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**AMALGAMATED REFERENCE DATA FOR SIZE-ADJUSTED BONE
DENSITOMETRY MEASUREMENTS IN 3598 CHILDREN AND YOUNG ADULTS
– THE ALPHABET STUDY**

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1 **Abstract**

2 The increasing use of dual energy X-ray absorptiometry (DXA) in children has led to the
3 need for robust reference data for interpretation of scans in daily clinical practice. Such data
4 need to be representative of the population being studied and be ‘future-proofed’ to software
5 and hardware upgrades. The aim was to combine all available paediatric DXA reference data
6 from seven UK centres to create reference curves adjusted for age, sex, ethnicity and body
7 size to enable clinical application, using in-vivo cross calibration and making data back- and
8 forward- compatible.

9 Seven UK sites collected data on GE-Lunar or Hologic Scanners between 1996 and 2012.
10 Males and females aged 4 to 20 years were recruited (n=3598). The split by ethnic group
11 was: White Caucasian 2887; South Asian 385; Black Afro-Caribbean 286; mixed heritage 40.
12 Scans of the total body and lumbar spine (L1-L4) were obtained. The European Spine
13 Phantom was used to cross-calibrate the 7 centres and 11 scanners. Reference curves were
14 produced for L1-L4 bone mineral apparent density (BMAD) and total body less head (TBLH)
15 and L1-L4 areal bone mineral density (aBMD) for GE Lunar Prodigy and iDXA (sex-and
16 ethnic-specific) and for Hologic (sex-specific). Regression equations for TBLH BMC were
17 produced using stepwise linear regression. Scans of 100 children were randomly selected to
18 test backwards and forwards compatibility of software versions, up to version 15.0 for GE
19 Lunar, and Apex 4.1 for Hologic.

20 For the first time, sex and ethnic- specific reference curves for lumbar spine BMAD, aBMD
21 and TBLH aBMD are provided for both GE-Lunar and Hologic scanners. These curves will
22 facilitate interpretation of DXA data in children using methods recommended in ISCD
23 guidelines. The databases have been created to allow future updates and analysis when more

24 definitive evidence for the best method of fracture prediction in children is agreed.

25

26 **Keywords: DXA; paediatric; BMD; BMC; reference; lean mass**

27

28 **Introduction**

29 The increasing availability and use of dual energy X-ray absorptiometry (DXA) technology
30 in children has brought to the fore the need for robust reference data for all DXA
31 manufacturers. Although manufacturer reference databases are available, they are often not
32 population based nor representative of the individual population being studied (1). Such
33 databases may also have wide variability due to small numbers, with limited power to model
34 rapid skeletal changes during different phases of growth. A further limitation for their use in
35 daily practice is the widespread use of multiple generations of hardware and acquisition and
36 analysis software that may distort the output. There is a need to enable transition between
37 them when monitoring skeletal health in individual patients or undertaking longitudinal
38 research studies.

39 In 2013 the International Society for Clinical Densitometry (ISCD) updated their 2007
40 Pediatric Bone Densitometry Guidelines for bone assessment in children (1-3). The
41 committee concluded that DXA is the preferred method for assessment of areal bone mineral
42 content (BMC) and density (aBMD) and that estimating aBMD should be part of the overall
43 assessment for children at elevated risk of a clinically significant fracture (1-3).

44 Measurements of total body less head (TBLH) and/ or posterior-anterior lumbar spine aBMD
45 or BMC are recommended; in conjunction with a history of clinically significant fractures,
46 these can be used to indicate the diagnosis of osteoporosis in children and adolescents (1-3).

47 In children with short stature or growth delay, the measurements should be size-corrected
48 using appropriate methods (4-7). The guidelines also acknowledge that adjustment for soft-
49 tissue measurements may be useful in children with malnutrition or in those with muscle and/
50 or skeletal deficits, as has been shown previously (8-11). Despite these guidelines, there are
51 still inconsistencies in the management of children with low BMD and bone fragility around

52 the world. The lack of robust reference data in a format that permits the diagnostic
53 application of ISCD recommendations is a source of inconsistency. .
54 The primary aim of the current study was to combine all available paediatric DXA reference
55 data from seven UK centres to create age-, sex-, ethnic- and size-corrected reference curves
56 for use in clinical practice and prediction equations for the assessment of the muscle and bone
57 relationship, and a database which is in-vivo cross calibrated and back- and forward-
58 compatible.

59

60 **Methods**

61 *Subjects*

62 Three thousand five hundred and ninety eight healthy, community dwelling children aged 4
63 to 20 years were recruited from 7 UK centres (Birmingham, Leeds, London, Glasgow,
64 Sheffield, Middlesbrough, Manchester) using centre-specific protocols, from 1996 to
65 2012(**Supplementary Table 1**). Participants were a self-selected convenience sample from
66 across each study region, recruited through advertisement in local schools and colleges,
67 general practice surgeries and youth groups. Children of White Caucasian, South-Asian and
68 Black Afro-Caribbean /African descent were included in the study, depending on centre-
69 specific inclusion and exclusion criteria. Ethnicity was defined by participants' self-reporting
70 both parents being of identical ethnic origin; where this was not the case, data were excluded.
71 All centres recruited healthy children without known metabolic bone disease, confirmed
72 through centre-specific screening questionnaires (**Supplementary Table 1**); abnormal results
73 were followed-up and excluded if metabolic bone disease was suspected. Children were
74 included who had had one or more moderate or high trauma fractures (12). At all centres,
75 local research Ethics Committees approved the studies. All research was carried out in
76 accordance with the Declaration of Helsinki.

77

78 *Anthropometric measurements*

79 Height and weight were measured according to centre specific protocols and body mass index
80 (BMI) calculated as weight/height² (kg/m²). To describe the population at each centre,
81 height, weight and BMI measurements were transformed to standard deviation scores (Z-
82 Scores) using the 1990 British growth reference data (13-15).

83

84 *Scan acquisition*

85 Children were scanned at each centre on either a GE Lunar™ DPX-L, Prodigy or iDXA
86 scanner (GE Medical Systems, Madison, Wisconsin, US) in Birmingham, Leeds, London,
87 Glasgow, Sheffield, Middlesbrough or on a QDR Discovery Hologic™ scanner (Hologic,
88 Bedford, MA, US) in Manchester. Total body, lumbar spine and proximal femur scans were
89 obtained; since the femur is not currently a recommended site according to the current ISCD
90 guidelines (2) only total body and lumbar spine are reported. Standard operating procedures
91 were followed in each centre. All scans were analysed centrally in Birmingham by two
92 Clinical Scientists and were scored for quality of scan acquisition and analysis. DPX-L scans
93 were analysed using software version 4.6c, Prodigy and iDXA scans using Encore version
94 15.0 (Basic and Enhanced) and Hologic scans using Apex 4.1. Spine bone mineral apparent
95 density (BMAD) was calculated using an adapted method of Carter et al. (g/cm³) (4, 16, 17).

96 Lumbar spine BMAD (g/cm³) =
$$\frac{(BMC_1 + BMC_2 + BMC_3 + BMC_4)}{(V_1 + V_2 + V_3 + V_4)}$$

97 Where V_n is the volume of the nth individual vertebra = AP_n^{1.5} (AP_n = Projected vertebral
98 area of the nth vertebra)

99 BMC_n is the bone mineral content of the nth vertebrae

100

101 Prediction equations were generated for GE Lunar (Prodigy, iDXA) and Hologic (Discovery)
102 for predicted total body less head bone mineral content (TBLH-BMC) by linear regression
103 analysis of log transformed, lean mass, fat mass, height and age (9, 18).

104

105 *Centre cross-calibration:*

106 The European Spine Phantom (ESP) was used to cross-calibrate bone measurements at 7
107 centres and 11 scanners. (19, 20). The phantom was measured once at each centre 10 times
108 without repositioning. For practical purposes this process was not repeated and therefore we
109 relied on local monitoring of scanner operation to verify machine stability. Birmingham was
110 used as the reference centre and all sites cross-calibrated to these measurements.

111 Additional measurements were taken on the iDXA and Hologic scanners using the Leeds
112 Paediatric Spine Phantom, developed by The University of Leeds (in-house).

113

114 *In-vivo cross calibration:*

115 *In-vivo* cross calibration was performed in Birmingham, firstly for DPX-L to Prodigy in
116 healthy children (n=105) and then for Prodigy to iDXA in children undergoing scans for
117 clinical purposes (n=70) . Both studies were approved by South Birmingham Ethics
118 Committees. Cross-calibration equations were produced using linear regression analysis of
119 absolute values. Machine differences were tested using paired t-test and machine bias with
120 Bland and Altman (**Supplementary table 2**). The equations were used to transform data from
121 the other GE-Lunar centres to Birmingham for lumbar spine DPX-L to Prodigy Basic and
122 iDXA; and for total body DPX-L to Prodigy basic, Prodigy enhanced and iDXA ¹. *In-vivo*
123 cross-calibration was not performed between Hologic and GE-scanners for bone or soft tissue
124 measurements.

^a Prodigy Enhanced is an option only available for total body scans.

125

126 *Back- and forward compatibility*

127 Scans of 100 children were selected from each of the GE Lunar and Hologic databases to
128 create equations for back- and forwards-compatibility of the reference curves. Within each
129 cohort of 100 children, 20 children per age-band (5-7, 8-10, 11-13, 14-16, 17-19 years) were
130 selected at random (10 male, 10 female) from each of the manufacturer specific datasets.
131 Total body and lumbar spine scans were analysed on software versions: GE-Lunar 10, 11, 13,
132 14, 15; Hologic 12.4, Apex 2.4, 3.1, 4.1. This sub-set of scans remains available for analysis
133 for future software versions.

134

135 *Statistical analysis*

136 The Lambda-Mu-Sigma (LMS) method was used to produce age reference curves for Lumbar
137 Spine BMAD, L1-L4 aBMD and TBLH BMD. The LMS curves were generated using the
138 method described by Cole and Green (21) (LMSchartmaker Pro version 2.54 © 1997-2011
139 Medical Research Council, UK). In brief, reference centile curves describe the distribution of
140 the dependent variable as it varies with the independent predictor covariate, here being age.
141 The curves are fitted using the parametric approach of the penalised log likelihood method as
142 cubic splines by non-linear regression. The degree of smoothing required for the curves is
143 expressed in terms of the equivalent degrees of freedom (edf) (21). The resulting model for
144 the dependent variable, generated from the raw data, is summarised by three parameters,
145 namely: L the Box-Cox power transformation needed to remove any skewness from the
146 distribution, M the median, and S the coefficient of variation. The LMS models were fitted
147 using the “Loop” analysis function in the software, setting the maximum edf’s for the cubic

148 splines at 3, 6 and 3 and the minimum edf's at 0,1 and 1, for L, M and S respectively. The
149 reference model choice was guided by the Schwarz Bayesian Criterion and visual inspection
150 of the curves, resulting in a parsimonious model. Goodness of fit was investigated using the
151 detrended Q-Q plots and ensuring the Q-test statistic was less than 2 (22-24). Standardized
152 residuals were tested for normality and the distribution of subjects within the expected
153 centiles was calculated.

154 Figures 1-3 and Supplemental Figures 3-5 highlight the age-related mean with the 5th and 95th
155 confidence intervals with each sex and ethnic group fitted separately. Standard deviation
156 scores (Z-scores) are calculated from the LMS parameters using the equation;

$$Z = ((\frac{y}{M})^L - 1) / L * S$$

157 *Z = Z- score, y = measured value, M = estimated mean, L = skewness, S = distribution*

158 The need for ethnic specific curves was tested using a one-sided t-test of the Z-scores
159 calculated from the gender specific white data. Where, a significant difference from zero was
160 observed, ethnic specific curves were generated. The goodness of fit of the curves is
161 described by comparing expected versus observed Z -score centile distributions in
162 **Supplemental Tables 7a-j.**

163 Regression equations for TBLH-BMC were produced using stepwise linear regression;
164 covariates in the initial model were log-transformed total body lean, total body fat, height and
165 age, only significant covariates were used. Residual plots were inspected for normality to
166 check for skewness and bias in the prediction models.

167 **Results**

168 A total of 3598 scans from children and young adults aged 4 to 20 years-old were included in
169 this study (1820 female, 1778 male). The split by ethnic group was: White Caucasian 2887;

170 South Asian 385; Black African/ Afro Caribbean 286 and 40 mixed heritage. One hundred
171 and one subjects were excluded (61 extreme body size [either height, weight or BMI SDS < -
172 3.5 or > 3.5SD]; 40 mixed heritage), leaving a total of 3497 subjects for the generation of
173 reference data (Table 1). Descriptive data by centre are shown in Table 2. There were small,
174 significant centre differences in height, weight and BMI SDS. Subjects were generally taller,
175 heavier with greater BMI than the 1990 UK-reference population (13-15).

176

177 *Manufacturer differences*

178 Phantom cross calibration: Using the ESP and with Birmingham as the reference centre there
179 were no significant differences between all 11 scanners in phantom BMC and aBMD
180 (including Hologic). In contrast, BA was more variable between the centres but the only
181 significant difference was observed between the Hologic scanner and all GE scanners
182 ($p=0.010$) (Supplemental Figure 1).

183 We explored these differences further using the Leeds Paediatric Spine Phantom scanned on
184 a Hologic Discovery and GE-Lunar iDXA scanners. There were no significant differences in
185 aBMD however BMC and BA were significantly different between the two ($p<0.001$), with
186 Hologic giving increasingly higher values compared to the iDXA with increasing BMC and
187 BA. Therefore, transformation equations were produced. However, when we applied these to
188 the *in-vivo* data there were still systematic differences between the Hologic and GE-Lunar
189 datasets. Consequently, we could not combine different manufacturer scan data and thus
190 needed to generate brand-specific reference data for use in clinical practice.

191 In-vivo cross-calibration: *In-vivo* cross-calibration data were only available for the GE-Lunar
192 scanners (25, 26). The strong linear relationships between scanners from a single
193 manufacturer enabled successful transformation of the *in-vivo* reference datasets collected
194 from three generations of GE-Lunar scanners. Once successfully transformed, the Bland

195 Altman tests showed no residual bias. Consequently, this allowed the pooling of all the GE-
196 Lunar data.

197

198 *Software differences – backwards and forwards compatibility*

199

200 For GE Lunar, there were no differences in any parameter measured using the basic analysis
201 from version 10 onwards (Prodigy). Version 14.0 included an enhanced total body analysis
202 to try and make Prodigy total body results comparable with the newly introduced iDXA.

203 Whilst there were no differences between the basic analysis, it is not surprising that there
204 were differences between the basic and enhanced total body analyses for all measured
205 parameters (aBMD, BMC, BA, lean and fat) (**Supplemental Figure 2**).

206 For Hologic there were no differences between software versions 12.4 through Apex 4.1. It is
207 important to note that this is only true if the same analysis option is used; for this study
208 NHANES BCA was selected throughout.

209

210 *Reference curve generation (Figures 1-3, Supplementary data S3-5)*

211 Because of the known differences in development between boys and girls their data were
212 separately analysed for BMAD , aBMD and TBLH-BMC.

213

214 *Size-adjusted lumbar spine (Supplemental tables 4a-c)*

215 Small, but significant differences were found for BMAD between White and Asian, and
216 White and Black children, (Figure 1). In girls, the mean difference in Z-score, calculated
217 using White as the referent group, was 0.25 (0.88), $p < 0.0001$ and 0.62 (1.18) $p < 0.0001$ for
218 South Asian and Black Caribbean girls respectively (Supplemental Table 7a-b). In boys, the
219 mean difference in Z-score, again calculated using White as referent group, was 0.24 (0.96),

220 $p=0.001$ and 0.46 (0.98) $p<0.0001$ for South Asian and Black Caribbean's respectively
221 (Supplemental Table 7a-b). When Z-scores were recalculated using ethnic-specific LMS data
222 they were no longer significantly different from 0. LMS data were therefore generated for
223 each ethnic group separately.

224 Figure 3 shows inter-scanner curve comparisons for males and females separately. Despite
225 cross-calibrating the Hologic BMC and BA values to GE Lunar using the ESP, highly
226 significant differences between the scanners remained confirming the differences described
227 earlier. The result of these differences was that calculated BMAD was lower from the
228 Hologic scanner. We explored whether this was due aBMD, BMC or BA. BMC and aBMD
229 were not different but BA was greater in Hologic. Using log-log transformation, (27) the
230 relationship between BA and BMC differed between scanners: for Prodigy, iDXA and DPX-
231 L this was $BA^{1.7}$ (expected $BA^{1.5}$ (4)), whereas for the QDR Discovery it was $BA^{1.9}$.

232

233 *Lumbar spine and total body less head areal BMD (Supplemental Tables S5-6)*

234 In contrast to the BMAD findings there were no significant differences in South Asian
235 children when compared to the white group. Differences remained for black compared to
236 white girls (lumbar spine 0.69 (1.14) $p<0.001$; TBLH 1.04 (1.08), $p<0.0001$) and boys
237 (lumbar spine 0.56 (0.97) $p<0.0001$; TBLH 0.93 (1.06), $p<0.0001$) (Supplemental Tables
238 S7d,e, 7e, h). We therefore combined the data for White and South Asian children, and re-
239 checked the distribution of Z-scores to check for normality and to ensure differences were not
240 significantly different from 0, they were not confirming the appropriateness of combining
241 data.

242

243 *Total body less head BMC (Tables 3-6)*

244 ANOVA was performed with TBLH-BMC as the dependent variable and lean body mass, fat
245 body mass, height, age, gender and ethnicity as co-variates or factors in the model.
246 Significant effects were noted for all covariates and factors. Total body lean mass was the
247 greatest predictor of TBLH-BMC, closely followed by total body fat mass, age and height.
248 Significant interactions were noted for all covariates between genders and ethnic groups
249 ($p < 0.001$). Girls had greater TBLH-BMC than males for the same lean mass, fat mass, height
250 and age. For the same gender, Afro-Caribbean children had greater TBLH-BMC for the same
251 covariate values (data not shown). Consequently, using stepwise linear regression analysis
252 with parsimonious variable selection of the log-transformed parameters, individual predictor
253 models were generated for each manufacturer, each ethnic group and each gender (Table 3a-
254 d). Individual Z-scores can be produced from by inputting age, height, lean and fat mass in to
255 the prediction equation. The predicted value can then be used to calculate the Z-score by
256 using the following equation:

$$Z - score = \frac{\text{Measured value} - \text{Predicted value}}{\text{Predicted value} \times SEE}$$

257

258

259 **Discussion**

260 For the first time, DXA measurements in children and young adults aged 4-20 years
261 combining data collected across multiple generations of GE-Lunar and Hologic DXA
262 scanners and software have been collated. Reference data are presented using some of the
263 recently recommended methods by ISCD for clinical use. We provide reference curves for
264 age- and size-adjusted lumbar spine and total body bone densitometry up to the age of 20
265 years. We also give prediction equations for size- and body composition-adjusted TBLH-
266 BMC measurements. These data enable calculation of sex-specific Z-scores for three ethnic

267 groups from 4 years-of-age through to the children switching to adult transition services.
268 Looking ahead, our random dataset of 100 healthy children provides forwards compatibility
269 of software, which allows us testing of future software updates.

270

271 *Scanner differences*

272 The strong linear relationships between the in-vivo cross-calibration of the reference datasets
273 enabled pooling of all of the GE-Lunar scanners after applying machine specific (i.e. Prodigy,
274 i-DXA) in-vivo transformation equations (**Supplementary Table 2a-b**). Unfortunately, only
275 data from *in-vitro* phantoms were available for cross-calibration between the two scanner
276 manufacturers. The observed BA differences were due to varying projectional errors of the
277 fan-beam (Hologic) versus narrow-fan (GE-Lunar) technology. Since the phantom consists
278 of an anthropomorphic spine set in a fixed position it cannot account for differences in body
279 thickness or spine depth which introduces significant errors in measurement when scanning
280 *in-vivo*. For this reason we were unable to cross-calibrate Hologic to GE-Lunar data. Our
281 findings confirm the inappropriate nature of using phantoms to cross-calibrate between
282 hardware with different properties, i.e. pencil → narrow-fan → fan beam (28,29).

283

284 *Software differences*

285 The data presented here are for the latest software version of each manufacturer; Encore 15.0
286 (GE Lunar) and Apex 4.1 (Hologic). With simple transformations it is possible to interpret
287 the DXA results using any version of software going back to GE Lunar Encore 10.0 and
288 Hologic 12.4. Our findings confirm that for both manufacturers it is necessary to always use
289 software specific reference data. It should be noted that for both, it is essential to ensure that
290 when comparing results from different software versions the same analysis options are
291 selected. For GE-Lunar this means selecting enhanced or basic analysis, and for Hologic

292 Apex software the NHANES BCA analysis should be switched on (30). For older, pre-Apex
293 versions of Hologic, the ‘auto whole body analysis’ should be used.

294

295 *Reference data and their use in fracture prediction*

296 Our study presents age- (TBLH-aBMD, spine aBMD) and size-adjusted data for bone
297 densitometric variables (BMAD, TBLH-BMC) previously shown to best predict fractures in
298 healthy or chronically ill children (31); these also represent some of the methods currently
299 recommended by ISCD (1, 2). In over 450 children with chronic disease the diagnostic odds
300 ratio for predicting vertebral fractures was 9.3 (5.3-14.9) for lumbar spine BMAD; for
301 predicting long bone fractures the odds ratio was 6.5 (4.1-10.2) for TBLH-BMC for lean
302 mass (31). BMAD has also been shown to be the best size-adjustment method for prediction
303 of fractures in healthy children (32). Current understanding is that when interpreting
304 paediatric bone density results it is preferable to use a size-adjustment method, such as
305 BMAD or a height-adjusted Z-score(1), however a firm consensus regarding the most
306 appropriate size-adjustment technique has yet to be established and for this reason the use of
307 age-adjusted aBMD is still recommended by ISCD (2). Unlike previous studies, some of
308 which are described below, that present reference data from a single manufacturer and using
309 one software version (7, 16, 33, 34) the data presented here can easily be applied to different
310 software versions and manufacturers. If necessary, data can be regenerated using newer size-
311 adjustment methodology.

312 The Bone Mineral Density Childhood Study (BMDCS) multi-center study generated robust
313 US-population-derived reference data for Hologic scanners (software version 12.3 for
314 baseline and Apex 2.1 for follow-up scans) from over 10 000 measurements in over 2000
315 individuals of TBLH and lumbar spine BMC and aBMD measurements in 5 to 20-year olds
316 (6, 6). Size-adjusted prediction equations using height for age Z-scores were also generated

317 and verified using an independent dataset. No data have yet been published to show whether
318 this method of adjustment significantly improves fracture prediction. Reference data were
319 also generated from the NHANES study; to date only LMS data for total body composition
320 have been published (33). It should be noted that both the NHANES and the BMDCS studies
321 generate Hologic reference data and are from much larger population samples than the UK
322 database presented here.

323 In contrast to the current study, NHANES data have been cross-calibrated from Hologic to
324 GE-Lunar. Data generated on Hologic 4500 scanners (software version Apex 3.0) were cross
325 calibrated to GE Lunar iDXA values (Software version 14.0) (29, 34). However, despite
326 being the largest published database (approximately 20 000 measurements), only data for
327 total body measurements were presented. Since reductions in TBLH-BMC only predict long
328 bone and not vertebral fracture risk (31), isolated total body data may have limited clinical
329 use. Another possible limitation of the NHANES reference database translation to GE
330 measurements is that pragmatic cross-calibration was performed using data from a native
331 Chinese population and then applied to transform a much larger dataset of a North American
332 US population (34).

333

334 *Limitations*

335 There are several limitations to this study. The previously discussed differences in phantom
336 measurements between the scanners due to projection error and table height differences
337 (Figure 3) and subsequent lack of in-vivo data for cross-calibration meant that we were
338 unable to create a single combined dataset, applicable to both manufacturers' scanners. The
339 data were all collected in UK centres, but are applicable for use worldwide provided the same
340 software and scan protocols are used. Caution should be applied when using the data in

341 populations in which there may be differences in growth rates or body habitus and robust
342 testing should be employed. In our study the sample size for the South Asian and Afro-
343 Caribbean populations were considerably smaller than the White population and recruited
344 mostly from one centre and as such we cannot be certain that this is fully representative of the
345 population. We cannot rule out recruitment bias in any of the centres but as can be seen from
346 **Supplementary Table 1** protocols and sampling strategies were broadly the same.
347 Although we cannot confirm that the differences between GE Lunar and Hologic reference
348 data were not due to population differences, it is likely that the differences are due to
349 differences in scanner technology. We believe the cross-calibration procedure is as robust as
350 it can be, since collecting repeated measurements on scanners across the country is neither
351 ethical nor feasible. Because only one centre collected Hologic data, in one ethnic group,
352 there are fewer subjects and the Hologic dataset did not include different ethnic groups.
353 Despite this, we have made this Hologic dataset robust to software updates and increased the
354 utility of the data previously published in 2007 (16). Finally, we have focussed on testing
355 the data based on bone measurements only, clearly repeating this work for body composition
356 would be an advantage (29, 34).

357

358 *Conclusion*

359 In conclusion, we present backwards- and forward- compatible ethnic- and sex specific
360 reference data for size-adjusted bone density in children and young adults, generated from
361 measurements in over 3500 individuals using GE and Hologic scanners. These data have
362 been produced using methods included in the most recent ISCD guidelines and for the first
363 time present curves for lumbar spine BMAD and prediction equations for TBLH-BMC taking
364 into account lean mass and body size, together with age-and gender- specific curves for

365 lumbar spine and TBLH aBMD. This reference database data has been specifically designed
366 to allow future updates and analysis when more definitive evidence for the best method of
367 fracture prediction in children is agreed.

368

369

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389

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391 collection: All authors and the ALPHABET study team. Data analysis: NJC Data
392 interpretation: NJC, KW Drafting manuscript: NJC, KW. Revising manuscript content: All
393 authors. Approving final version of manuscript: All authors; Data integrity of manuscript:
394 NJC

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491

492 **Figure legends**

493 **Figure 1** Comparison of GE Lunar iDXA™ lumbar spine BMAD LMS reference curves
494 between the three different ethnic groups. (A) BMAD (g/cm^3) for girls; (B) BMAD (g/cm^3)
495 for boys. Solid black line represents the mean for White Caucasian Children ($\pm 95\%$
496 Confidence interval -dotted black line). Dark grey dashed line represents the mean for Black
497 Afro-Caribbean Children; Dashed light grey line represents the mean for South Asian
498 Children.

499

500 **Figure 2** Comparison of lumbar spine BMAD LMS reference curves between males and
501 females (A) GE Lunar iDXA; (B) Hologic Discovery. Solid black line represents males
502 (mean $\pm 95\%$ Confidence interval). Dashed line represents females (mean $\pm 95\%$ Confidence
503 interval).

504

505 **Figure 3** Comparison of lumbar spine BMAD LMS reference curves between manufacturers,
506 GE Lunar iDXA™ compared to Transformed Hologic Discovery (Hologic data transformed
507 using cross calibration equations generated from the European Spine Phantom). (A) Females;
508 (B) Males. Solid black line represents GE Lunar iDXA™ (mean $\pm 95\%$ Confidence interval).
509 Dashed line represents Hologic Discovery (mean $\pm 95\%$ Confidence interval).

510

511 TABLES

512 **Table 1** Distribution of subjects used for the generation of reference data

GE Lunar Prodigy	2547	Male	1245	White Caucasian	925
				South Asian	192
				Black Afro Caribbean	128
		Female	1302	White Caucasian	970
				South Asian	184
				Black Afro Caribbean	148
GE Lunar iDXA (including transformed Prodigy)	2910	Male	1411	White Caucasian	1091
				South Asian	192
				Black Afro Caribbean	128
		Female	1499	White Caucasian	1167
				South Asian	184
				Black Afro Caribbean	148
Hologic Discovery	587	Male	325	White Caucasian	325
		Female	262	White Caucasian	262

513

514

515 **Table 2** Patient anthropometric data. Mean (SD)

Centre	Number	Mean (SD) Height Z-score	Mean (SD) Weight Z-score	Mean (SD) BMI Z-score
Birmingham	935	0.20 (1.09)	0.45 (1.24)	0.46 (1.25)
Middlesbrough	390	0.35 (0.97)	0.41 (0.96)	0.31 (1.00)
Leeds	171	0.34 (1.00)	0.42 (1.10)	0.31 (1.11)
Glasgow	212	0.15 (1.02)	0.34 (1.07)	0.36 (1.02)
London	372	0.11 (1.03)	0.29 (1.10)	0.27 (1.12)
Sheffield	830	0.40 (1.05)	0.59 (1.11)	0.51 (1.15)
Manchester	587	0.30 (0.96)	0.47 (1.01)	0.41 (1.03)
TOTAL	3497	0.28 (1.03)	0.46 (1.11)	0.42 (1.14)
Centre Differences (p value)		<0.001	0.001	0.003

516

517 Using a one-sided t-test all Z-scores were significantly ($p < 0.0001$) greater than zero. Centre
 518 differences were compared using ANOVA.

519

Table 3a Prediction Equations for Total body less head bone mineral content (TBLH-BMC (g)) for lean mass (g), fat mass (g), height (cm) and age (1decimal place) for the GE Lunar Prodigy™- Software version Encore 15.0.

		GE Prodigy	r ²	SEE
Girls	White Caucasian	$TBLH-BMC = 3.77 \times 10^{-4} \times LEAN^{0.845} \times FAT^{0.130} \times Height^{0.928} \times Age^{0.179}$	0.966	0.0988
	South Asian	$TBLH-BMC = 2.24 \times 10^{-4} \times LEAN^{0.603} \times FAT^{0.122} \times Height^{1.535} \times Age^{0.216}$	0.970	0.0935
	Black Afro-Caribbean	$TBLH-BMC = 1.02 \times 10^{-3} \times LEAN^{0.941} \times FAT^{0.100} \times Height^{0.543} \times Age^{0.311}$	0.967	0.1002
Boys	White Caucasian	$TBLH-BMC = 2.93 \times 10^{-4} \times LEAN^{0.939} \times FAT^{0.073} \times Height^{0.930} \times Age^{0.079}$	0.972	0.0976
	South Asian	$TBLH-BMC = 1.47 \times 10^{-4} \times LEAN^{0.978} \times FAT^{0.060} \times Height^{1.060}$	0.978	0.0932
	Black Afro-Caribbean	$TBLH-BMC = 1.94 \times 10^{-3} \times LEAN^{0.983} \times FAT^{0.048} \times Height^{1.018}$	0.973	0.0883

Total body less head BMC = TBLH-BMC; Total body lean mass = LEAN; Total body fat mass = FAT; SEE = Standard error estimate

Z-Score = (Measure Value – Predicted Value) / (Predicted Value x SEE)

Table 3b Prediction Equations for Total body less head bone mineral content (TBLH-BMC (g)) for lean mass (g), fat mass (g), height (cm) and age (1dp) for the GE Lunar Prodigy™ using the ENHANCED analysis mode - Software version Encore 15.0.

		GE Prodigy-Enhanced	r ²	SEE
Girls	White Caucasian	$TBLH-BMC = 4.24 \times 10^{-3} \times LEAN^{0.682} \times FAT^{0.079} \times Height^{0.905} \times Age^{0.122}$	0.967	0.0818
	South Asian	$TBLH-BMC = 6.04 \times 10^{-3} \times LEAN^{0.511} \times FAT^{0.106} \times Height^{1.110} \times Age^{0.185}$	0.937	0.0809
	Black Afro- Caribbean	$TBLH-BMC = 9.01 \times 10^{-3} \times LEAN^{0.744} \times FAT^{0.103} \times Height^{0.545} \times Age^{0.234}$	0.961	0.0910
Boys	White Caucasian	$TBLH-BMC = 1.47 \times 10^{-3} \times LEAN^{0.813} \times FAT^{0.055} \times Height^{0.949}$	0.974	0.0839
	South Asian	$TBLH-BMC = 5.06 \times 10^{-3} \times LEAN^{0.883} \times FAT^{0.044} \times Height^{0.586}$	0.979	0.0775
	Black Afro- Caribbean	$TBLH-BMC = 3.81 \times 10^{-3} \times LEAN^{0.856} \times FAT^{0.047} \times Height^{0.692}$	0.974	0.0735

Total body less head BMC = TBLH-BMC; Total body lean mass = LEAN; Total body fat mass = FAT; SEE = Standard error estimate

Z-Score = (Measure Value – Predicted Value) / (Predicted Value x SEE)

Table 3c Prediction Equations for Total body less head bone mineral content (TBLH-BMC (g)) for lean mass (g), fat mass (g), height (cm) and age (1dp) for the GE Lunar iDXA™ - Software version Encore 15.0.

		GE Lunar iDXA	r ²	SEE
Girls	White Caucasian	$TBLH-BMC = 1.85 \times 10^{-3} \times LEAN^{0.736} \times FAT^{0.077} \times Height^{0.950} \times Age^{0.135}$	0.965	0.0843
	South Asian	$TBLH-BMC = 2.58 \times 10^{-3} \times LEAN^{0.538} \times FAT^{0.110} \times Height^{1.210} \times Age^{0.192}$	0.967	0.0836
	Black Afro- Caribbean	$TBLH-BMC = 4.27 \times 10^{-3} \times LEAN^{0.787} \times FAT^{0.105} \times Height^{0.594} \times Age^{0.239}$	0.962	0.0931
Boys	White Caucasian	$TBLH-BMC = 5.88 \times 10^{-4} \times LEAN^{0.827} \times FAT^{0.055} \times Height^{1.095}$	0.974	0.0849
	South Asian	$TBLH-BMC = 2.01 \times 10^{-3} \times LEAN^{0.906} \times FAT^{0.047} \times Height^{0.708}$	0.980	0.0798
	Black Afro- Caribbean	$TBLH-BMC = 1.78 \times 10^{-3} \times LEAN^{0.887} \times FAT^{0.051} \times Height^{0.765}$	0.975	0.0754

Total body less head BMC = TBLH-BMC; Total body lean mass = LEAN; Total body fat mass = FAT; SEE = Standard error estimate

Z-Score = (Measure Value – Predicted Value) / (Predicted Value x SEE)

Table 3d Prediction Equations for Total body less head bone mineral content (TBLH-BMC (g)) for lean mass (g), fat mass (g), height (cm) and age (1dp) for the Hologic Discovery – Software version Apex 4.1.

		Hologic Discovery	r ²	SEE
Girls	White Caucasian	$TBLH-BMC = 1.20 \times 10^{-2} \times LEAN^{0.704} \times Height^{0.717} \times Age^{0.235}$	0.954	0.0871
Boys	White Caucasian	$TBLH-BMC = 4.77 \times 10^{-3} \times LEAN^{1.041} \times FAT^{-0.046} \times Height^{0.398}$	0.960	0.0962

Total body less head BMC = TBLH-BMC; Total body lean mass = LEAN; Total body fat mass = FAT; SEE = Standard error estimate

Z-Score = (Measure Value – Predicted Value) / (Predicted Value x SEE)

