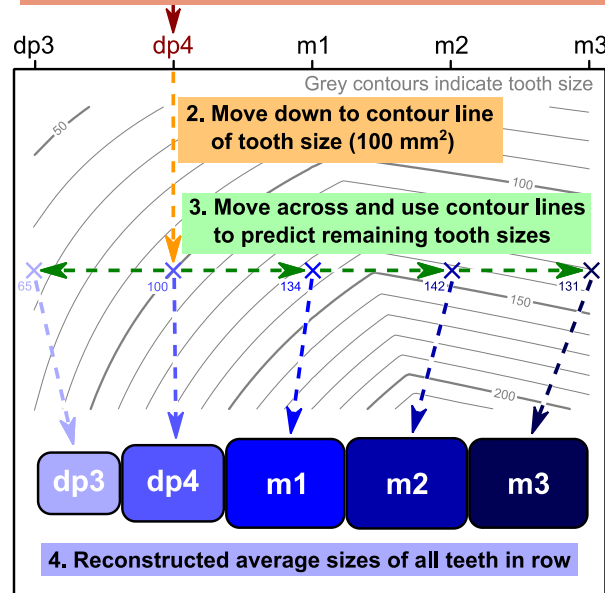


Schematic diagram of results

Predict fossil hominin tooth row from single tooth position

1. Calculate average area for a sample of fossil teeth, e.g. **dp4** with average size of 100 mm²



Supplementary Methods

Equivalencies of Inhibitory Cascade Pattern

The original formulation of the inhibitory cascade by Kavanagh *et al.*⁶ established that the ratio of the $m2/m1$ varies with the ratio of the $m3/m1$ as:

$$\frac{m3}{m1} = 2 \times \left(\frac{m2}{m1}\right) - 1$$

Kavanagh *et al.*^{6:430} showed that this is mathematically equivalent to $m2$ being 1/3 of the total area of the row. In addition, it can be shown that $m2$ is the average of $m1$ and $m3$ when the inhibitory cascade pattern is being followed:

$$\frac{m3}{m1} = 2 \times \left(\frac{m2}{m1}\right) - 1$$

$$2 \times \left(\frac{m2}{m1}\right) = \frac{m3}{m1} + 1$$

$$2 \times m2 = m1 \left(\frac{m3}{m1} + 1\right)$$

$$m2 = (m1 + m3)/2$$

Therefore, $m2$ is the sum of $m1$ and $m3$ divided by 2, which is the average of $m1$ and $m3$.

We can also show that the pattern is linear when the inhibitory cascade is being followed, i.e. that the change from $m1$ to $m2$ (given by $m2 - m1$) is the same as from $m2$ to $m3$ (given by $m3 - m2$):

$$m2 = (m1 + m3)/2$$

$$2 \times m2 = m1 + m3$$

$$m2 - m1 = m3 - m2$$

Therefore, these four statements are mathematically equivalent, and all express the expected pattern according to the inhibitory cascade:

1. $m_3/m_1 = 2 \times (m_2/m_1) - 1$
 2. m_2 is 1/3 of the total of the tooth row.
 3. m_2 is the average of m_1 and m_3 .
 4. The change in sizes across three teeth, m_1 , m_2 and m_3 , is linear.
- All of these statements are true whether the absolute tooth size is used (e.g. in mm^2) or if they are some proportion (e.g. percentage of largest tooth in the row, or percentage of m_1).

Derivation of Prediction Surface

Variables:

PropArea, area of tooth as a percentage of largest tooth in the row. *PropAreaA* and *PropAreaB* represent proportional area of tooth for regions A and B respectively.

T, tooth position (1 to 5 for dp_3 to m_3)

AM1, area of first molar (m_1), tooth position 3

Ta, tooth position slope coefficient for *PropArea*

Aa, area of m_1 slope coefficient for *PropArea*

Pb, intercept coefficient for *PropArea*

MaxArea, absolute area of largest tooth in the row

Ma, slope coefficient for *MaxArea*

Mb, intercept coefficient for *MaxArea*

Area, area of teeth for given tooth position and area of m_1 . *AreaA* and *AreaB* represent proportional area of tooth for regions A and B respectively.

AreaAStA area of teeth in region A for given tooth position and area of m_1 , standardized by *AreaA* at m_1 .

AreaBStA area of teeth in region B for given tooth position and area of m_1 , standardized by *AreaA* at m_1 .

We use the planar regressions that describe the proportional size of each tooth for each tooth position and the size of the m_1 in that row.

General regression equation for multiple linear regression of *PropArea* vs *T* and *AM1*:

$$\text{PropArea} = Ta \times T + Aa \times AM1 + Pb \quad \text{Eq. 1}$$

The hominin tooth size data are divided into two regions, A and B, where Region A is where tooth positions are 1, 2 and 3, while Region B is where tooth positions are 4 and 5.

OLS regression equations for Plane A (data in Region A) calculated from *Homo* data based on Eq. 1:

$$\text{PropAreaAH} = 0.2378 \times T + -0.001658 \times AM1 + 0.4406 \quad \text{Eq. 2}$$

OLS regression equations for Plane B (data in Region B) calculated from *Homo* data based on Eq. 1:

$$\text{PropAreaBH} = -0.08217 \times T + 0.0006890 \times AM1 + 1.226 \quad \text{Eq. 3}$$

Extended Data Fig. 10a shows the two planes. The tooth position at which the maximum tooth size occurs is the 'reversal' position, which is the intersection between Planes A and B. The intersection between Planes A and B can be found by setting Eq. 2 equal to Eq. 3:

$$\text{PropAreaA} = \text{PropAreaB} \quad \text{Eq. 4}$$

$$0.2378 \times T + -0.001658 \times AM1 + 0.4406 = -0.08217 \times T + 0.0006890 \times AM1 + 1.226 \quad \text{Eq. 5}$$

Solve Eq. 5 for *AM1*:

$$AM1H = -426.1(0.7852 - 0.3199 \times T) \quad \text{Eq. 6}$$

MaxArea gives the expected size of the largest tooth for a given m1 size:

$$MaxArea = Ma \times AM1 + Mb \quad \text{Eq. 7}$$

Regression equation for *Homo* data based on Eq. 7 (see Extended Data Fig. 6):

$$MaxAreaH = 1.312 \times AM1 - 30.44 \quad \text{Eq. 8}$$

The area of each tooth in a row (mm^2) can be estimated by multiplying the proportional size of each tooth in the row (*PropArea*) by the expected maximum area for the size of m1 in that row (*MaxArea*).

$$Area = PropArea \times MaxArea \quad \text{Eq. 9}$$

$$= (Ta \times T + Aa \times AM1 + Pb) \times (Ma \times AM1 + Mb) \quad \text{Eq. 10}$$

Formula for *AreaAH* for *Homo* (substituting equation for Plane A (Eq. 2) and Eq. 8 into Eq. 10):

$$AreaAH = (0.2378 \times T + -0.001658 \times AM1 + 0.4406) \times (1.312 \times AM1 - 30.44) \quad \text{Eq. 11}$$

Formula for *AreaBH* for *Homo* (substituting equation for Plane B (Eq. 3) and Eq. 8 into Eq. 10):

$$AreaBH = (-0.08217 \times T + 0.0006890 \times AM1 + 1.226) \times (1.312 \times AM1 - 30.44) \quad \text{Eq. 12}$$

The resulting surfaces show how the areas of all teeth in the row can be calculated given the tooth position (*T*) and the area of the m1 (*AMI*) in that row (Extended Data Fig. 10b).

The expected relationship between predicted m1 size and actual m1 size is linear, where slope = 1. However, when the *Area* formulas are used to predict m1 size, the relationship between predicted m1 area and actual m1 area is not linear because the cross-section through the surface (i.e. *AreaAH*) at $T = 3$ (i.e. at m1) is not a linear function with slope = 1 and intercept = 0. For example, if the *Area* formula is used to predict the area of m1 (*AMI*), given *Area* = 80 and $T = 3$, then the predicted area of m1 is $AMI \approx 76.1$, not the expected value of 80. The non-linear relationship is illustrated in Extended Data Fig. 10c, which is a cross-section of the cyan surface in Extended Data Fig. 10b at $T = 3$. The expected 1:1 relationship is shown in black, while *AreaAH* diverges from the 1:1 relationship.

To ensure that the cross-section at $T = 3$ is always 1:1, *Area* can be standardized by dividing by the ratio of calculated m1 area (*AMI* when $T = 3$; Eq. 13) to m1 area (Eq. 14). This forces the prediction surface at m1 to be a 1:1 linear function of *AMI*, so m1 size is always predicted correctly. The standardized area is *AreaSt*.

$$AreaAtM1 = (Ta \times 3 + Aa \times AM1 + Pb) \times (Ma \times AM1 + Mb) \quad \text{Eq. 13}$$

$$AreaSt = \frac{Area}{\left(\frac{AreaAtM1}{AMI}\right)} \quad \text{Eq. 14}$$

$$= \frac{(Ta \times T + Aa \times AM1 + Pb) \times (Ma \times AM1 + Mb)}{\left(\frac{(Ta \times 3 + Aa \times AM1 + Pb) \times (Ma \times AM1 + Mb)}{AMI}\right)} \quad \text{Eq. 15}$$

Substituting Eq. 11 and Eq. 13 into Eq. 15 gives *AreaAH* standardized by the value of m1 on *AreaAH*.

$$AreaAHStA = \frac{AreaAH}{\left(\frac{AreaAHAtM1}{AM1}\right)} \quad \text{Eq. 16}$$

$$AreaAHStA = \frac{(0.2378 \times T + -0.001658 \times AM1 + 0.4406) \times (1.312 \times AM1 - 30.44)}{\left(\frac{(0.2378 \times T + -0.001658 \times 3 + 0.4406) \times (1.312 \times AM1 - 30.44)}{AM1}\right)} \quad \text{Eq. 17}$$

AreaB standardized by the value of m1 on *AreaA* is:

$$AreaBHStA = \frac{AreaBH}{\left(\frac{AreaAHAtM1}{AM1}\right)} \quad \text{Eq. 18}$$

$$AreaBHStA = \frac{(-0.08217 \times T + 0.0006890 \times AM1 + 1.226) \times (1.312 \times AM1 - 30.44)}{\left(\frac{(0.2378 \times T + -0.001658 \times 3 + 0.4406) \times (1.312 \times AM1 - 30.44)}{AM1}\right)} \quad \text{Eq. 19}$$

To calculate the predicted sizes for a given tooth position and size, the *Area* formulas were solved for *AM1* using Wolfram Mathematica 8.0, taking the quadratic solution that corresponds with the tooth area data.

$$AM1AreaAHStA = 4.011 \times 10^{-29} \times \left(3.313 \times 10^{30} + 1.247 \times 10^{28} AreaAH - \sqrt{(-4.327 \times 10^{59} AreaAH + (-3.313 \times 10^{30} - 1.247 \times 10^{28} AreaAH - 1.788 \times 10^{30} T)^2} + 1.788 \times 10^{30} T\right) \quad \text{Eq. 20}$$

$$AM1AreaBHStA = 8.380 \times 10^{-24} \times \left(-1.062 \times 10^{26} - 1.436 \times 10^{23} AreaBH + 7.116 \times 10^{24} T + \sqrt{(2.385 \times 10^{49} AreaBH + (-1.062 \times 10^{26} - 1.436 \times 10^{23} AreaBH + 7.116 \times 10^{24} T)^2}\right) \quad \text{Eq. 21}$$

Eq. 20 and Eq. 21 are used in Supplementary Spreadsheet 1 to predict the size of the m1 in the row of Tooth X (the mean size of a sample of teeth of a given position). From the predicted size of the m1, the sizes of the remainder of the teeth in the row are predicted in Supplementary Spreadsheet 1 using Eq. 17 and Eq. 19 given the region of each tooth on the prediction surface.

OLS regression equations for Plane A (data in Region A) calculated from australopith data based on Eq. 1:

$$PropAreaAA = 0.2304 \times T + 2.379 \times 10^{-6} \times AM1 + 0.08096 \quad \text{Eq. 22}$$

OLS regression equations for Plane B (data in Region B) calculated from australopith data based on Eq. 1:

$$PropAreaBA = 0.009631 \times T + 0.0001680 \times AM1 + 0.9062 \quad \text{Eq. 23}$$

$$AM1A = -6037(0.8253 - 0.2208 \times T) \quad \text{Eq. 24}$$

Max area for australopiths:

$$MaxAreaA = 1.298 \times AM1 + 0.1495 \quad \text{Eq. 25}$$

Following the same steps as above using the australopith data give the following prediction equations:

$$\begin{aligned}
 AM1AreaAAS\tau A &= 1.973 \times 10^{-12} && \text{Eq. 26} \\
 &\times \left(-8.625 \times 10^{15} + 2.534 \times 10^{11} AreaAA \right. \\
 &+ \left. \sqrt{(8.341 \times 10^{28} AreaAA + (-8.625 \times 10^{15} + 2.534 \times 10^{11} AreaAA - 2.455 \times 10^{16} T)^2} - 2.455 \right. \\
 &\left. \times 10^{16} T \right)
 \end{aligned}$$

$$\begin{aligned}
 AM1AreaBAS\tau A &= 5.587 \times 10^{-17} && \text{Eq. 27} \\
 &\times \left(-4.827 \times 10^{19} + 1.267 \times 10^{14} AreaBA \right. \\
 &+ \left. \sqrt{(1.473 \times 10^{26} AreaBA + (-4.827 \times 10^{19} + 1.267 \times 10^{14} AreaBA - 5.130 \times 10^{17} T)^2} - 5.130 \right. \\
 &\left. \times 10^{17} T \right)
 \end{aligned}$$

Calculation of Confidence Intervals for Predictions in Supplementary Spreadsheet 1

The calculation of the 95% confidence intervals cannot be done using the usual approach³² since the predicted means are estimated as the product of two independent distributions. The regression of the maximum tooth size is done using the model matrix (X_m), a new point (x_m) at which a prediction is made, a standard error (s_m) and the regression coefficients (β_m), and yields the distribution of the prediction (m)

$$\frac{m - \beta_m x_m}{s_m \sqrt{x_m^T (X_m^T X_m)^{-1} x_m}} \sim t(df_m).$$

The two relative size planes are regressed using a model matrix (X_p), the standard error (s_p), and regression coefficients (β_p). Once the appropriate plane is determined, the prediction (p) is distributed

$$\frac{p - \beta_p x_p}{s_p \sqrt{x_p^T (X_p^T X_p)^{-1} x_p}} \sim t(df_p).$$

A derivation of the formulas is available³². The product distribution of the product of these two estimates can be calculated, but has no closed form solution. As such, a numerical approach is used. For each new point, mp is estimated by taking a random sample of 100,000 student t variables on the appropriate number of degrees of freedom and the t variables are substituted in each equation above, and the product mp is taken. The 2.5th quantile and the 97.5th quantile are then taken from this resulting sample from the product distribution.

This approach works for any new point at which a model prediction is made. However, to implement this in Excel, a simplification was made to ease computational complexity. A cubic spline was fit to each of the tooth positions at intervals of 1 mm² in m1 size, on the range of existing m1 sizes.

Phylogenetic Tree

The phylogeny in Fig. 4 is modified from Dembo *et al.*²⁸ to include only taxa represented in this study. The divergence date for *P. pygmaeus* was set at 17.5 mya³³ and the split between *P. paniscus* and *P. troglodytes* at 2.1 mya³⁴. The placement of *A. deyiremeda* (***) is based on the date of 3.3–3.5 mya and the phylogenetic position suggested in Haile-Selassie *et al.*¹⁷.

Supplementary Tables

Supplementary Table 1 | *Homo sapiens* population-level mean rectangular area (mesiodistal length \times maximum buccolingual width, mm²) of lower deciduous premolars and molars. These studies have data for both deciduous premolar and at least two molars from the same population, sometimes published in separate references. Sex: B, both; F, female; M, male.

Population	Sex	dp3	dp4	m1	m2	m3	Source
Icelanders	F			118.41	108.21		35
Icelanders	M			125.03	116.14		35
Icelanders	F	56.74	89.77				36
Icelanders	M	58.66	91.87				36
Chicago Whites	B	55.15	86.90				37
Chicago Whites	M			116.06	113.19	104.35	37
Jats (Haryana, India)	F	52.74	86.39	107.50	94.47		38
Jats (Haryana, India)	M	55.76	90.44	112.96	102.11		38
Yuendumu Aborigines	F	60.66	101.42				39
Yuendumu Aborigines	M	65.19	107.27				39
Yuendumu Aborigines	F			132.86	124.54	127.43	40
Yuendumu Aborigines	M			142.29	132.03	135.99	40
Michigan Whites	F	56.51	84.75	103.31	89.97		41
Michigan Whites	M	57.95	87.91	112.03	99.50		41
Burlington Growth Centre, Toronto	F	48.41	81.54	108.56	98.68		42
Burlington Growth Centre, Toronto	M	50.73	84.79	115.59	106.29		42
Late Archaic/Glacial Kame	B	54.27	93.60				43
Late Archaic/Glacial Kame	B			126.55	118.55	107.70	44
South Australian caucasoid	F	59.35	85.90	117.70	113.73		45
South Australian caucasoid	M	57.62	88.98	123.34	117.06		45

Supplementary Table 2 | *Homo sapiens* population-level mean rectangular area (mesiodistal length × buccolingual width, mm²) of lower molars. Sex: B, both; F, female; M, male. Comp: Compilation source.

Region	Population	Sex	m1	m2	m3	Source	Comp.
Africa	American Negros	B	128.52	130.80	129.60	46	23
Africa	Negro (S. Africa)	F	118.22	112.45	111.53	47	29
Africa	Negro (S. Africa)	M	123.70	119.00	116.70	47	29
Africa	Negro (W. Africa, Teso)	B	119.45	112.72	126.28	48	29
Africa	Nubian Agriculturalist (3300-1100BC)	F	113.18	104.53	98.30	49	29
Africa	Nubian Agriculturalist (3300-1100BC)	M	124.25	116.30	109.50	49	29
Africa	Nubian Intensive Agriculturalist (1-1400AD)	F	109.65	102.89	93.98	49	29
Africa	Nubian Intensive Agriculturalist (1-1400AD)	M	120.26	112.33	105.26	49	29
Africa	Nubian Mesolithic (10000-7000BC)	F	125.10	118.81	113.83	49	29
Africa	Nubian Mesolithic (10000-7000BC)	M	139.24	130.19	124.88	49	29
Africa	San (Bushman, Kalahari nomadic)	F	111.37	110.09	95.65	50	29
Africa	San (Bushman, Kalahari nomadic)	M	114.14	110.77	102.11	50	29
Africa	San (Bushman, skeletal)	B	111.18	107.06	95.04	51	29
Africa	San (Bushman, skeletal)	F	117.72	106.05	90.24	52	29
Africa	San (Bushman, skeletal)	M	94.50	110.21	101.00	52	29
Africa	San (Bushmen, Kalahari settlers)	F	111.62	107.32	98.01	50	29
Africa	San (Bushmen, Kalahari settlers)	M	101.57	114.49	99.50	50	29
Africa	Wadi Halfa Mesolithic	B	139.15	135.70	135.70	53	23
Asia	Chinese	M	120.89	111.27	109.71	54 in 47	29
Asia	Chinese (Bronze Age Shang Dynasty)	B	120.33	112.35	108.64	55	29
Asia	Chinese (Hong Kong)	B	113.42	106.28	99.77	55	29
Asia	India (Bronze Age Harappans)	F	106.58	91.57	83.06	56	29
Asia	India (Bronze Age Harappans)	M	111.41	100.28	94.02	56	29
Asia	India (Inagaon 1700-700BC)	B	123.56	107.95	109.44	57	29
Asia	India (Ramapuram)	M	114.13	101.88	104.75	58	29
Asia	IndiaPunjabi	B	115.19	105.03	104.48	59	
Asia	Indonesians (Gilimanuk)	B	127.60	118.77	113.36	60	23
Asia	Japanese (Ainu Hokkaido)	F	111.28	97.00	89.24	61	29
Asia	Japanese (Ainu Hokkaido)	M	122.10	106.05	97.92	61	29
Asia	Japanese (Early Jomon)	F	120.99	105.04	98.94	61	29
Asia	Japanese (Early Jomon)	M	128.82	113.40	107.06	61	29
Asia	Japanese (Early Yayoi)	F	124.30	111.28	108.12	61	29
Asia	Japanese (Early Yayoi)	M	135.70	125.40	114.40	61	29
Asia	Japanese (Hokoriku)	F	116.63	108.12	91.20	62 in 47	29
Asia	Japanese (Hokoriku)	M	120.96	115.50	102.00	62 in 47	29
Asia	Japanese (Korean descent)	F	117.70	111.28	96.03	61	29
Asia	Japanese (Korean descent)	M	127.68	122.04	114.40	61	29
Asia	Japanese (Kyoto)	F	117.66	113.36	97.92	61	29

Region	Population	Sex	m1	m2	m3	Source	Comp.
Asia	Japanese (Kyoto)	M	127.65	120.91	106.05	61	29
Asia	Japanese (Tokyo, Edo Period)	F	118.77	113.36	103.95	61	29
Asia	Japanese (Tokyo, Edo Period)	M	126.50	119.78	108.07	61	29
Asia	Malay	B	125.06	115.28	106.30		23
Asia	South-east (Bronze Age Java)	B	127.20	121.03	117.71	55	29
Asia	South-east (Bronze Age Thai)	B	125.85	112.25	105.74	55	29
Asia	South-east (Java)	B	125.93	110.96	104.71	55	29
Asia	South-east (Java)	F	119.84	106.08	109.14	63	29
Asia	South-east (Java)	M	126.50	114.45	113.36	63	29
Asia	South-east (Thai)	B	123.48	112.85	113.82	55	29
Asia	Tibet	F	100.86	78.86	62.07	64	29
Asia	Tibet	M	103.50	79.94	56.40	64	29
Australia	Broadbeach	F		133.28	117.52	65	29
Australia	Broadbeach	M	142.68	151.04	139.15	65	29
Australia	Western Australia	F	119.74	126.00	119.46	66	29
Australia	Western Australia	M	139.32	136.89	133.02	66	29
Australia	Yuendumu	F	132.93	124.98	126.33	67,68	29
Australia	Yuendumu	M	142.67	132.82	132.47	67,68	29
Europe	American Whites	B	126.14	110.09	114.13		23
Europe	Caucasoid	B	120.40	115.11	111.57		23
Europe	Lapps (Skolt)	F	116.95	109.30	104.03	69	29
Europe	Lapps (Skolt)	M	123.18	115.23	108.04	69	29
Europe	Norwegian Lapps	F	111.72	96.78	88.04	22	29
Europe	Norwegian Lapps	M	113.88	105.84	94.25	22	29
North America	Dickson Mound	B	130.13	124.65	119.77	23	23
North America	Eskimo (Hall Beach)	M	131.04	122.21		70	29
North America	Eskimo (Igloodik)	F	120.99	115.56	114.49	70	29
North America	Eskimo (Igloodik)	M	132.21	126.56	128.80	70	29
North America	Eskimo, Aleut	F	115.25	121.87	117.75	71	29
North America	Eskimo, Aleut	M	122.07	118.39	112.97	71	29
North America	Indian Knoll (4160-2558BC)	F	125.96	117.16	117.07	72	29
North America	Indian Knoll (4160-2558BC)	M	133.80	123.40	120.27	72	29
North America	Ohio Valley (Adena)	F	120.58	113.14	103.94	44	29
North America	Ohio Valley (Adena)	M	124.17	119.01	108.37	44	29
North America	Ohio Valley (Hopewell)	F	120.73	112.47	111.57	44	29
North America	Ohio Valley (Hopewell)	M	132.28	124.12	119.72	44	29
North America	Pecos Pueblo (800-1100AD)	B	128.45	119.46	112.41	73	29
North America	Tennessee Archaic (6000-500BC)	B	119.77	115.67	114.17	74	29
North America	Tennessee Mississippi (1300-1550AD)	B	120.59	114.02	107.20	74	29
North America	Tennessee Woodland (700-1150AD)	B	124.65	119.23	118.26	74	29
Oceania	Bougainville (Nasioi)	F	122.49	112.78	106.49	75	29
Oceania	Bougainville (Nasioi)	M	130.74	122.10	115.53	75	29
Oceania	New Britain	B	133.48	123.52	125.91	76	23
Oceania	New Britain (West Nakanai)	M	120.75	105.00	113.30	77	29
Oceania	New Guinea (Goroka)	M	139.95	130.98	125.18	78	29
Oceania	New Guinea (Lufa)	M	139.88	130.04	120.19	78	29

Region	Population	Sex	m1	m2	m3	Source	Comp.
South America	Lengua	F	125.53	110.58	104.54	79	29
South America	Lengua	M	132.26	119.65	115.85	79	29
South America	Peruvian (2500-1000BP)	B	122.04	114.48	104.04	80	29
South America	Peruvian (4000-2500BP)	B	115.56	111.28	112.35	80	29
South America	Peruvian (6500-4000BP)	B	117.66	111.24	99.96	80	29

Supplementary Table 3 | Fossil hominin specimen-level rectangular area (mesiodistal length × buccolingual width, mm²) of lower deciduous premolars and molars. Side: L, left; R, right; U, unknown.

Species	Specimens	Side	dp3	dp4	m1	m2	m3	Source
<i>Ardipithecus ramidus</i>	ARA-VP-1/128	L			111.1	154.7		81
<i>Ardipithecus ramidus</i>	ARA-VP-1/128	R			115.36		139.7	81
<i>Ardipithecus ramidus</i>	ARA-VP-1/129	R	35.77					81
<i>Ardipithecus ramidus</i>	ARA-VP-1/200	L			113.3			81
<i>Ardipithecus ramidus</i>	GWM16/P10	R					149.345	82
<i>Ardipithecus ramidus</i>	GWM3/P1	R			102	147.32		82
<i>Ardipithecus ramidus</i>	GWM3W/P185	L			107.625			82
<i>Ardipithecus ramidus</i>	GWM5SW/P56	L			118.8	148.59		82
<i>Ardipithecus ramidus</i>	GWM5SW/P56	R			113.42	148.59	144.64	82
<i>Ardipithecus ramidus</i>	GWM9N/P50	L			102.82	140.12		82
<i>Ardipithecus ramidus</i>	GWM9N/P50	R			107			82
<i>Ardipithecus ramidus</i>	KNM-TH 13150	R			115.44	149.34		83
<i>Australopithecus afarensis</i>	AL 128-23	L			124.32	151.25		84
<i>Australopithecus afarensis</i>	AL 145-35	L			174.2	218.68		84
<i>Australopithecus afarensis</i>	AL 188-1	R					234.08	84
<i>Australopithecus afarensis</i>	AL 198-1	L				153.76	176.66	84
<i>Australopithecus afarensis</i>	AL 200-1b	R			162.5			84
<i>Australopithecus afarensis</i>	AL 207-13	L				165		84
<i>Australopithecus afarensis</i>	AL 241-14	L			197.1			84
<i>Australopithecus afarensis</i>	AL 266-1	L			149.94			84
<i>Australopithecus afarensis</i>	AL 266-1	R			152.32	182	207	84
<i>Australopithecus afarensis</i>	AL 277-1	L				218.95		84
<i>Australopithecus afarensis</i>	AL 288-1i	L					173.24	84
<i>Australopithecus afarensis</i>	AL 288-1i	R			136.4	161.04	172.02	84
<i>Australopithecus afarensis</i>	AL 333-30	R	68					84
<i>Australopithecus afarensis</i>	AL 333-43	L	75.05	109.98				84
<i>Australopithecus afarensis</i>	AL 333-43	R	72.38	112.52				84
<i>Australopithecus afarensis</i>	AL 333-74	L			182.25		195.96	84
<i>Australopithecus afarensis</i>	AL 333w-1	L			159.72	168.75		84
<i>Australopithecus afarensis</i>	AL 333w-1	R			158.6	180.7		84
<i>Australopithecus afarensis</i>	AL 333w-12	R			166.37			84
<i>Australopithecus afarensis</i>	AL 333w-27	L				217.14		84
<i>Australopithecus afarensis</i>	AL 333w-32,60	L			174.24	211.7	204.48	84
<i>Australopithecus afarensis</i>	AL 333w-32,60	R					200.22	84
<i>Australopithecus afarensis</i>	AL 333w-48	R				152.46		84
<i>Australopithecus afarensis</i>	AL 333w-57	L				173.03	180	84
<i>Australopithecus afarensis</i>	AL 333w-59	L				201.6	183.4	84
<i>Australopithecus afarensis</i>	AL 400-1a	L			161.2	221.92	202.5	84
<i>Australopithecus afarensis</i>	AL 400-1a	R			168.91	213.15	212.52	84
<i>Australopithecus afarensis</i>	AL 417-1d	L			143.84	170.3	198.17	85
<i>Australopithecus afarensis</i>	LH 2	L		131.44				86
<i>Australopithecus afarensis</i>	LH 2	R	69.16	133.56	194.6			86
<i>Australopithecus afarensis</i>	LH 3	R	80.64		180.88			86
<i>Australopithecus afarensis</i>	LH 4	L				201.28		86
<i>Australopithecus afarensis</i>	LH 4	R			167.58		225.78	86
<i>Australopithecus afarensis</i>	MAK-VP-1/12	L			159.82	186.2	205.02	87
<i>Australopithecus afarensis</i>	MAK-VP-1/12	R			157.3	188.86	203.68	87
<i>Australopithecus afarensis</i>	MAK-VP-1/2	R			168.64	195	202.8	87

Species	Specimens	Side	dp3	dp4	m1	m2	m3	Source
<i>Australopithecus africanus</i>	MLD 2	U		138.43	212.8			88,89
<i>Australopithecus africanus</i>	MLD 5	R		125.66				89
<i>Australopithecus africanus</i>	Sts 18	R		123.6	209.89			90
<i>Australopithecus africanus</i>	Sts 24	L	62.05					91
<i>Australopithecus africanus</i>	Sts 24	R	60.48	98.1	150.29			89
<i>Australopithecus africanus</i>	Sts 4	U				188.76		90
<i>Australopithecus africanus</i>	Sts 41	U					197.81	90
<i>Australopithecus africanus</i>	Sts 52b	L				196.84	173.99	90
<i>Australopithecus africanus</i>	Sts 52b	R			167.7	192.96	174.15	90
<i>Australopithecus africanus</i>	Sts 6	U				214.2		90
<i>Australopithecus africanus</i>	Sts 7	L				223.38	221.76	90
<i>Australopithecus africanus</i>	Sts 9	U			193.7			90
<i>Australopithecus africanus</i>	Stw 106	L	69.92	129.54				89
<i>Australopithecus africanus</i>	Stw 123	L		106.7				89
<i>Australopithecus africanus</i>	Taung	L		124.02	185.9			88,89
<i>Australopithecus africanus</i>	Taung	R	69.6	123.05	189			90
<i>Australopithecus africanus</i>	TM 1515	U				257.04		90
<i>Australopithecus africanus</i>	TM 1518	U					247.16	90
<i>Australopithecus africanus</i>	TM 1519	U					218.23	90
<i>Australopithecus africanus</i>	TM 1520	U					220.8	90
<i>Australopithecus anamensis</i>	ARI-VP-1/352	R					170.17	92
<i>Australopithecus anamensis</i>	ARI-VP-3/176	R				265.68		92
<i>Australopithecus anamensis</i>	ARI-VP-3/80	R				178.92		92
<i>Australopithecus anamensis</i>	KNM-ER 20422	L			126.44			93
<i>Australopithecus anamensis</i>	KNM-ER 20428	L					207.7	93
<i>Australopithecus anamensis</i>	KNM-KP 29286	L			146.37		188.76	94
<i>Australopithecus anamensis</i>	KNM-KP 29286	R			145.14	203	188.94	94
<i>Australopithecus anamensis</i>	KNM-KP 30500	L			179.55	238.58		94
<i>Australopithecus anamensis</i>	KNM-KP 30500	R			180.78	233.73	227.8	94
<i>Australopithecus anamensis</i>	KNM-KP 34725	L	63.92		167.14	224		95
<i>Australopithecus anamensis</i>	KNM-KP 29281	L			151.13	172.62	175.68	94
<i>Australopithecus anamensis</i>	KNM-KP 29281	R			151.2	175.14	170.17	94
<i>Australopithecus anamensis</i>	KNM-KP 31712	L		83.43	127.05			95
<i>Australopithecus anamensis</i>	KNM-KP 31717	L					169.88	95
<i>Australopithecus anamensis</i>	KNM-KP 31729	R		82.45				95
<i>Australopithecus anamensis</i>	MSD-VP-3/24	L					168.74	92
<i>Australopithecus anamensis</i>	MSD-VP-5/16	L			143.99	173.24		92
<i>Australopithecus deviremeda</i>	BRT-VP-3/14	R			158.76	204.24	211.72	17
<i>Australopithecus sediba</i>	MH1	R			145	185.76	205.62	96
<i>Australopithecus sediba</i>	MH2	R			130.98	172.02	180.34	96
<i>Australopithecus sediba</i>	MH2	L					176.25	96
<i>Homo erectus</i> (Asia)	D 211	L			162.5	133.4	113.42	97
<i>Homo erectus</i> (Asia)	D 211	R			162.36	141.45	118.72	97
<i>Homo erectus</i> (Asia)	D 2600	L				180.7	187.5	98
<i>Homo erectus</i> (Asia)	D 2600	R				186.26	208.81	98
<i>Homo erectus</i> (Asia)	D 2735	L			147.84	138.6		98
<i>Homo erectus</i> (Asia)	D 2735	R			154.28	145.52		98
<i>Homo erectus</i> (Asia)	Hexian PA 831	U					120.91	99
<i>Homo erectus</i> (Asia)	Hexian PA 834	U			163.75	180.88		99
<i>Homo erectus</i> (Asia)	Hexian PA 838	U				189.04		99
<i>Homo erectus</i> (Asia)	Hexian PA 839	U				191.62		99
<i>Homo erectus</i> (Asia)	Lantian	R			144.9	163.8		99

Species	Specimens	Side	dp3	dp4	m1	m2	m3	Source
<i>Homo erectus</i> (Asia)	PCG2	L		107.67				100
<i>Homo erectus</i> (Asia)	Sangiran 9	R				179.07	175.26	101 in ⁹⁷
<i>Homo erectus</i> (Asia)	Sangiran 1B	U			162.5	171.6	181.25	102
<i>Homo erectus</i> (Asia)	Sangiran FS67	R	66.24					103
<i>Homo erectus</i> (Asia)	Sangiran FS72	R		134				103
<i>Homo erectus</i> (Asia)	Zhoukoudian G1-6	L			165	158.75	147.6	104 in ⁹⁷
<i>Homo erectus</i> (Asia)	Zhoukoudian G1-7	R				162.54	159.96	104 in ⁹⁷
<i>Homo erectus</i> (Asia)	ZKD 100	L			165			104
<i>Homo erectus</i> (Asia)	ZKD 101	U			118.65			104
<i>Homo erectus</i> (Asia)	ZKD 102	L			163.8			104
<i>Homo erectus</i> (Asia)	ZKD 106	U				135.66		104
<i>Homo erectus</i> (Asia)	ZKD 110	U				158.75		104
<i>Homo erectus</i> (Asia)	ZKD 111	U				158.76		104
<i>Homo erectus</i> (Asia)	ZKD 114	U					100	104
<i>Homo erectus</i> (Asia)	ZKD 115	U					159.96	104
<i>Homo erectus</i> (Asia)	ZKD 116	U					147.6	104
<i>Homo erectus</i> (Asia)	ZKD 117	U					102	104
<i>Homo erectus</i> (Asia)	ZKD 123	U	50.82					104
<i>Homo erectus</i> (Asia)	ZKD 125, ZKD 128	R	68.6	98.28				104
<i>Homo erectus</i> (Asia)	ZKD 126	U		94.5				104
<i>Homo erectus</i> (Asia)	ZKD 127	R		123.22				104
<i>Homo erectus</i> (Asia)	ZKD 129	R		110.88				104
<i>Homo erectus</i> (Asia)	ZKD 131	U					137.16	104
<i>Homo erectus</i> (Asia)	ZKD 134	U					113.42	104
<i>Homo erectus</i> (Asia)	ZKD 136	U					117.72	104
<i>Homo erectus</i> (Asia)	ZKD 137	U			152.1			104
<i>Homo erectus</i> (Asia)	ZKD 138	U				129.95		104
<i>Homo erectus</i> (Asia)	ZKD 147	U			143.91			104
<i>Homo erectus</i> (Asia)	ZKD 34	U			132.98			104
<i>Homo erectus</i> (Asia)	ZKD 35	R			157.5			104
<i>Homo erectus</i> (Asia)	ZKD 36	L			180.48			104
<i>Homo erectus</i> (Asia)	ZKD 38	L			99.99			104
<i>Homo erectus</i> (Asia)	ZKD 43	U				142.08		104
<i>Homo erectus</i> (Asia)	ZKD 44	U				150.65		104
<i>Homo erectus</i> (Asia)	ZKD 45	U				145.2		104
<i>Homo erectus</i> (Asia)	ZKD 51	U					147.62	104
<i>Homo erectus</i> (Asia)	ZKD 52	U					139.08	104
<i>Homo erectus</i> (Asia)	ZKD 96	R			132.16			104
<i>Homo erectus</i> (Asia)	ZKD 97	R			132.09			104
<i>Homo erectus</i> (Asia)	ZKD 98	R			171.36			104
<i>Homo erectus</i> (Asia)	ZKD 99	R			163.48			104
<i>Homo erectus</i> (Africa)	KNM-ER 1507	L	53.04	102.83				89
<i>Homo erectus</i> (Africa)	KNM-ER 1808	R					163.2	105
<i>Homo erectus</i> (Africa)	KNM-ER 820	L	71.1	100.58	131.61			106
<i>Homo erectus</i> (Africa)	KNM-ER 820	R	70.07	103.4	131.76			106
<i>Homo erectus</i> (Africa)	KNM-ER 992	L			138.43	158.6	164.82	101 in ⁹⁷
<i>Homo erectus</i> (Africa)	KNM-WT 15000	L			132.98	140.3		107
<i>Homo erectus</i> (Africa)	KNM-WT 15000	R			132.09	141.36		107
<i>Homo erectus</i> (Africa)	OH 22	R			153.4	147.32		108
<i>Homo erectus</i> (Africa)	OH 51	L			172.62			108
<i>Homo erectus</i> (Africa)	Ternifine I	R			165	171.6	152.5	109 in ¹¹⁰
<i>Homo erectus</i> (Africa)	Ternifine II	L			182	189	167.5	109 in ¹¹⁰

Species	Specimens	Side	dp3	dp4	m1	m2	m3	Source
<i>Homo erectus</i> (Africa)	Ternifine III	R			153.75	150.06	139.2	109 in 110
<i>Homo erectus</i> (Africa)	Thomas I	L			182	198	146.37	111 in 110
<i>Homo erectus</i> (Africa)	Rabat	U			143	141.25	127.2	102
<i>Homo floresiensis</i>	LB1	R			98.04	107	95	112
<i>Homo floresiensis</i>	LB6-1	U			82	84.39	70.31	112
<i>Homo habilis</i>	OH 13	L			150.8		180.56	113
<i>Homo habilis</i>	OH 13	R			149.64	168.19	179.58	113
<i>Homo habilis</i>	OH 16	R			187.96	246.13	228.8	113
<i>Homo habilis</i>	OH 27	R					201.96	113
<i>Homo habilis</i>	OH 37	L			141.24	199.5		113
<i>Homo habilis</i>	OH 4	R					201.5	113
<i>Homo habilis</i>	OH 7	L			180.56	210.6		113
<i>Homo habilis</i>	OH 7	R			183.52			113
<i>Homo heidelbergensis</i>	Arago 1	L		100.7				89
<i>Homo heidelbergensis</i>	Arago 1	R	82.82					89
<i>Homo heidelbergensis</i>	Arago 13	U			184.96	204.4	174.2	97
<i>Homo heidelbergensis</i>	Arago 2	U			119.9	129.71	101.85	110
<i>Homo heidelbergensis</i>	Arago 22	R	68.25					89
<i>Homo heidelbergensis</i>	Arago 5	L		100.44				89
<i>Homo heidelbergensis</i>	Arago 6	R	80					114 in 115
<i>Homo heidelbergensis</i>	Arago 8	R			179.4	230.74	165	110
<i>Homo heidelbergensis</i>	AT 1	L			125.4	117.42	96.6	110
<i>Homo heidelbergensis</i>	AT 1	R			131.08	123.12	107.91	110
<i>Homo heidelbergensis</i>	AT 100	L					109	110
<i>Homo heidelbergensis</i>	AT 101	R			111.24			110
<i>Homo heidelbergensis</i>	AT 11	L				107		110
<i>Homo heidelbergensis</i>	AT 13	L					143.51	110
<i>Homo heidelbergensis</i>	AT 2	R			124.02			110
<i>Homo heidelbergensis</i>	AT 21	L			136.8			110
<i>Homo heidelbergensis</i>	AT 22, AT 75	L			111.28	106.92	98.44	110
<i>Homo heidelbergensis</i>	AT 30	R					107.67	110
<i>Homo heidelbergensis</i>	Mauer	L					129.95	102
<i>Homo heidelbergensis</i>	Mauer	R			129.92	152.4	132.98	102
<i>Homo neanderthalensis</i>	Amud 1	L			117.72	113.4	127.44	116
<i>Homo neanderthalensis</i>	Amud 1	R			118.8	116.39	121.8	116
<i>Homo neanderthalensis</i>	Amud III	R		95.68				116
<i>Homo neanderthalensis</i>	Archi	L	66.6	95.68				117 in 115
<i>Homo neanderthalensis</i>	Archi	R	69.3	98.58				117 in 115
<i>Homo neanderthalensis</i>	Barakai	U		111.1				118 in 119
<i>Homo neanderthalensis</i>	Couvin	R		87				119
<i>Homo neanderthalensis</i>	Cova Negra	R		87.87				120
<i>Homo neanderthalensis</i>	Dederiyeh 1	L	66.3	102.6	120.75			121
<i>Homo neanderthalensis</i>	Dederiyeh 1	R	68.8					121
<i>Homo neanderthalensis</i>	Dederiyeh 2	L	64.38	94.94				122
<i>Homo neanderthalensis</i>	Engis 2	R	58.8	90.9				123 in 115
<i>Homo neanderthalensis</i>	Fate 12	L				126.44		124
<i>Homo neanderthalensis</i>	Fate 2	L			113			124
<i>Homo neanderthalensis</i>	Fate 3	R					139.7	124
<i>Homo neanderthalensis</i>	Fate 5	L		85.85				124
<i>Homo neanderthalensis</i>	Fate 6	R			138.75			124
<i>Homo neanderthalensis</i>	Gibraltar 2	L	72.54	108.78				125,126
<i>Homo neanderthalensis</i>	Gibraltar 2	R	76.56	105.06				125,127

Species	Specimens	Side	dp3	dp4	m1	m2	m3	Source
<i>Homo neanderthalensis</i>	Grotte du Bison I Q5.1	R		81.7				128
<i>Homo neanderthalensis</i>	Grotte du Bison I S6	R		87				128
<i>Homo neanderthalensis</i>	Grotte du Renne 16	R				81.84		129
<i>Homo neanderthalensis</i>	Grotte du Renne 18	R	72.96					129
<i>Homo neanderthalensis</i>	Grotte du Renne 21	R				146.16		129
<i>Homo neanderthalensis</i>	Grotte du Renne 25	R	63					129
<i>Homo neanderthalensis</i>	Grotte du Renne 29	R		89				129
<i>Homo neanderthalensis</i>	Grotte du Renne 30	R			71.44			129
<i>Homo neanderthalensis</i>	Grotte du Renne 33	R	60.59					129
<i>Homo neanderthalensis</i>	Grotte du Renne 35	R			133.2			129
<i>Homo neanderthalensis</i>	Grotte du Renne 5	R				141.52		129
<i>Homo neanderthalensis</i>	Grotte du Renne 6	R					122.04	129
<i>Homo neanderthalensis</i>	Hortus 15	L		83.72				130 in 115
<i>Homo neanderthalensis</i>	Hortus 2	L		83.72				130 in 125
<i>Homo neanderthalensis</i>	KMH 1	L	65.52	92.92	84.15			131
<i>Homo neanderthalensis</i>	KMH 1	R	67.5	91.35				131
<i>Homo neanderthalensis</i>	KMH 14	L				140		131
<i>Homo neanderthalensis</i>	KMH 2	L				117.72	118.77	131
<i>Homo neanderthalensis</i>	KMH 2	R				120.99	123.17	131
<i>Homo neanderthalensis</i>	KMH 32	U				127.6		131
<i>Homo neanderthalensis</i>	Krapina 1, Krapina 7, Krapina 79	R			174.795	172.36	122.5	132
<i>Homo neanderthalensis</i>	Krapina 10, Krapina 77, Krapina 108	R			154.24	154.945	140.065	132
<i>Homo neanderthalensis</i>	Krapina 104	R			112.32			132
<i>Homo neanderthalensis</i>	Krapina 105	R			170			132
<i>Homo neanderthalensis</i>	Krapina 107	L				170.8		132
<i>Homo neanderthalensis</i>	Krapina 2, Krapina 84	R			125.97	141.45		132
<i>Homo neanderthalensis</i>	Krapina 3, Krapina 5, Krapina 82	L			154.2	142.945	140.79	132
<i>Homo neanderthalensis</i>	Krapina 4	L					128.26	132
<i>Homo neanderthalensis</i>	Krapina 51	L	77.9					132
<i>Homo neanderthalensis</i>	Krapina 52, Krapina 62	L		85.2025	116.28			132
<i>Homo neanderthalensis</i>	Krapina 53	R		109.545	151.875	160.475		132
<i>Homo neanderthalensis</i>	Krapina 54, Krapina 83	L			123.585	112.7		132
<i>Homo neanderthalensis</i>	Krapina 55, Krapina 106	L			157.3	150.93	136.53	132
<i>Homo neanderthalensis</i>	Krapina 57	R			143.125	143.22	129.34	132
<i>Homo neanderthalensis</i>	Krapina 58	L			127.53	135.09	128.4	132
<i>Homo neanderthalensis</i>	Krapina 58	R			129.6	136.23	132.37	132
<i>Homo neanderthalensis</i>	Krapina 59	L			130.98	140.3		132
<i>Homo neanderthalensis</i>	Krapina 59	R					133.34	132
<i>Homo neanderthalensis</i>	Krapina 6	L				135.85		132
<i>Homo neanderthalensis</i>	Krapina 63	L		115.575				132
<i>Homo neanderthalensis</i>	Krapina 64	L	77.42	91.91	130.68			132
<i>Homo neanderthalensis</i>	Krapina 65	L			102.46			132
<i>Homo neanderthalensis</i>	Krapina 66	L			106.5			132
<i>Homo neanderthalensis</i>	Krapina 68	R			112.86			132
<i>Homo neanderthalensis</i>	Krapina 78	L					116.48	132

Species	Specimens	Side	dp3	dp4	m1	m2	m3	Source
<i>Homo neanderthalensis</i>	Krapina 8	R					155.68	132
<i>Homo neanderthalensis</i>	Krapina 80	R			146.05			132
<i>Homo neanderthalensis</i>	Krapina 85	L					126.42	132
<i>Homo neanderthalensis</i>	Krapina 86	L				145.41		132
<i>Homo neanderthalensis</i>	Krapina 9	L					125.19	132
<i>Homo neanderthalensis</i>	La Ferrassie 4 bis	L		76.8				133 in 115
<i>Homo neanderthalensis</i>	La Ferrassie 8	L	62.16	101.37				133 in 115
<i>Homo neanderthalensis</i>	La Ferrassie 8	R	62.25					133 in 115
<i>Homo neanderthalensis</i>	La Quina 761	R		99.75				134 in 119
<i>Homo neanderthalensis</i>	Molare	L		118.65				115
<i>Homo neanderthalensis</i>	Molare	R	79.2	110.74				115
<i>Homo neanderthalensis</i>	Pech de l'Aze	R	65.25	92.92				125
<i>Homo neanderthalensis</i>	Roc de Marsal infant	L	66	96.46				135
<i>Homo neanderthalensis</i>	Roc de Marsal infant	R	65.7	99.64				135
<i>Homo neanderthalensis</i>	Salemas	L		110				136 in 135
<i>Homo neanderthalensis</i>	Scladina 4A-13	R		89.3				119
<i>Homo neanderthalensis</i>	Shanidar 1	L			105	117.72	124.2	137
<i>Homo neanderthalensis</i>	Shanidar 1	R			109.2	121	126.44	137
<i>Homo neanderthalensis</i>	Shanidar 2	L			126.44	135.6	131.04	137
<i>Homo neanderthalensis</i>	Shanidar 2	R			124.32	129.92	120.96	137
<i>Homo neanderthalensis</i>	Shanidar 6	R				148.68	156.16	137
<i>Homo neanderthalensis</i>	Shanidar 7	L	67.64	91				138
<i>Homo neanderthalensis</i>	Shanidar 7	R	61.92					138
<i>Homo neanderthalensis</i>	Tabun series IV	L		105.28				139:209 in 119
<i>Homo neanderthalensis</i>	Teshik-Tash	L	78.3	96.9				140 in 125
<i>Homo neanderthalensis</i>	Teshik-Tash	R	76.5	105.6				140 in 125
<i>Paranthropus boisei</i>	KNM-ER 1477	L	97.9	168.51				141
<i>Paranthropus boisei</i>	KNM-ER 1477	R	101.2	161.46				141
<i>Paranthropus boisei</i>	KNM-ER 3230	L			262.4	372.13	332.1	105
<i>Paranthropus boisei</i>	KNM-ER 3230	R			246.33	377.88	358.28	105
<i>Paranthropus boisei</i>	KNM-ER 729	R			248	351	418	142
<i>Paranthropus boisei</i>	OH 26	R					290.5	113
<i>Paranthropus boisei</i>	OH 30	L			252			113
<i>Paranthropus boisei</i>	OH 38	R				316.8		113
<i>Paranthropus robustus</i>	DNH 10	R					230.79	143
<i>Paranthropus robustus</i>	DNH 18	R					270.04	143
<i>Paranthropus robustus</i>	DNH 19	L				252.32		143
<i>Paranthropus robustus</i>	DNH 2	L		117.16				143
<i>Paranthropus robustus</i>	DNH 21	L				212.67	195.91	143
<i>Paranthropus robustus</i>	DNH 44	R	94.16					143
<i>Paranthropus robustus</i>	DNH 46	R			198.45			143
<i>Paranthropus robustus</i>	DNH 51	R				233.52	236.3	143
<i>Paranthropus robustus</i>	DNH 56	L		113.85				143
<i>Paranthropus robustus</i>	DNH 56	R		116				143
<i>Paranthropus robustus</i>	DNH 60	R	72.38	123.22	161.84	188.5		143
<i>Paranthropus robustus</i>	DNH 67	R			178.12			143
<i>Paranthropus robustus</i>	DNH 7	L			168.84	180.9	206.72	144
<i>Paranthropus robustus</i>	DNH 7	R				191.7	206.36	144
<i>Paranthropus robustus</i>	DNH 75	R					231.82	143
<i>Paranthropus robustus</i>	DNH 8	L			227.65	238.5	309.42	144

Species	Specimens	Side	dp3	dp4	m1	m2	m3	Source
<i>Paranthropus robustus</i>	DNH 8	R			223.44	229.4	291.4	144
<i>Paranthropus robustus</i>	KB 5223	L	74.26	128.75	179.8			88
<i>Paranthropus robustus</i>	KB 5223	R		128.75	178.75			88
<i>Paranthropus robustus</i>	SK 1	U				260.1		90
<i>Paranthropus robustus</i>	SK 104	U			205.5			90
<i>Paranthropus robustus</i>	SK 12	U					246.24	90
<i>Paranthropus robustus</i>	SK 1586	L				207		145
<i>Paranthropus robustus</i>	SK 1586	R				223.5	250.5	145
<i>Paranthropus robustus</i>	SK 1587	L			189.63			145
<i>Paranthropus robustus</i>	SK 1587	R				196.08		145
<i>Paranthropus robustus</i>	SK 1588	L			172.2			145
<i>Paranthropus robustus</i>	SK 1648	R					227.65	145
<i>Paranthropus robustus</i>	SK 23	L			217.56	224.96	208	90
<i>Paranthropus robustus</i>	SK 23	R			214.62	222	249.12	90
<i>Paranthropus robustus</i>	SK 25	L			205.72			90
<i>Paranthropus robustus</i>	SK 25	R			203	250.5		90
<i>Paranthropus robustus</i>	SK 34	U			208.8	270.54	297	90
<i>Paranthropus robustus</i>	SK 37	U				246.5		90
<i>Paranthropus robustus</i>	SK 3974	R			199.66			145
<i>Paranthropus robustus</i>	SK 3976	L			276.8			145
<i>Paranthropus robustus</i>	SK 3978	L	79.56	136.5				145
<i>Paranthropus robustus</i>	SK 3978	R	81.18	138.03				145
<i>Paranthropus robustus</i>	SK 438	L			141.7			89
<i>Paranthropus robustus</i>	SK 5	U				213		90
<i>Paranthropus robustus</i>	SK 55b	L		138.99	195.91			90
<i>Paranthropus robustus</i>	SK 55b	R			197.28	224.51	212.35	90
<i>Paranthropus robustus</i>	SK 6	L			243.2	275.31	292.3	90
<i>Paranthropus robustus</i>	SK 6	R			235.06	281.88	289.6	90
<i>Paranthropus robustus</i>	SK 61	L	100.7	158.27				90
<i>Paranthropus robustus</i>	SK 61	R	105.45	160.8	224.84			90
<i>Paranthropus robustus</i>	SK 62	L	85.05	143.88				90
<i>Paranthropus robustus</i>	SK 62	R			160.8			90
<i>Paranthropus robustus</i>	SK 63	L	72.54	127.2	199.8			90
<i>Paranthropus robustus</i>	SK 63	R	84.6	127.2	201.15			90
<i>Paranthropus robustus</i>	SK 64	R	92.22	134.62				90
<i>Paranthropus robustus</i>	SK 81	U					251.37	90
<i>Paranthropus robustus</i>	SK 828	U			223.08			90
<i>Paranthropus robustus</i>	SK 838	U			182			90
<i>Paranthropus robustus</i>	SK 839, SK 852	R	81					89
<i>Paranthropus robustus</i>	SK 840	U					206.4	90
<i>Paranthropus robustus</i>	SK 841	U			116.4		210.98	90
<i>Paranthropus robustus</i>	SK 842	U			125.24			90
<i>Paranthropus robustus</i>	SK 843	U			197.1	219.62		90
<i>Paranthropus robustus</i>	SK 844	U					215.28	90
<i>Paranthropus robustus</i>	SK 845	R			217.5	233.6		90
<i>Paranthropus robustus</i>	SK 846	U			200.1			90
<i>Paranthropus robustus</i>	SKW 4767	U			233.28			146
<i>Paranthropus robustus</i>	SKW 5	L				231.84		147
<i>Paranthropus robustus</i>	SKW 5	R			179.52	233.28	233.8	147
<i>Paranthropus robustus</i>	SKX 4446	U			221.65	270.18		148
<i>Paranthropus robustus</i>	SKX 4446	U			221.65	270.18		146
<i>Paranthropus robustus</i>	SKX 5002	L					247.42	149

Species	Specimens	Side	dp3	dp4	m1	m2	m3	Source
<i>Paranthropus robustus</i>	SKX 5013	U			163.2			148
<i>Paranthropus robustus</i>	SKX 5013	U			163.2			146
<i>Paranthropus robustus</i>	SKX 5014	U					258	148
<i>Paranthropus robustus</i>	SKX 5023	U			180.48			148
<i>Paranthropus robustus</i>	SKX 5023	U			180.48			146
<i>Paranthropus robustus</i>	TM 1536	L	79.38					89
<i>Paranthropus robustus</i>	TM 1536	R	76.8	116.62	149.86			88,89
<i>Paranthropus robustus</i>	TM 1601	R	85					89

Supplementary Table 4 | Hominin mean rectangular area (mesiodistal length × buccolingual width, mm²) of lower deciduous premolars and molars, and number of individuals sampled at each tooth position for fossil hominin species in parentheses.

Species	dp3	dp4	m1	m2	m3
<i>Homo sapiens</i>	55.96	90.11	122.46	114.45	109.95
<i>Ardipithecus ramidus</i>	35.77 (1)	(0)	110.37 (7)	148.01 (5)	144.56 (3)
<i>Australopithecus afarensis</i>	72.88 (4)	121.88 (2)	165.11 (17)	186.28 (18)	199.28 (13)
<i>Australopithecus africanus</i>	66.93 (3)	120.80 (7)	186.97 (6)	215.66 (5)	213.31 (6)
<i>Australopithecus anamensis</i>	(0)	(0)	148.82 (7)	207.84 (7)	186.58 (7)
<i>Australopithecus deyiremeda</i>	(0)	(0)	158.76 (1)	204.24 (1)	211.72 (1)
<i>Australopithecus sediba</i>	(0)	(0)	137.99 (2)	178.89 (2)	191.96 (2)
<i>Homo erectus</i> (Asia)	61.89 (3)	111.43 (6)	150.69 (19)	160.07 (18)	141.49 (16)
<i>Homo erectus</i> (Africa)	61.81 (2)	102.41 (2)	155.44 (10)	162.08 (8)	151.54 (7)
<i>Homo floresiensis</i>	(0)	(0)	90.02 (2)	95.70 (2)	82.66 (2)
<i>Homo habilis</i>	(0)	(0)	165.37 (4)	206.11 (4)	203.08 (4)
<i>Homo heidelbergensis</i>	77.02 (3)	100.57 (2)	136.20 (9)	150.21 (7)	125.93 (9)
<i>Homo neanderthalensis</i>	68.61 (17)	96.52 (34)	130.50 (24)	137.74 (24)	131.57 (19)
<i>Paranthropus boisei</i>	99.55 (1)	164.99 (1)	251.46 (3)	347.60 (3)	351.23 (3)
<i>Paranthropus robustus</i>	84.02 (11)	131.00 (14)	199.79 (29)	232.09 (20)	240.40 (21)

Supplementary Table 5 | Mean, maximum and standard deviation of prediction error rates (%) for *Homo* and australopith fossil individuals. The observed size of the tooth in the ‘Predicted using’ column was used to predict the sizes of each other tooth in the fossil. The average prediction errors for all tooth positions and specimens were 10.3% and 7.9% for *Homo* and australopith specimens respectively. * indicates values that include *Homo neanderthalensis* KMH 1 where m1 is incompletely developed and so does not fit predictions.

Mean Prediction Error

Taxon	Predicted using	Predicted tooth				
		dp3	dp4	m1	m2	m3
<i>Homo</i>	dp3		12.0	30.1*		
	dp4	8.6		12.7*	1.8	
	m1	17.1*	9.4*		10.2	15.5
	m2		1.2	7.0		8.7
	m3			10.1	8.5	
Australopiths	dp3		7.8	10.0	20.0	
	dp4	7.1		6.6	14.5	9.6
	m1	9.2	7.0		8.1	7.9
	m2	16.3	12.3	7.5		7.4
	m3		7.9	7.5	7.8	

Maximum Prediction Error

Taxon	Predicted using	Predicted tooth				
		dp3	dp4	m1	m2	m3
<i>Homo</i>	dp3		36.7	62.2*		
	dp4	26.7		46.0*	1.8	
	m1	32.8*	30.1*		34.8	54.5
	m2		1.2	21.1		29.2
	m3			26.9	22.5	
Australopiths	dp3		19.1	29.1	20	
	dp4	16.1		16.1	17.4	19.2
	m1	22.6	19.2		22.8	23.5
	m2	16.3	14.5	18.0		21.4
	m3		15.7	29.2	27.2	

SD of the Prediction Error

Taxon	Predicted using	Predicted tooth				
		dp3	dp4	m1	m2	m3
<i>Homo</i>	dp3		10.9	22.8*		
	dp4	7.4		16.9*		
	m1	11.1*	10.7*		7.2	12.2
	m2			4.7		7.6
	m3			6.4	6.7	
Australopiths	dp3		6.0	8.5		
	dp4	5.1		4.7	4.1	13.5
	m1	6.7	5.3		5.9	7.0
	m2		3.1	4.9		5.0
	m3		11.0	6.7	5.8	

Supplementary Table 6 | Great ape mean rectangular area (mesiodistal length × buccolingual width, mm²) of lower deciduous premolars and molars. Number of individuals is in parentheses. Sex: F, female; M, male.

Species	Sex	dp3	dp4	m1	m2	m3	Source
<i>Gorilla gorilla</i>	F	80.24 (20)	128.55 (16)	198.74 (35)	245.75 (28)	226.61 (17)	¹⁵⁰
<i>Gorilla gorilla</i>	M	89.53 (16)	144.83 (13)	222.63 (30)	280.18 (23)	260.68 (17)	¹⁵⁰
<i>Pan paniscus</i>	F	36.00 (2)	55.08 (2)	86.24 (40)	92.82 (25)	78.26 (13)	³⁰
<i>Pan paniscus</i>	M	36.00 (3)	55.89 (4)	87.22 (33)	90.16 (17)	75.60 (12)	³⁰
<i>Pan troglodytes</i>	F	41.09 (17)	63.8 (16)	103.79 (48)	113.35 (36)	103.44 (32)	¹⁵⁰
<i>Pan troglodytes</i>	M	40.53 (19)	66.1 (18)	106.37 (30)	113.76 (17)	102.43 (10)	¹⁵⁰
<i>Pongo pygmaeus</i>	F	56.32 (5)	94.28 (5)	150.56 (5)	166.06 (3)	159.29 (3)	¹⁵¹
<i>Pongo pygmaeus</i>	M	67.6 (9)	104.49 (9)	161.17 (12)	179.54 (6)	183.37 (3)	¹⁵¹

Supplementary Table 7 | 2D and 3D measures of tooth size for six fossil hominin specimens. Rectangular area (mesiodistal length × maximum buccolingual width, MDBLArea), 3D area of the enamel-dentine junction (EDJ3DArea), cross-sectional area of the tooth at the cervix (CervixArea) and outline area of the outer enamel surface (OES2DArea) for each tooth position.

Species	Specimen	Tooth	Cervix Area	EDJ3D Area	MDBL Area	OES2D Area
<i>Australopithecus anamensis</i>	KNM-KP 29286	m1	115.3	226.7	148.9	129.3
<i>Australopithecus anamensis</i>	KNM-KP 29286	m2	160.6	279.6	198.8	168.7
<i>Australopithecus anamensis</i>	KNM-KP 29286	m3	148.5	255.9	193.7	159.3
<i>Homo erectus</i>	Sangiran 1B	m1	114.5	265.4	169.0	141.6
<i>Homo erectus</i>	Sangiran 1B	m2	128.3	277.7	180.8	153.9
<i>Homo erectus</i>	Sangiran 1B	m3	120.7	253.7	180.0	147.2
<i>Homo neanderthalensis</i>	Scladina 4A I	m1	73.9	216.1	124.0	102.4
<i>Homo neanderthalensis</i>	Scladina 4A I	m2	84.0	211.1	128.3	106.3
<i>Homo neanderthalensis</i>	Scladina 4A I	m3	80.5	189.2	126.5	102.3
<i>Paranthropus boisei</i>	KNM-ER 15930	m1	125.6	281.8	181.9	154.0
<i>Paranthropus boisei</i>	KNM-ER 15930	m2	164.6	320.2	244.2	201.8
<i>Paranthropus boisei</i>	KNM-ER 15930	m3	176.7	342.3	277.5	220.0
<i>Paranthropus robustus</i>	DNH 8	m1	125.8	316.0	215.9	185.5
<i>Paranthropus robustus</i>	DNH 8	m2	150.9	317.4	242.8	203.0
<i>Paranthropus robustus</i>	DNH 8	m3	163.1	370.0	304.3	243.7
<i>Paranthropus robustus</i>	SK 6	m1	162.0	352.7	246.2	207.3
<i>Paranthropus robustus</i>	SK 6	m2	190.7	373.6	286.9	234.2
<i>Paranthropus robustus</i>	SK 6	m3	200.4	374.5	301.3	245.3

Supplementary Table 8 | Results for multiple linear regression of proportion of maximum in tooth row (PropRowMaxP) vs tooth position (ToothNoP) and area of m1 (AreaM1) for *Homo* species.

Plane A

Coefficients:

	Estimate	Std. Error	t value	Pr(> t)	
(Intercept)	0.4405763	0.0940624	4.684	0.000352	***
ToothNoP	0.2377533	0.0134270	17.707	5.56e-11	***
AreaM1	-0.0016578	0.0006297	-2.633	0.019684	*

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 0.04591 on 14 degrees of freedom

(14 observations deleted due to missingness)

Multiple R-squared: 0.9593, Adjusted R-squared: 0.9535

F-statistic: 165 on 2 and 14 DF, p-value: 1.848e-10

Plane B

Coefficients:

	Estimate	Std. Error	t value	Pr(> t)	
(Intercept)	1.2257523	0.1115384	10.990	2.86e-07	***
ToothNoP	-0.0821669	0.0207012	-3.969	0.0022	**
AreaM1	0.0006890	0.0004452	1.548	0.1500	

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 0.03873 on 11 degrees of freedom

(10 observations deleted due to missingness)

Multiple R-squared: 0.6226, Adjusted R-squared: 0.554

F-statistic: 9.075 on 2 and 11 DF, p-value: 0.004701

Supplementary Table 9 | Results for multiple linear regression of proportion of maximum in tooth row (PropRowMaxP) vs tooth position (ToothNoP) and area of m1 (AreaM1) for australopithecines.

Plane A

Coefficients:

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	8.096e-02	7.664e-02	1.056	0.309
ToothNoP	2.304e-01	1.652e-02	13.950	1.32e-09 ***
AreaM1	2.379e-06	3.358e-04	0.007	0.994

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 0.0576 on 14 degrees of freedom

(17 observations deleted due to missingness)

Multiple R-squared: 0.9345, Adjusted R-squared: 0.9251

F-statistic: 99.8 on 2 and 14 DF, p-value: 5.197e-09

Plane B

Coefficients:

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	0.9062316	0.0813763	11.136	5.09e-08 ***
ToothNoP	0.0096313	0.0162932	0.591	0.565
AreaM1	0.0001680	0.0002022	0.831	0.421

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 0.03259 on 13 degrees of freedom

(10 observations deleted due to missingness)

Multiple R-squared: 0.07408, Adjusted R-squared: -0.06837

F-statistic: 0.5201 on 2 and 13 DF, p-value: 0.6064

Supplementary Table 10 | Results for multiple linear regression of proportion of maximum in tooth row (PropRowMaxP) vs tooth position (ToothNoP) and area of m1 (AreaM1) for great apes.

Plane A

Coefficients:

	Estimate	Std. Error	t value	Pr(> t)	
(Intercept)	0.1729463	0.0432668	3.997	0.00312	**
ToothNoP	0.2677833	0.0135136	19.816	9.85e-09	***
AreaM1	-0.0007271	0.0002288	-3.178	0.01121	*

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 0.03822 on 9 degrees of freedom

(10 observations deleted due to missingness)

Multiple R-squared: 0.9781, Adjusted R-squared: 0.9733

F-statistic: 201.4 on 2 and 9 DF, p-value: 3.374e-08

Plane B

Coefficients:

	Estimate	Std. Error	t value	Pr(> t)	
(Intercept)	1.2875205	0.1459685	8.821	0.000311	***
ToothNoP	-0.0836563	0.0307087	-2.724	0.041567	*
AreaM1	0.0003374	0.0003184	1.060	0.337646	

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 0.04343 on 5 degrees of freedom

(10 observations deleted due to missingness)

Multiple R-squared: 0.6309, Adjusted R-squared: 0.4832

F-statistic: 4.272 on 2 and 5 DF, p-value: 0.08279

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