# EVOLUTIONARY AND BIOGEOGRAPHICAL IMPLICATIONS OF PHYLOGENETIC ANALYSIS OF THE LATE PALAEOZOIC TRILOBITE PALADIN

by david K. Brezinski

ABSTRACT. Phylogenetic analysis of 19 species assigned to the Carboniferous and Permian trilobite genus *Paladin* produced a single most parsimonious tree with four distinct branches termed Clades A to D. Clade A is made up of six species that are geographically restricted to the interior shelf regions of the United States and are assigned to *Kaskia*. Clade B of five species, which includes *Paladin morrowensis* (Mather), is known from the southern and western United States; these five species conform morphologically to the original diagnosis of *Paladin*. Clade C consists of five Viséan to Namurian species from western and eastern Europe and at least three of the species should be assigned to *Weberides*. Clade D consists of two late Carboniferous and one early Permian species from the Ukraine, western Russia, and Spitsbergen. This clade is distinct from all the others analysed and probably represents a yet unnamed genus.

KEY WORDS: Trilobita, Carboniferous, evolution, phylogenetics, palaeobiogeography.

PALADIN is one of the most widespread and long-lived late Palaeozoic trilobite genera. It ranges from the early Carboniferous (early Viséan) to Late Permian (Guadalupean), a duration of nearly 100 million years. Examination of systematic literature indicates that approximately 50 species are assigned to this genus worldwide (Hahn and Hahn 1970; Osmólska 1970; Kobayashi and Hamada 1980). A number of these species from the late Carboniferous and Permian can be reassigned a priori to other genera. Some clear examples of this are P. gruenewaldi (Möller, 1867), P. roemeri (Möller, 1867), and P. pyriformus Chamberlain, 1969. The first two species, by lacking an anterior border, are more appropriately assigned to Triproetus Kobayashi and Hamada, 1979; the last, which displays a medial preoccipital lobe, should be assigned to Pseudophillipsia (Carniphillipsia) Hahn and Brauckmann, 1975. It is not clear from the current analysis whether the Permian species previously assigned to Paladin and herein assigned to Neokaskia and Triproetus are derived from the early Carboniferous Paladin stock. This unresolved question may warrant future study. Even with the reassignments considered above, there are still approximately 40 species that can be attributed to Paladin, all of which are from the Carboniferous.

Although the revised stratigraphical range of *Paladin* spans most of the Carboniferous, North American representatives are restricted to the Viséan and Namurian (Osagean–Morrowan) as outlined as 'Stage 2' of Brezinski (1999). The relatively complete marine record for the Carboniferous and Permian of North America notwithstanding, it is puzzling why this long-ranging and cosmopolitan genus has never been recognized outside the Stage 2 limits documented by Brezinski (1999). This perplexing stratigraphical and geographical restriction spurred the current study. Because this genus is such an important component of the Carboniferous trilobite faunas of the United States, a taxonomic re-evaluation was undertaken in an attempt to understand better the phylogeny and systematic relationships of *Paladin*.

## Taxonomic confusion

Weller (1936) erected *Paladin* on a Namurian species from the southern United States, *Griffithides morrowensis* Mather, 1915. Characterised by a forwardly expanding glabella, large posteriorly located palpebral lobes, anterior border, and bordered pygidium, according to Weller, *Paladin* was similar to

Kaskia, which he erected in the same publication. He noted that Kaskia differed from Paladin by lacking the distinct anterior border, and had more forwardly located palpebral lobes, but that there were several species which exhibited morphologies intermediate between the two type species. This gradation in morphology has through time posed a considerable amount of confusion when subsequent species were assigned to either genus. Whittington (1954) recognized this gradation in morphology between the two and chose to consider Kaskia a subgenus of Paladin. Whittington (1954, p. 5) also noted that some of the European species assigned to Weberides by Reed (1942) might also be assignable to Paladin. Cisne (1967) also noted the difficulty in separating Kaskia from Paladin and considered the two to be congeneric. The lumping of Kaskia and Weberides under Paladin was accepted by Hahn and Hahn (1970), Osmólska (1970), and most subsequent workers, with the exception of Kobayashi and Hamada (1980) who continued to separate Weberides as a distinct genus.

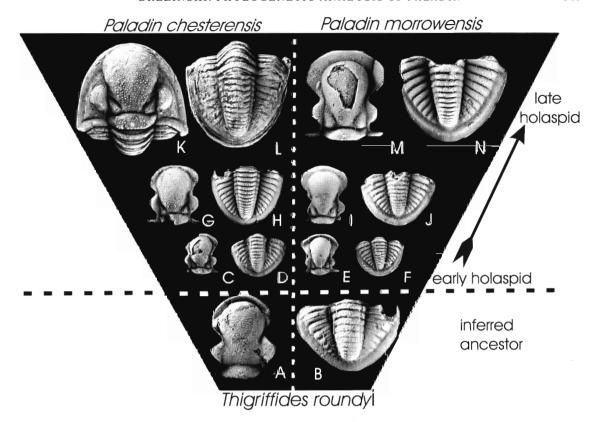
## PUTATIVE ANCESTRAL MORPHOLOGY

Throughout the earliest Carboniferous (Tournaisian) the characteristic phillipsiid morphology consisted of a forwardly tapering to parallel-sided glabella, narrow, forward-located, palpebral lobes, and borderless pygidium. The few exceptions include *Griffithides*, *Bollandia*, *Griffithidella*, and *Thigriffides*. It can be surmised that the widespread Viséan and Namurian *Paladin* probably originated from one of these possible ancestors. Although Weller (1936), in his original remarks on the genus, proposed that *Paladin* was derived from *Griffithides*, this is not a likely ancestral candidate since it is characterized by narrow palpebral lobes located forward on a cranidium that lacks an anterior border, and has a long pygidium that is weakly bordered. Furthermore, it is unlikely that *Griffithides* served as a precursor to *Paladin* since both genera have the same stratigraphical ranges. *Bollandia* also is characterized by small, forward-located palpebral lobes, and lacks an anterior cranidial border, consequently, as with *Griffithides*, it is not a likely ancestor to *Paladin*. The large, posteriorly located palpebral lobes, forwardly expanding glabella, anterior cranidial border, and distinctly bordered pygidium are pleisiomorphic characters that point to only two other potential ancestral genera, *Griffithidella* and *Thigriffides*. The former has ten thoracic segments, a very short pygidium, and a narrow glabella that, in some species, is forwardly converging or medially constricted. These characters tend to suggest that *Griffithidella* is also an improbable ancestor to *Paladin*.

Hahn and Hahn (1967, fig. 5) suggested that *Paladin* was derived from the early Carboniferous *Thigriffides*. This was agreed by Owens (*in* Owens and Hahn 1993), but alternatively, Owens and Hahn (1993) suggested that *Rhenogriffides* Hahn *et al.*, 1987 may have supplied the ancestral stock for *Paladin*. Although *Rhenogriffides* shares a superficial resemblance to *Paladin*, it is characterized by transversely narrow palpebral lobes, a poorly defined anterior border, and glabella with parallel to slightly forward-tapering sides. Clearly, *Thigriffides*, with its large posteriorly located palpebral lobes, forwardly expanding glabella, and anterior border, has a closer resemblance to *Paladin*, and is more tenable as the ancestor. These presumed pleisiomorphic characters, although widespread in the middle to late Carboniferous, are unusual in the Tournaisian.

The dearth of ontogenetic studies for late Palaeozoic trilobites hinders phylogenetic reconstruction based on ontogenies. However, comparison of variation through the holaspid stage of several species aids reconstruction of ancestral morphologies. Text-figure 1 illustrates such holaspid variation of *P. morrowensis* (Mather, 1915) and *P. chesterensis* (Weller, 1936). Notwithstanding the marked morphological divergence exhibited by larger holaspid specimens, smaller, and presumably younger, individuals show a marked interspecific morphological convergence. This is interpreted to suggest that these species share a common ancestral history. The putative ancestor which will be used in subsequent analysis is *Thigriffides roundyi* (Girty, 1926) (Text-fig. 1).

While there have been no detailed ontogenetic studies of North American species of *Paladin*, Clarkson and Zhang (1991) provided detailed documentation of morphological changes shown by *P. eichwaldi shunnerensis* (King, 1914) from Great Britain. The marked similarity between the early holaspid individuals of *P. chesterensis* (Weller, 1936), *P. eichwaldi* (Fischer v. Waldheim, 1825), and *P. morrowensis* (Mather, 1915) allows a reasonable interpretation that the European taxa assigned to



TEXT-FIG. 1. A-B, morphology in inferred ancestor *Thigriffides roundyi* (CFM 51952, US National Museum 3523, respectively). C-N, morphological variation of cranidia and pygidia observed in individuals of increasing size through the holaspid stage of *Paladin chesterensis* (Weller, 1936) and *P. morrowensis* (Mather, 1915). C-D, small holaspid individuals of *P. chesterensis* (Carnegie Museum of Natural History (CM) 41720–41721). E-F, small holaspid individuals of *P. morrowensis* (CM 41724–41725). G-H, middle holaspid individuals of *P. chesterensis* (CM 41722–41723). I-J, middle holaspid individuals of *P. morrowensis* (CM 41726–41727). K-L, larger holaspid individuals of *P. chesterensis* (Chicago Field Museum (CFM) 34437). M-N, larger holaspid individuals of *P. morrowensis* (CM 41728–41729). All photographs ×2·0.

*Paladin* and the North American representatives share a close common ancestry (see Clarkson and Zhang, 1991, figs 12–13; text-fig. 3).

## PHYLOGENETIC ANALYSIS

The plethora of apparently gradational morphologies displayed by species assigned to *Paladin*, and exemplified by the *Paladin-Kaskia* conundrum, is confusing both to traditional taxonomy and phylogeny reconstruction. Because of this, modern phylogenetic procedures were chosen to analyse a number of species assigned to *Paladin*. The initial phylogenetic study, which concentrated on 11 North American species, produced a single tree with a distinct two-branch topology which reflects a dichotomy of taxa into two separate clades. Because these results are mirrored in later analyses they will not be discussed here. The topology produced in this early analysis raised a question regarding the relationship between these North American clades and the European representatives of the genus. This question led to the inclusion in the analysis of an additional eight species known from Europe, five from the Viséan and Namurian and three from Westphalian to Stephanian. The Viséan–Namurian species are: *P. cuspidatus* (Reed, 1943),

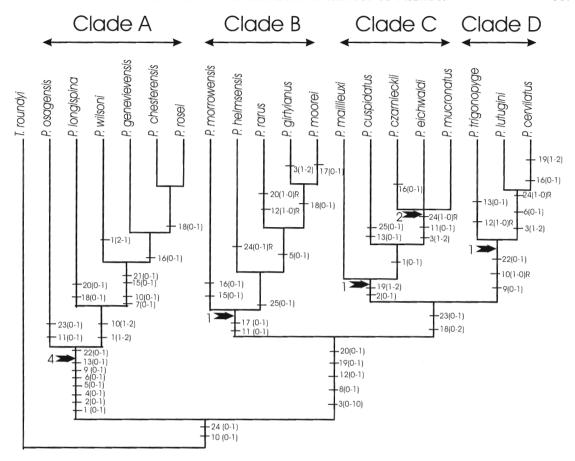
TABLE 1. List of characters and character-states used in phylogenetic analysis of *Paladin* ingroup using *Thigriffides* roundyi as an outgroup.

- 1. Anterior border. Distinct and wide (0). Narrow (1). Indistinct to obsolete (2).
- 2. Anterior border furrow. Shallow (0). Absent (1).
- 3. Shape of the base of glabella. Quadrate (0). Subrounded (1). Acutely rounded (2).
- 4. Palpebral midline position. Less than or equal to 0.35 cranidial length (0). Greater than 0.35 cranidial length (1).
- 5. Palpebral outline. Nearly semicircular (0). Crescentic (1). Narrow (2).
- 6. Character of S1. Shallow (0). Narrow, deeply incised (1).
- 7. Lateral border. Sharply furrowed (0). Indistinct furrow rounded at margin (1).
- 8. Occipital furrow longitudinal profile. Narrowly incised (0). V-shaped profile (1).
- 9. L2-3 present. No (0). Yes (1).
- 10. Posterior facial sutures. Short curved (0). With a short straight section (1). With a long straight section (2).
- 11. L1 outline. Suboval (0). Subtriangular (1).
- 12. L1 position with respect to glabella base. Sub-even (0). Behind glabella base (1).
- 13. S1 trace. Straight (0). Recurved posteriorly (1).
- 14. Pygidial/Axial width ratio. Less than or equal to 0.33 (0). Greater than 0.33 (1).
- 15. Pleural furrow. Well incised (0). Shallow-narrow (1).
- 16. Interpleural furrows. Distinct, except on most posterior ribs (0). Distinct on only anteriormost ribs (1).
- 17. Axial terminus outline. Broadly rounded (0). Sharply rounded with tapered extension (1).
- 18. Pygidial outline. Parabolic (0). Semi-elliptical (1). Lanceolate (2).
- 19. Pygidial border. Evenly wide (0). Slightly wider posterior (1). Posteriorly extended (2).
- 20. Pleural field convexity (transverse). Arched (strongly convex) (0). Flattened exsagittal (low convexity) (1).
- 21. Pygidial border slope. Less than lateral part of pleural field slope (0). Equal to pleural fields (1). Flanged (2).
- 22. L1 depressed in height relative to posterior of glabella. Yes (0). No (1).
- 23. Pygidium axial transverse profile. Rounded (0). Subtrapezoidal (1).
- 24. Cranidial width at  $\Delta$  relative to  $\beta$ .  $\Delta$  greater than  $\beta$  (0).  $\Delta$  equal to or less than  $\beta$  (1).
- 25. Shape of sutures at  $\beta$ . Broadly rounded (0). Acutely rounded (1).

P. czarnieckii Osmólska, 1970, P. eichwaldi (Fischer v. Waldheim, 1825), P. maillieuxi (Demanet, 1938), and P. mucronatus (M°Coy, 1844). The three late Carboniferous species are: P. cervilatus (Weber, 1933), P. lutugini (Weber, 1933), and P. trigonopyge Osmólska, 1968.

The 19 Paladin species comprising the ingroup were compared using 25 morphological characters. These taxa were examined using PAUP 4.0 in beta version employing a heuristic search command with simple addition. All characters were unordered and of equal weight, and are summarised in Table 1. The list of character states for each ingroup species is given in the Appendix. A single most parsimonious tree was retained by the analysis (Text-fig. 2). This most parsimonious tree, of length 62 steps, produced a consistency index of 0.48 and a retention index (RI) of 0.80. The topology of the single most parsimonious tree displayed four separate branches, labelled Clades A to D. The sister clades, Clades A and B, were produced by the initial analysis of North American species; these two groups maintained their integrity during the more comprehensive analysis. Bremer indices, an indication of branch stability, were calculated for this topology and are plotted on their respective branches on Text-figure 2. The total support index (ti) of 0.15 was calculated from the branch supports, and suggests a fairly stable topology. Fifty per cent consensus trees also were produced by bootstrap and jack-knife analyses to test the stability of the four-branch topology. Although the four-branch topology was retained in the bootstrap analysis, Clade B was lost during jack-knife analysis. However, during both analyses several of the taxa were removed from Clades C and D. The possible significance of these omissions is discussed below.

Clade A consists of six species: *P. osagensis* Cisne, 1967, the oldest representative of the genus; *P. chesterensis* (Weller, 1936); *P. genevievensis* (Walter, 1924); *P. longispina* (Strong, 1872); *P. rosei* Cisne, 1967; and *P. wilsoni* (Walter, 1924). This clade ranges from early to late Viséan (late Osagean—middle Chesterian). The large number of characters that define this group (see Text-fig. 2) directly reflects that it was the ingroup defined in the earlier analysis. The result of the large number of synapomorphies is a



TEXT-FIG. 2. Single most parsimonious cladogram of 19 ingroup taxa assigned to *Paladin*. Characters 1–25 given in Table 1; assigned character states given in Appendix. Distribution of characters given on branches; R denotes a reversal to an ancestral state. Bremer branch support index number given in dark numbers to left of arrows at branch locations. Cladogram generated by Treeview (Page 1996).

relatively high Bremer branch index of 4. This branch is also retained on the consensus trees produced by bootstrap and jack-knife analyses.

Clade B is made up of the remaining five North American species: *P. girtyianus* Hahn and Hahn, 1970; *P. helmsensis* Whittington, 1954; *P. moorei* (Branson, 1937); *P. morrowensis* (Mather, 1915); and *P. rarus* Whittington, 1954. Species of this clade range in age from the late Viséan to late Namurian (middle Chesterian–middle Morrowan). Although this branch is defined by a low Bremer branch index of 1, both sister clades A and C have much higher indices, which serve to isolate the low stability branch between them. Although this branch is retained on the consensus tree from the bootstrap analysis, it is lost during the jack-knife analysis.

The branch defining Clade C is created by the five Viséan–Namurian species from Europe. This group contains *P. cuspidatus* (Reed, 1943), *P. czarnieckii* Osmólska, 1970, *P. eichwaldi* (Fischer v. Waldheim, 1825), *P. maillieuxi* (Demanet, 1938), and *P. mucronatus* (M°Coy, 1844). *Paladin czarnieckii* Osmólska, 1970, *P. eichwaldi* (Fischer v. Waldheim, 1825), and *P. mucronatus* (M°Coy, 1844) produce an unresolved polytomy. Bremer indices on this branch occur at the node at the base of the branch and between the three polytomous species and their sister species. This later index may suggest that the three unresolved species represent a separate clade from the two non-polytomous species. This is further suggested by the bootstrap

and jack-knife consensus trees, which each maintain the three polytomous European species as a separate branch. This polytomy suggests that the characters used in the analysis of North American species is not sufficiently discriminating for all European species. Clearly, further analysis of this clade is warranted to evaluate whether the three unresolved species indeed represent a separate group that is not differentiable by the current characters utilized. The three unresolved species are characterized by the shape of the glabellar base, a narrow anterior cranidial border, broadly rounded anterior facial suture, and posterior elongation of the pygidial border.

The fourth clade, termed here Clade D, is made up of the three species from upper Carboniferous—lower Permian strata. Both *P. cervilatus* (Weber, 1933) and *P. lutugini* (Weber, 1933) are known from the Westphalian of the Donets Basin of Ukraine, whereas *P. trigonopyge* (Osmólska, 1968) is known from lower Permian (Asselian) strata of Spitsbergen. Although this branch has a Bremer index of only 1, it is maintained on the consensus trees for both the bootstrap and jack-knife analyses, albeit without *P. trigonopyge* (Osmólska, 1968). The removal of *P. trigonopyge* (Osmólska, 1968) from this branch during both of the variance analyses raises questions about its relationship to this clade as well as other *Paladin* clades.

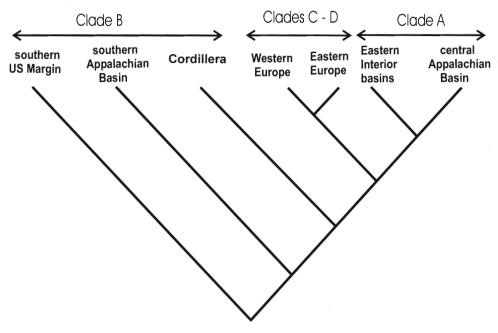
## PALAEOGEOGRAPHICAL IMPLICATIONS

The four clades delineated by phylogenetic analysis of 19 ingroup species assigned to *Paladin* reflect subgroups that are not only differentiatable on a phylogenetic basis but also on palaeogeographical grounds. In an effort to evaluate possible vicariance patterns within the ingroup species, Brookes Parsimony Analysis (BPA) was conducted (Text-fig. 3).

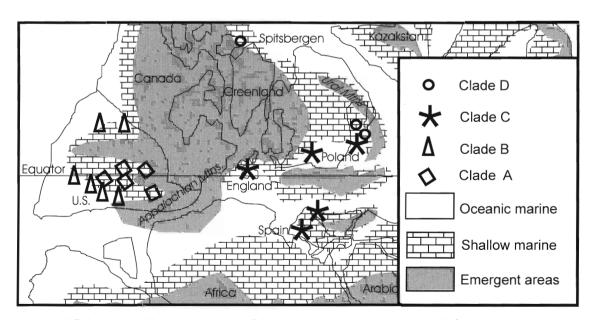
The ingroup species are present in seven separate geographical areas (Text fig. 4). Clade A is present in the central Appalachian Basin (Area 1), the Illinois and Michigan basins, and upper Mississippi Valley area (Area 2). Clade B is present in the southern Appalachian Basin (Area 3), Cordilleran region (Area 4), and the southern United States (Arkansas, Oklahoma, and Texas) (Area 5). Clade C is present in western Europe, including the United Kingdom, Belgium, Spain, and France (Area 6), as well as eastern Europe (Germany, Poland, Czech Republic, Ukraine, and Spitsbergen) (Area 7). Clade D is known only from the Ukraine and Spitsbergen.

Utilizing BPA with the above-defined areas, a single area cladogram was derived (Text-fig. 3), with a consistency index of 0.92 and 38 steps long. This single area cladogram reveals an interesting vicariance pattern, but one that might be intuitively expected. On the retained tree, Area 5, the southern margin of the United States, represents the ancestral area. Sister areas to this ancestral area are both the southern Appalachian Basin and the Cordillera. These three areas are the known sites where species of Clade B occur. Western Europe and eastern Europe are sister areas to one another, as they form a branch which is a sister area to both the Cordillera and eastern Interior areas of the United States. The European branch contains species of Clades C and D. Perhaps the most derived areas are 1 and 2 which contain species of Clade A.

Because the area cladogram produces relationships that are consistent with the cladogram in Text-figure 2, a number of palaeogeographical inferences may be proposed (Text-fig. 4). The restriction of Clade B to Areas 3, 4, and 5 is consistent with both phylogenetic and environmental occurrences. Area 5, the area where the type species of *Paladin* is known, juxtaposes the known area of occurrence of *Thigriffides roundyi* (Girty, 1926), the interpreted ancestor of *Paladin*. Brezinski (1998) found that this ancestral species inhabited presumed deep water, starved basin deposits off the southern margin of the United States during the Tournaisian. *Paladin*, the interpreted descendant of *Thigriffides*, is known from shelf edge and deeper off-shelf deposits of Arkansas and Oklahoma (Sutherland and Manger 1979). The environments that produced these deposits represent a variety of water depths from shallow, shelf edge, sand shoals, to relatively deeper, off-shelf shales. Although these rocks were deposited in a range of water depths the environments that produced them appear to be relatively long-lived and stable through time. This is indicated by the lack of interbedded lithologies and facies shifts (Sutherland and Manger 1979). The other areas where Clade B is known can also be attributed to similar environmental settings.



TEXT-FIG. 3. Area cladogram derived from phylogenetic cladogram illustrated in Text-figure 2 using Brookes Parsimony Analysis (BPA) (see Wiley 1988). Geographical areas are described in text. Cladogram generated by Treeview (Page 1996).



TEXT-FIG. 4. Palaeogeographical reconstruction of the pan-equatorial region during the Viséan. Continental and shelf area reconstructions from Scotese and McKerrow (1990). Symbols represent approximate locations where species of *Paladin* are known and clade in which they are assigned.

Another inference suggested from Text-figures 2 and 3 is that western Europe served as an ancestral area for eastern Europe. Although this inference is based on the area cladogram, it may be premature to make such an interpretation based on the limited number of species used in the current analysis.

Lastly, the most derived areas, the Eastern Interior areas of the United States and the central Appalachian Basin, were the sites where Clade A is present. Clade A occurrences are within deposits of interbedded marine and non-marine strata; the former were deposited in relatively shallow marine environments during cyclothemic iterations of sea level (Swann 1964; Brezinski 1983, 1989). It is suggested here that these interior basins allowed for the allopatric isolation of these taxa, permitting this lineage to evolve with little exchange with the ancestral southern areas.

## TAXONOMIC SIGNIFICANCE

While there is a dearth of information about the ontogenetic development of species of *Paladin*, morphological similarities within the early holaspid stage of a number of the species of *Paladin* argue for a shared ancestry (Text-fig. 5). The divergence of the 19 ingroup species into four, phylogenetically, palaeogeographically, and in some cases palaeoecologically, distinct groups suggests that a taxonomic separation may be warranted and that the current procedure of lumping the taxa together is inadvisable. The branch stability of Clade A along with its palaeogeographical segregation in the central

The branch stability of Clade A along with its palaeogeographical segregation in the central Appalachian, Illinois, and Michigan basins argues that this clade has significant divergence from its proposed ancestor. Interestingly, the position of these species on this branch of the cladogram is congruent with the stratigraphical occurrence of these species. *Paladin osagensis* Cisne, 1967, the earliest representative known (early Viséan), is succeeded, stratigraphically, by *P. longispina* (Strong, 1872) (middle Viséan strata), *P. wilsoni* (Walter, 1924) and *P. genevievensis* (Walter, 1924) (late middle Viséan), and finally by *P. rosei* Cisne, 1967, and *P. chesterensis* (Weller, 1936) (late Viséan). All of these factors point strongly towards Clade A being a distinct monophyletic group. One of the most derived species on this branch, *P. chesterensis* (Weller, 1936), was originally designated the type species of the genus *Kaskia* by Weller (1936). Considering the factors discussed above, it is herein recommended that the genus *Kaskia* be re-erected and that the species discussed herein as Clade A be assigned to it.

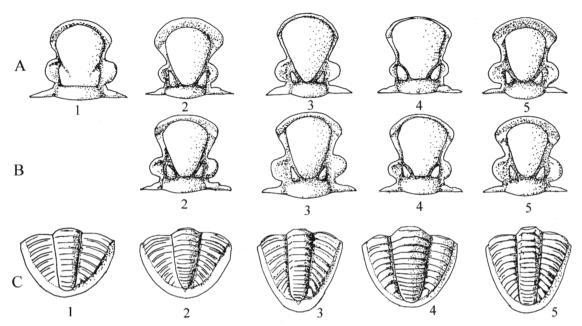
Some morphological attributes that should be used to characterise large holaspid individuals of *Kaskia* include a narrow, indistinct, anterior border, a subquadrate base of the glabella, long straight section of the posterior branch of the facial sutures, narrow, crescentic palpebral lobes, a broad, indistinct, trough-like, lateral border furrow, and a semielliptical pygidium, with shallow pleural furrows.

The branch stability of Clade C is manifested with two separate support values. Although the entire branch has a Bremer branch support number of 3, the unresolved polytomy of *P. czarnieckii* Osmólska, 1970, *P. eichwaldi* (Fischer v. Waldheim, 1825), and *P. mucronatus* (M<sup>c</sup>Coy, 1844) has an additional support value of 2 and is retained in both the bootstrap and jack-knife analyses. This trichotomy points to the possibility that Clade C, when examined more thoroughly, might provide a separation of the western European species into two separate and recognizable groups. Clade C is present in Europe and there is currently no known overlap of geographical distribution with it and members of clades A or B.

An important taxon within Clade C is *Paladin mucronatus* (M<sup>c</sup>Coy, 1844). Reed (1942) erected *Weberides* based on this taxon, and although other species within this clade do not show the terminal mucro on the pygidium, they do exhibit a broadening of the posterior pygidial border. Furthermore, the triangular outline to the pygidium is not seen in either Clades A or B. It is herein recommended that the unresolved species of Clade C be returned to the resurrected *Weberides*. Further analysis is needed to determine if *P. cuspidatus* (Reed, 1943) and *P. maillieuxi* (Demanet, 1938) should be assigned to *Weberides* or to a separate genus.

Characters which redefine this genus include the pygidial character mentioned above, in addition to the acutely rounded posterior base of the glabella, large palpebral lobes that are commonly broader than the cranidal width at  $\beta$ , and narrow anterior border that lacks a border furrow. While the current analysis suggests that perhaps more than one clade is represented by these species, sufficient information is currently not available to delineate such a separation.

While Clade B has a relatively low Bremer index, it is retained by bootstrap analysis, but lost during



TEXT-FIG. 5. Sketches of general cranidial morphologies shown by holaspid individuals of the various *Paladin* clades as well as that of the presumed ancestral morphology of *Thigriffides roundyi* (Girty, 1926) cranidium (A1) and pygidium (C1). Smaller holaspid cranidial morphologies (row B), larger holaspid cranidial morphologies (row A), and holaspid pygidial morphologies (row C). Column 2, *Paladin morrowensis* (Mather, 1915); column 3, *P. chesterensis* (Weller, 1936); column 4, *P. eichwaldi* (Fischer v. Wildhelm, 1825); and column 5, *P. lutugini* (Weber, 1933).

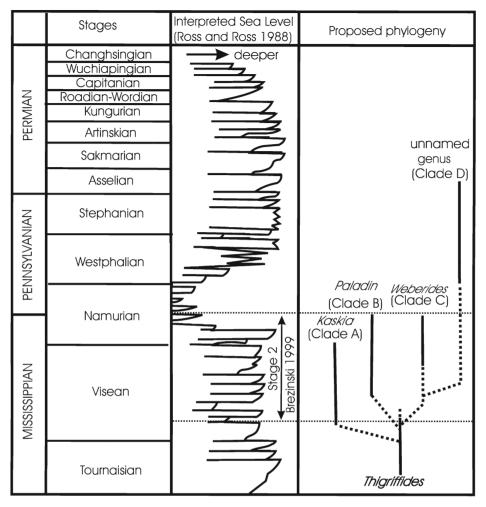
jack-knife analysis. This clade contains the type species of *Paladin, P. morrowensis* (Mather, 1915), as well as some of the more definitively assigned *Paladin* species (e.g. *P. helmsensis* Whittington, 1954, *P. girtyianus* Hahn and Hahn, 1970). Because the species of this clade exhibit a morphological separation from the other species examined, it should be considered a separate genus from either of its sister clades. Therefore, the characteristics of Clade B should be considered as those that typify the true *Paladin* morphology.

Refined morphological characteristics that typify *Paladin* are the subrounded base to the glabella, parabolic outline to the pygidium, sharply tapered terminus to the pygidial axis, and distinct anterior border, commonly separated from the front of the glabella by a shallow, narrow border furrow.

Clade D, like Clade B, has a relatively low Bremer support index of 1. However, a branch formed by two of the taxa, *P. cervilatus* (Weber, 1933) and *P. lutugini* (Weber, 1933), is retained on both the bootstrap and jack-knife consensus trees. Notwithstanding that this clade (or at least part of it) forms a derived branch separate from the other clades, the current analysis, based only on photographs of the type material, is insufficient to be certain that these species are truly a separate and distinct group. Therefore, it would be imprudent with the current information to erect a separate genus to include these species. Further analysis of these species as well as other late Carboniferous species such as *P. subbakewellensis* Osmólska, 1970 is needed.

## **EVOLUTIONARY INTERPRETATIONS**

The phylogenetic analysis presented here suggests that many of the taxa currently assigned to *Paladin* can be referred to a number of other genera. Similarity of morphology within the early holaspid stages, especially among the mid-Carboniferous genera, indicates their shared ancestry. A proposed phylogeny, consistent with the cladogram illustrated in Text-figure 2, is presented in Text-figure 6. The interpreted



TEXT-FIG. 6. Hypothesis of phylogeny based on phylogenetic and palaeogeographical reconstructions presented in Text-figures 2–4. Sea level curve from Ross and Ross (1988), and dashed lines delineate Stage 2 of North American trilobite evolution as outlined in Brezinski (1999).

ancestor, *Thigriffides*, ranged through most of the Tournaisian, and inhabited deeper water, basinal deposits. During the early Viséan, the early representatives of the *Kaskia* lineage appeared and migrated into the depositional basins of the shallow shelf interior of the central and eastern United States. In these quasi-isolated locations, this lineage evolved into the derived morphology seen in *Kaskia chesterensis* Weller, 1936. During the middle Viséan, a second lineage with morphologies consistent with true *Paladin* individuals was derived from the putative ancestral *Thigriffides* stock. The *Paladin* lineage inhabited areas along the margin of the late Mississippian and earliest Pennsylvanian shelves of the southern and western United States. Coincident with this lineage was the derivation of another lineage (or group of lineages) that inhabited the periphery of the Rheic seaway of Europe and western Russia and Ukraine and are manifested in the genus *Weberides*. This lineage, or group of lineages, arose in the middle to late Viséan and survived into the early Namurian.

The termination of the Kaskia, Paladin, and Weberides lineages by the middle Namurian indicates an extinction episode that spanned a considerable geographical area. This is consistent with the

interpretations of Brezinski (1999) that the Stage 2 of North American trilobite evolution was terminated during the mid-Carboniferous eustatic fall in sea level (Saunders and Ramsbottom 1986). The appearance of a *Paladin*-like morphology appears after this extinction may be the result of the Donets Basin of eastern Ukraine serving as a refuge during the mid-Carboniferous regression. Thus, the Westphalian and Stephanian taxa currently assigned to *Paladin* may represent Lazarus taxa (Jablonski 1986) that appeared during late Carboniferous submergence. Alternatively, these species may represent a convergent, late Carboniferous morphology.

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#### DAVID K. BREZINSKI

Carnegie Museum of Natural History
4400 Forbes Avenue
Pittsburgh, PA 15213, USA
e-mail brezinskid@carnegiemuseums.org
and
Maryland Geological Survey
Baltimore, MD 21218, USA

e-mail dbrezinski@dnr.state.md.us

# APPENDIX

Character states for ingroup relative to outgroup species

	12344	67890	11111 12345	11112 67890	22222 12345
Thigriffides roundyi	00000	00000	00000	00000	00000
Paladin chesterensis	21011	11012	00111	10100	11011
Paladin wilsoni	21011	11012	00111	00000	11011
Paladin genevievensis	11011	11012	00111	10000	11011
Paladin rosei	21011	11012	00111	10100	11011
Paladin longispina	21011	10012	00100	001?1	01?11
Paladin osagensis	11011	10011	10100	0???0	01110
Paladin morrowensis	00100	00101	11001	11011	01010
Paladin girtyianus	00201	00101	11000	01111	00011
Paladin helmsensis	00101	00101	11000	01011	00000
Paladin rarus	01101	00101	10000	01010	00011
Paladin moorei	00101	00101	11000	00111	00011
Paladin czarnieckii	11200	00101	11000	10221	00100
Paladin eichwaldi	11200	00101	11000	00221	00100
Paladin mucronatus	11200	00101	11000	00221	00100
Paladin cuspidatus	11100	00101	01100	00221	00111
Paladin maillieuxi	01100	00101	01000	00221	00110
Paladin lutugini	00200	1?110	01000	00211	01100
Paladin cervilatus	00200	1?11?	01000	10221	011?0
Paladin trigonopyge	00100	0?110	00100	00211	01110