

The added value of biomarker analysis to the genesis of Plaggic Anthrosols; the identification of stable fillings used for the production of plaggic manure.

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Abstract.

Plaggic Anthrosols are the result of historical forms of land management in cultural landscapes on chemically poor sandy substrates. Application of plaggic manure was responsible for the development of the plaggic horizons of these agricultural soils. Pollen diagrams reflect aspects of the environmental development but the interpretation of the pollen spectra is complicated due to the mix of the aeolian pollen influx of crop species and species in the surroundings, and of pollen occurring in the used stable fillings. Pollen diagrams and radiocarbon dates of plaggic Anthrosols suggested a development period of more than a millennium. *Calluna* is present in almost all the pollen spectra, indicating the presence of heath in the landscape during the whole period of soil development. Optically stimulated luminescence dating of the plaggic horizon made clear that the deposition of plaggic covers started in the 16th century and accelerated in the 18th century. The stable fillings, used for the production of plaggic manure and responsible for the rise of the soil surface, cannot be identified with pollen diagrams alone. Biomarker analyses provide more evidence about the sources of stable fillings. The oldest biomarker spectra of the plaggic horizons of three typical plaggic Anthrosols examined in this study, were dominated by biomarkers of forests species as *Quercus* and *Betula* while the spectra of middle part of the plaggic horizons were dominated by biomarkers of stem tissue of crop species as *Secale* and *Avena*. Only the youngest spectra of the plaggic horizons were dominated by biomarkers of *Calluna*. This indicates that the use of heath sods as stable filling was most likely introduced very late in the development of the Anthrosols. Before the 19th century the mineral component in plaggic manure cannot be explained by the use of heath sods. We conclude that other sources of materials, containing mineral grains must have been responsible for the raise of the plaggic horizon.

Key words

Plaggic Anthrosols, plaggic manure, radiocarbon/luminescence dating, palynology, biomarkers, Netherlands.

1. Introduction.

Plaggic Anthrosols occur in cultural landscapes, developed on coversands. These chemical poor Late-glacial aeolian sand deposits dominate the surface geology of an extensive area in northwestern Europe. Plaggic Anthrosols are the characteristic soils that developed on ancient arable fields, fertilized with plaggic stable manure. Plaggic Anthrosols have a complex genesis and are valuable records of environmental and agricultural history (van Mourik et al., 2011).

In previous palaeopedological studies of such soil records in The Netherlands (van Mourik et al, 2011, 2012, 2013a), information was unlocked by application of pollen analysis, radiocarbon (¹⁴C) and Optically Stimulated Luminescence (OSL) dating. Radiocarbon dates of soil organic carbon, extracted from humic horizons from plaggic Anthrosols, suggested the start of sedentary agriculture between 3000 and 2000 BP but are not indicative for the age of the plaggic sediments due to the complexity of soil organic carbon in plaggic sediments (Mook & Streurman, 1983; van Mourik et al., 1995). It was assumed that farmers used organic sods as stable filling, firstly dug on forest soils and later on heaths for the production of stable manure to fertilize the fields. The mineral fraction of the sods was supposed to be responsible for the development of the plaggic horizon and the raise of the land surface. OSL dating applied on quartz grains extracted from plaggic sediments provides more reliable ages of the plaggic sediments. The OSL dates suggested that the rise of the plaggic horizons started in the 16th century and accelerated in the 18th century (Bokhorst et. al., 2005). This is rather well in line with historical data, as presented by Spek, (2004, p 965).

The use of ectorganic matter from forest soils in the Dutch coversand area, must have been strongly reduced in the 11th-13th century, due to commercial forest clear cuttings as recorded in archived documents (Vera, 2011).

58 These deforestations resulted in a regional extension of sand drifting and the managers of the heaths had to
59 protect their valuable ecotopes against this ‘historical environmental catastrophe’ (Vera, 2011).
60 Heaths were already present in the Late Paleolithic landscape (Doorenbosch, 2013) and played a ceremonial
61 role in the society of our ancestors. People already had the knowledge to manage the heath as sustainable
62 grazing areas for cattle (Doorenbosch, 2013).
63 The use of heath for sheep grazing and other purposes as honey and oil production could continue until the
64 middle of the 18th century (Vera, 2011). In SE-Netherlands sustainable use of the heaths was promoted by
65 many management rules and laws (van Mourik, 1978; Veera 2011). Over the course of the 18th century, the
66 population growth resulted in an increasing food demand. In the course of the 18th century, the deep stable
67 economy was introduced and the booming demand for manure resulted in intensification of manure production.
68 Farmers started with the use of heath sods as (additional) stable filling (Spek, 2004). This caused heath
69 degradation and initiated the second extension of sand drifting. The use of sods finished at the end of the 19th
70 century after the introduction of chemical fertilizers (Spek, 2004).
71 Through the combination of OSL and ¹⁴C dating, historical records and the conventional paleoecological proxy
72 of fossil pollen analysis we have a good impression of the paleoecological environment and the age of such
73 deposits. However, it remains problematic to reconstruct the combination of crop residues and various
74 materials used by farmers as stable filling to produce the stable manure, together responsible for the rise of
75 the surface of Anthrosols. This is also hindering a detailed interpretation of the agricultural practices and shifts
76 therein related to the plaggic agriculture system, and specifically the timing of the onset of the intensive heath
77 sod driven deep stable agriculture with which plaggic Anthrosols are most commonly associated. To address
78 this issue, in the present study we expanded our paleoecological toolset with an adapted application of the
79 recently developed biomarker approach (Jansen et al., 2010). This biomarker approach consists of a
80 combination of analytical chemical analysis and modelling with the VERHIB model to unravel concentration
81 patterns of higher chain length (C₂₀-C₃₆) *n*-alkanes of higher plant origin preserved in a soil or sedimentary
82 archive into the (groups of) species responsible for their production (Jansen et al., 2010). The approach was
83 originally developed to unravel past local vegetation composition. Upon successful application in a tropical
84 ecosystem setting (Jansen et al., 2013), its applicability in palaeopedology was explored (van Mourik & Jansen,
85 2013b). This pilot application concerned a polycyclic soil sequence in driftsand deposits. It showed that the
86 comparison of pollen and biomarker spectra allowed us to indicate the plant species responsible for carbon
87 sequestration in the humic horizons (van Mourik & Jansen, 2013b). Important conclusion was that biomarker
88 analysis showed promise not only in the reconstruction of past local vegetation composition of a specific site,
89 but also in studies where the emphasis lies not on the vegetation per se, but rather on reconstructing various
90 sources of soil organic matter input (van Mourik & Jansen, 2013b).
91 Goal of the present study was to further explore the applicability of biomarker analysis as part of a multi-proxy
92 reconstruction aimed at unraveling the sources of stable fillings used for the production of plaggic manure in
93 the context of the historic development of the plaggic agriculture ecosystem. For this, we applied biomarker
94 analysis on three previous investigated plaggic Anthrosol.

95 **Materials and methods.**

96 **2. Profile selection**

100 Fig. 1

101 Fig. 2 a,b,c

102 The distribution area of plaggic Anthrosol in NW-Europe is indicated in fig.1. Pape (1972) published the first
103 map of the distribution of plaggic agriculture in NW-Europe. Bastiaens & van Mourik (1995) found traces of
104 intensification and extension of this area in Vlaanderen (Belgium) while van Mourik (1999b) also reported
105 plaggic Anthrosols in Schleswig (Germany). Beside this area with ‘real’ plaggic Anthrosols, Spek (2004, p. 724)
106 summarized information about the occurrence of soils with some evidence of application of plaggic manure in
107 the Atlantic coastal zones of Norway, Denmark, France, Galicia, Scotland and Ireland.

108 For this pilot study we selected three previously investigated plaggic Anthrosols in the Netherlands with an
109 undisturbed plaggic horizon: Valenakker, Nabbegat and Posteles (fig.2). Pollen diagrams, ¹⁴C and OSL dates of
110 these profiles were available and previously published separately in various articles. Here we combined these,
111 and re-sampled the plaggic horizons of the profiles for biomarker analysis and new fossil pollen analysis to
112 allow for comparison. Vertical sampling resolution was: Valenakker 20 cm, Nabbegat and Posteles 10 cm.

113 Valenakker (van Mourik et al. 2012) is situated southwest of the city Weert (middle Limburg) on the sport fields
114 of a former college. As a result, during the 20th century the soil has never been ploughed or subjected to land
115 consolidation. This profile has never been affected by roots of *Zea mays*, introduced in The Netherlands in the
116 middle of the 20th century (van Mourik & Horsten, 1995).

117 Nabbegat (van Mourik et al. 2013a) is situated on the Maashorst (eastern North-Brabant). The plaggic deposits
118 were buried by drift sand around 1800 AD. Consequently, the plaggic deposits have perfectly been protected
119 against damage by land consolidation or pollution afterwards (van Mourik et al. 2013a). The site is now
120 vegetated by oak and birch trees. Roots of these trees may have caused input of organic matter by
121 decomposed roots in the upper part of the plaggic horizon (fig.3).

122
123 Fig. 3

124
125 Posteles (van Mourik et al., 2011) is situated in Twente (eastern Overijssel). The landowner informed us that
126 during the last three generations this land was never subjected to deep ploughing or land consolidation but
127 since 1960 *Zea mays* was regular sowed. In contrast to Valenakker and Nabbegat we can expect biomarkers of
128 this deep rooting cultivated plant.

129 130 131 **2.1. Pollen analysis.**

132
133 Pollen diagrams of plaggic Anthrosols provide paleoecological information about plant species, present on site
134 and in the region during the formation of the plaggic horizon. Previous research showed that pollen grains,
135 infiltrated in soils and incorporated in plaggic deposits, are well preserved in the anaerobic and acid
136 microenvironment of excremental aggregates (van Mourik, 1999a, 2001) (fig 4,5).

137
138 Fig. 4

139
140 Fig. 5

141
142 Samples for pollen extraction were collected in 10 ml tubes in profile pits. For a correct matching of pollen and
143 biomarker spectra of the plaggic deposits, the same samples were treated for both pollen and biomarker
144 extraction and analysis. The pollen extractions were carried out using the tufa extraction method (Moore et al.,
145 1991, p. 50). For the identification of pollen grains, the pollen key of Moore et al. (1991, p. 83-166) was applied.
146 Pollen scores were based on the total pollen sum of arboreal and non-arboreal plant species. For the
147 estimation of the pollen concentrations of the various soil horizons, the exotic marker grain method was
148 applied (Moore et al., 1991, p. 53).

149 150 151 **2.2. ¹⁴C and OSL dating.**

152
153 The determination of the age of plaggic deposits is subjected to various complications (Spek, 2004). Pollen
154 stratification is disturbed by bioturbation and ploughing. Besides, the pollen content is a mix of the regular
155 pollen influx and pollen in stable fillings, used for the production of stable manure (van Mourik et al., 2011).
156 The ages of humic horizons of buried Podzols cannot be correctly determined by ¹⁴C dating due to the complex
157 composition of soil organic carbon (van Mourik et al., 1995). During a period of active soil formation, hard
158 decomposable organic carbon can accumulate in the Ah horizon, especially in the humin fraction but also in the
159 humic acid fraction. Especially the accumulation of charcoal fragments in the organic aggregates is responsible
160 for overestimation of the ¹⁴C ages (fig.4). During the Early Holocene small amounts of charcoal fragments were
161 released after (natural) forest fires, but the amount increased drastic in the iron time due to the charcoal
162 production for the melting of iron from placic horizons and iron stone (Beukenkamp and Sevink, 2005). The age
163 of the humic acid fraction was considered the best estimate of the moment of fossilization of the Ah horizon
164 after burying by driftsand. The difference between humin and humic acids ages was interpreted as a measure
165 for the period of soil activity and humin accumulation. Later, OSL dating confirmed that radiocarbon dates, not
166 only of the humin fraction but also of the humic acids, overestimate the true ages (Bokhorts et al., 2005).
167 Conventional radiocarbon dating of humin and humic acids showed in presented diagrams, extracted from
168 plaggic deposits, was performed in the CIO (Centre for Isotope Research of the University of Groningen).

169 OSL dates provide reliable information about the moment of fossilization of plaggic material under the rising
170 furrow because the quartz grain were perfectly bleached during active ploughing (Bokhorts et al., 2005). OSL
171 dating of quartz grains, extracted from plaggic deposits, was performed in the NCL (Netherland Centre for
172 Luminesce Dating, Wageningen University).

173
174

175 2.3. Biomarker analysis.

176
177

178 2.3.1. The application of the VERHIB model

179 A detailed description of the biomarker approach using the VERHIB method is presented in our previous
180 publications (Jansen et al., 2010; Jansen et al., 2013; Van Mourik & Jansen 2013). Briefly, the basis of the
181 method lies in the unraveling of the preserved concentration patterns of C₂₀-C₃₆ *n*-alkanes, which are exclusive
182 to the epicuticular wax layers on leaves and roots of higher plants (Kolattukudy et al., 1976). While such an
183 application in itself is not new (e.g. Pancost et al., 2002; Hughen et al., 2004) the novelty of our approach lies in
184 the application of the VERHIB model that we specifically developed to unravel the mixed *n*-alkane signal
185 encountered in soil or sedimentary archives (Jansen et al., 2010). The VERHIB model consists of a linear
186 regression model that describes how a certain input of plant derived compounds such as *n*-alkanes over time in
187 a certain archive at a certain location, results in accumulation of these compounds. An inversion of the forward
188 model is used to reconstruct the accumulation encountered with depth into its most likely vegetation origin
189 (Jansen et al., 2010). An important aspect of biomarker analysis using VERHIB is that it is an indirect
190 reconstruction. While the biomarker patterns, in the present study the *n*-alkanes, are directly measured, the
191 reconstruction into the most likely combination of vegetation biomass input responsible for the observed
192 pattern is inferred by the model. For this, several parameters must be inputted into the model (Jansen et al.,
193 2010) the most important of which is the selection of the expected plant species that have been responsible for
194 the input of biomass in the archive in question, and subsequent inclusion of their *n*-alkane signature in the
195 VERHIB reference base. In the present study, the selection of species to include was based on the (expected)
196 crop history of the sites under study, as well as the anticipated origin of the stable fillings used as manure. An
197 important matter of debate when using *n*-alkane patterns to reconstruct past vegetation input is the genotypic
198 plasticity of the *n*-alkane patterns, in particular in relation to prevailing environmental factors such as climate
199 (e.g. Shepherd and Griffiths, 2006). In a previous study focusing on vegetation of relevance for reconstructions
200 in ecosystems in North-Western Europe where plaggic agriculture occurred, we found that while genotypic
201 plasticity related to climatic factors may influence the signal, such influence does not eradicate the different
202 vegetation origins (Kirkels et al., 2013). To limit external influences as much as possible, the vegetation selected
203 for inclusion in the VERHIB reference base was sampled in close vicinity to the three study sites as much as
204 possible. The first group of selected plant species concerned the main sources of stable fillings, used for the
205 manure production: fermented litter from deciduous forest soils (*Quercus robur*, *Betula pendula*), grass sods
206 from brook valleys (*Molinia caerulea*) and heath sods (*Calluna vulgaris*).

207 The second group concerned crop species. Close to the educational Field Study Centre Orvelte (Drenthe) is a
208 traditional plaggic field where they continued with the cultivation of traditional crop species. There we sampled
209 *Fagopyrum esculentum*, *Spergula arvensis*, *Avena sativa*, *Secale cereal*, *Spergula arvensis*. The modern crop
210 species *Zea mays* corn was sampled on the Posteles.

211 The concentration patterns of the *n*-alkanes with carbon numbers 20-36 in the selected vegetation samples
212 and in the soil samples were subsequently used as input for the VERHIB model (see 2.3.2 for a description of
213 the extraction and analysis of the biomarkers).

214 A second parameter that must be considered in the application of VERHIB, is input of leaf and root material.
215 VERHIB considers the species specific *n*-alkane patterns in plant roots separately from the patterns in plant
216 leaves, and uses this to deal with the input of young root material at depth (Jansen et al., 2010). A first
217 selection criterion here concerns whether or not leaf and root material can be expected to have entered the
218 soil at all. For the deciduous forest soil material potentially used as stable fillings (*Quercus robur*, *Betula
219 pendula*), exclusively leaf derived biomass input is expected as the trees did not grow on-site. In contrast, for
220 the crop species *Zea Mays* and *Spergula Arvensis* only root material is expected to have entered the soil in
221 appreciable amounts as the leaf material is mostly removed during harvest. For the other species considered,
222 both leaf and root material must be taken into account. A selection of root and/or leaf derived *n*-alkane
223 patterns to be considered in the VERHIB reference base was made in accordance with the previous. With
224 respect to the ratio of input of leaf vs. root biomass as required by the model, no exact information is available
225 for the soil profile under study. Therefore, for those species where both leaf and root material is considered to
226 have possibly entered the soil, in line with the exploratory nature of the present study, we applied an assumed

226 leaf/root biomass input ratio of 1.0 and assumed that while input of leaf material always occurred at the top of
227 the soil profile, root input also occurred with depth. Since our pilot study in polycyclic driftsand deposits
228 showed that VERHIB was unable to filter out root input sufficiently (Van Mourik & Jansen, 2013), when
229 interpreting the occurrence of a certain species with depth in the profiles under study as modelled by VERHIB,
230 the possibility of young root input being responsible for the signal was explicitly taken into account.
231 Figure 6 shows a flow diagram that illustrates the functioning of the VERHIB modelling as well as the selection
232 of parameters and reference base species as described above.
233

234 Fig. 6

236 2.3.2. Extraction and analysis of the biomarkers.

237 Approximately 0.1 g of each of the freeze-dried and ground vegetation and soil samples was extracted by
238 accelerated Solvent Extraction (ASE) using a Dionex 200 ASE extractor. The extraction temperature was 75°C
239 and the extraction pressure 17×10^6 Pa, employing a heating phase of 5 min and a static extraction time of 20
240 min. Dichloromethane/methanol (DCM/MeOH) (93:7 v/v) was used as the extractant (Jansen et al., 2006). The
241 extracts were subsequently fractionated into three fractions containing the *n*-alkanes, the esters and the
242 combination of alcohols and fatty acids respectively. For this, a silica column consisting of extracted cotton
243 wool and silica gel was used, followed by elution with hexane, hexane/DCM (4:1) and DCM/Methanol (9:1)
244 respectively. Separation of the *n*-alkanes took place by on-column injection of 1.0 µl of the first fraction on a 30
245 m Rtx-5Sil MS column (Restek) with an internal diameter of 0.25 mm and film thickness of 0.1 µm, using He as a
246 carrier gas. Temperature programming was: 50°C (hold 2 min); 40°C/min to 80°C (hold 2 min); 20°C/min to
247 130°C; 4°C/min to 350°C (hold 10 min). Subsequent MS detection in full scan mode used a mass-to-charge
248 ratio (*m/z*) of 50-650 with a cycle time of 0.65 s and followed electron impact ionization (70 eV). The *n*-alkanes
249 were identified from the total ion current (TIC) signal by their mass spectra (dominant fragment ion
250 represented by *m/z* = 57) and retention times and quantified using a deuterated internal standard (*d*₄₂-*n*-C₂₀
251 alkane (Jansen et al. 2010) as well as a conventional external *n*-alkane standard.
252

253 Fig.7

254
255 Figure 7 presents the *n*-alkane biomarker distribution in the leaves and/or roots of the species, inserted in the
256 reference base. The results show the odd-over-even chain-length predominance typical of higher plants
257 (Kolattukudy et al., 1976). The observed variation in patterns and concentrations is in line with the variation
258 found in other species in previous work (e.g. Jansen et al., 2006).
259

260 3. The vertical distribution of biomarkers and pollen in the analysed profiles.

261 3.1. Profile Valenakker

262
263 Profile Valenakker is a plaggic Anthrosol (Aan), overlying a ploughed umbric Podzol (2ABp, 2Bs). The pollen
264 diagram (fig.3) and the absolute dates (table 1) reflects a soil development of ≈ 1400 year.
265 The post sedimentary pollen spectra in the 2BS show percentages of tree species as *Corylus* and *Quercus* of the
266 Middle Subatlantic. The presence of Poaceae, Cyperaceae, *Rumex* and Ranunculaceae reflects a period of
267 pasture. The high scores of Cerealia in ploughed 2ABp and even the 2B indicate a form of sedentary agriculture
268 before the start of plaggic agriculture. ¹⁴C dating indicate a carbon age of the base (60 cm) of the Aan horizon
269 of ≈ 600 AD. The OSL age of the lower part of the plaggic horizon is 800-900 year younger, ≈ 1560 AD.
270

271 Fig.8.

272 Table 1.

273 Fig.9.

274 Micromorphological observations (fig.5ab) of the plaggic deposits show the complexity of soil organic matter.
275 There are various sources of organic carbon as plants roots, tissue of table fillings and sods. Also the

276 composition of pollen spectra is complex, a mix of the regular pollen influx of plants on the fields and in the
277 surrounding infiltrating into the soil and pollen, and pollen present in various stable fillings.
278 In previous studies the origin of stable fillings, used in plaggic agriculture, was reconstructed on the base of
279 pollen diagrams (Spek, 2004; van Mourik et al., 2012a, 2012b). The pollen spectra of the Aan horizon show very
280 low scores of arboreal trees but reasonable scores of Ericaceae and Poaceae. Ericaceae pollen may indicate the
281 use of heath sods, Poaceae pollen the use of grassland sods, the combination of sods from degrading heath and
282 the rise of the land surface by plaggic manure is caused by the mineral fraction in such sods. However, the rise
283 of the plaggic horizon of ≈ 60 cm cannot be explained by the use of heath sods if it is true that the use of heath
284 sods (with a mineral fraction) was introduced in the course of the 18th century when better construction
285 materials enabled the farmers to build deep stables (Vera, 2011). In fact, the sources of stable fillings cannot be
286 satisfactorily detected with pollen diagrams.
287 The biomarker spectrum of the base is dominated by *Quercus*. Despite the low percentages *Quercus* pollen it is
288 very likely that the farmers used forest litter as stable filling. The middle spectrum is dominated by markers of
289 *Avena* and *Secale*. This points to the use of straw from these crop species as stable filling. Pollen of *Cerealia* is
290 present in the whole diagram. In the upper spectrum biomarkers of *Calluna* are present together with *Avena*
291 and *Secale*. This points to the use of heath sods as additional stable filling during the last phase in the
292 development of the plaggic horizon.
293

294 3.2. Profile Nabbegat

295

Fig. 10.

296

Table 2.

297

Fig. 11.

298

299 Profile Nabbegat is a haplic Arenosol (with Mormoder humus form), overlying a plaggic Anthrosol, overlying a
300 ploughed umbric Podzol. The pollen diagram (fig.6.) and the absolute dates (table 2) reflect a soil development
301 of ≈ 3000 year.

302 The post sedimentary pollen spectra of the 3ABp reflect the start of agriculture (increase of *Cerealia*) on a
303 former heath (decrease of *Ericaceae*) in a surrounding with coppice hedges (*Quercus*, *Corylus*). Based on
304 radiocarbon dates, the agricultural activities started before ≈ 1000 BC, the OSL dates point to deposition of
305 plaggic material after ≈ 1500 AD.

306 The radiocarbon ages indicate that the farmers used organic matter with very few mineral 'contamination' for
307 a long time. The OSL ages indicate that the rise of the plaggic horizon started ≈ 1500 AD due to mineral grains
308 as part of the manure. The plaggic horizon developed between 1500 and 1800 AD. Around 1800 AD, short after
309 the introduction of the deep stable economy (Vera, 2011), the plaggic Anthrosol was overblown by driftsand.
310 Apparently, the use of heath sods resulted in heath degradation, sand drifting and acceleration of the rise of
311 the plaggic horizon (van Mourik et al., 2012a). The sand drifting stabilized under planted *Quercus* trees; the
312 roots of these trees reached the buried Anthrosol and may have contributed the scores of biomarkers in the
313 upperpart of the buried plaggic horizon. The composition of the pollen spectra of the plaggic horizon is rather
314 uniform, dominated by Ericaceae and *Cerealia*.

315 Fig.7. shows the results of biomarker analysis. Biomarkers of *Quercus* were present in all the spectra, dominant
316 in the lower spectra, regular in the other spectra. This points to the use of forest litter as stable filling during
317 the development of the lower part of the plaggic horizon. The main crop species during this time was *Spergula*.
318 The middle part is dominated by markers of *Avena* and *Secale*, indicating the use of straw. Only in the upper
319 spectrum *Calluna* was found, indicating the use of heath sods during the last phase of the development of the
320 plaggic horizon.

321

322

3.3. Profile Posteles

Fig.12.

Table 3.

Fig.13.

Profile Posteles is a plaggic Anthrosol, overlying a ploughed umbric Podzol. The pollen diagram (fig.8) and the absolute dates (table 3) reflect a soil development of at least 1200 year.

The pollen content of the buried ploughed Podzol (2Ap, 2B) is post-sedimentary infiltrated in Late-Glacial coversand by bioturbation and agriculture. Characteristic is the sharp decrease of pollen concentrations with depth, shown by the pollen density curve. The spectra of the 2B horizon already reflect evidence of agriculture (Cerealia) in a deforested landscape (low percentages of *Alnus*, *Quercus*, *Fagus*). The spectra of the 2Ap horizon show increasing percentages of Cerealia.

The radiocarbon age of the base of the plaggic deposits (95 cm) is ≈ 850 AD, The OSL age ≈ 1500 AD. The OSL age of the 2Ap (105 cm) is 2035 ± 450 BC, ≈ 3500 year older than sample 95. In this part of the profile we see the effect of bioturbation on the age of the coversand. Grains from the base of the Aan were transported to the 2Ap and reversed, which explains the large standard deviation of the OSL of sample 105 cm.

The actual Ap horizon (the active plough horizon) is palynologically characterized by peak percentages of Cerealia, a slight extension of *Pinus* (planted on the abandoned heath after 1900 AD) and the appearance of *Zea mays* (introduced in Dutch agriculture after 1950 AD). Pollen of Cerealia, Ericaceae and Poaceae were found in all the spectra of the Aan.

The lowest spectrum (80) is dominated by the crop species *Spergula* and the score of *Quercus* indicates the use of forest litter during the development of this part of the Aan.

The spectra 10, 20, 40, 60 are dominated by biomarkers from roots of *Zea mays*. This crop species was introduced around 1950 AD, but the markers of the decomposed *Zea* roots seem to suppress all the others (this was not the case in the profiles Valenakker and Nabbegat). Spectrum 50 is dominated by *Avena* and *Secale*, spectrum 30 by *Zea* and *Secale* and spectrum 0 by *Zea* and *Calluna*. Again the use of heath sods seems restricted to the youngest part of the plaggic horizon.

4. Discussion

Pollen diagrams of plaggic Anthrosols provide valuable paleoecological information to reconstruct the soil dynamics during the plaggic agriculture. However, interpretation of pollen diagrams is complicated. Pollen grains, extracted from plaggic deposits, may originate from two sources (van Mourik et al., 2011). The first source concerns the regional pollen influx from flowering species and local flowering crop species. Pollen grains precipitate on the soil surface and may infiltrate into the Anthrosols by ploughing and bioturbation. This pollen influx will be mixed with the pollen content of materials, used as stable fillings to produce manure.

Pollen will be preserved in plaggic deposits in the anaerobic and acid micro environment of humic aggregates, produced by worms and micro arthropods (van Mourik, 1999b, 2001). In general it is not possible to make a clear separation between pollen grains originating from the regular pollen influx or from materials as sods. Therefore, the identification of the various sources of stall fillings cannot be based on pollen analysis alone. Additional information, acquired by biomarker analysis proved very useful for this purpose.

In the pollen diagrams *Fagopyrum* is found in almost all the spectra of the plaggic deposits and in Valenakker and Nabbegat even in the top spectra of the buried ploughed Podzol, probably as result of pollen infiltration. *Fagopyrum* as crop species on sandy soils was introduced after 1350 AD (Leenders, 1996). Based on this palynological time marker, plaggic deposition started around 1350 AD.

The radiocarbon ages of plaggic deposits are much older. This is caused by (1) older organic carbon, present in the applied stable fillings (as forest litter) for the manure production and (2) accumulation of hardly decomposable organic carbon during active soil formation. Consequently the radiocarbon dates overestimate the ages of the plaggic sediments, but approach the age of the introduction of agricultural soil management (van Mourik et al., 1995, 2011, 2012a, 2012b). Manuring of infertile soils came already in use in the Bronze Age

374 and also the Celtic fields are an example of a prehistorical agricultural system based on manure management
375 ([Spek, 2004](#)).

376 The mineral component of stable manure, applied on the fields, was responsible for the thickening of the
377 plaggic horizon. Ploughing of the furrow will bleach the OSL signal of the mineral grains until the moment that
378 the grains are no longer part of the active soil furrow. For that reason, OSL dating of the plaggic horizon provide
379 reliable ages of the plaggic deposits ([Bockhorst et al., 2005](#)). The OSL dates of the profiles Valenakker,
380 Nabbegat and Posteles indicate a start of the thickening \approx 1550 AD.

381 It was not possible to determine the sources of stable fillings palynologically. Possible stable fillings were forest
382 litter, sods from moist grass lands and heath sods. But in almost all spectra of the pollen diagrams Ericaceae,
383 Poaceae and arboreal pollen occur. Biomarkers extracted from plaggic deposits, originate from two sources.
384 The first source concerns biomarkers from decomposed roots of crop species, the second source of organic
385 material as straw and sods, used as stable filling for manure production.

386 In the three diagrams we find *Quercus* as dominant marker in the lowest part of the Aan-horizon, indicating the
387 use of forest litter. In Nabbegat, *Quercus* markers can also originate from roots of the planted *Quercus* forest
388 after the stabilization of the sand drifting. This is not the case on Valenakker and Posteles. The middle part of
389 the Aan-horizon is dominated by markers of *Avena* and *Secale*, indicating the use of straw as stable filling.

390 Only in the top of the Aan-horizon markers of *Calluna* are present, indicating the use of heath sods as stable
391 filling. Based on the results of the biomarker analysis we can conclude that heath sods were used as stable
392 filling only in the 18th and 19th century. This fits with the observations about the use of heaths in historical
393 archives [Vera \(2011\)](#).

394 So the question rises about heath management before the introduction of the deep stable economy. Some
395 researchers point to careful heath management before the 19th century. In interviews with farmers, born
396 before 1950, [Burny \(1999\)](#) collected essential information about historical heaths management in the Belgian
397 Kempen. A historical study of land use in the Campina also indicated carefully maintenance and sustainable use
398 of valuable common fields ([de Keyzer, 2014](#)). Before the 19th century, heath sods were never dug on the dry
399 *Calluna* heath, only on the moist *Erica* heath. These organic sods were not used as stable filling but as fuel for
400 the furnace. Burning of *Calluna* heaths was the most important management action to rejuvenate the heath.
401 Juvenile heath is food for cows. Sods digging was a bad action due to the resistance and incoherence of these
402 dry sods and also the long recovery period. Mowing of older *Calluna* shrubs took place. Twigs were used for
403 roofs, burning and brooms. ([Burny, 1999](#)). Because of the very low nutrient contribution to the manure of
404 mowed *Calluna*, the farmers preferred the use of twigs of broom (*Genista*). When in the course of the 18th the
405 authority relationships changed and the population growth and the demand for food increased, farmers
406 started to intensify their production ([Vera, 2011](#)). They needed more manure and started with the deep stable
407 economy and the use of *Calluna* heath sods.

408 An important factor may be the presence of pollen and biomarkers content in sheep droppings. According to
409 [Simpson et al., \(1999\)](#) biomarkers survive the congestion process and stay in the manure. But what do sheep
410 consume? Grazing sheep are very selective in collecting food ([Oom et al., 2008](#); [Smits & Noordijk, 2013](#)). They
411 prefer grasses (*Molinia*, *Festuca* and *Corynephorous*). Only in years that there is insufficient grass available at
412 the end of the summer, they eat shoots of *Calluna*, at that time nourishing with high concentrations Ca, Mg and
413 but no P. Pollen extractions from sheep droppings showed that only in droppings, collected during the summer
414 season *Calluna* pollen is present. During the flowering season of *Calluna*, the animals consume pollen,
415 precipitated on the grasses. That explains the presence of *Calluna* pollen and the absence of *Calluna*
416 biomarkers in the lower parts of the plaggic horizons.

417 If it is true that *Calluna* heath sods were dug only in the 18th and 19th century, how can we explain the mineral
418 component in the plaggic manure, responsible of the rise of the land surface before that time?

419 According to [Smits and Noordijk \(2013\)](#) there are several sources of minerals. Firstly, a small amount of mineral
420 grains will be incorporated in the manure during emptying out the manure of the stable. Secondly, farmers had
421 the knowledge that the addition of sand could improve the fertility of the soil. Not the leached and acid sand
422 from heath sods but not leached sand, dug on sheep walks and in blown out depressions in nearby drift sand
423 landscapes.

424

425

426 5. Conclusions

427

428 • The vertical zoning of biomarkers and pollen in plaggic horizons are different. Palynologically, the plaggic
429 horizon is a homogenous, the biomarker diagrams show differentiation.

430 • We can identify various stable fillings used, based on the vertical distribution of biomarkers.

- 431 • The biomarker spectra of the base layer of the plaggic horizon are dominated by biomarkers of deciduous
 432 trees litter (dominated by *Quercus*), indicating the use of organic matter from the forest floor.
 433 • The biomarker spectra of the middle part of the plaggic deposits are dominated by crop species (*Avena*,
 434 *Secale*), indicating the use of straw from these species as stable filling during a relatively long time.
 435 • Only the top spectra of the plaggic horizons are dominated by *Calluna*, indicating that heath sods were
 436 used as stable filling only during the last phase in the development of the plaggic horizon.
 437 • Profile Posteles shows the impact of the contribution of biomarkers of roots of *Zea mays*, introduced
 438 around 1950 AD, suppressing the other species.
 439 • The negligible percentages of *Calluna* in biomarker spectra of plaggic deposits with exception of the top,
 440 suggest an overestimating of the use of heath sods in the traditional interpretation of the genesis of
 441 plaggic horizons, the dominance of crop species in biomarker spectra of plaggic deposits suggests
 442 underestimating of the use of straw as source material for the production of organic stable manure to
 443 fertilize ancient arable fields.
 444

445
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 448 agriculture. We are grateful to Annemarie Philip (IBED, University of Amsterdam) for the preparation of the
 449 pollen slides, Hans van der Plicht (CIO, University Groningen) for production of the radiocarbon dates and Jakob
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 451 produced by Jan van Arkel (IBED, University of Amsterdam).
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453
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575 **Tables (including table captions)**

576

Table 1. ¹⁴ C and OSL dates of the plaggic deposits of Valenakker.				
Horizon	Depth (cm)	Calendric ¹⁴ C ages humin	Calendric ¹⁴ C ages humic acids	Calendric OSL ages
Aan	20	–	–	1775 ± 20 AD
Aan	40	771 ± 92 AD	1049 ± 78 AD	1635 ± 30 AD
Aan	60	595 ± 61 AD	698 ± 54 AD	1565 ± 30 AD

577

Table 2. ¹⁴ C and OSL dates of the plaggic deposits of Nabbegat.				
Horizon	Depth (cm)	Calendric ¹⁴ C ages humin	Calendric ¹⁴ C ages humic acids	Calendric OSL ages
C	70	–	–	1803 ± 12 AD
2An	80	428 ± 107 AD	626 ± 45 AD	1770 ± 11 AD
2An	105	37 ± 133 BC	3 ± 101 AD	–
2An	130	1182 ± 139 BC	811 ± 101 BC	1676 ± 14AD
3ABp	140	–	1299 ± 78 BC	–
3ABp	150	–	1385 ± 72 BC	–

578

Table 3. ¹⁴ C and OSL dates of the plaggic deposits of Posteles.				
Horizon	Depth cm	Calendric ¹⁴ C ages humin	Calendric ¹⁴ C ages Humic acids	Calendric OSL ages
Aan	45	–	–	1758 ± 14 AD
Aan	59	–	–	1711 ± 20 AD
Aan	70	1132 ± 68 AD	1172 ± 51 AD	1651 ± 31 AD
Aan	82	–	–	1626 ± 20 AD
Aan	95	884 ± 82 AD	861 ± 85 AD	1517 ± 31 AD
2ABp	105	–	–	2035 ± 450 BC

579

580

581 **Figure Captions**

582

583 Fig. 1. The location of sampled profiles Valenakker, Nabbegat and Posteles in the distribution area of plaggic
584 agriculture.

585

586 Fig. 2. The plaggic Anthrosols Valenakker, Nabbegat and Posteles. The location of the OSL samples are indicated
587 in the white circles (depth in cm); the locations of the profiles are indicated in fig. 1.

588

589 Fig. 3. Cross-section of a (living) tree root in the thin section of the 2 Aan of Nabbegat (70-80cm). Characteristic
590 is the double fringing of the root tissue. Such roots were only found in the upper part of the 2Aan of Nabbegat.
591 Roots of crop species were not found in the thin sections of the three profiles; they decompose rather fast
592 compared with tree roots.

593

594 Fig. 4. Distribution pattern of organic aggregates in a thin section of the Aan of Valenakker (40-50 cm). In the
595 fabric of the aggregates are charcoal particles visible

596

597 Fig. 5. Pollen grains, visible in a welded aggregate of the same thin sections. Pollen grains in thin sections are
598 observable as not double fringing, empty spheroidal objects. The palynological characteristics as sculpture and
599 aperture are not visible without the chemical treatments during pollen extraction.

600

601 Fig. 6. Flow diagram of the methodology of biomarker analysis.

602

603 Fig. 7. The *n*-alkane biomarker distribution in leaves and/or roots of species sampled, for the reference base of
604 this pilot study.

605

606 Fig. 8. Pollen diagram Valenakker. Pollen density in k.grain/ml.

607

608 Fig. 9. Biomarker diagram Valenakker.

609

610 Fig. 10. Pollen diagram Nabbegat. Log D = pollen density in log k.grain/ml.

611

612 Fig. 11. Biomarker diagram Nabbegat.

613

614 Fig. 12. Pollen diagram Posteles; Pollen density in k.grain/ml.

615

616 Fig. 13. Biomarker diagram Posteles.

617

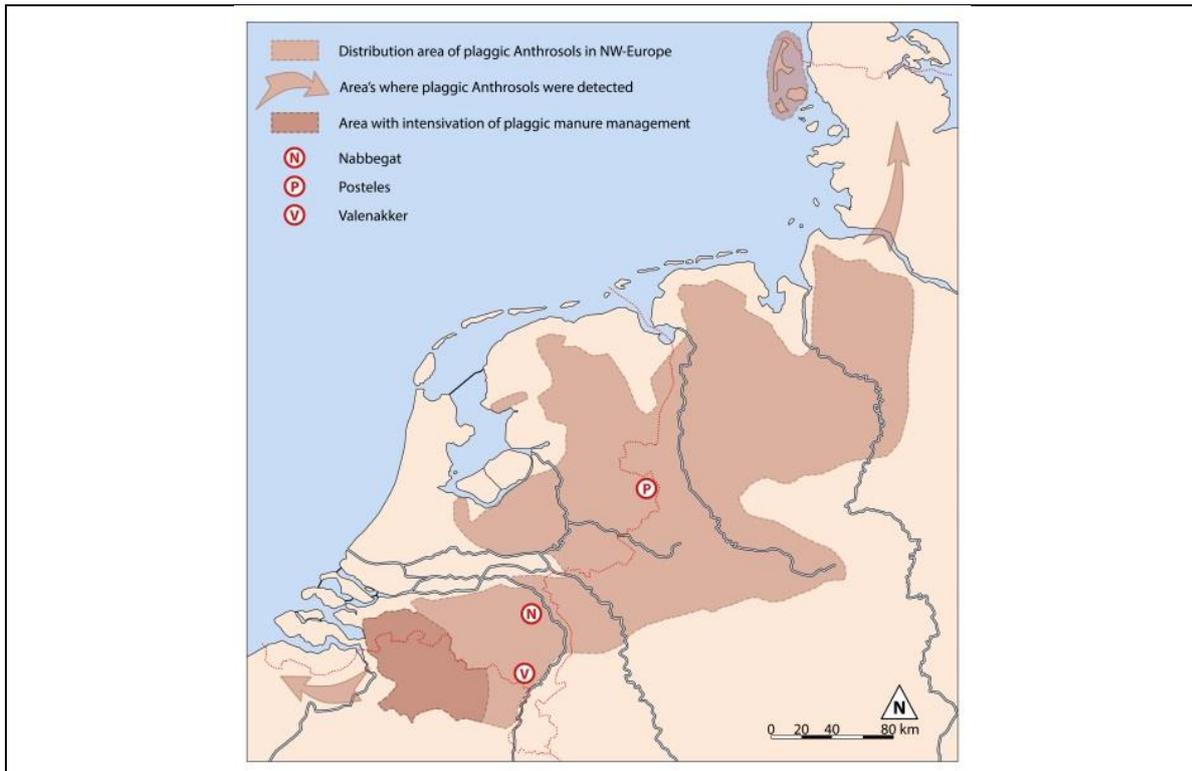


Fig. 1. The location of sampled profiles Valenakker, Nabbegat and Posteles in the distribution area of plaggic agriculture.

619

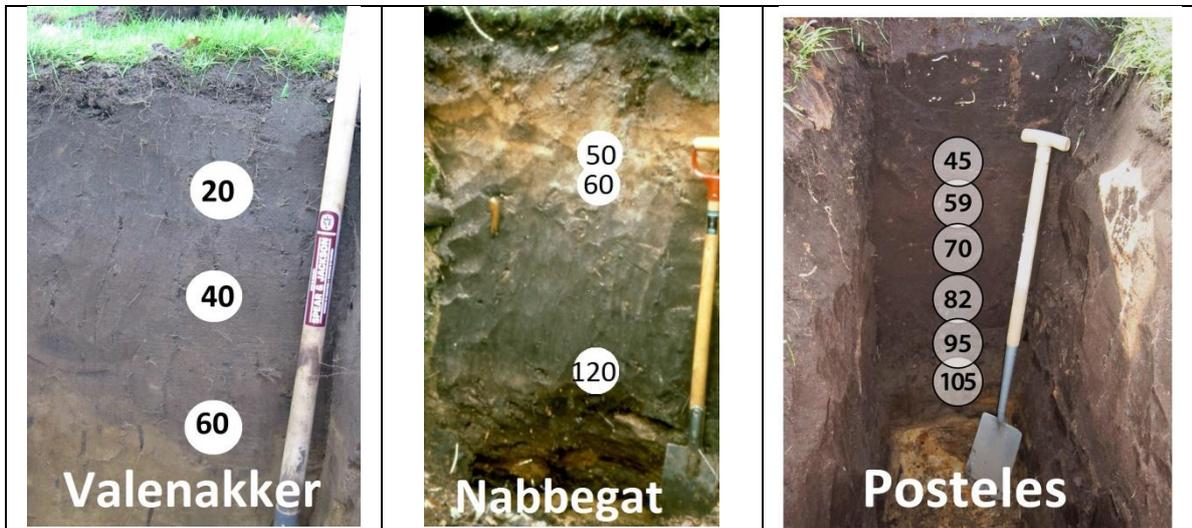


Fig. 2. The plaggic Anthrosols Valenakker, Nabbegat and Posteles. The location of the OSL samples are indicated in the white circles (depth in cm); the locations of the profiles are indicated in fig. 1.

620



Fig. 3. Cross-section of a tree root in a thin section of the 2 An of Nabbegat (70-80cm). Characteristic is the double fringing of the root tissue. Such roots were only found in the upper part of the 2An of Nabbegat. Roots of crop species were not found in thin sections of the three profiles. The decompose rate of such roots is fast.

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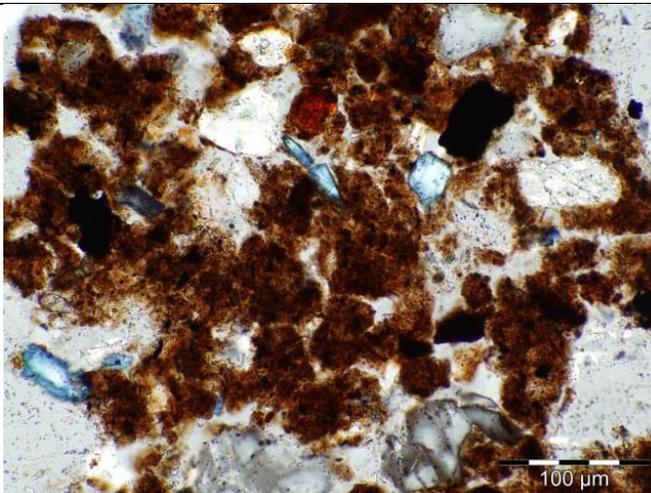


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622

623

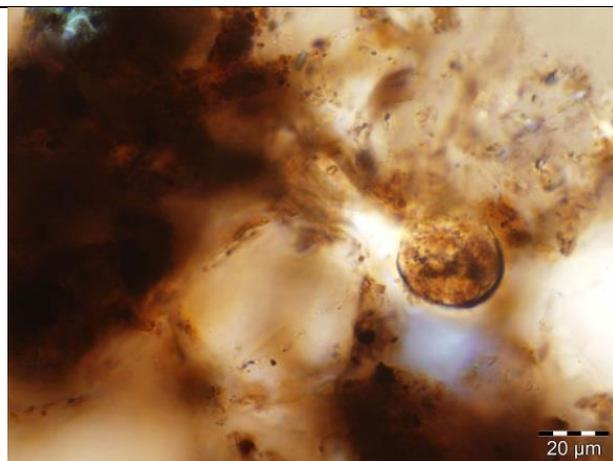


Fig. 5. Pollen grains, visible in an aged aggregate of the same thin sections. Pollen grains are in thin sections observable as not double fringing and empty spheroidal objects. The palynological characteristics as sculpture and aperture are not visible without the chemical treatments during pollen extraction.

624

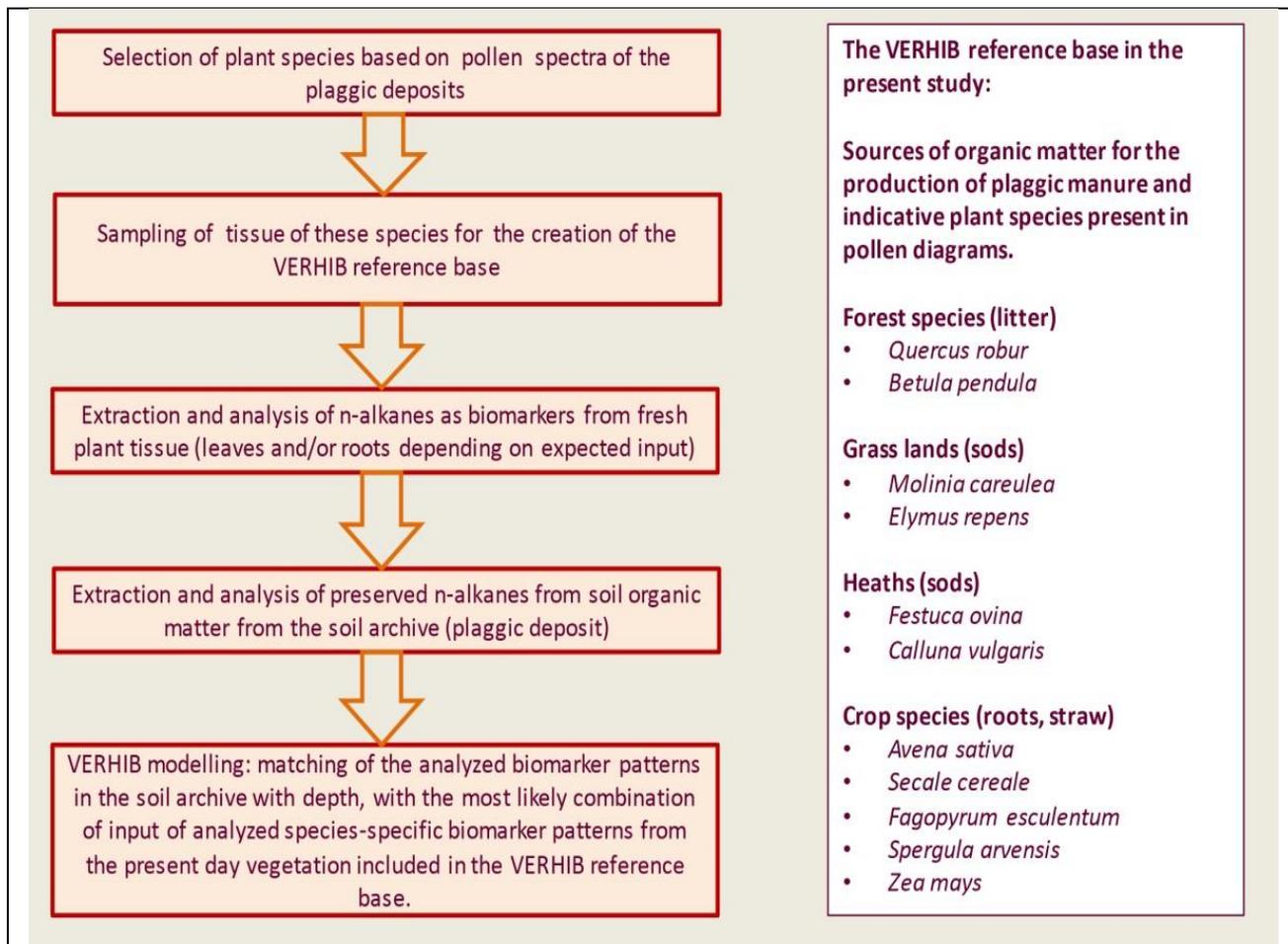


Fig. 6. Flow diagram of the methodology of biomarker analysis.

625

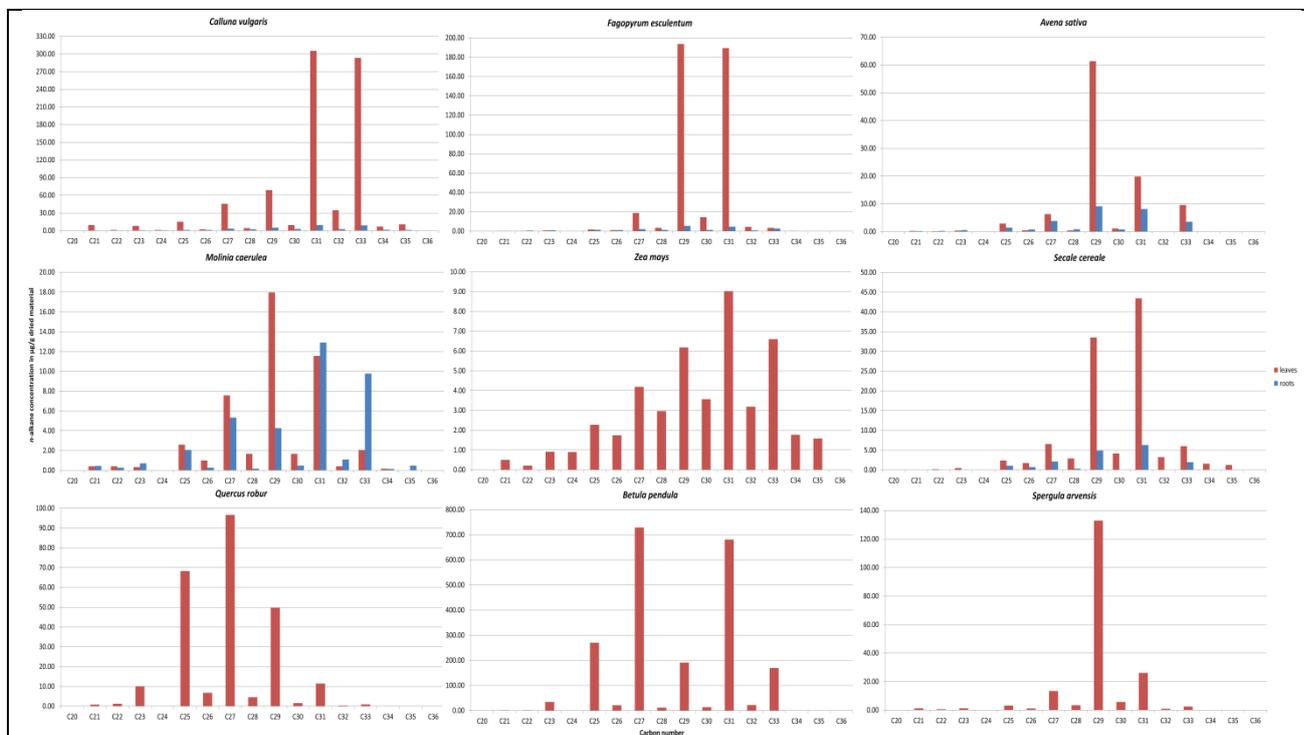


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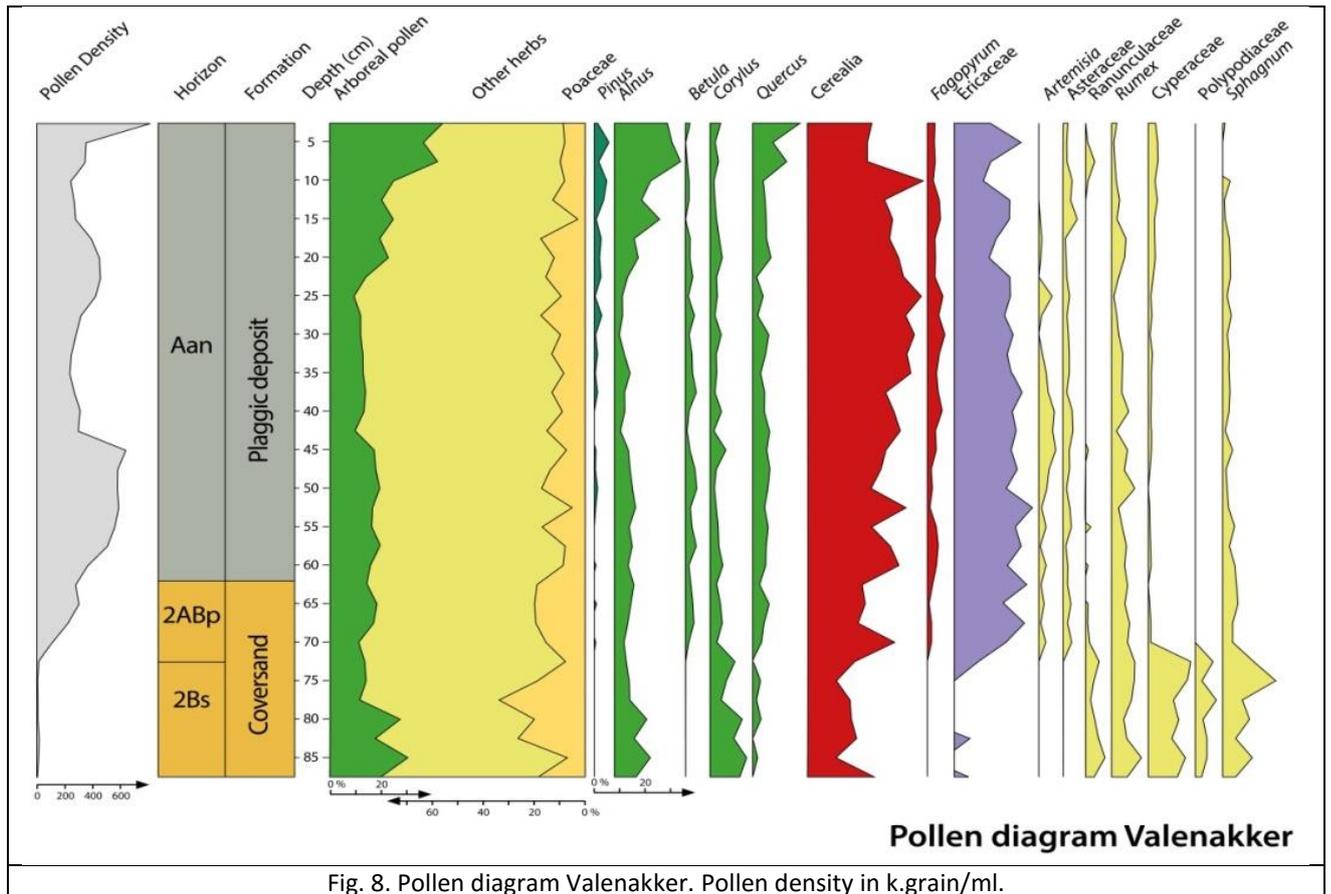


Fig. 8. Pollen diagram Valenakker. Pollen density in k.grain/ml.

627

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Horizon	Depth (cm)	Calendric ¹⁴ C ages humin	Calendric ¹⁴ C ages humic acids	Calendric OSL ages
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Aan	40	771 ± 92 AD	1049 ± 78 AD	1635 ± 30 AD
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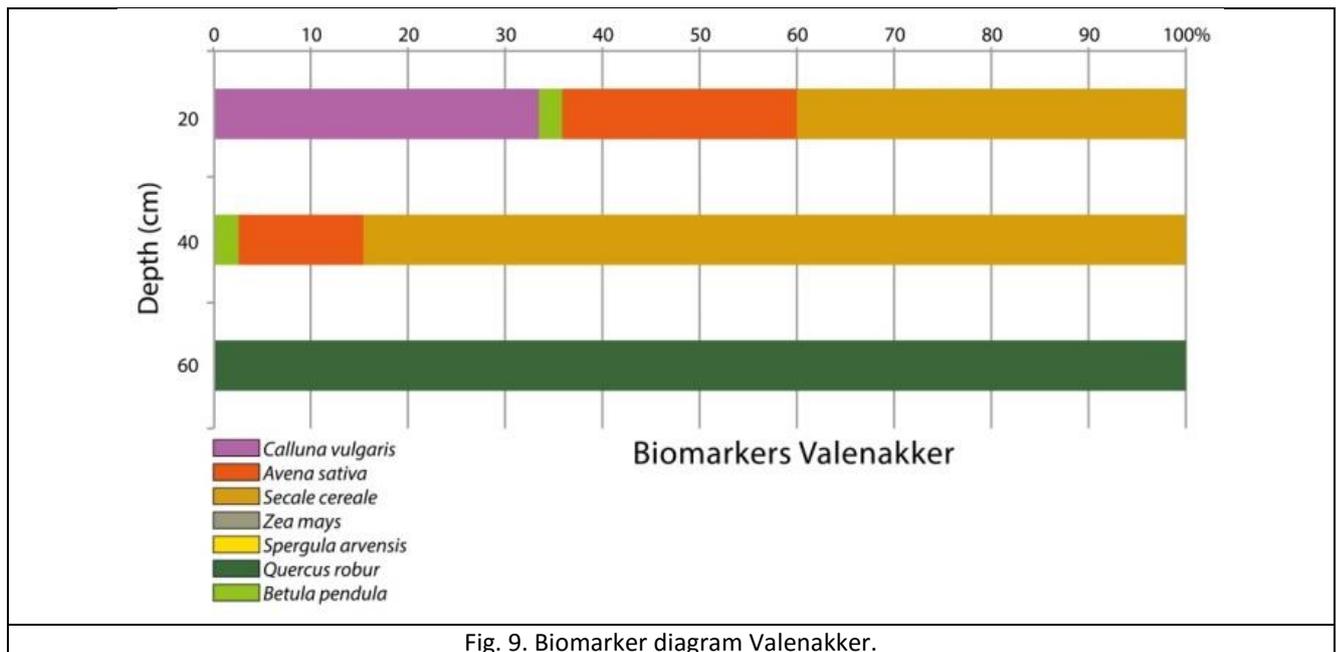


Fig. 9. Biomarker diagram Valenakker.

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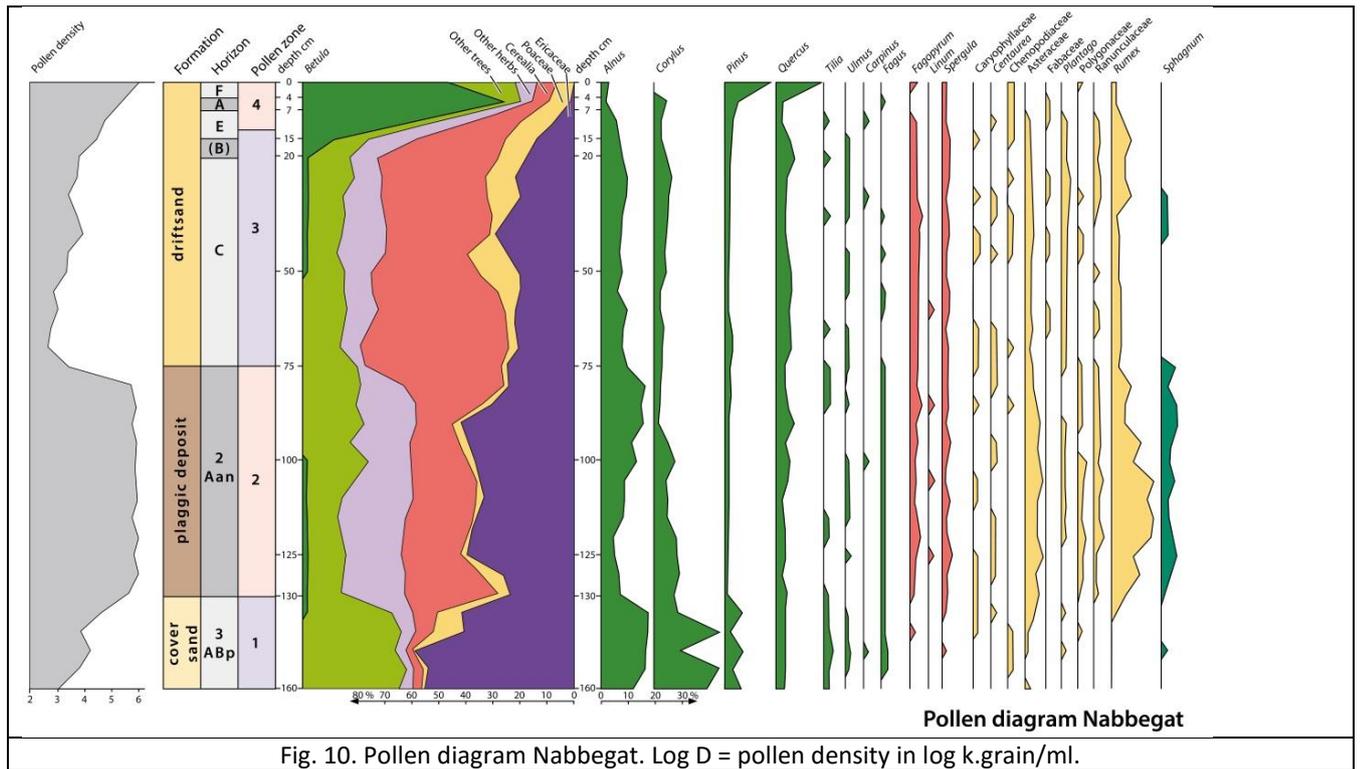


Fig. 10. Pollen diagram Nabbegat. Log D = pollen density in log k.grain/ml.

630

Table 2. ¹⁴C and OSL dates of the plaggic deposits of Nabbegat.

Horizon	Depth (cm)	Calendric ¹⁴ C ages humin	Calendric ¹⁴ C ages humic acids	Calendric OSL ages
C	70	—	—	1803 ± 12 AD
2An	80	428 ± 107 AD	626 ± 45 AD	1770 ± 11 AD
2An	105	37 ± 133 BC	3 ± 101 AD	—
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3ABp	150	—	1385 ± 72 BC	—

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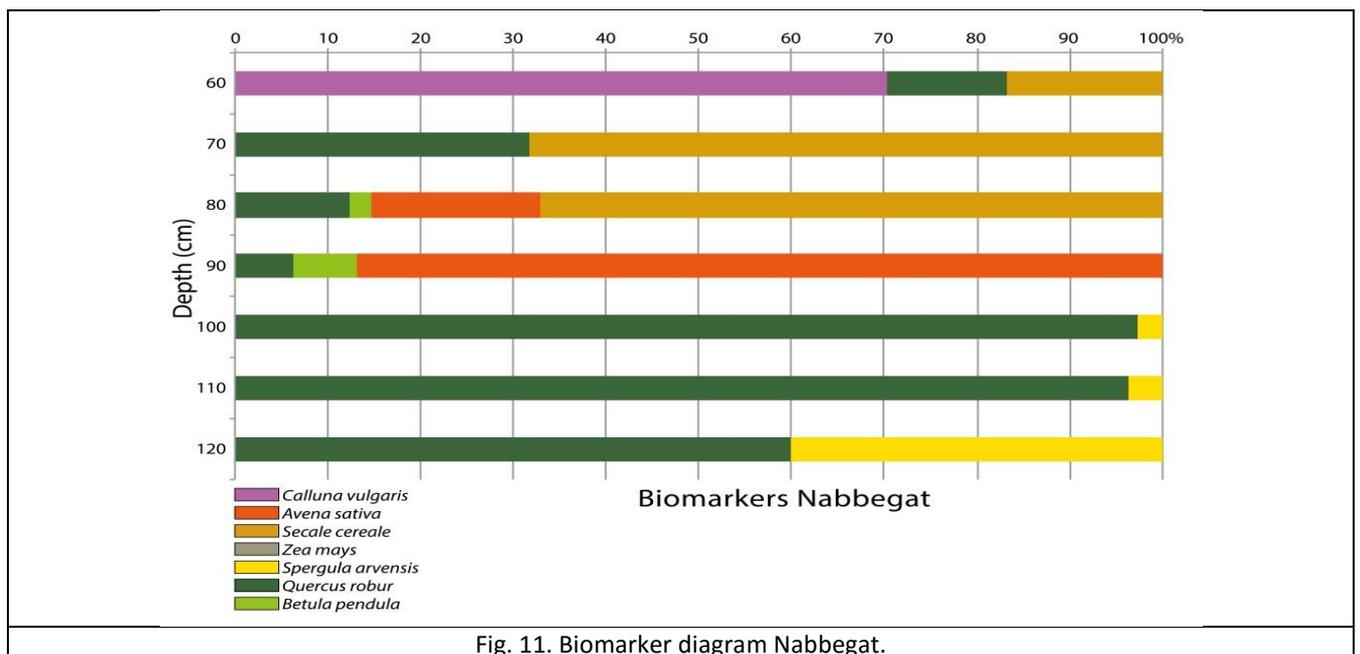
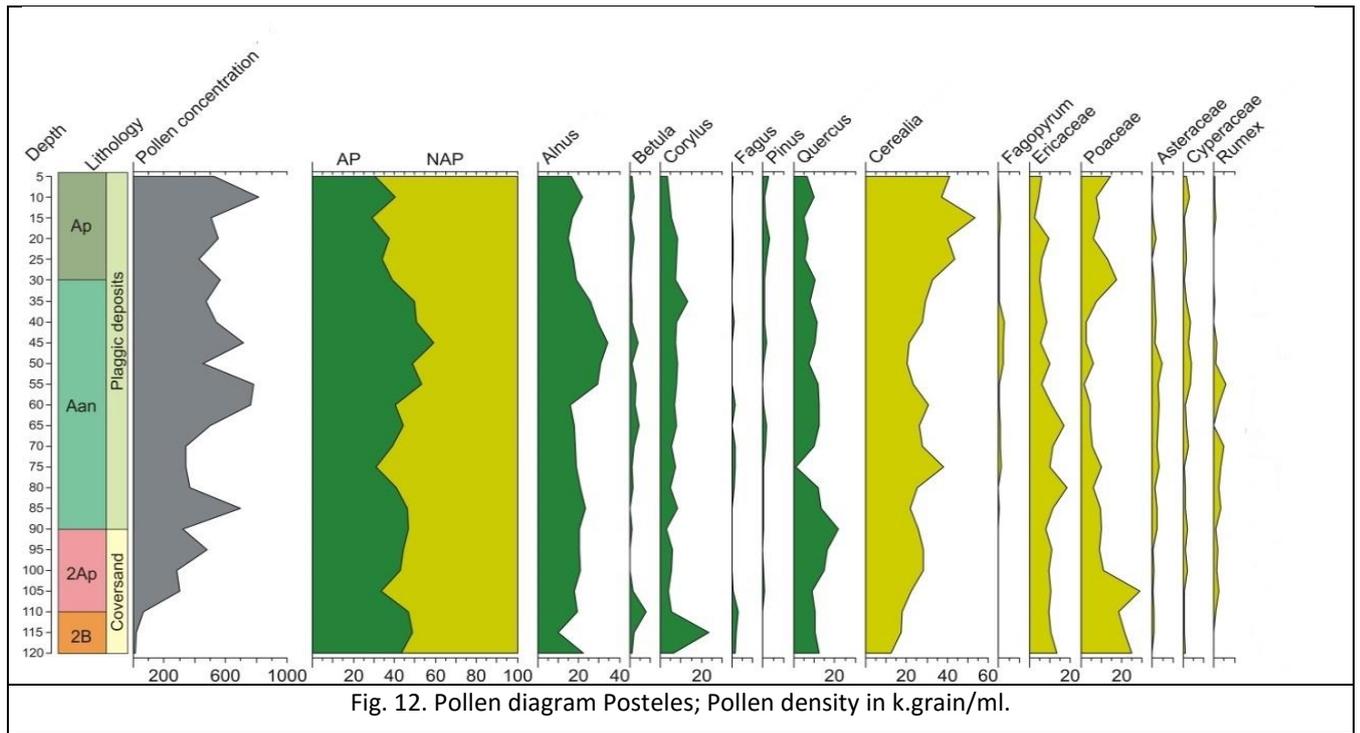


Fig. 11. Biomarker diagram Nabbegat.



632

Table 3. ¹⁴C and OSL dates of the plaggic deposits of Posteles.

Horizon	Depth cm	Calendric ¹⁴ C ages humin	Calendric ¹⁴ C ages Humic acids	Calendric OSL ages
Aa	45	–	–	1758 ± 14 AD
Aa	59	–	–	1711 ± 20 AD
Aa	70	1132 ± 68 AD	1172 ± 51 AD	1651 ± 31 AD
Aa	82	–	–	1626 ± 20 AD
Aa	95	884 ± 82 AD	861 ± 85 AD	1517 ± 31 AD
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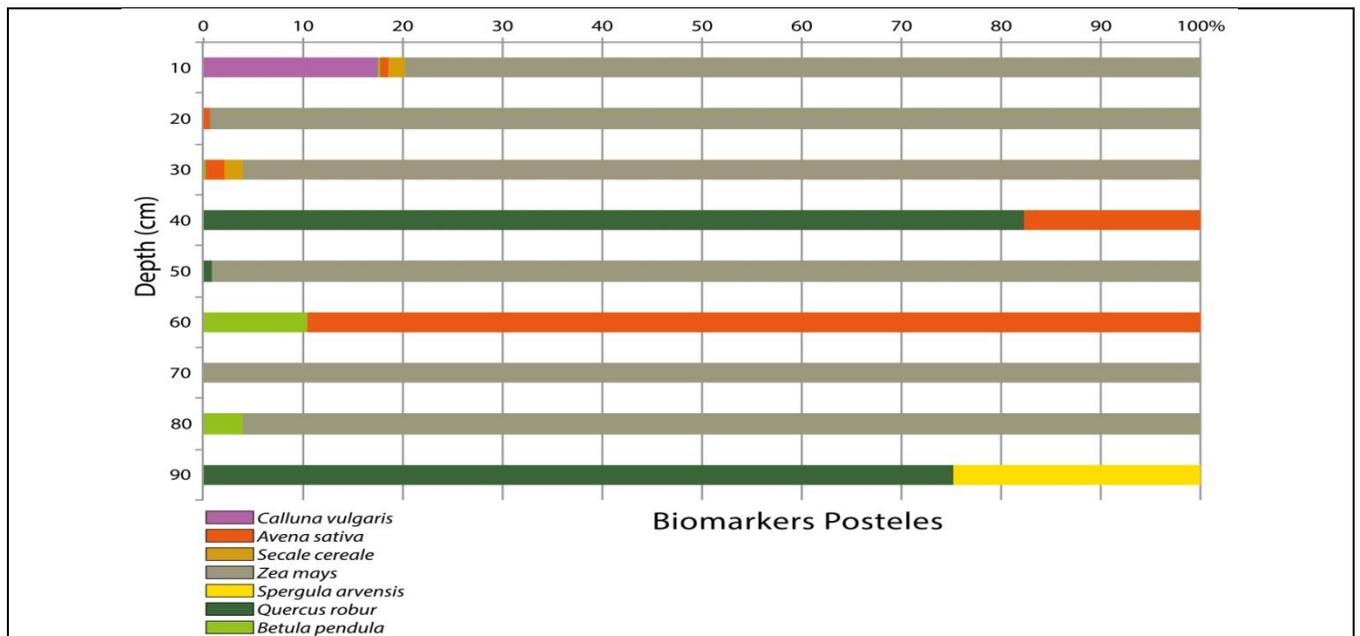


Fig. 13. Biomarker diagram Posteles.

634 **Figures 1 – 13 (including figure captions).**
635
636