] upper third premolar area. (E) Square root of calculated ( $\mathrm{MD} \times \mathrm{BL}$ ) upper second molar area. ( $\mathbf{F}$ ) Square root of calculated ( $\mathrm{MD} \times \mathrm{BL}$ ) lower second molar area. Measures were taken on original specimens by D.J.D. for Au. africanus, Au. robustus,
and Au. sediba. Measurements for Au. afarensis, H. habilis, H. rudolfensis, and $H$. erectus are from (6). $\mathrm{P}^{4}$ is not fully erupted on the right side of MH1, therefore measures of the maxillary postcanine dentition are presented for the left side only. Dental metrics for Au. sediba are as follows (MD, BL, in millimeters): Maxillary: MH1: RI1 10.1, 6.9; LI2 7.7 (damaged), 5.1; RC 9.0, 8.8; LP3 9.0, 11.2; LP4 9.2, 12.1; LM1 12.9, 12.0; LM2 12.9, 13.7; LM3 13.3, 14.1; MH2: RM3 11.3, 12.9. Mandibular: MH1: LC 8.0, 8.5; RM1 12.5, 11.6; RM2 14.4, 12.9; RM3 14.9, 13.8; MH2: RM1 11.8, 11.1; RM2 14.1, 12.2; RM3 14.2, 12.7; LM3 14.1, 12.5.


Fig. 4. Representative ossa coxae, in lateral view, from left to right, of $A u$. afarensis (AL 288-1), Au. africanus (Sts 14), Au. sediba (MH1), and H. erectus (KNM-WT 15000). The specimens are oriented so that the iliac blades all lie in the plane of the photograph (which thus leads to differences between specimens in the orientation of the acetabula and ischial tuberosities). MH1 possesses derived, Homo-like morphology compared to other australopithecines, including a relative reduction in the weight transfer distance from the sacroiliac (yellow) to hip (circle)
joints; expansion of the retroauricular surface of the ilium (blue arrows) (determined by striking a line from the center of the sphere representing the femoral head to the most distant point on the posterior ilium; the superior arrow marks the terminus of this line, and the inferior arrow marks the intersection of this line with the most anterior point on the auricular face); narrowing of the tuberoacetabular sulcus (delimited by yellow arrows); and pronouncement of the acetabulocristal (green arrows) and acetabulosacral buttresses.
(table S2 and fig. S5). The calcaneus is markedly primitive in its overall morphology: the bone is strongly angled along the proximodistal axis, with the point of maximum inflexion occurring at an enlarged peroneal trochlea; the lateral plantar tubercle is lacking; the calcaneal axis is set about $45^{\circ}$ to the transverse plane; and the calcaneocuboid facet is vertically set and lacks an expanded posterior projection for the beak of the cuboid (23).

Discussion. The age and overall morphology of $A u$. sediba imply that it is most likely descended from Au. africanus, and appears more derived toward Homo than do Au. afarensis, Au. garhi, and Au. africanus. Elsewhere in South Africa, the Sterkfontein cranium Stw 53, dated to 2.0 to 1.5 Ma , is generally considered to represent either $H$. habilis (10, 24, 25) or perhaps an undiagnosed form of early Homo (26). It played an important role in the assignment of OH 62 to H. habilis (27). However, the derived craniodental morphology of Au. sediba casts doubt on the attribution of Stw 53 to early Homo [see also (28)]: Stw 53 appears to be more primitive than MH1 in retaining closely spaced temporal lines; marked postorbital constriction; a weakly developed supraorbital torus; narrow, nonprojecting nasal bones; anterior pillars; marked nasoalveolar prognathism; medial and lateral expansion of the frontal process of the zygomatic bone; and laterally flared zygomatics. If Stw 53 instead represents $A u$. africanus, the assignment of OH 62 to H. habilis becomes tenuous. Attribution of the partial skeleton KNM-ER 3735 to $H$. habilis was tentatively based, in part, on a favorable comparison with OH 62 and on the hypothesis that there were no other contemporaneous non-
robust australopith species to which it could be assigned in East Africa (29). As a result, the interpretation of KNM-ER 3735 as $H$. habilis also becomes uncertain.

The phylogenetic significance of the cooccurrence of derived postcranial features in $A u$. sediba, H. erectus, and a sample of isolated fossils generally referred to Homo sp. indet. (table S2) is not clear: The latter might represent early $H$. erectus, it might sample the postcranium of $H$. rudolfensis (which would then imply an evolutionary pathway from Au. sediba to $H$. rudolfensis to $H$. erectus), or it might represent the postcranium of H. habilis [which would suggest that OH 62 and KNM-ER 3735 (two specimens with ostensibly more primitive postcranial skeletons) do not belong in this taxon]. If the latter possibility holds, it could suggest a phylogenetic sequence from $A u$. sediba to $H$. habilis to H. erectus. Conversely, although the overall postcranial morphology of Au. sediba is similar to that of other australopiths, a number of derived features of the os coxa align the Malapa hominins with later Homo (H. erectus) to the exclusion of other australopiths. Additionally, Au. sediba shares a small number of cranial traits with $H$. erectus that are not exhibited in the $H$. habilis-H. rudolfensis hypodigm, including slight postorbital constriction and convexity of the infraorbital region (18). Following on this, MH1 compares favorably with SK 847 (H. erectus) in the development of the supraorbital torus, nasal bones, infraorbital region, frontal process of the zygomatic, and subnasal projection. However, MH1 differs from SK 847 in its relatively smaller size, the robust glabelar region, the weakly developed supratoral sulcus, the steeply inclined zygomaticoalveolar crests with a
high masseter origin, and the moderate canine juga, all features aligning MH1 with Australopithecus. It is thus not possible to establish the precise phylogenetic position of $A u$. sediba in relation to the various species assigned to early Homo. We can conclude that combined craniodental and postcranial evidence demonstrates that this new species shares more derived features with early Homo than does any other known australopith species (Table 1 and table S2) and thus represents a candidate ancestor for the genus, or a sister group to a close ancestor that persisted for some time after the first appearance of Homo.

The discovery of a <1.95-million-year-old (10) australopith that is potentially ancestral to Homo is seemingly at odds with the recovery of older fossils attributed to the latter genus (5) or of approximately contemporaneous fossils attributable to $H$. erectus $(6,30)$. However, it is unlikely that Malapa represents either the earliest or the latest temporal appearance of $A u$. sediba, nor does it encompass the geographical expanse that the species once occupied. We hypothesize that $A u$. sediba was derived via cladogenesis from $A u$. africanus ( $\approx 3.0$ to 2.4 Ma ), a taxon whose first and last appearance dates are also uncertain (31). The possibility that $A u$. sediba split from Au. africanus before the earliest appearance of Homo cannot be discounted.

Although the skull and skeleton of Au. sediba do evince derived features shared with early Homo, the overall body plan is that of a hominin at an australopith adaptive grade. This supports the argument, based on endocranial volume and craniodental morphology, that this species is most parsimoniously attributed to the genus Australopithecus. The Malapa specimens dem-
onstrate that the evolutionary transition from a small-bodied and perhaps more arboreal-adapted hominin (such as Au. africanus) to a largerbodied, possibly full-striding terrestrial biped (such as $H$. erectus) occurred in a mosaic fashion. Changes in functionally important aspects of pelvic morphology, including a reduction of the sacroacetabular weight-bearing load arm and enhanced acetabulosacral buttressing (reflecting enhancement of the hip extensor mechanism), enlargement of the iliofemoral ligament attachment (reflecting a shift in position of the line of transfer of weight to behind the center of rotation of the hip joint), enlargement of the acetabulocristal buttress (denoting enhancement of an alternating pelvic tilt mechanism), and reduction of the distance from the acetabulum to the ischial tuberosity (reflecting a reduction in the moment arm of the hamstring muscles) $(20,32)$ occurred within the context of an otherwise australopith body plan, and seemingly before an increase in hominin encephalization [in contrast to the argument in (33)]. Relative humeral and femoral diaphyseal strength measures (table S2) also suggest that habitual locomotor patterns in Au. sediba involved a more modern human-like mechanical load-sharing than that seen in the H. habilis specimen OH $62(34,35)$. Mosaic evolutionary changes are mirrored in craniodental morphology, because the increasingly wide spacing of the temporal lines and reduction in postorbital constriction that characterize Homo first appeared in an australopith and before significant cranial expansion. Moreover, dental reduction, particularly in the postcanine dentition, preceded the cuspal rearrangement (wide spacing of postcanine tooth cusps) that marks early Homo.

The pattern of dental eruption and epiphyseal fusion exhibited by MH1 indicates that its age at death was 12 to 13 years by human standards, whereas in MH2 the advanced degree of occlusal attrition and epiphyseal closure indicates that it had reached full adulthood (SOM text S1). Although juvenile, MH1 exhibits pronounced development of the supraorbital region and canine juga, eversion of the gonial angle of the mandible, and large rugose muscle scars in the skeleton, all indicating that this was a male individual. And, although fully adult, the mandible and skeleton of MH2 are smaller than in MH1, which, combined with the less rugose muscle scars and the shape of the pubic body of the os coxa, suggests that MH2 was a female. In terms of dental dimensions, MH1 has mandibular molar occlusal surface areas that are $10.7 \%\left(\mathrm{M}_{1}\right)$ and $8.1 \%\left(\mathrm{M}_{2}\right)$ larger than those of MH2. Dimorphism in the postcranial skeleton likewise is not great, though the juvenile status of MH1 tends to confound efforts to assess adult body size. The diameter of the proximal epiphysis for the femoral head of MH1 ( 29.8 mm ) is approximately $9.1 \%$ smaller than the superoinferior diameter of MH2's femoral head ( 32.7 mm ). It is likely that MH1 would have experienced some appositional increase in joint size before maturity, thus this disparity would probably have de-
creased somewhat. The distal humeral epiphysis of MH 1 is fully fused and its articular breadth $(35.3 \mathrm{~mm})$ is only marginally larger than that of MH2 ( 35.2 mm ). Thus, although the dentition and postcranial skeleton are at odds in the degree of apparent size differences, the overall level of dimorphism, if these sex attributions are correct, appears slight in the Malapa hominins and was probably similar to that evinced by modern humans.

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18. Rak (36) describes a feature in the infraorbital region of Au. boisei that he refers to as a nasomaxillary basin: a concave depression that is surrounded by a more elevated topography. We see a similar concavity in the infraorbital region of specimens of $H$. habilis-H. rudolfensis (KNM-ER 1470, KNM-ER 1805, KNM-ER 1813, and OH 24 ), although it is not clear whether they represent homologous structures. In specimens of $A u$. africanus, $A u$. sediba, and $H$. erectus, we recognize a slight convexity in this area.
19. Some humeri that are probably best attributed to Australopithecus lack marked development of the trochlear/capitular keel [or "lateral crest": see (37)], and thus the absence of a marked crest does not reliably differentiate Australopithecus from Homo. However, although some specimens of early Homo (such as KNM-WT 15000) have crests that are more strongly developed than those of modern humans, none exhibit the marked crests of the australopiths. Thus, the marked crest seen in the Malapa humeri can be seen to be shared with Australopithecus rather than Homo.
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41. We thank the South African Heritage Resources Agency for the permits to work at the Malapa site; the Nash family for granting access to the Malapa site and continued support of research on their reserve; the South African Department of Science and Technology, the South African National Research Foundation, the Institute for Human Evolution, the Palaeontological Scientific Trust, the Andrew W. Mellon Foundation, the AfricaArray Program, the U.S. Diplomatic Mission to South Africa, and Sir Richard Branson for funding; the University of the Witwatersrand's Schools of Geosciences and Anatomical Sciences and the Bernard Price Institute for Palaeontology for support and facilities; the Gauteng Government, Gauteng Department of Agriculture, Conservation and Environment and the Cradle of Humankind Management Authority; E. Mbua, P. Kiura, V. Iminjili, and the National Museums of Kenya for access to comparative specimens; Optech and Optron; Duke University; the Ray A. Rothrock Fellowship of Texas A\&M University; and the University of Zurich 2009 Field School. Numerous individuals have been involved in the ongoing preparation and excavation of these fossils, including C. Dube, B. Eloff, C. Kemp, M. Kgasi, M. Languza, J. Malaza, G. Mokoma, P. Mukanela, T. Nemvhundi, M. Ngcamphalala, S. Jirah, S. Tshabalala, and C. Yates. Other individuals who have given significant support to this project include B. de Klerk, C. Steininger, B. Kuhn, L. Pollarolo, B. Zipfel, J. Kretzen, D. Conforti, J. McCaffery, C. Dlamini, H. Visser, R. McCrae-Samuel, B. Nkosi, B. Louw, L. Backwell, F. Thackeray, and M. Peltier. T. Stidham helped construct the cladogram in fig. S3. J. Smilg facilitated computed tomography scanning of the specimens. R. Clarke and F. Kirera provided valuable discussions on these and other hominin fossils in Africa.

## Supporting Online Material

www.sciencemag.org/cgi/content/full/328/5975/195/DC1
SOM Text 1 to 4
Figs. S1 to S5
Tables S1 and S2
References

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## ERRATUM

Research Articles: "Australopithecus sediba: A new species of Homo-like Australopith from South Africa" by L. R. Berger et al. (9 April, p. 195). In the legend of Fig. 3, the mesiodistal diameter of the RM2 of the mandible of the adult individual MH 2 should be 13.1 mm (not 14.1 mm ). In Fig. 4, the specimen number of the pelvis of Australopithecus afarensis (Lucy) should be A.L. 288-1 (not A.L. 228-1). These errors do not affect the Research Article's conclusions.

## COMMENTARY

Connections and health


Intellectual property and genetic tests

PERSPECTIVES

## LETTERS

## China's Plan Flawed But Courageous

IN HER NEWS FOCUS STORY "HAS CHINA OUTgrown the one-child policy?" (17 September, p. 1458), M. Hvistendahl explores the consequences of China's fertility policies. Overall, the story presents the one-child policy in a negative light, buttressed by statistical data, such as the current gender ratio of 119:100 at birth and the projected graying of the Chinese population. These data are accurate but misleadingly incomplete. The single most relevant statistic is the one that drove the original decision to implement the policy: the total population of China as a function of time. Even with the reduced population growth that came with the one-child policy and with China's rapid shift toward

an urbanized and export-driven economy, those numbers are sobering. In 1960, China had a population of 646 million; in 1980, it was 981 million; by 2000, it had grown to 1.267 billion; and in 2010 , it is projected to be 1.354 billion (1). In the past 50 years, China's population has increased by an amount equal to or greater than the population of all of Central and North America ( $\sim 500$ million). Without defending or criticizing the one-child policy, we can at least recognize that it stands as a brave attempt by
the inhabitants of an overcrowded planet to create a more livable future for our children and grandchildren.

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## Reference

1. United Nations, Department of Economic and Social Affairs, Population Division, Population Estimates and Projections Section, World Population Prospects: The 2008 Revision (http://esa.un.org/unpd/wpp2008/ tab-sorting_population.htm).

## science Readers' Picks TALK

## Your Breakthrough Nominations

EVERY YEAR, THE EDITORS AT SCIENCE GET TOGETHER TO LOOK back and decide which scientific advances merit the title of Breakthrough of the Year. This year, we also asked you, our readers, to weigh in with your nominations on our Web site. You responded immediately with a wide range of worthy ideas, and your comments were still pouring in as the magazine went to press.

As it turned out, almost all of your nominations overlapped with our selections. However, you felt that our number-one pick, the first quantum machine, was a distant dark horse to the synthetic-biology results we deemed a runner-up.

You also pointed to some results that didn't make our list. Of these nominations, the overwhelming favorite was the arsenic-based bacterium that Science published online on 2 December. The News editors who selected our breakthroughs agree that the idea of a life form with arsenic in its DNA is, as one of you put it, "effing badass!" But because the paper ran late in the year, we feel it is prudent to allow for further analysis before giving it a nod. Other recent results you felt warranted attention, such as the apparent reversal of aging in genetically engineered mice by researchers at Harvard University, fell into our wait-and-see category as well.

CERN physicists' success in trapping atoms of antihydrogen, reported in November, also justifiably impressed readers. We passed it over only because it's sure to be quickly overshadowed by follow-up experiments to measure critical properties of the mysterious antimatter.

The rest of your suggestions were all over the scientific map. They include basic research (the crystal structure of the eukaryotic ribosome and quantum entanglement in solid-state circuits); far-flung results such as the detection of free oxygen on Saturn's moon Rhea ("totally amazing"); various transformations of stem cells; and mind-boggling speculations from theoretical physics (the holographic universe). Only a couple of inventions crept in, notably a mechanical walker for paralyzed people. One techminded reader asked, "Is there any way we can sneak the commercial realization of memristors in here?" (That's not Science's kind of breakthrough, but you could try Business Week.)

We thank the many readers who participated in our Breakthrough discussion. You can read all the suggestions sent by your fellow readers, including those sent after our press deadline, at Science's Talk page (talk.sciencemag.org) and the Facebook page for our online news outlet, ScienceNOW.

SCIENCE NEWS STAFF

## Regulating Genetic Tests: Account for Benefits

THE POLICY FORUM "REGULATING DIRECT-TOconsumer personal genome testing" (A. L. McGuire et al., 8 October, p. 181) perpetuates the myth that risk-based stratification represents the best way to regulate direct-toconsumer genome tests. In fact, risk-based regulation fails to account for tolerability of risk ( 1 ), where the greater the expected benefit, the greater should be the tolerability of risk. The proposed method would likely delay access to all test results deemed high risk, regardless of the potential benefit of those results to the patient.

Sophisticated regulation based on tolerability of risk reduces the approval time and evidentiary requirements for the products with the highest potential for benefit. The U.S. Food and Drug Administration (FDA) already applies this strategy by granting an accelerated approval process for life-saving drugs, yet the agency often appears to take the opposite approach for diagnostic tests. This is an important issue as FDA considers expanding its reach into tests developed in laboratories. For many direct-to-consumer genetic tests, it is the absence of actionable benefit that appropriately reduces the tolerability of risk, not a high level of risk in absolute terms. The Policy Forum should have more explicitly stated that for patients, the harm caused by delaying a new test can be greater than the benefit of the regulation itself.

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## Reference

1. F. Bouder, D. Slavin, R. Lofstedt, The Tolerability of Risk: A New Framework for Risk Management (Earthscan, London, 2007).

## Regulating Genetic Tests: Who Owns the Data?

DIRECT-TO-CONSUMER (DTC) GENETIC TESTing is not as dangerous as A. L. McGuire et al. suggest ("Regulating direct-to-consumer personal genome testing," Policy Forum, 8 October, p. 181). The belief that genetic information will compel consumers to behave unreasonably is unfounded.

Unlike the classical genetic tests that look for rare genetic abnormalities (such as Huntington's disease), genetic tests sold directly to consumers cannot diagnose a disease. They merely provide information about DNA sequence variations, or singlenucleotide polymorphisms (SNPs). Certain SNPs can be found more often among indi-
viduals with a particular disease or condition. For example, one particular combination of two SNPs in the APOE gene occurs 3 to 20 times more frequently in individuals with Alzheimer's (1). Consequently, a person with this particular combination of SNPs could be at a greater risk to develop Alzheimer's; the mere presence of these SNPs is not diagnostic.

There is no reason to require government approval before allowing a person to see his or her own genetic information. After receiving results that indicate a healthrelated risk, a person will likely see a doctor to determine the next appropriate step. This action would be no different from a person's response to learning that he or she has increased blood pressure, cholesterol, or weight. Should DTC sales of weight scales be regulated too?

The real question is what will happen if governments, big pharma, committees, and other faceless bodies gain control of our genetic information. College admissions based on genotype? Advertising geared to those with a particular SNP? This is a question for the ethicists to ponder.

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## Reference

1. L. Bertram et al., Nat. Genet. 39, 17 (2007).

## Response

IN OUR POLICY FORUM "REGULATING DIRECT-to-consumer personal genome testing" (8 October, p. 181), we recommend a riskstratified regulatory approach for direct-toconsumer (DTC) genetic tests. If tests are low risk (as existing data indicate for the vast majority of tests sold DTC ) then the oversight should be minimal and should focus on ensuring that consumers have accurate and truthful information. Only high-risk tests would be subject to greater oversight and enhanced scrutiny by agencies such as the U.S. Food and Drug Administration (FDA).

Williams argues that this type of riskbased strategy fails to account for the impact of benefit on the tolerability of risk. We agree that benefit is an important consideration. A high-risk test that has the potential to benefit consumers significantly, and thus has proven therapeutic utility, should be cleared or approved for distribution. In fact, the sort of external data-driven review that would be required to assess benefit is precisely what the FDA would provide when evaluating a high-risk new test product. However, the primary goal of regulatory oversight is
to ensure consumer safety, so regardless of the potential for benefit, a test that poses significant risk ought to be subject to external review. The data required for the review in this circumstance allow regulators to take into account both risks and benefits.

Williams also argues that delayed access to a new test may prevent the introduction of life-saving diagnostics. We are not aware of any evidence that life-saving genetic tests, such as those used in newborn screening, are being delayed, but we agree that waiting for robust effectiveness data could effectively regulate DTC companies out of existence and should be avoided. We therefore believe that the FDA's premarket oversight should focus on the legitimacy of claims made and that there should be enhanced postmarket data collection and surveillance. This approach is quite flexible in terms of the amount and type of evidence required for a test to move to market. It is more hospitable to innovative tests than is the case in many other countries, such as Germany, where DTC testing has been banned.

Baskys believes that all individuals have a right to receive their genetic information. We do not necessarily disagree. We simply argue that consumers need access to accurate and truthful information in order to make informed decisions. For tests that have the potential to influence important medical decisions, this includes contextual information about the individual's other risk factors, including family history, environmental exposures, and biological symptoms. We agree that there is a dearth of scientific evidence on the potential risks associated with receiving genetic information DTC; there is also a lack of evidence of the potential benefits. As noted by Baskys, there are few, if any, DTC tests that have as much predictive value as a blood pressure test or a weight scale. Indeed, available evidence tells us that genetic risk information does little to motivate healthy behavior change (1). Given this reality, it seems appropriate that the vast majority of

## Letters to the Editor <br> Letters ( $\sim 300$ words) discuss material published in Science in the previous 3 months or issues of general interest. They can be submitted through the Web (www.submit2science.org) or by regular mail (1200 New York Ave., NW, Washington, DC 20005, USA). Letters are not acknowledged upon receipt, nor are authors generally consulted before publication. Whether published in full or in part, letters are subject to editing for clarity and space.

DTC tests will not require significant oversight, as per our stratified approach.

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## Reference

1. T. M. Marteau et al., "Effects of communicating DNA-based disease risk estimates on risk-reducing behaviours" (Cochrane Database of Systematic Reviews, issue 10, 10.1002/14651858.CD007275.pub2, 2010).

## CORRECTIONS AND CLARIFICATIONS

Books et al.: "How neuromythologies support sex role stereotypes" by D. F. Halpern (3 December, p. 1320). An error was introduced during the production process. Reference 5 should read: R. A. Lippa, M. L. Collaer, M. Peters, Arch. Sex. Behav. 39, 997 (2010).

Reports: "A low-magnetic-field soft gamma repeater" by N. Rea et al. (12 November, p. 944). Paolo Esposito's
affiliation contained an error. He is at the Osservatorio Astronomico di Cagliari. The affiliation has been corrected in the HTML version online.

Random Samples: "What, us bitter?" (8 October, p. 157). The accompanying photograph mistakenly showed a scene in the Swiss Alps, not the Pamir Mountains.

Research Articles: "Australopithecus sediba: A new
species of Homo-like Australopith from South Africa" by L. R. Berger et al. (9 April, p. 195). In the legend for Fig. 3, the mesiodistal diameter of the RM2 of the mandible of the adult individual MH 2 should be 13.1 mm (not 14.1 mm ). In Fig. 4, the specimen number of the pelvis of Australopithecus afarensis (Lucy) should be A.L. 288-1 (not A.L. 228-1). These errors do not affect the Research Article's conclusions.

## TECHNICAL COMMENT ABSTRACTS

## Comment on "The Incidence of Fire in Amazonian Forests with Implications for REDD"

Jennifer K. Balch, Daniel C. Nepstad, Paulo M. Brando, Ane Alencar

Aragão and Shimabukuro (Reports, 4 June 2010, p. 1275) reported that fires increase in agricultural frontiers even as deforestation decreases and concluded that these fires lead to unaccounted carbon emissions under the United Nations climate treaty's tropical deforestation and forest degradation component. Emissions from postdeforestation management activities are, in fact, included in these estimates-but burning of standing forests is not. Full text at www.sciencemag.org/cgi/content/ful//330/6011/1627-b

## Response to Comment on "The Incidence of Fire in Amazonian Forests with Implications for REDD"

Luiz E. O. C. Aragão and Yosio E. Shimabukuro

Balch et al. suggest that the increased fire frequency reported in our study is principally due to post-deforestation activities. We present a new analysis demonstrating that for the majority of grid cells with positive fire trends, there is a low likelihood that these trends have resulted exclusively from post-deforestation activities. We therefore confirm that fires pose a growing threat to reducing emissions from deforestation and degradation (REDD) policies. Full text at www.sciencemag.org/cgi/content/full/330/6011/1627-c

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