

FORAMINIFERAL CHARACTERISATION OF MID-UPPER JURASSIC SEQUENCES IN THE WESSEX BASIN (UNITED KINGDOM)

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Received September 19, 2002; accepted October 6, 2003

Key words: Jurassic, foraminifera, sequence stratigraphy, Dorset Coast, United Kingdom.

Abstract. The use of foraminifera in the characterisation of sequences (systems tracts, maximum flooding surfaces, etc.) has developed over the last decade. Much of this work has been based in the Cenozoic successions of the Gulf of Mexico, although there is a growing application of such data in the Middle East and the North Sea Basin. The easiest surface to characterise has been the maximum flooding surface with its high diversity and high(er) abundance faunas; the characterisation of individual systems tracts has been less successful.

Using the well-known mid-Upper Jurassic successions of the Dorset coastal sections, we have investigated a number of high resolution (para)sequences for their foraminiferal content. Using data of foraminiferal diversity and standing crops from a range of modern substrates we have investigated the potential faunas available after deposition, taphonomy, compaction, groundwater dissolution and modern weathering. By understanding the processes involved we have identified the key foraminiferal features of typical mid-Upper Jurassic sequences and indicated how this work may help in the correlation of successions in North Dorset and Normandy.

Riassunto. L'uso dei foraminiferi nella caratterizzazione delle sequenze (system tracts, superfici di massimo allagamento, ecc.) si è sviluppato durante l'ultimo decennio. La maggior parte di questo lavoro è basata sulle successioni cenozoiche del Golfo del Messico, sebbene ci sia una crescente applicazione di tali dati nei bacini del Medio Oriente e del Mare del Nord. La superficie più facile da caratterizzare è stata la superficie di massimo allagamento, con alta diversità e alte abbondanze faunistiche; la caratterizzazione dei singoli systems tracts ha avuto meno successo.

Usando le ben note successioni costiere del Dorset della parte centrale del Giurassico superiore si è indagato il contenuto in foraminiferi di un certo numero di (para)sequenze ad alta risoluzione. Usando i dati sulla diversità e produttività fissa dei foraminiferi da una gamma di substrati moderni, si sono investigate le faune potenzialmente disponibili dopo la deposizione, la tafonomia, la compattazione, la dissoluzione delle acque circolanti e l'alterazione meteorica moderna. Capendo i processi coinvolti, è stato possibile identificare le chiavi caratteristiche dei forami-

niferi delle tipiche sequenze della parte centrale del Giurassico superiore, indicando come questo lavoro possa essere d'aiuto nella correlazione delle successioni nel Dorset settentrionale ed in Normandia.

Introduction

Sequence stratigraphy developed in the 1970s and was based, initially, on the interpretation of seismic profiles. Vail et al. (1977) expanded these initial concepts to include both borehole and outcrop data and closely related the resulting stratigraphy to sea level changes. Short-term fluctuations in sea level were shown to generate "sequences" or "genetically related strata bounded by unconformities or their correlative conformities" (Van Wagoner et al. 1988, p. 39). Sequences soon developed a nomenclature, with the three main subdivisions being represented (in ascending order) by the Lowstand Systems Tract (LST), Transgressive Systems Tract (TST) and Highstand Systems Tract (HST). The Maximum Flooding Surface (MFS) is an important part of sequence stratigraphic interpretation and separates the TST and the HST. In some successions (see below) the 'surface' cannot be located precisely and we have adopted the concept of a Zone of Maximum Flooding following the work of Montanez & Osleger (1993), Strasser et al. (1994, 1999) and Oliver (1998). This "Exxon Model" of a sequence must not be confused with the "Galloway Model", which uses the MFS as they key to sequence identification (Galloway 1989a,b; Reading 1996, pp. 25-26, fig.2.10), and the "Einsele Model" which is based on the recognition of transgressive/regressive cycles (Einsele & Bayer 1991).

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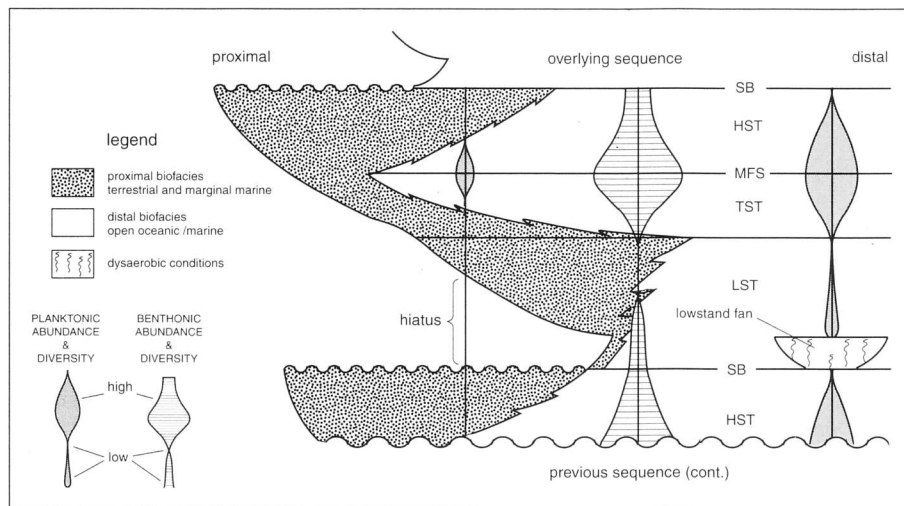


Fig. 1 - The "Exxon model" of a sequence (based on Emery & Myers 1996, fig. 6.14a) and the hypothetical distribution of benthonic and planktonic foraminifera. The planktonic foraminifera are most abundant (and diverse) in the region of the MFS with deeper-water morphotypes being found in more distal regions while only surface-water morphotypes are to be found in more proximal environments. The abundance (and diversity) of benthonic foraminifera mirrors that of the planktonic fauna, although benthonic taxa are found in a much wider range of proximal environments. SB = Sequence Boundary; LST = Lowstand Systems Tract; TST = Transgressive Systems Tract; MFS = Maximum Flooding Surface; HST = Highstand Systems Tract.

The "Exxon Model" is used by Emery & Myers (1996) in their description of how palaeontology and biostratigraphy contribute to sequence stratigraphy (Fig. 1). Emery & Myers (*op. cit.*, pp. 101, 104, fig. 6.14) describe how the MFS can be identified by the abundance and diversity of benthonic foraminifera. It also coincides with the most landward distribution of diverse, open marine, plankton. This approach has been used effectively in younger rocks, most notably in the Cenozoic of the Gulf of Mexico (Shaffer 1987, 1990; Armentrout 1987, 1991; Armentrout & Clement 1990; Armentrout *et al.* 1990, 1999; Pacht *et al.* 1990; Vail & Wornardt 1990; Van der Zwan & Brugman 1999). More recently, a number of authors have extended this approach to the interpretation of Mesozoic successions (Olsson 1988; Cubaynes *et al.* 1990; Simmons *et al.* 1991; Powell 1992; Hart 1997, 2000; Henderson 1997; Henderson & Hart 2000; Oxford *et al.* 2000; Sharland *et al.* 2001) with varying degrees of success. In their work on the Jurassic to Lower Cretaceous successions of the North Sea Basin, Partington *et al.* (1993) used the distinctive assemblages associated with the maximum flooding events to generate a biostratigraphical framework for correlation. One of the problems associated with more ancient successions is the loss of diagnostic taxa or even complete faunas as a result of taphonomy, burial and other diagenetic processes. In this account we report on our research into the preservation and identification of microfaunas within Jurassic sequences.

Mid-Upper Jurassic of the Dorset Coast

The dominantly siliciclastic succession of the Dorset Coast has been the subject of a number of investigations many of which have directly, or indirectly, related

the sedimentary succession to changes in sea level (Arkell 1933, 1956; Wilson 1968a,b; Talbot 1973, 1974; Brookfield 1973a,b, 1978; Fürsich 1975; Sun 1989, 1990a,b; Rioult *et al.* 1991; Coe 1992, 1995; Oliver 1998; Newell 2000; Henderson & Hart 2000; Oxford *et al.* 2000).

In many of these interpretations (Fig. 2) of the sequence stratigraphy there are different, if not conflicting, conclusions especially between Wilson (1991), Rioult *et al.* (1991), Coe (1992, 1995), Oliver (1998) and Newell (2000). In attempting to resolve some of these differences, and to explore the use of foraminifera in the recognition of Jurassic sequences, Oxford *et al.* (2000) used a part of the succession for a pilot investigation. The Nothe Grit Formation and the Redcliff Formation are well exposed on the Dorset Coast between Weymouth and Ringstead (Fig. 3). In this succession 96 samples were collected and prepared using the solvent method of Brasier (1980). All samples were washed on a 63 μm sieve, dried gently and stored ready for investigation. A minimum of 300 foraminifera were picked from the 500-250 μm size fraction with counts of the other size fractions (250-125 μm and 125-63 μm) also being recorded for further analysis (not presented here). Fig. 4 records the results of this preliminary investigation with the percentage of dry sediment retained on the 63 μm sieve plotted alongside the generic diversity recorded in each sample.

The most clay-rich samples are located in the Nothe Clay Member (samples between O1 and O6) and it is these same samples that record the highest levels of generic (and specific) diversity. This would appear to be the "zone of maximum flooding" recorded by Oliver (1998) and would clearly show up as a gamma-ray peak on a wireline log. Unfortunately planktonic foraminifera are not recorded at this level in the UK Jurassic succession and one cannot fully test all the features of the Emery & Myers (1996, fig.

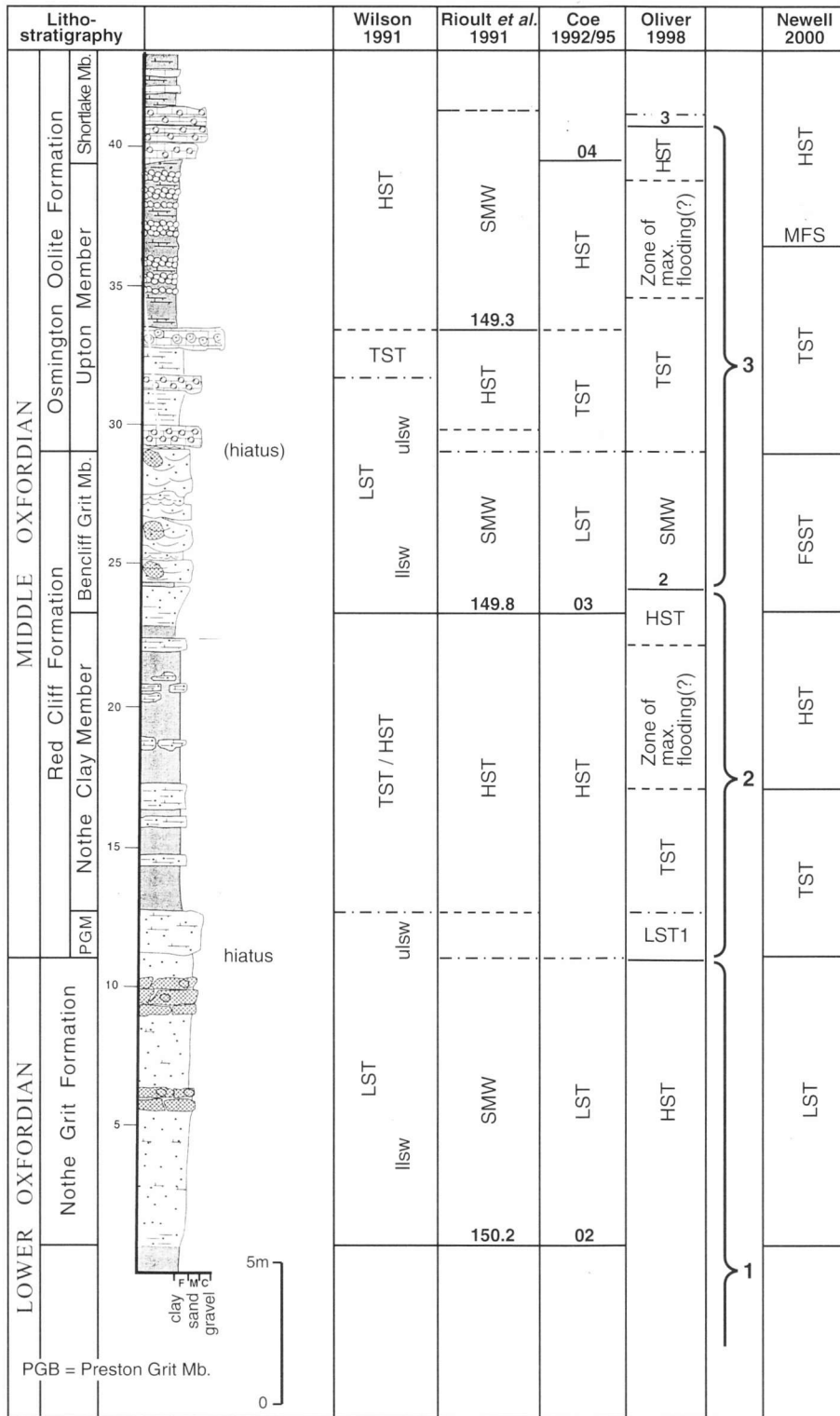


Fig. 2 - The Jurassic succession of the Weymouth to Ringstead Bay section (see Fig. 3) with the major lithostratigraphical units identified. The various sequence stratigraphical interpretations are also indicated. Additional terms include: llsw = lower lowstand wedge; ulsw = upper lowstand wedge; SMW = Shelf Margin Wedge; FSST = Falling Stage Systems Tract. O2/O3/O4 are the unconformities identified by Coe (1995). The radiometric dates (in millions years) are the ages assigned to Sequence Boundaries by Rioult et al. (1991).

Interpretation of the fauna

The faunas we record from geological samples are the product of:

- the initial fauna;
- post-mortem taphonomy;
- diagenesis and compaction;
- changes during burial and "geological time";
- modern weathering and exhumation (if not collected from the sub-surface);
- errors introduced during sampling, processing and analysis.

Work on modern-day faunas and their relationship with the substrate are important in helping to understand the death assemblages preserved in the geological record. In a recent survey of the foraminifera of Plymouth Sound (Fig. 6), Castignetti (1997) sampled a series of siliciclastic locations at monthly intervals over a yearly cycle and was able to determine

6.14) predictions. The genera recorded in this part of the succession are indicated in Fig. 5. The pattern is not clear, but samples O3 and O4 appear to be close to the zone of 'maximum flooding' and a number of important genera (*Ophthalmidium*, *Nubeculina*, *Vaginulina*, *Fronicularia*, *Citharina*, *Lingulina* and *Epistomina*) are restricted to this interval. The miliolids (with a porcellanous wall) and the epistominids (with an aragonitic wall) point to a preservational control of the fauna that characterises this interval.

the standing crop of all taxa from a wide range of environments. Fig. 7 shows the total annual production and the total annual diversity for these sites, all but one of which were located within Plymouth Sound. The most diverse and productive fauna is at Location 9 (mud) with the lowest values recorded at Location 11 (sand waves). The controls on the fauna are complex (and the subject of a further publication by the authors) but it is clear that in the nutrient-rich clays there is an abundant, di-

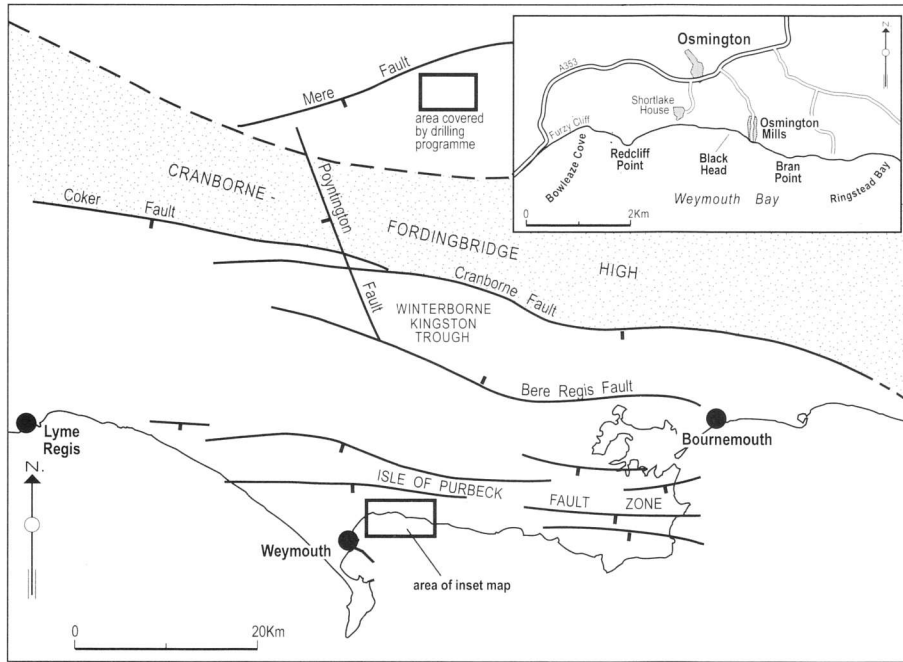


Fig. 3 - Outline structural map of Dorset showing the locations mentioned in the text. The area of North Dorset indicated in the box marks the location of the 5 boreholes used by Henderson (1997) and Henderson & Hart (2000).

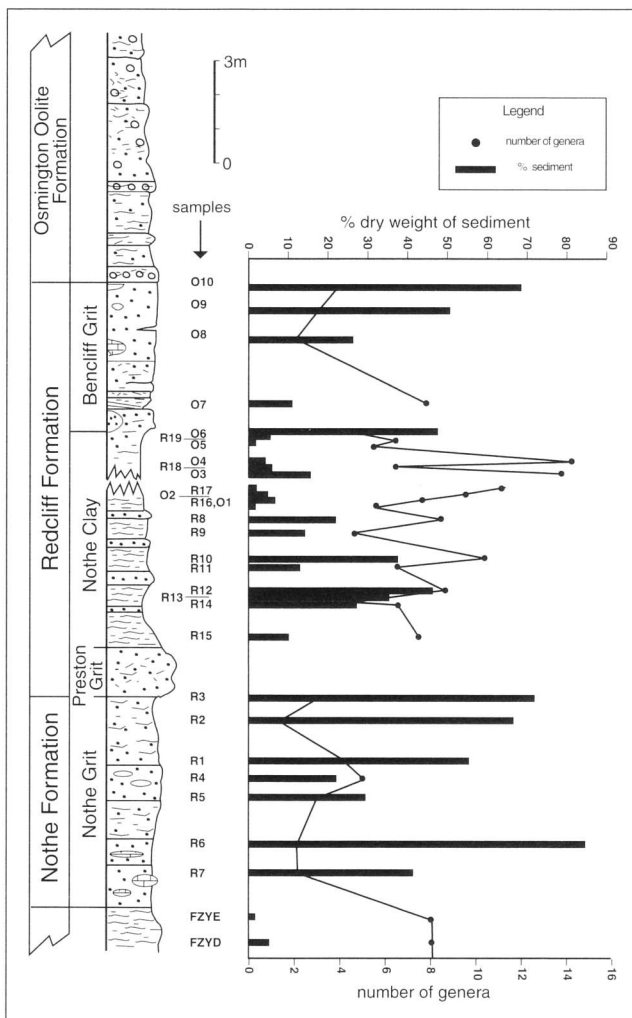


Fig. 4 - Sample locations within the Nothe Grit Formation, Preston Grit Member, Nothe Clay Member and the Benclyff Grit Member. Plotted against these samples are the percentages of dry sediment (by weight) retained on the 63µm sieve and the generic diversity.

verse fauna with the nutrient-poor sands supporting only a poor, in-situ, living fauna.

It is evident that the faunas in the clay-rich environments (Fig. 8) will be further enhanced by the slow rate of sedimentation (= several yearly standing crops in an average micropalaeontological sample) compared to the dilution effect of the higher rates of sedimentation in the more sand-rich environments. Compaction will further accentuate this situation with clays often being subjected to ~50% reduction. This is often detected by the presence of compressed agglutinated foraminifera where the flexible chitinous inner lining of the test has allowed an inward collapse of the chambers (Henderson, pers. comm.). This is particularly seen in genera such as *Trochammina*, *Haplophragmoides*, *Ammobaculites* and *Reophax*.

In recent years a number of investigations of the preservation potential of foraminifera have been undertaken by Murray (1989), Boltovskoy & Totah (1992), Alve & Murray (1995) and Murray & Alve (2000). It is particularly noted by Alve & Murray (1995) and Murray & Alve (2000) that high diversity agglutinated assemblages can be derived from high diversity assemblages apparently dominated by calcareous taxa through the selective dissolution of the calcareous component over time. This will particularly effect the aragonitic taxa and, in the Jurassic and Cretaceous, the epistominids will often only be found in the dense, clay-rich parts of the succession that have acted as aquacludes (Fig. 8). In Jurassic sequences, therefore, the aragonitic fauna of epistominids and planktonic foraminifera (Oxford et al. 2002) are the most likely indicators of maximum flooding surfaces or “zones”.

In samples collected at outcrop, even the best preserved faunas can still be removed by modern processes. A sure sign of this will be the presence of selenite (gypsum)

sample codes	FD	FE	R7	R6	R5	R4	R1	R2	R3	R15	R14	R13	R12	R11	R10	R9	R8	R16	O1	O2	R17	O3	R18	O4	O5	R19	O6	O7	O8	O9	O10
<i>Ammobaculites</i> spp.																															
<i>Haplophragmoides</i> spp.																															
<i>Textularia</i> spp.																															
<i>Trochammina</i> spp.																															
<i>Lenticulina</i> spp.																															
<i>Citharina</i> spp.																															
<i>Dentalina</i> spp.																															
<i>Nodosaria</i> spp.																															
<i>Planularia</i> spp.																															
<i>Vaginulina</i> spp.																															
<i>Nubeculina bigoti</i> Cushman																															
<i>Lagena</i> spp.																															
<i>Epistomina</i> spp.																															
<i>Eoguttulina</i> spp.																															
<i>Pseudonodosaria</i> spp.																															
<i>Triplasia</i> spp.																															
<i>Marginulina</i> spp.																															
<i>Lingulina</i> spp.																															
<i>Frolicularia</i> spp.																															
<i>Ophthalmidium</i> spp.																															
<i>Suboidellina</i> spp.																															
<i>Ramulina</i> spp.																															
<i>Trocholina</i> spp.																															
<i>Tristix</i> spp.																															

Fig. 5 - Distribution of foraminifera in the Nothe Grit and Redcliff Formations. *Nubeculina bigoti* Cushman is identified as this is the only species recorded within that genus in this succession.

crystals on bedding and fracture surfaces, or processed residues containing an abundance of such crystals.

The dense, impermeable clays will, therefore, contain the most abundant and diverse faunas from deposition, through compaction, to preservation in an aquaclude. These dense clays should also record the highest levels of gamma-rays in wireline logs and be readily identified

as a zone of maximum flooding by both palaeontologists and sedimentologists.

The Oxfordian succession of the Dorset Coast

Using the form of analysis shown in Fig. 4, the complete Oxfordian succession of the Dorset Coast

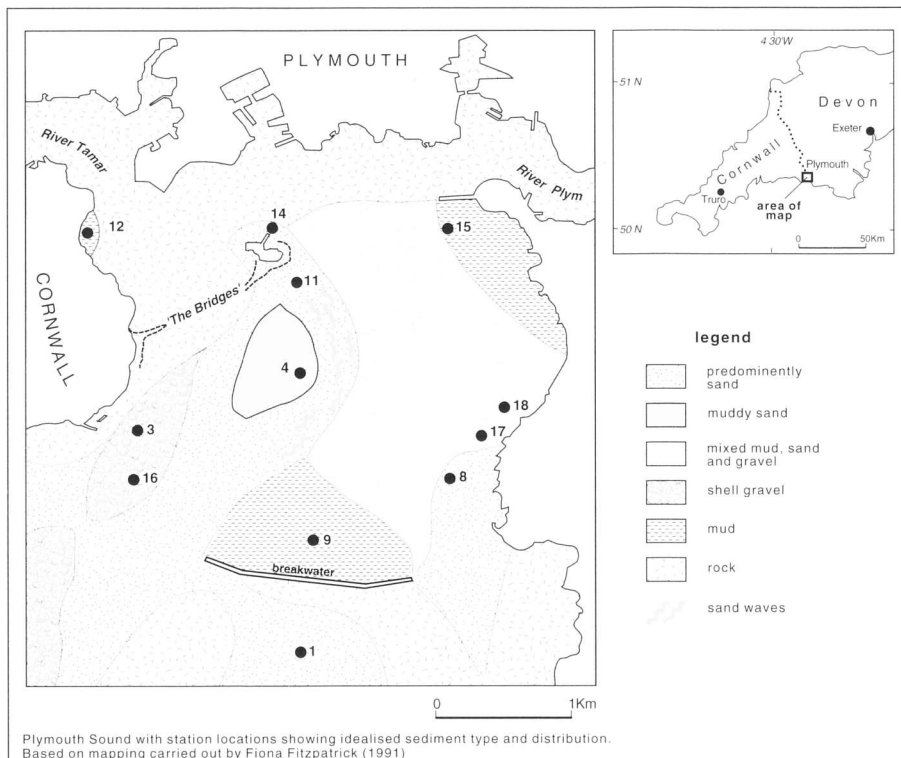


Fig. 6 - Sediment distribution map of Plymouth Sound and the location of the sampling sites used in the monthly analysis (1994-1996) of the benthonic foraminifera (after Castignetti et al. 2000).

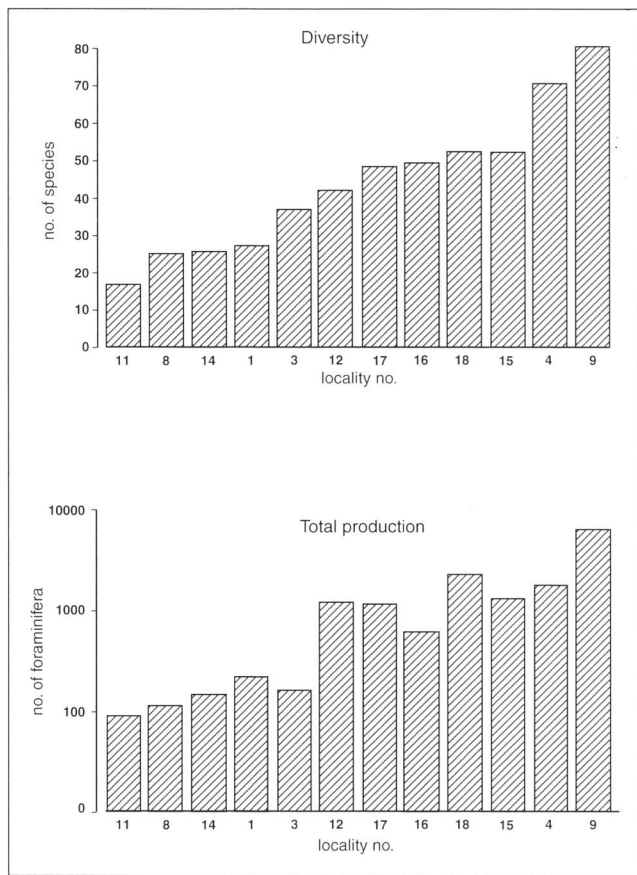


Fig. 7 - Plots of annual specific diversity and annual total production for the sites indicated in Fig. 6 (after Castignetti 1997).

(Weymouth to Ringstead) has been analysed (Fig. 9). The 'cycle' shown in Fig. 4 can be seen in the middle of this diagram, picked out by a number of diversity peaks (around sample OSM3). These pick out the zone of maximum flooding described above. This is within the upper part of the Nothe Clay Member of the Redcliff Formation. Lower in the succession, below sample FZ3, there are reduced percentages of sediment retained on the 63µm sieve and moderate levels of foraminiferal diversity. This is typical of the upper part of the Oxford Clay Formation (Mariae Zone). It is the Mariae Zone, on the shores of the Fleet west of Weymouth (Fig. 3), that has yielded planktonic foraminifera (Oxford et al. 2002) and these assemblages are coeval with the occurrence of planktonic foraminifera in the Marnes de Villers Formation on the Normandy Coast (Samson et al., 1992; Oxford et al. 2002, fig. 2). In North Dorset (Fig. 3) Henderson & Hart (2000, fig. 3) have identified several floods of epistominids at this level which probably records the same maximum flooding event.

Above the Nothe Clay 'cycle' (Figs 4, 9) there is a marked drop in diversity and an increase in sediment retained on the 63µm sieve. This appears to indicate a significant break (= sequence boundary) within the lower part of the Bencliff Grit Member and, although the exact

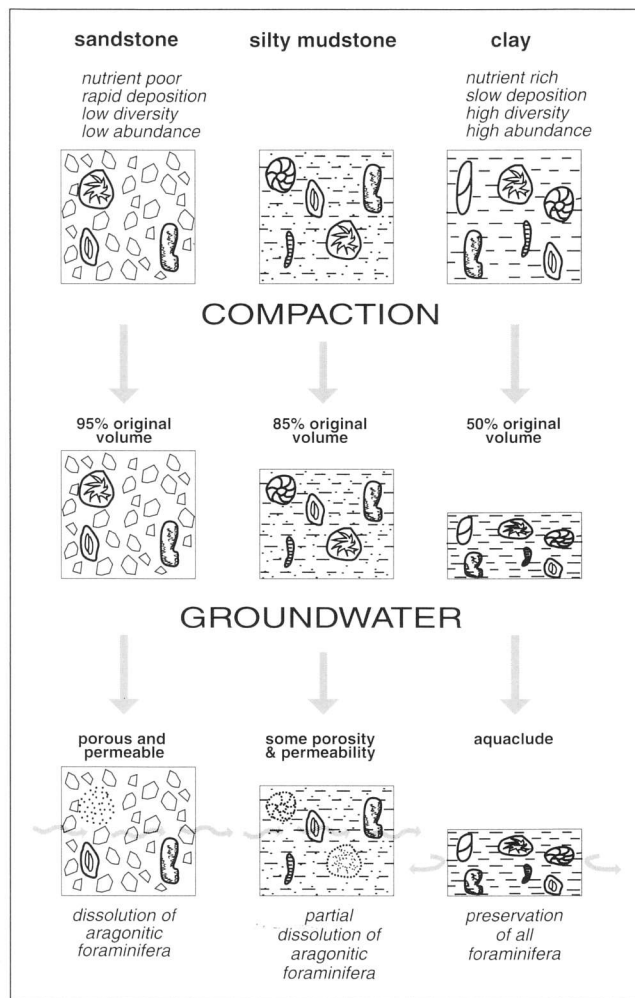


Fig. 8 - Theoretical model for the history of a "sample" from deposition to final collection, showing the effects of taphonomy, compaction and dissolution by groundwater flow.

position of this is disputed by the various workers who have studied this succession (Fig. 2), all show this event at about this level. In the overlying Upton Member of the Osmington Oolite Formation (around sample BH1 on Fig. 9) we record a major diversity peak. Above this level there are four more diversity peaks in descending order of magnitude until, at about the level of sample R2 there appears to be a major sequence boundary before the onset of the major cycle at the base of the Kimmeridge Clay Formation. If all the diversity peaks between the top of the Oxford Clay Formation and the base of the Kimmeridge Clay Formation are considered, it is possible to detect a large 'cycle' with a peak at about the level of the mid-Upton member. Is this a 3rd Order sequence in the Van Wagoner et al. (1988) terminology? If this is the case, then the regularly spaced minor peaks would be 4th Order sequences (or parasequences). In both North Dorset and Normandy our work has detected comparable diversity fluctuations with floods of diverse assemblages (with epistominids) at regular intervals (see Henderson & Hart 2000).

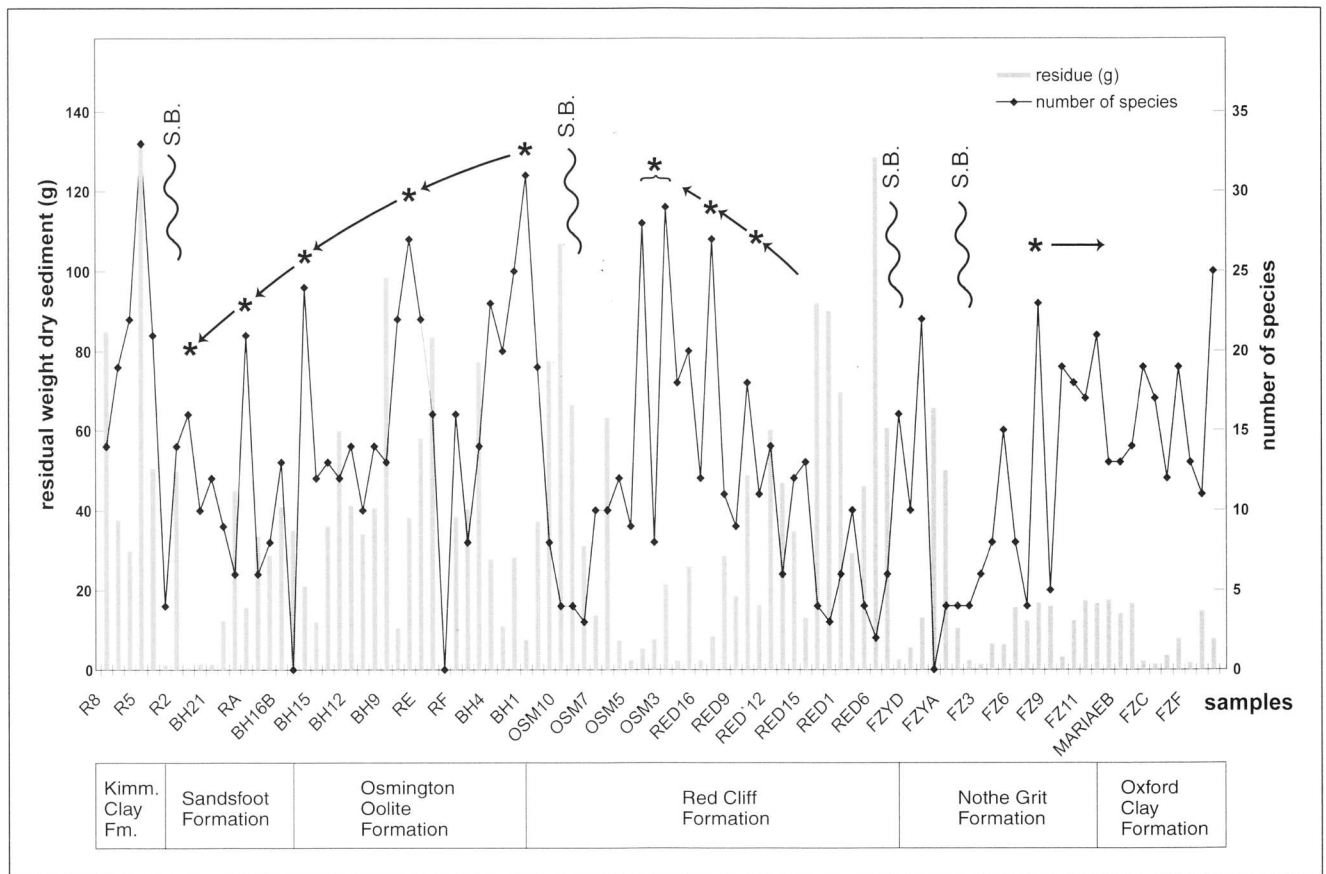


Fig. 9 - Analysis of the Oxfordian succession of the Dorset Coast following the methodology used in Figure 4. Locations of potential Maximum Flooding Surfaces (or Zones) are indicated by an asterisk; some of these are sequences while others may be parasequences. Potential sequence boundaries are indicated by the 'wavy' lines and initials SB. The sample codes are as follows:

- FZ / FZY ~ Furzy Cliff [Upper Oxford Clay - Nothe Grit Formation]
- RED ~ Redcliff [Nothe Grit Formation - Nothe Clay Member]
- OSM ~ Osmington [Nothe Clay Member - Osmington Oolite Formation]
- BH ~ Black Head [Osmington Oolite and Sandsfoot Formations]
- R ~ Ringstead Bay [Sandsfoot Formation and base of Kimmeridge Clay Formation]

Only an outline assessment of our results have been presented here as there is clearly a great deal of detailed information to process on all the faunas we have recovered in our samples. This work is in hand, including that based on the material available from North Dorset and Normandy.

Summary

Using data from the analysis of foraminifera living in modern siliciclastic environments we have developed a model for the categorization of Jurassic assemblages and their interpretation in a sequence stratigraphic context. Preliminary work comparing the Dorset Coast with North Dorset and Normandy indicate that the foraminifera

can provide significant assistance in the development of correlations based on sequence stratigraphy. Maximum flooding events in the Jurassic appear to be identified by high diversity assemblages in which epistominids are a significant component. Planktonic foraminifera (which at this stratigraphic level may also have been aragonitic) have also been shown to occur at maximum flooding surfaces within the mid-Upper Jurassic (Oxford et al. 2002; Hart et al. 2002).

Acknowledgements. M.J.O. acknowledges receipt of a studentship part-funded by the Faculty of Science and Amerada Hess (London). Mr. John Abraham is thanked for his assistance with the final versions of the figures. Comments given by two reviewers, S. Feist-Burkhardt and F. Fursich, were helpful in the preparation of the final version of the paper.

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