

## ON A NEW INDEX OF HOST SPECIFICITY

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### Summary « On a new index of host specificity »

To facilitate consistent comparisons of host specificities among disparate taxa, a new index of host specificity is proposed. This index takes into account host range and host taxonomic rank, but not prevalence or intensity of infection. The central premise of this index is that a parasite lacking specificity at a higher taxonomic level should be considered to be less host specific than one lacking specificity at a lower taxonomic level. A ranking scheme is adopted that accommodates this notion. Host associations are ranked in order beginning with 1 for the most host specific scenario, corresponding to a taxon parasitizing 1 species, 1 genus, 1 family, 1 order, 1 class of host, and ending with a rank of 11,795,988,501, for the least host specific scenario, corresponding to a taxon parasitizing the greatest number of hosts allowed by this scheme (i.e., 1,000 species, 500 genera, 150 families, 75 orders, 5 classes). These unwieldy rank values are converted to more intuitive index values, which range from 1 to approximately 10, by taking the log (base 10) of the rank value. A Windows program to assist with the computation of rank and index value is available at <http://darwin.eeb.uconn.edu/specification/specification.html>. Standardization of the terminology applied to host specificity is proposed using this new index

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as follows : *oioxeny* for parasites restricted to a single species of host (index value of 0), *mesostenoxeny* for parasites restricted to species in a single genus of hosts (index values between 0 and 3.0004), *metastenoxeny* for parasites restricted to a single family of hosts (index values greater than or equal to 3.0004 and less than 5.574322), and *euryxeny* for parasitizing host in greater than one family (index values equal to or greater than 5.574322). In addition, ranges of index values are suggested to delimit the ecological terms 'specialist' and 'generalist.' The potential utility of the index was examined using data gathered from the literature on host ranges for approximately 700 species representing five phyla of parasitic organisms, and six common generalizations about host specificity.

#### **Key words**

Specificity Index - Host Range - Specialist - Generalist - Oioxeny - Stenoxeny - Euryxeny

#### **Résumé « Sur un nouvel indice de spécificité pour l'hôte »**

Afin de faciliter des comparaisons cohérentes des spécificités pour l'hôte entre des taxons quelconques, un nouvel index de spécificité pour l'hôte est proposé. Cet index prend en compte le spectre d'hôtes et la taxonomie de l'hôte mais ni la prévalence ni l'intensité de l'infestation. Le principe autour duquel s'articule cet index est qu'un parasite dépourvu de spécificité à un niveau taxonomique élevé doit être considéré comme moins spécifique pour l'hôte qu'un parasite dépourvu de spécificité à un niveau taxonomique moins élevé. Un système de rangs est adopté pour répondre à cette notion.

Les associations d'hôtes sont rangées dans un ordre qui commence à 1 pour le scénario le plus spécifique, correspondant à un taxon qui parasite 1 espèce, 1 genre, 1 famille, 1 ordre, 1 classe d'hôte, et se termine à la valeur 11.795.988.501 pour le scénario le moins spécifique, correspondant à un taxon qui parasiterait le plus grand nombre d'hôtes permis par le système (à savoir 1000 espèces, 500 genres, 150 familles, 75 ordres, 5 classes). Ces valeurs de rang peu maniables sont converties en valeurs d'indice plus intuitives qui vont de 1 à approximativement 10 en prenant le logarithme de base 10 de la valeur de rang. Un programme sur Windows pour le calcul du rang et de l'indice est disponible à <http://darwin.eeb.uconn.edu/specification/specification.html>.

Une standardisation de la terminologie appliquée à la spécificité d'hôte est proposée en utilisant le nouvel indice comme suit : *oioxénie* pour les parasites dont le spectre est réduit à une seule espèce d'hôte (indice = 0), *mésosténoxénie* pour les parasites se limitant aux espèces d'un même genre

d'hôtes (indice entre 0 et 3,0004), *métasténoxénie* pour les parasites n'exploitant que les membres d'une même famille d'hôtes (indice entre 3,0004 et 5,574322) et *euryxénie* pour les parasites se trouvant dans plus d'une famille d'hôtes (indice égal ou supérieur à 5,574322). Des intervalles des valeurs de l'indice sont suggérées pour délimiter les termes écologiques de « spécialiste » ou « généraliste ». L'utilité potentielle de l'indice a été testée en utilisant des données prises dans la littérature sur des spectres d'hôtes d'environ 700 espèces représentant cinq phylums de parasites, et six généralisations habituelles relatives à la spécificité.

### Mots-clés

Spécificité d'hôte - Indice de spécificité - Spectre d'hôtes - Spécialiste - Généraliste - Oioxénie - Sténoxénie - Euryxénie

The concept of host specificity is among the most intriguing aspects of Parasitology. The literature exploring the various facets of this concept is extensive. Generalizations about the specificities of parasite groups abound in the literature (e.g., Rohde, 1978 ; Noble *et al.*, 1989 ; Holmes, 1990 ; Poulin, 1992 ; Sasal *et al.*, 1999). Within the last 50 years alone, two major symposia have been held to address this topic ; both of these resulted in interesting compilations of papers (Baer, 1957 ; Chabaud, 1982). By now it is apparent that all parasites exhibit some degree of specificity for the kinds of organisms they parasitize at any single life-stage. It is also apparent that the extensive diversity of known parasites collectively exhibits a wide spectrum of host associations (e.g., Rohde, 1978 ; Noble *et al.*, 1989 ; Poulin, 1992 ; Combes, 2001). Despite the continuous nature of this spectrum of host associations, the utility of distinguishing between parasites falling at either of its extremes is widely recognized. However, neither the specific boundaries of the extremes, nor the terminology for characterizing host specificity at either extreme are universally agreed upon.

At one end of this spectrum are taxa that parasitize only a single species of host. A number of terms have been used to describe species exhibiting such restricted host specificity. Sandground (1929) considered such taxa to be 'monoxenous,' but noted that this term could also be applied to parasites associated with several very closely related hosts. Marshall (1981) restricted use of 'monoxeny' to parasites known only from one host species. Hargis (1957) referred to such taxa as exhibiting 'species-specificity.' Ludwig (1982) used the term 'stenohospitalic.' Euzet and Combes (1980) suggested the term 'oioxenous' for parasites restricted to one species of host ; Lambert (1982) followed suit. Such taxa have also been described as 'host-specific' by many authors (e.g., Moss & Wojcik, 1978), and as 'specialists' in much of the ecological literature (e.g., Dusek *et al.*, 1998 ; Sasal *et al.*, 1999). The

latter two terms have also been applied to parasite species exhibiting slightly more relaxed host specificities ; most commonly, those parasitizing a suite of closely related hosts (e.g., Esch & Fernandez, 1993). Cumming (1998 ; pg. 380) very precisely defined specialists as « those species that have been collected more than ten times and where 90% or more of collections are from a single taxon. » Some of the problems associated with the lack of precision of terms used to characterize ‘species specificity’ in specific taxonomic groups have been articulated by others (e.g., Williams, 1986).

The boundary at the other end of the spectrum is less well defined. A diversity of terms has been proposed to describe a lack of specificity, and the exact concept to which each of these terms has been applied has also varied. Parasites designated as lacking in specificity have included : species parasitizing two or more closely related species of hosts (e.g., Ludwig, 1982), species parasitizing many species of hosts (e.g., Sandground, 1929), and species parasitizing many distantly related species of hosts (e.g., Esch & Fernandez, 1993). Sandground (1929) referred to taxa parasitizing a diversity of hosts as ‘polyxenous.’ Marshall (1981) more precisely defined ‘polyxenous’ taxa as those parasitizing more than one family of host. Euzet and Combes (1980) proposed the term ‘euryxenous’ for species parasitizing a number of distantly related hosts. Hoogstraal and Aeschlimann (1982) developed an elaborate scheme of terms that incorporated specificity of both immature and adult tick stages ; taxa exhibiting the least amount of host specificity (i.e., in which both adults and immatures « are both catholic in host acceptability » [p. 157]), were referred to as ‘nonparticular’ or ‘telotropic.’ Ludwig (1982) termed species exhibiting wide host specificity ‘euryhospitalic.’ Among ecologists, such taxa are most commonly referred to as ‘generalists.’ However, application of the latter term has also varied. To some (e.g., Dusek *et al.*, 1998), a generalist is a parasite occurring on several host species ; to others (e.g., see Esch & Fernandez, 1993), it is a parasite associated with a broad range of unrelated host taxa. Cumming (1998) very precisely defined generalists as any species collected from greater than 50 species of host.

Many authors have sought to recognize one or more categories of specificity for parasites exhibiting host associations falling between the two extremes described above. For example, Sandground (1929) referred to parasites whose potential hosts are few in number as ‘oligoxenous.’ A category for parasites restricted to a closely related group of hosts is widely recognized. But, again, a number of different terms have been applied. For example, a natural group of monogeneans parasitizing a natural group of fish species was described as exhibiting ‘supraspecific specificity’ by Hargis (1957). Hoogstraal and Aeschlimann (1982) proposed the terms ‘ditropic-A,’ ‘ditropic-B,’ ‘telotropic-moderate,’ etc. for such taxa, depending on how

taxon-limited the specificity exhibited by the parasite was. Euzet and Combes (1980) proposed the term 'stenoxenous' for parasites restricted to a small group of related hosts, usually the same genus or family. Marshall (1981) recognized two categories of moderately host-specific taxa ; he termed parasites associated with two or more congeners 'oligoxenous' and those associated with two or more genera in the same family of host 'pleioxenous.' Llewellyn (1982) advocated a somewhat different approach that emphasized degree of host relationship over number of hosts. He proposed the term 'phylogenetic specificity' for related parasites occurring on related hosts and 'ecological specificity' for parasites occurring on unrelated hosts, sharing a common habitat.

Although some of this terminology has subsequently been adopted by other authors (e.g., Lambert, 1982 followed Euzet & Combes, 1980), no set of terms or concepts has been universally accepted. It remains commonplace for authors to characterize parasites as exhibiting, for example, « little host specificity » (Steinlein *et al.*, 2001, p. 611), « very broad host specificity » (Podlipaev, 2001, p. 648), or « a rather high level of host specificity » (Telford *et al.*, 2001, p. 904). Clearly these types of characterizations limit the power of comparisons across taxa.

A metric or index of host specificity would greatly facilitate precise, reproducible specificity comparisons. However, the quantitative measures available are surprisingly few in number. In addition, as noted by Lymbery (1989), there is variation in the meaning intended by different authors using the term 'host specificity.' In much of the ecological literature this term is used in the sense of 'host preference' or 'host specificity' sensu Lymbery (1989). The available indices primarily reflect this concept of 'host specificity.' Rohde (1980) evaluated three host specificity indices and their potential applications. The first of these relies to a large extent on intensities of infection. The second relies heavily on prevalence of infection. The third is based on probability theory and relies to some extent on number of host species infected relative to the number of host species examined, although one of the variations of this index accommodates host range. Price (1980) suggested that an index of specificity for a higher taxon, such as family, might be calculated as the percent of species in that category utilizing only a single host. May's (1990) method for calculating effective specialization is considered by some (e.g., Mawdsley & Stork, 1997) to be a measure of host specificity. However, that metric was developed primarily in response to the interest in determining accurate estimates of global arthropod diversity (e.g., May, 1979 ; Erwin, 1982 ; Mawdsley & Stork, 1997 ; Ødegaard, 2000) and, as a consequence, is a host-centric rather than parasite-centric metric. This is also true of the index described by Marshall (1981). Alternatively, Poulin (1992, 1997) used number of known host species (or 'host range' sensu

Lymbery, 1989) to estimate host specificity at the species level. To estimate host specificity for higher taxa, Poulin (1992) used percentage of parasite species in the taxon known from only a single host, and mean number of known host species per parasite species. With the exception of the metrics utilized by Poulin and one version of Rohde's third index, all of the above indices are essentially measures of either host preference, or host specificity in its more ecological sense (i.e., sensu Lymbery, 1989). None of the above metrics take host relationships into account.

It is 'host specificity' in the sense of 'host range' that is of interest to us here. However, we prefer a slightly more comprehensive concept combining information on host range (i.e., number of host species parasitized) and the phylogenetic limits of this host range. Because none of the above indices were designed to account for any element of host relationships, all of these metrics fail to distinguish between the specificity of a taxon parasitizing a group of related hosts from that of a taxon parasitizing a suite of more distantly related hosts, and thus are inappropriate indices of the concept of host specificity we are striving to measure here.

The development of an index of host specificity that incorporates data on both number and basic taxonomic position of hosts is the subject of this chapter. This seemed an appropriate endeavor to undertake in a book honoring Professor Louis Euzet, for Professor Euzet has not only contributed to the formalization of the terminology of host specificity, but has also contributed enormously to the taxonomy and systematics of two of the most host-specific parasite groups known. This index was designed to measure host specificity sensu Euzet and Combes (1980) rather than, for example, host specificity as defined by Lymbery (1989). Thus, it does not take into account prevalence or intensity of infection. In order to distinguish the index described here from other specificity indices, we have adopted the abbreviation *HS*.

### Index of Host Specificity (*HS*)

The issue of how best to incorporate basic host phylogenetic information was perhaps the biggest challenge in the development of the index. Although a strictly tree-based system might reflect the phylogenetic components of the hosts most effectively, it would only be useful for comparing those parasites for which well-corroborated host phylogenies were available and only for comparing parasites in which the tree of host relationships for one was a subset of the tree of host relationships for the other. In addition, if the comparisons were to be meaningful, the host trees utilized would need to include all species in each host group, because, barring extensive polytomy, the number of nodes in a tree is generally correlated with number of species in the tree and thus missing taxa would significantly affect the assessment of relative phylogenetic specificity. As a result, a tree-based index would also

be severely biased by differences in the extent of extinction among host groups because host range information is generally available only for extant taxa. However, even if fully resolved trees of all host taxa were available, the issue of which nodes to consider equivalent when comparing trees generated for parasites of host taxa consisting of different numbers of included species would be difficult to resolve. In short, although a tree-based index might be useful for analyses of co-evolution, it is not clear that such a system could be used to compare host specificities of distantly related taxa.

Instead, the Linnaean hierarchy from species through class, was adopted as the criterion to be used to represent host relationships. This hierarchy was incorporated by requiring that each parasite be scored for the number of species, genera, families, orders and classes of taxa it parasitizes. Taxonomic categories above class were not included because examination of the host associations of a disparate suite of 700 parasite taxa (see Appendix I) failed to identify a single taxon parasitizing more than 1 phylum (or kingdom) at any single life-stage; although, several taxa parasitizing more than a single class of hosts were identified. If taxa parasitizing more than a single phylum or kingdom, at the same life-stage, exist, they are rare and are best treated on an individual basis when they arise. It should be noted, however, that the trichodinid ciliates may prove to be just such an exception (Van As & Basson, 1987). In basing our index of specificity on the rank at which specificity is detected, we are implicitly assuming that taxonomic categories across host groups are comparable, that higher taxa represent monophyletic units, and that phyletic distance is a more important component of specificity than the number of species parasitized. It was our hope that at least discrepancies in taxonomic rank among disparate host taxa would be buffered by our focus on major taxonomic categories only. There seemed little reason to believe that differences in classification among host groups were major enough to transcend the basic major taxonomic categories of classification. Thus, for example, even if genera of fish and mammals are not quite equivalent, they are not so different that a genus of fish is the equivalent of a family of mammals. Our index is a move in the direction of a tree-based index of specificity that, we hope, represents a useful compromise between incorporating relevant phylogenetic information on hosts and allowing broad comparisons among parasites infecting distantly related host taxa.

One of the primary features we wished the index to reflect was the premise that a parasite lacking specificity at a higher taxonomic level (e.g., a parasite hosted by two genera) should be considered less host specific than one lacking specificity at a relatively lower taxonomic level (e.g., a parasite hosted by any number of species in a single genus). The higher the taxonomic category at which this lack of specificity is exhibited, the less host

specific the taxon. A basic ranking scheme was adopted that would reflect this premise. In this scheme, all possible numbers of hosts in each taxonomic category are uniquely ranked, in sequence, beginning with 1 species in 1 genus in 1 family in 1 order in 1 class of host (i.e., the most host specific scenario) and ending with the maximum number of species, genera, families, orders and classes (i.e., the least host specific scenario).

It is easiest to understand how a parasite species is assigned a unique rank by describing how data on host ranges is used to determine the rank assigned to a species. Let  $S$ ,  $G$ ,  $F$ ,  $O$ , and  $C$  be the number of species, genera, families, orders, and classes, respectively, in which a particular taxon occurs. We can then represent the host distribution of any parasite species as a decimalized index, namely  $C.O.F.G.S$ . We can then list all possible decimalized indices by enumerating all combinations of  $S$ ,  $G$ ,  $F$ ,  $O$ , and  $C$ , provided that we specify the maximum number of taxa a parasite can infect at any hierarchical level. Let  $S_{\text{Max}}$ ,  $G_{\text{Max}}$ ,  $F_{\text{Max}}$ ,  $O_{\text{Max}}$ , and  $C_{\text{Max}}$  be those maximum numbers. We can then sort the decimalized indices from most specific (1.1.1.1.1) to least specific ( $C_{\text{Max}}.O_{\text{Max}}.F_{\text{Max}}.G_{\text{Max}}.S_{\text{Max}}$ ) by sorting first on the number of classes in which a species is found, then on the number of orders, then on the number of families, then on the number of genera, then on the number of species. Any parasite species will have a unique position in this sorted list ranging from 1 for the most specific taxa to some very large number that depends on  $S_{\text{Max}}$ ,  $G_{\text{Max}}$ ,  $F_{\text{Max}}$ ,  $O_{\text{Max}}$ , and  $C_{\text{Max}}$  for the least specific. That unique position is the rank assigned to a parasite species. The C++ code for a more computationally efficient algorithm is included as Appendix II.

However, before the total number of possible ranks could be determined, it was necessary to determine the maximum number of host taxa to allow at each of the five taxonomic levels (i.e.,  $S_{\text{Max}}$ ,  $G_{\text{Max}}$ ,  $F_{\text{Max}}$ ,  $O_{\text{Max}}$ , and  $C_{\text{Max}}$ ). Of primary concern was that these values be set large enough to accommodate the specificity exhibited by even the least host specific parasite system; the larger the maximum values allowed, the more likely this is to be the case. However, the larger these values, the greater the number of possible ranks, and the more unwieldy the index. We initially considered setting these numbers to the total number of known taxa in each category. So, for example, these values might be set to 2 million species, 100,000 genera, etc. However, this strategy results in an outrageously large number of ranks (possibly as many as  $10^{15}$ ), and, given that no parasite taxon is known to parasitize all species, this strategy was determined to be unnecessarily conservative. It seemed more appropriate to assign these maximum values following examination of published host data. Thus, data from the literature were assembled on numbers of species, genera, families, orders, and classes

parasitized by each of 700 species representing a diversity of host/parasite systems, at least some of which are generally considered to be notoriously nonhost-specific. These data are summarized in Appendix I.

One additional consideration played a role in the determination of these maximum values. Although host specificity is most commonly investigated at the level of parasite species, it has been addressed at higher taxonomic categories such genus, family, or order (e.g., Hargis, 1957 ; Moss, 1979 ; Rohde, 1979 ; Marshall, 1981 ; Kim, 1982 ; Ludwig, 1982 ; Cumming, 1998 ; Caira & Jensen, 2001). If the index were to have utility for addressing questions relating to higher taxonomic categories of parasites, the maximum values would need to be sufficiently high so as to accommodate all host taxa at each taxonomic category parasitized by, for example all species in a genus, or all species in a family of parasite. The maximum numbers of host taxa parasitized by any single species of parasite in each of the five taxonomic categories seen in Appendix I are : 58 species, 37 genera, 17 families, 12 orders and 2 classes. The maximum numbers of host taxa parasitized by any single genus of parasite in each of the five taxonomic categories seen in Appendix I are : 100 species, 71 genera, 45 families, 18 orders and 2 classes. The maximum numbers of host taxa parasitized by any single family of parasite in each of the 5 taxonomic categories seen in Appendix I are : 139 species, 97 genera, 55 families, 19 orders and 2 classes. Given that these data represent only a tiny fraction of all parasite taxa, and less host specific taxa are likely to exist, it seemed appropriate to set the maximum values substantially higher than any seen among these data. We chose maximum values of : 1,000 species, 500 genera, 150 families, 75 orders and 5 classes. These numbers result in a total number of 11,795,988,501 ranks. The highest rank represents the least host specific scenario allowed under this configuration and would be assigned to any taxon parasitizing 1,000 species, 500 genera, 150 families, 75 orders and 5 classes of hosts. So, a taxon parasitizing 1 species in 1 genus in 1 family in 1 order would receive the rank of 1. A taxon parasitizing 1,000 species in 500 genera in 100 families in 50 orders receive the rank of 11,795,988,501. Taxa exhibiting all other possible combinations of host associations receive ranks falling between these two values. For example, a taxon parasitizing 1,000 species in 1 genus in 1 family in 1 order in 1 class would receive a rank of 1,000. A taxon parasitizing 2 species in 2 genera in 1 family in 1 order in 1 class would receive a rank of 1001, etc. The corresponding decimalized index value for the most host specific parasite would be : 1.1.1.1.1 ; the decimalized index value for the least host specific parasite allowed by this system would be : 1000.500.150.75.5.

Unfortunately, the extensive range of the raw rank values and the com-

plex representation of the decimalized index make them awkward to use and difficult to compare. Perhaps more importantly, the difference in specificity between a species that parasitizes only 1 species and one that parasitizes 2 species in 1 genus seems far greater to us than the difference between a species that parasitizes 10 species in 5 genera in 1 family and one that parasitizes, 11 species in 5 genera in 1 family. By taking the logarithm (base 10) of the raw rank value we change our metric from ranks of 1-11,795,988,501, to an index of 0 to ~10 (or more precisely, 10.07173434). A taxon parasitizing 1 species in 1 genus in 1 family in 1 order in 1 class (i.e., with a rank of 1) would be assigned an index value of  $\log_{10}(1)$ , or 0. A taxon parasitizing 1,000 species in 500 genera in 150 families in 75 orders in 5 classes (i.e. with the rank of 11,795,988,501) would be assigned the index value of  $\log_{10}(11,795,988,501)$ , or 10.07173434. The lower the index value, the higher the degree of specificity exhibited by a taxon.

The conversion to a  $\log_{10}$  scale has several consequences that are important to note. Perhaps most importantly, this conversion results in the index being of much finer scale at its lower values. Thus, for example, taxa parasitizing hosts in 1 order in 1 class and any number of genera or species (below 500 and 1,000, respectively) will be assigned index values between 0 and 7.65957204. Whereas, those parasitizing any combination of numbers of taxonomic categories in 2 to 75 orders in 2 to 5 classes will receive values between 7.65957204 and 10.0717343. This attribute reflects our decision to emphasize proportional differences in degree of specificity rather than absolute differences in specificity rank. The detailed comparison of taxa exhibiting greater specificities (lower indices) may be more interesting than the probing of those with lesser specificities (higher indices). An appreciation for the degrees of host specificity represented by the various index values can be obtained by examining Table I, in which the host associations represented by each whole number index value are presented. In order to illustrate the variation in scale noted above, index values of 7.5, 8.5, and 9.5 are also included.

To facilitate computation of the rank and its corresponding index value from the number of host species, genera, families, orders and classes parasitized, a Windows program (including source code in C++) is available from <http://darwin.eeb.uconn.edu/specification/specification.html>. The program uses our choice of values for maximum number of species, genera, families, orders and classes by default (i.e., 1,000, 500, 100, 75, and 5, respectively), but the algorithm is completely general (see Appendix II). Although users can modify the default values of these maximums by selecting appropriate options, we encourage users to resist changing the default values unless they encounter a taxon that exceeds the maximum values described above. Changes in these default values will require recalculation of all index values

HOSTS						Rank	Index (HS)
No. species	No. genera	No. families	No. orders	No. classes			
1	1	1	1	1	1	1	0.00000000
10	1	1	1	1	1	10	1.00000000
100	1	1	1	1	1	100	2.00000000
1000	1	1	1	1	1	1000	3.00000000
55	11	1	1	1	1	10000	4.00000000
565	106	1	1	1	1	100000	5.00000000
455	297	3	1	1	1	1000000	6.00000000
877	283	28	1	1	1	10000000	7.00000000
778	114	97	1	1	1	31622777	7.5000001
944	41	28	3	1	1	100000000	8.00000000
906	96	20	8	1	1	316227766	8.50000000
325	171	41	25	1	1	1000000000	9.00000000
633	178	122	18	2	1	3162277660	9.50000000
573	120	38	16	5	1	10000000000	10.00000000
1000	500	150	75	5	1	11795988501	10.07173434
<b>Host specificity categories</b>							
oioxeny	1	1	1	1	1	1	0.00000000
lower boundary of mesostenoxeny	2	1	1	1	1	2	0.30103000
lower boundary of metastenoxeny	2	2	1	1	1	1001	3.00043408
lower boundary of euryxeny	2	2	2	1	1	375251	5.57432186

Table 1. Host specificity index landmarks.

obtained using our original default values if comparisons are to be made across datasets. We recommend that investigators using this index present at least the rank and index value for each parasite taxon investigated. However, it would be even more useful to present the number of species, genera, families, orders and classes of hosts in each case. This could be done in a concise manner using the decimalized index scheme described above. These host data are necessary if the maximum value of any taxonomic category is altered and recalibration of the index is required. Although these values can be retrieved from the raw rank, the retrieval process is very tedious.

So as to increase the precision of the terminology used to characterize the host associations of parasites (Combes, 2001), we suggest that quantitative definitions based on the index developed here would be very appropriate. We propose a system that allows the basic terms suggested by Euzet and Combes (1980) to be applied with only slight modification. We prefer these terms over those of, for example, Sandground (1929) and Marshall (1981), because the terms used by the latter authors have a history of alternate usage associated with the number of hosts parasitized over the course of the life cycle of a parasite (i.e., monoxenous = one host; polyxenous = greater than one host). We think it advisable not to confuse the issue of number of hosts in the complete life-cycle of a taxon with number of hosts parasitized by one life-stage of a taxon. We propose the following:

- 1) An *oioxenous* taxon is restricted to a single species of host and has an index value of 0.

2) Two subcategories of stenoxenous specificity that reflect the two intermediate categories of specificity of Marshall (1981) should be recognized :

a) A *mesostenoxenous* taxon parasitizes greater than 1 species of host, but is restricted to a single genus of host ; such taxa have index values greater than 0 and less than 3.000434077. Note that because the highest possible index value of a parasite restricted to a single genus of host is 3.00000000 (i.e. 1,000 species, 1 genus, 1 family, 1 order and 1 class), it is necessary to calculate and present index values to at least four decimal places if the difference between this and the next rank is to be detected.

b) A *metastenoxenous* taxon parasitizes more than one genus of host, but is restricted to a single family of host ; such taxa have index values greater than or equal 3.00043400 and less than 5.574321858. Once again, because the highest possible index value of a parasite restricted to a single family of host is 5.574320701 (i.e. 1,000 species, 500 genera, 1 family, 1 order and 1 class), it is necessary to calculate and present index values to at least 6 decimal places if the difference between this and the next rank is to be detected.

3) A *euryxenous* taxon parasitizes more than one family of host ; such taxa have index values equal to or greater than 5.574322.

Because ecologists so commonly use the terms 'specialist' and 'generalist' to describe host associations, we would also like to suggest index values to delimit these terms. Although, as noted above, there is not complete consensus among ecologists as to the boundary between these two terms, we suggest that taxa exhibiting oioxeny or mesostenoxeny should be considered specialists and those exhibiting metastenoxeny or euryxeny should be considered generalists. Thus, specialists would be taxa with index values less than 3.0004 and generalists would those with index values of 3.0004 or greater.

### **Exploring the Index**

The index and the data provided in Appendix I are used below to explore several of the major generalizations about parasite specificity that exist in the literature. This allows us to demonstrate the utility of the index. In addition, we attempt to gain additional insight into patterns of specificity by comparing an index value calculated from a rank generated for a higher parasite taxon with an index value that represents a mean of the index values of the species associated with that higher parasite taxon.

The host association data provided in Appendix I were obtained from a variety of sources. While it was beyond the scope of this work to attempt to verify the specific identities of the host or parasite taxa, we have attempted to standardize the higher categories to which each of the host species of record was assigned. The classification schemes followed were : Fishbase

(2002) for fishes, Nowak (1991) for mammals, Clements (2000) for birds, and Stevens (2001) for plants. The host data presented here include natural infections only; data from experimental infections were excluded. This is particularly relevant to the mermithid nematodes. Hosts suspected to be the result of accidental infections were also excluded. In general, the hosts presented are only those utilized by the adult parasite.

The 700 species in Appendix I were chosen to represent the major groups of metazoan parasites, while also representing a diversity of endoparasites and ectoparasites, and a number of distinct host groups. As noted above, a special point was made to include groups generally considered to be lacking in host specificity. This was the primary factor that led to the inclusion of the mycorrhizal fungi. Groups generally perceived to be very host specific are also represented. Whenever possible, data were collected for multiple species in multiple genera of parasites in a family so as to allow comparisons of index calculations for parasites of different taxonomic levels, as well as to allow these numbers to be compared to mean index calculations at these levels. In deference to the taxonomic interests of Professor Euzet, there is a slight emphasis on monogeneans.

The overall organization of Appendix I is by parasite phylum. Data provided include parasite family, genus and species identity. For each species of parasite the number of species, genera, families, orders and classes of hosts parasitized, the source of these data, the rank, and the index value assigned to each species based on these associations are provided. The endoparasites consist of: 1) Nematoda (12 mermithids, 1 diplogasterid, 4 steinernematids, 2 heterorhabditids, 1 neotylenchid, 1 allantonematid, and 26 cystidicolids), 2) Digenea (49 hemiurids, 49 leptocreadiids, 18 allocreadiids), 3) Cestoda (3 cephalochlamydids, 8 litobothriids, and 35 onchobothriids), and 4) Acanthocephala (19 neoechinorhynchids). The ectoparasites consist of: 1) Monogenea (34 gyrodactylids, 11 dactylogyrids, 39 monocotylids, 2 capsalids, 4 hexabothriids, 5 mazocreadiids, 4 discocotylids, 5 diclydophorids, 15 microcotylids, 8 gastrocotylids, 2) Copepoda (79 caligids and 28 panderids), 3) Acari : 'ticks' (19 ixodids), 4) Acari : 'mites' (114 harpyrhynchids), 5) Siphonaptera (53 ceratophyllids), and 6) Mallophaga (8 phyllopterids). In addition, 46 mycorrhizal fungi of the order Aphyllonales were included.

#### I. Parasite species indices ( $HS_s$ )

Index values for individual species (designated ) are plotted in Figure 1. Shaded regions indicate the recommended index ranges for oioxenous, mesostenoxenous, metastenoxenous, and euryxenous taxa, as described above. Several basic trends are apparent from these data. For example, species in some groups, such as the onchobothriid tapeworms and many of the monogeneans, all exhibit similar degrees of specificity. These species

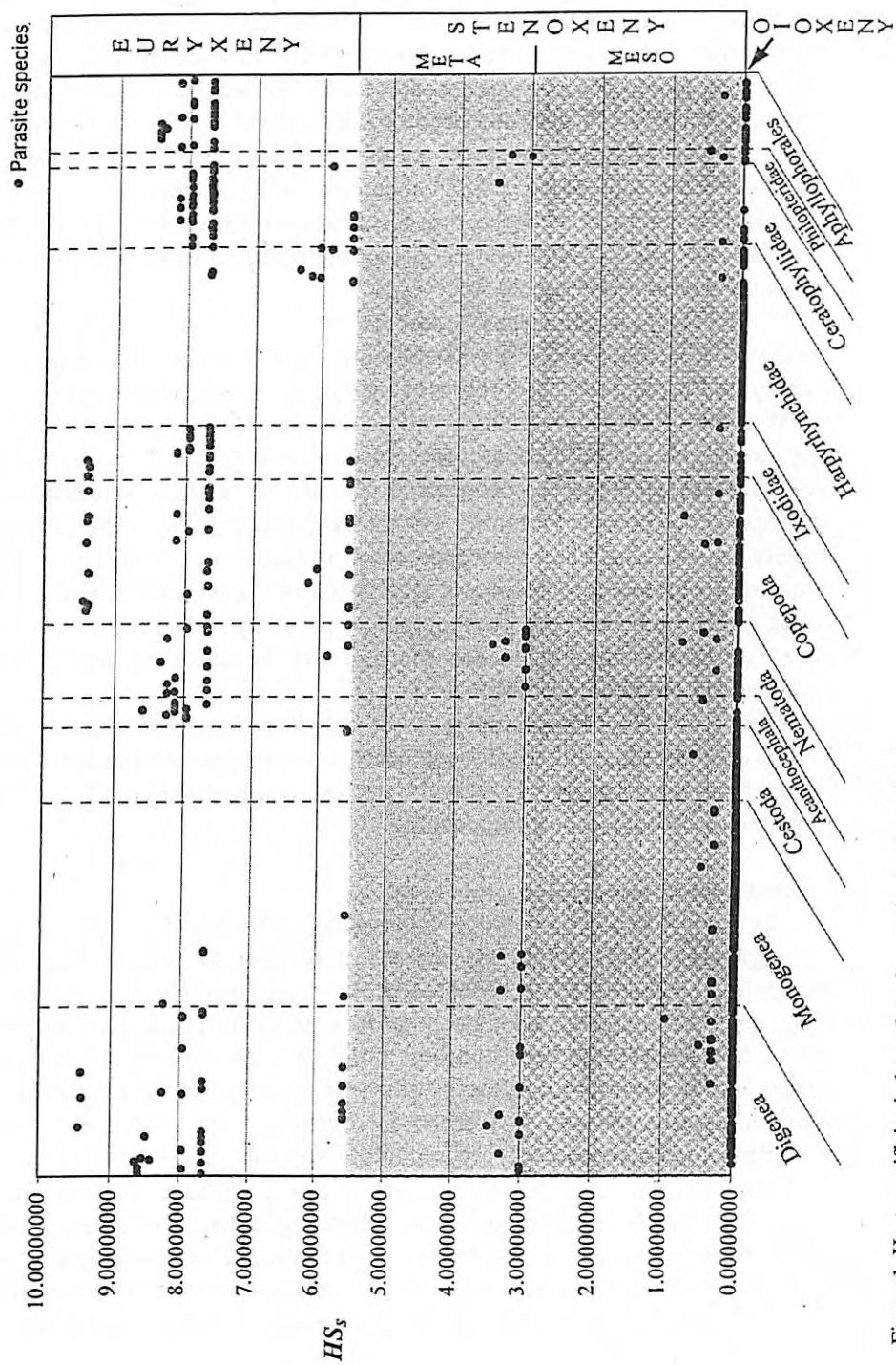


Figure 1. Host specificity index values for species in Appendix 1.

tend to be very host specific ; most are oioxenous (i.e., have index values of 0). However, there are other groups, such as the copepods and the digeneans, that collectively include species with a wide diversity of host specificities. Both of these groups include many species that are oioxenous, but many more species that are among the most euryxenous of the species included in this study. Both groups also include a number of stenoxenous species. However, whereas the digeneans include both mesostenoxenous and metastenoxenous species, the non-oioxenous copepods are entirely metastenoxenous. The mycorrhizal fungi are interesting in that species are either oioxenous or euryxenous.

Oioxenous specificity was exhibited by one or more species in all but 4 of the 31 families in the 5 phyla examined. In the higher parasite taxa (i.e., family or above) that are represented by 8 or more species in Appendix I, the greatest numbers of oioxenous taxa were seen among the Platyhelminthes in the monogenean families Dactylogyridae, Monocotylidae, and Microcotylidae, and in the tapeworm families Onchobothriidae and Litobothriidae, as well as in the arthropod family Harpyrhynchidae ; in each case, approximately 90% or greater of the species had an index value of 0. The smallest number of oioxenous species was seen in the arthropod families Ceratophyllidae (fleas) and Ixodidae (ticks), and in the Platyhelminthes, in the digenetic family Allocreadiidae. Less than 30% of the species in each of these families had index values of 0.

The least host specific species were caligid and pandarid copepods, and hemiurid digeneans. Of the 12 species that exhibited index values greater than 9.3, 5 were caligid copepods, 4 were pandarid copepods, and the remaining 3 were hemiurid digeneans.

## II. Generic indices

Two types of generic index values can be obtained from this system. Mean generic values can be calculated by averaging the index values of all species in a genus ( $HS_s$ ). Alternatively, an index value for the genus itself ( $HS_g$ ) can be generated by determining the rank for the genus from the number of host species, genera, families, orders and classes parasitized by all species in the genus. Both of these values are presented in Table II for the 79 non-monotypic (metazoan) genera from Appendix I for which data on specific identifications of hosts were available. Note that these two index values are conspicuously different for a number of genera. Instances in which these values are similar are indicative of substantial host overlap among the species in the genus. For example, in the ceratophyllid flea genus *Monopsyllus*, the mean generic index value ( $HS_s$ ) for the six included species was  $7.89 \pm 0.08$ , and the index value for the genus ( $HS_g$ ) was 8.14. Indeed, the 80

Family	n*	Genus	Mean Generic Index ( $\bar{HS}_s$ )	HOSTS						Rank	Generic Index ( $HS_s$ )	Generic Index Ratio ( $\bar{HS}_s / HS_s$ )
				No. species	No. genera	No. familles	No. orders	No. classes				
ALLOCREADIIDAE	4	Bunodera	6.80520758	39	28	12	9	1	355973905	8.55141816	0.7	
	12	Crepidostomum	4.60066378	100	41	20	15	2	3011437755	9.47877389	0.4	
HEMIURIDAE	4	Aponurus	1.91489301	4	4	4	3	1	90953349	7.95881869	0.2	
	3	Brachyphallus	6.38497096	20	16	11	8	1	312908095	8.49541680	0.7	
	4	Derogenes	4.27683193	50	28	16	10	2	2803855722	9.44775566	0.4	
	3	Dinomia	3.85839706	5	4	4	2	1	45665799	7.65959106	0.5	
	6	Dinurus	2.40827895	7	7	4	1	1	1125740	6.03143810	0.3	
	5	Ectenurus	1.11486437	4	4	3	1	1	750499	5.87535012	0.1	
	7	Genolinea	4.63702690	29	24	13	6	1	227158274	8.35632856	0.5	
	4	Gonocerca	1.91489539	4	3	2	2	1	45664801	7.65958157	0.2	
	2	Gonocerella	0.00000000	2	2	2	2	1	45663801	7.65957206	0.0	
	4	Hemurus	2.35496561	20	15	8	6	2	2629434956	9.41986243	0.2	
LEPOCREADIIDAE	2	Dihemistophenus	0.00000000	1	1	1	1	1	1	0.00000000	0.0	
	2	Labifer	0.00000000	2	2	2	1	1	375251	5.57432186	0.0	
	12	Lepidapedon	1.30370611	17	11	6	4	1	136615131	8.13549880	0.1	
	8	Lepocreadium	0.00000000	5	5	5	3	1	91697855	7.96235918	0.0	
	4	Neolepidapedon	0.31381813	13	3	3	2	1	46038061	7.66311702	0.0	
	8	Opechona	3.90503962	21	8	8	6	1	225316034	8.35279210	0.4	
	35	Gyrodactylus	1.30460880	40	26	24	10	1	402589229	8.60486215	0.1	
	4	Antcyrocephalus	0.00000000	4	4	4	3	1	91325602	7.96059254	0.0	
DACTYLOGYRIDAE	2	Pseudohalitrema	0.00000000	2	2	2	1	1	375251	5.57432186	0.0	
	4	Callicoryle	0.00000000	3	3	3	2	1	46038051	7.66311693	0.0	
MONOCOTYLIDAE	3	Decacotyle	0.00000000	2	2	2	1	1	375251	5.57432186	0.0	
	2	Dendramonocotyle	2.78716151	4	3	2	1	1	376251	5.57457766	0.4	
	2	Dianthus	0.00000000	2	2	2	1	1	375251	5.57432186	0.0	
	3	Espruthotrema	0.00000000	3	3	3	2	1	46038051	7.66311693	0.0	
	4	Heterocotyle	0.00000000	3	3	2	1	1	376250	5.57347651	0.0	
	3	Merizocotyle	0.00000000	3	3	3	1	1	749501	5.87477222	0.0	
	6	Monocotyle	0.10034333	5	3	1	1	1	2002	3.30146407	0.0	
HEXABOTHRIIDAE	4	Neheterocotyle	0.00000000	3	3	3	1	1	749501	5.87477222	0.0	
	3	Thaumatoctyle	0.00000000	3	2	2	1	1	375252	5.57432302	0.0	
	2	Squadronchacotyle	0.00000000	2	2	2	2	1	45663801	7.65957206	0.0	
MAZOCRAEIDAE	2	Clapicecotyle	0.00000000	1	1	1	1	1	1	0.00000000	0.0	
	2	Mesocraeoides	0.00000000	2	2	1	1	1	1001	3.00043408	0.0	
DISCOCOTYLIDAE	3	Togia	0.00000000	3	3	2	1	1	376250	5.57547651	0.0	
	4	Chorictyle	0.07523750	5	4	3	2	1	46039050	7.66312633	0.0	
MICROCOTYLIDAE	3	Axonoides	0.00000000	2	1	1	1	1	2	3.01030000	0.0	
	2	Heteroxazine	0.00000000	2	2	2	1	1	375251	5.57432186	0.0	
	7	Microcopule	0.00000000	7	6	4	1	1	1124746	6.05105446	0.0	
	11	Acanthobothrium	0.00000000	5	3	3	2	1	46038053	7.66311695	0.0	
LITOBOTHRIIDAE	6	Callichothrium	0.00000000	6	1	1	1	1	6	0.77815125	0.0	
	9	Pedilobothrium	0.00000000	4	4	2	1	1	377248	5.57662695	0.0	
	6	Playbothrium	0.00000000	5	4	1	1	1	2999	3.47697647	0.0	
	8	Litobothrium	0.00000000	4	3	2	1	1	376251	5.57547766	0.0	
PSEUDOPHYLLIDAE	2	Cephalochlamys	5.57432359	6	2	2	1	1	375255	5.57432649	1.0	
	14	Neoechinorhynchus	5.78121852	56	36	15	10	2	280303121	9.4470104	0.6	
CYSTIDICOLIDAE	6	Ascarophis	4.00894550	30	17	12	7	1	270215379	8.43171006	0.45	
	4	Cystidicola	3.97918832	23	12	7	6	1	224951727	8.35208933	0.47	
	2	Cystidicoloides	4.06656281	16	9	5	5	1	180411685	8.25626466	0.45	
	3	Salvelinema	1.10077698	6	2	1	1	1	1005	3.00216606	0.36	
	7	Spinitectus	2.28809292	26	23	14	8	1	314004917	8.49693645	0.22	
HETERORHABDITIDAE	2	Heterorrhabdites	0.00000000	2	2	1	1	1	1001	3.00043408	0.0	
	2	Hydromermis	4.52630004	11	9	3	1	1	755476	5.87822067	0.71	
MERMITIDAE	2	Isomeris	2.03980766	9	2	1	1	1	1008	3.00046053	0.67	
	4	Naeplectana	6.05148415	17	15	10	3	1	93459132	7.97103976	0.75	
CALIGIDAE	2	Anurites	0.00000000	2	2	2	2	1	45663801	7.65957206	0.0	
	41	Caligus	2.79633133	85	71	45	18	2	313939080	9.49684653	0.25	
	29	Lepocyclopus	2.96787712	80	56	33	12	2	2893405801	9.46140933	0.31	
	2	Pupulina	0.00000000	1	1	1	1	1	1	0.00000000	N	
	2	Demoleus	3.82978603	2	2	2	2	1	45663801	7.65957206	0.56	
PANDARIDAE	3	Dinemoura	5.21143516	7	6	5	2	1	46784552	7.67010248	0.67	
	2	Echthrogaleus	8.53575874	8	8	7	3	2	2496552365	9.39734068	0.96	
	5	Nestipus	2.23019176	8	5	3	2	1	46040049	7.66313578	0.29	
	8	Pandarus	4.26642669	18	14	9	6	2	2629801246	9.41992293	0.45	
	2	Perisopus	7.49709281	14	10	9	6	2	2629797288	9.41992227	0.79	
	3	Dermacentor	7.98077969	39	30	13	5	1	183366585	8.26332020	0.96	
	15	Ixodes	5.22084675	42	28	13	5	1	183364645	8.26331560	0.63	
	2	Amalaraeus	7.81187831	20	14	6	3	1	92077049	7.96415139	0.98	
	7	Ceratophyllus	3.02871424	17	14	6	3	1	92077046	7.96415138	0.38	
	2	Dactylophylla	3.82978603	2	2	2	2	1	45663801	7.65957206	0.50	
EXODIDAE	3	Malereus	7.76169532	15	11	5	3	1	91703820	7.96238743	0.97	
	9	Megabothris	6.51968398	55	29	10	4	1	138103918	8.41020600	0.80	
	6	Monoplyllus	7.89157796	42	29	12	4	1	138833505	8.14249429	0.96	
	3	Opisocrotis	7.75907477	14	10	5	3	1	91702829	7.96238273	0.97	
	3	Opisodassy	7.76463049	17	11	6	3	1	92074082	7.96413740	0.97	
	5	Oreopexas	6.88754953	38	26	9	4	1	137734696	8.13904335	0.84	
	4	Oropsylla	7.89018204	33	23	9	3	1	93190714	7.96937264	0.99	
	4	Thrassis	7.21441970	21	13	5	2	1	46791503	7.67016700	0.94	
	8	Catlingcola	0.88405192	11	8	1	1	1	6983	3.84404204	0.23	

Number of species in genus.

Table 2. Generic host specificity indices for non-monotypic metazoan genera from Appendix I. (N/A, 1 applicable).

PARASITES		HOSTS						Rank	Familial Index ( $HS_f$ )	Familial Index Ratio ( $HS_s / HS_f$ )
Family	n*	Mean Familial Index ( $HS_s$ )	No. species	No. genera	No. families	No. orders	No. classes			
Allocreadiidae	18	5.05716660	117	53	24	16	1	648892693	8.81217288	0.573884179
Hemiuroidae	50	2.76484776	100	65	34	13	2	2934983639	9.46760568	0.292032416
Leptocreadiidae	41	1.31744914	48	28	21	9	1	399208274	8.60119953	0.153170396
Gyrodactylidae	35	1.30460080	40	26	24	10	1	402589229	8.60486215	0.151612051
Dactylogyridae	12	0.00000000	12	12	10	6	2	2630165551	9.41998309	0
Monocotylidae	40	0.15440958	20	15	12	4	2	2542931784	9.40533471	0.016417233
Capsalidae	2	0.23856063	4	2	2	1	1	375253	5.57432417	0.042796332
Hexabothriidae	4	0.00000000	4	4	4	2	1	46411302	7.66662375	0
Mazocracidae	5	0.00000000	2	2	1	1	1	1001	3.00043408	0
Discocotylidae	4	0.00000000	4	4	3	1	1	750499	5.87535012	0
Diclydophoridae	5	0.06020600	6	5	3	2	1	46040047	7.66313576	0.007856575
Microcotylidae	15	0.00000000	12	9	6	2	1	47157796	7.67355350	0
Gastrocotylidae	8	0.11288625	6	5	2	1	1	378246	5.57777434	0.020238583
Onchobothriidae	36	0.01672389	26	15	8	4	1	137356626	8.13784961	0.002055075
Litobothriidae	8	0.00000000	4	3	3	1	1	749502	5.87477280	0
Pseudophyllidea	3	3.71621573	6	3	3	1	1	749504	5.87477395	0.632571697
Neoechinorhynchidae	19	4.28495687	58	39	17	11	1	442069917	8.64549096	0.495629096
Cystidicolidae	26	2.70857567	77	49	29	16	2	3054748735	9.48497549	0.285564857
Heterorhabdiidae	2	0.00000000	2	2	1	1	1	1001	3.00043408	0
Mermithidae	12	3.48708946	107	52	16	6	1	228269326	8.35844756	0.417193436
Steinernematidae	4	6.05148415	16	14	10	3	1	93548145	7.97103518	0.759184224
Caligidae	80	2.50897526	139	97	55	19	2	3181753992	9.50266660	0.264028548
Panderidae	28	4.14172096	28	21	14	8	2	2718114962	9.43426782	0.439008202
Ixodidae	19	5.80650774	65	34	16	5	1	184454509	8.26588928	0.702466189
Ceratophyllidae	53	6.49238163	116	44	15	4	1	139935024	8.14592643	0.797009608
Philopteridae	8	0.88495192	11	8	1	1	1	6983	3.84404204	0.230213901

\* Number of species in family.

Table 3. Familial host specificity indices for non-monotypic metazoan families from Appendix I.

host records for these 6 flea species include only 42 unique host species. Instances in which these values differ are indicative of little host overlap among the species in the genus. For example, the mean generic index value ( $HS_g$ ) for the four included species of diclydophorid monogeneans in the genus *Choricotyle* was 0.075, whereas the index value for this genus ( $HS_g$ ) was 7.66. Examination of the host lists for these four species reveals no overlap in host species ; each parasitizes different species of teleosts, and collectively, these hosts belong to three different families of fishes. Calculation of the ratio of  $HS_s$  to  $HS_g$  provides a more convenient way to assess the proportion of the generic host range accounted for by the host range of an average species. A ratio of 1 indicates complete overlap in the host species parasitized by all species in the genus ; a ratio of 0 indicates no overlap in the host species parasitized by species in the genus. So for example, in the ceratophyllid flea genus *Monophyllus* this ratio (i.e.,  $HS_s / HS_g$ ) is 0.97, whereas this ratio is 0.0098 for the diclydophorid monogenean genus *Choricotyle*.

### III. Familial indices

Similarly, two types of familial index values can be obtained from this system. Mean familial values can be calculated by averaging the index values of all species in all genera in a family. Alternatively, an index value for the family ( $HS_f$ ) itself can be determined by calculating a rank for the family from the number of host species, genera, families, orders and classes parasitized by all species in all genera in the family. Both of these values are presented in Table III for most of the non-monotypic (metazoan) families from Appendix I. Again, it is important to note that these two index values are conspicuously different for many families. In instances in which these values are similar, there exists much host overlap among the species in the family. In instances in which the index value for the family is conspicuously higher than the mean familial index value, there is little host overlap among the species in the family.

One of the advantages of this system is that mean index values can be calculated for essentially any suite of parasites so as to allow a diversity of comparisons to be made. We use such calculations below to examine several common generalizations about parasite specificity. In each case, the mean is followed by the standard error and the p value obtained from a Wilcoxon signed-rank test for each comparison.

1) « ...[A]mong parasites of fish, acanthocephalans and nematodes show little host specificity, whereas monogeneans seem to be highly specific. » (e.g., Sasal *et al.*, 1999 ; pg. 437).

Mean index values for the parasites of fish in Appendix I support this statement. The 19 species of acanthocephalans had a mean index value of  $4.28 \pm 0.95$ , and the 26 species of nematodes of fishes had a mean index value of  $2.71 \pm 0.73$ . By comparison, the 130 species of monogeneans had a mean index value of  $0.41 \pm 0.13$ . Wilcoxon signed-rank tests suggest that monogeneans exhibit significantly greater host specificity (at p = 0.05 level) than both acanthocephalans and nematodes (p = 7.71e-07, p = 0.0034, respectively).

2) « In both marine and freshwater fish, monogeneans as a group are possibly the most host-specific of all fish parasites. » (Poulin, 1992 ; pg. 753).

The mean specificity index values for the groups of fish parasites included in Appendix I support this statement with only one exception. The mean index value for the 109 species of digenleans was  $2.60 \pm 0.33$ . The mean index value for the 108 copepod species was  $2.93 \pm 0.37$ . As noted above, the mean index values for acanthocephalans was  $4.28 \pm 0.95$ , that for nematodes was  $2.71 \pm 0.73$ , and that for monogeneans was  $0.41 \pm 0.13$ . Individual Wilcoxon signed-rank tests comparing monogeneans to digenleans, copepods, acanthocephalans and nematodes were all significant at the

$p = 0.05$  level ( $p = 4.595e-09$ ,  $p = 1.494e-07$ ,  $7.7111e-07$ ,  $p = 0.000343$ , respectively). The 44 species of fish cestodes however, had a mean index value lower than that of the monogeneans ( $0.01 \pm 0.01$  versus  $0.41 \pm 0.13$ ). The results of the Wilcoxon signed-rank test comparing the index values of monogeneans and cestodes suggest the cestodes are significantly more host specific ( $p = 0.02$ ) than the monogeneans. It is important to note, however, that, whereas the majority of the other fish parasite groups in Appendix I parasitize teleosts, the cestode groups included parasitize elasmobranchs. It would be very interesting to investigate this question using cestodes of teleosts because they are generally perceived to be less host specific than the cestodes of elasmobranchs. In which case, elasmobranchs should not be included as 'fishes' in these kinds of comparisons.

3) Ectoparasites [of fishes] are more host-specific than endoparasites. (Rohde & Heap, 1998).

The mean index values for the species in Appendix I that fall into these two groups support this statement. The mean index value for all ectoparasites of fishes (copepods and monogeneans) was :  $1.56 \pm 0.20$  ( $n = 238$ ). The mean index value for all endoparasites of fishes (i.e., acanthocephalans, nematodes, and digeneans) was  $2.83 \pm 0.29$  ( $n = 154$ ). The Wilcoxon signed-rank test was significant ( $p = 3.507e-05$ ). Cestodes were not included in this comparison for the reasons described in comparison 2, above.

4) « ...[P]arasitic worms with direct life cycles...are more host specific than the worms that have an indirect cycle. » (Noble *et al.*, 1989 ; pg. 501).

This statement also appears to be supported by the data in Appendix I. The mean index value for the species of worms presumed to possess direct life-cycles, which from the data in Appendix I include the monogeneans and the entomogenous nematodes only, is  $0.90 \pm 0.17$  ( $n = 151$ ). Whereas the mean index value for the species of worms presumed to possess indirect life cycles (acanthocephalans, digeneans, cystidicolid nematodes and cestodes) is  $2.22 \pm 0.24$  ( $n = 201$ ). The difference between these two groups was significant using the Wilcoxon signed-rank test ( $p = 0.00158$ ).

5) Monogeneans exhibit a high degree of host specificity at the species level, but rather low specificity at the generic level. (see Llewellyn, 1957)

Comparison of the mean generic specificity index value for each of the 21 non-monotypic genera of monogeneans with the generic index value based on the full spectrum of hosts of all species in each genus allows Llewellyn's observation to be addressed. In 16 of the 21 genera the mean generic index values were at least 5 index points lower than the index value calculated for the genus as a whole ; the generic values for six genera are greater than 7 index points above the mean generic index values. As is apparent from Table II, these differences are more marked in the monogeneans than in any of the other parasite groups, providing support for

Llewellyn's observation. If it were not for the fact that so many of the monogenean species exhibit oioxenous specificity, and thus have index values of 0, this question could be more directly addressed by comparing the mean generic index value to the generic index value.

6) The degree of host specificity exhibited by 'ticks' has been controversial (e.g., Hoogstraal & Aeschlimann, 1982 ; Klompen *et al.*, 1996 ; Cumming, 1998).

The tick data analyzed here (which, admittedly, are very limited) result in a mean species host specificity index value of  $5.81 \pm 0.81$  ( $n = 19$ ). Although we were not able to include the tick data of Cumming (1998) in Appendix I, because of the lack of data on host species identities, the data provided indicate that a number of these species are less host specific than all of the 700 taxa included in Appendix I. At a minimum (i.e., when only data on order and class of host are included) the host specificity index values for the 14 species of ticks that parasitize hosts in 3 classes treated by Cumming (1998) exceed a specificity index value of 9.6. By comparison, the mean index values for the other groups of arthropods parasitizing vertebrates from Appendix I were : lice  $0.88 \pm 0.50$  ( $n = 8$ ), harpyrhynchid mites  $0.62 \pm 0.18$  ( $n = 114$ ), and fleas  $6.49 \pm 0.37$  ( $n = 53$ ). The results of Wilcoxon signed-rank tests suggest ticks are significantly less host specific than lice and mites, but not fleas ( $p = 0.00980$ ,  $p = 1.754e-13$ , and  $p = 0.0954$ ). These values are certainly consistent with the notion expressed previously by many authors (e.g., Marshall, 1981 ; Fain, 1994) that arthropods that are intimately associated with their hosts for extended periods of their lives tend to be much more host specific than those that are less intimately associated with their hosts for only brief periods of time.

Testing the above generalizations was not our main intention. Rather it was our intention to develop a mechanism that would facilitate the consistent comparisons of host specificity at different taxonomic levels and among disparate taxa. There currently exists disagreement about terms used to refer to host specificity. In some cases, even when terms are used routinely, their definitions are vague and there is disagreement about how they should be applied. We favor a quantitative index of specificity that increases the precision of the terms used to describe host specificity in parasitology, and perhaps also ecology, in general. The index described above allows quantitative comparisons of specificity patterns among distantly related groups of hosts and parasites. However the power of the index developed here depends on detailed and accurate host specificity data. If broader questions are to be addressed, generation of this type of data is strongly encouraged for as many parasite species as possible and with as broad geographic sampling as possible. It is our hope that this index will allow parasite systematists and ecolo-

gists to assess hypotheses about specificity in a more precise and reproducible way, and perhaps even to reveal patterns that would not otherwise have been detected.

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#### Appendix I. Host specificity of selected parasite species.

D I G E N E A	<i>Dinosma oregonensis</i>	Love & Moser, 1983	2	2	1	1	1	1001	3.00043408
	<i>Dinosma pectoralis</i>	Love & Moser, 1983	2	2	1	1	1	1001	3.00043408
	<i>Dinosma tortum</i>	Love & Moser, 1983	3	2	2	1	1	375252	5.57432302
	<i>Dinurus barbatus</i>	Love & Moser, 1983	2	2	2	1	1	375251	5.57432186
	<i>Dinurus breviductus</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	<i>Dinurus euthynni</i>	Love & Moser, 1983	3	3	1	1	1	2000	3.30103000
	<i>Dinurus longisinus</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	<i>Dinurus scombri</i>	Love & Moser, 1983	2	2	2	1	1	375251	5.57432186
	<i>Dinurus tornatus</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	<i>Dissosaccus laevis</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	<i>Ectenurus americanus</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	<i>Ectenurus lepidocibilis</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	<i>Ectenurus lepidus</i>	Love & Moser, 1983	2	2	2	1	1	375251	5.57432186
	<i>Ectenurus virgulus</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	<i>Ectenurus yamagutii</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	<i>Elytorphallus mexicanus</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	<i>Genarchopsis muellieri</i>	Love & Moser, 1983	13	10	5	4	2	2540355877	9.40489456
	<i>Genolinea aburame</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	<i>Genolinea anaura</i>	Love & Moser, 1983	7	6	5	3	1	91698852	7.96236390
	<i>Genolinea latilcauda</i>	Love & Moser, 1983	19	17	8	5	1	181527384	8.25894215
	<i>Genolinea lintoni</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	<i>Genolinea manteri</i>	Love & Moser, 1983	4	4	3	2	1	46039049	7.66312634
	<i>Genolinea montereyensis</i>	Love & Moser, 1983	2	2	1	1	1	1001	3.00043408
	<i>Genolinea tanyopa</i>	Love & Moser, 1983	2	2	2	1	1	375251	5.57432186
	<i>Glomeridurus ulmeri</i>	Love & Moser, 1983	2	1	1	1	1	2	0.30103000
	<i>Gonocerca haedrichi</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	<i>Gonocerca oregonensis</i>	Love & Moser, 1983	3	3	2	2	1	45664800	7.65958156
	<i>Gonocerca oshoro</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	<i>Gonocerca phycidis</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	<i>Gonocarella pacifica</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	<i>Gonocarella trachinoti</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	<i>Hemilurus communis</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	<i>Hemilurus levinseni</i>	Love & Moser, 1983	20	15	8	6	2	2629434956	9.41986243
	<i>Hemilurus luehei</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	<i>Hemilurus odhneri</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
LEPOCREADIIDAE	<i>Blanius plicatum</i>	Love & Moser, 1983	2	2	2	1	1	375251	5.57432186
	n (genera) = 11	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	n (species) = 41	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	<i>Cephalolepidapedon saba</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	<i>Dihemistophenus fragilis</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	<i>Dihemistophenus tydiae</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	<i>Echeneidocoelium indicum</i>	Love & Moser, 1983	2	2	1	1	1	2	0.30103000
	<i>Labifer balli</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	<i>Labifer secundus</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	<i>Lepidapedon antimorae</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	<i>Lepidapedon calli</i>	Love & Moser, 1983	2	2	1	1	1	1001	3.00043408
	<i>Lepidapedon cascadensis</i>	Love & Moser, 1983	2	1	1	1	1	2	0.30103000

		<i>Lepidapedon elongatum</i>	Love & Moser, 1983	2	1	1	1	1	2	0.30103000
		<i>Lepidapedon epinepheli</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Lepidapedon filiformis</i>	Love & Moser, 1983	7	6	5	3	1	91698852	7.96236390
		<i>Lepidapedon hancocki</i>	Love & Moser, 1983	2	2	1	1	1	1001	3.00043408
		<i>Lepidapedon luteum</i>	Love & Moser, 1983	3	1	1	1	1	3	0.47712125
		<i>Lepidapedon microyleum</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Lepidapedon nicolli</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Lepidapedon oregonensis</i>	Love & Moser, 1983	2	1	1	1	1	2	0.30103000
		<i>Lepidapedon yaquina</i>	Love & Moser, 1983	2	1	1	1	1	2	0.30103000
		<i>Lepocreadium areolatum</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Lepocreadium bimarginatum</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Lepocreadium bravae</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Lepocreadium californianum</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Lepocreadium ghanensis</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Lepocreadium manteri</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Lepocreadium scombri</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Lepocreadium seiferoides</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Neolabrifer bravae</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Neolepidapedon medialunae</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Neolepidapedon pugetense</i>	Love & Moser, 1983	2	1	1	1	1	2	0.30103000
		<i>Neolepidapedon retursum</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Neolepidapedon sebastisci</i>	Love & Moser, 1983	9	1	1	1	1	9	0.95424251
		<i>Opechona alaskensis</i>	Love & Moser, 1983	10	3	3	3	1	90952358	7.95881396
		<i>Opechona bacillaris</i>	Love & Moser, 1983	3	3	3	3	1	90952351	7.95881393
		<i>Opechona occidentalis</i>	Love & Moser, 1983	10	2	2	2	1	45663809	7.65957213
		<i>Opechona olsoni</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Opechona orientalis</i>	Love & Moser, 1983	3	3	3	2	1	46038051	7.66311693
		<i>Opechona parvasoma</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Opechona pharyngodactyla</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Opechona theragrae</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Pseudocreadium diodontis</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	GYRODACTYLIDAE	<i>Gyrodactylus abalonia</i>	Beverley-Burton, 1984	9	8	5	5	1	180410686	8.25626226
		<i>Gyrodactylus adpersi</i>	Beverley-Burton, 1984	1	1	1	1	1	1	0.00000000
		<i>Gyrodactylus aldrichi</i>	Beverley-Burton, 1984	1	1	1	1	1	1	0.00000000
		<i>Gyrodactylus alexanderi</i>	Beverley-Burton, 1984	1	1	1	1	1	1	0.00000000
		<i>Gyrodactylus aquilinus</i>	Beverley-Burton, 1984	1	1	1	1	1	1	0.00000000
		<i>Gyrodactylus atratuli</i>	Beverley-Burton, 1984	3	2	2	1	1	375252	5.57432302
		<i>Gyrodactylus bairdi</i>	Beverley-Burton, 1984	2	1	1	1	1	2	0.30103000
		<i>Gyrodactylus bullatarudis</i>	Beverley-Burton, 1984	1	1	1	1	1	1	0.00000000
		<i>Gyrodactylus cameroni</i>	Beverley-Burton, 1984	1	1	1	1	1	1	0.00000000
		<i>Gyrodactylus canadensis</i>	Beverley-Burton, 1984	3	3	1	1	1	2000	3.30103000
		<i>Gyrodactylus colemanensis</i>	Beverley-Burton, 1984	2	2	1	1	1	1001	3.00043408
		<i>Gyrodactylus commersoni</i>	Beverley-Burton, 1984	1	1	1	1	1	1	0.00000000
		<i>Gyrodactylus couesi</i>	Beverley-Burton, 1984	1	1	1	1	1	1	0.00000000
		<i>Gyrodactylus dechirali</i>	Beverley-Burton, 1984	2	1	1	1	1	2	0.30103000



M O N O G E N E A	<i>Dionchus rachycentris</i>	Hargis, 1957	1	1	1	1	1	1	0.00000000
	<i>Dionchus ramorae</i>	Hargis, 1957	1	1	1	1	1	1	0.00000000
	<i>Empruhotrema dasyatidis</i>	Chisholm et al., 2001	1	1	1	1	1	1	0.00000000
	<i>Empruhotrema raiiae</i>	Hargis, 1957	1	1	1	1	1	1	0.00000000
	<i>Empruhotrema quindecimta</i>	Chisholm et al., 2001	1	1	1	1	1	1	0.00000000
	<i>Heterocotyle astobatus</i>	Hargis, 1957	1	1	1	1	1	1	0.00000000
	<i>Heterocotyle americana</i>	Hargis, 1957	1	1	1	1	1	1	0.00000000
	<i>Heterocotyle capricornensis</i>	Chisholm et al., 2001	1	1	1	1	1	1	0.00000000
	<i>Heterocotyle pseudominima</i>	Hargis, 1957	1	1	1	1	1	1	0.00000000
	<i>Loimopapillosum dasyatidis</i>	Hargis, 1957	2	1	1	1	1	1	0.00000000
	<i>Loimosina sciodon</i>	Hargis, 1957	1	1	1	1	1	1	0.00000000
	<i>Loimosina wilsoni</i>	Hargis, 1957	1	1	1	1	1	1	0.00000000
	<i>Merizocotyle australensis</i>	Chisholm et al., 2001	1	1	1	1	1	1	0.00000000
	<i>Merizocotyle icopae</i>	Chisholm et al., 2001	1	1	1	1	1	1	0.00000000
	<i>Merizocotyle urolophii</i>	Chisholm et al., 2001	1	1	1	1	1	1	0.00000000
	<i>Monocotyle coralli</i>	Chisholm et al., 2001	1	1	1	1	1	1	0.00000000
	<i>Monocotyle deademalis</i>	Hargis, 1957	2	1	1	1	1	2	0.30103000
	<i>Monocotyle helicophallus</i>	Chisholm et al., 2001	1	1	1	1	1	1	0.00000000
	<i>Monocotyle multiparous</i>	Chisholm et al., 2001	1	1	1	1	1	1	0.00000000
	<i>Monocotyle pricei</i>	Hargis, 1957	2	1	1	1	1	2	0.30103000
	<i>Monocotyle spiremae</i>	Chisholm et al., 2001	1	1	1	1	1	1	0.00000000
	<i>Neoheterocotyle inpristi</i>	Hargis, 1957	1	1	1	1	1	1	0.00000000
	<i>Neoheterocotyle rhinobatidis</i>	Chisholm, et al 2001	1	1	1	1	1	1	0.00000000
	<i>Neoheterocotyle rhinobatis</i>	Chisholm, et al 2001	1	1	1	1	1	1	0.00000000
	<i>Neoheterocotyle rhynchobatus</i>	Chisholm, et al 2001	1	1	1	1	1	1	0.00000000
	<i>Thaumatocotyle longicirrus</i>	Hargis, 1957	1	1	1	1	1	1	0.00000000
	<i>Thaumatocotyle pseudodasybatis</i>	Hargis, 1957	1	1	1	1	1	1	0.00000000
	<i>Thaumatocotyle retorta</i>	Hargis, 1957	1	1	1	1	1	1	0.00000000
	<i>Troglocephalus rhinobatidis</i>	Chisholm et al., 2001	1	1	1	1	1	1	0.00000000
CAPSALIDAE	<i>Benedenia posterocarpa</i>	Hargis, 1957	1	1	1	1	1	1	0.00000000
n (genera) = 2; n (species) = 2	<i>Entobdella corona</i>	Hargis, 1957	3	1	1	1	3	0.47712125	
HEXABOTRIIIDAE	<i>Dasyochocotyle spiniphalus</i>	Hargis, 1957	1	1	1	1	1	1	0.00000000
n (genera) = 3	<i>Heterochocotyle leucas</i>	Hargis, 1957	1	1	1	1	1	1	0.00000000
n (species) = 4	<i>Squalonchocotyle inpristi</i>	Hargis, 1957	1	1	1	1	1	1	0.00000000
	<i>Squalonchocotyle liburonis</i>	Hargis, 1957	1	1	1	1	1	1	0.00000000
MAZOCRAEIIDAE	<i>Clupaeocotyle brevoortia</i>	Hargis, 1957	1	1	1	1	1	1	0.00000000
n (genera) = 3	<i>Clupaeocotyle megaconfibula</i>	Hargis, 1957	1	1	1	1	1	1	0.00000000
n (species) = 5	<i>Cunea brevoortea</i>	Hargis, 1957	1	1	1	1	1	1	0.00000000
	<i>Mesocraeoides georgiei</i>	Hargis, 1957	1	1	1	1	1	1	0.00000000
	<i>Mesocraeoides opisthonema</i>	Hargis, 1957	1	1	1	1	1	1	0.00000000
DISCOCOTYLIDAE	<i>Bicotylophora trachinoti</i>	Hargis, 1957	1	1	1	1	1	1	0.00000000
n (genera) = 2	<i>Talga bairdiella</i>	Hargis, 1957	1	1	1	1	1	1	0.00000000
n (species) = 4	<i>Talga cupida</i>	Hargis, 1957	1	1	1	1	1	1	0.00000000
	<i>Talga micropogone</i>	Hargis, 1957	1	1	1	1	1	1	0.00000000

DICLYDOPHORIDAE	<i>Choricyle aspnocorda</i>	Hargis, 1957	1	1	1	1	1	1	0.00000000
n (genera) = 2	<i>Choricyle lusitanensis</i>	Hargis, 1957	1	1	1	1	1	1	0.00000000
n (species) = 5	<i>Choricyle prionotii</i>	Hargis, 1957	1	1	1	1	1	1	0.00000000
	<i>Choricyle sinosynoscioni</i>	Hargis, 1957	2	1	1	1	1	2	0.30103000
	<i>Petocotyle minima</i>	Hargis, 1957	1	1	1	1	1	1	0.00000000
MICROCOTYLIDAE	<i>Axinoides gracilis</i>	Hargis, 1957	1	1	1	1	1	1	0.00000000
n (genera) = 6	<i>Axinoides raphidoma</i>	Hargis, 1957	1	1	1	1	1	1	0.00000000
n (species) = 15	<i>Axinoides truncatus</i>	Hargis, 1957	1	1	1	1	1	1	0.00000000
	<i>Cempocotyle carangus</i>	Hargis, 1957	1	1	1	1	1	1	0.00000000
	<i>Heteraxine carangus</i>	Hargis, 1957	1	1	1	1	1	1	0.00000000
	<i>Heteraxine xanthophyllus</i>	Hargis, 1957	1	1	1	1	1	1	0.00000000
	<i>Metamicrocotyla macracantha</i>	Hargis, 1957	1	1	1	1	1	1	0.00000000
	<i>Microcotyle archosargi</i>	Hargis, 1957	1	1	1	1	1	1	0.00000000
	<i>Microcotyle heteracantha</i>	Hargis, 1957	1	1	1	1	1	1	0.00000000
	<i>Microcotyle pogoniae</i>	Hargis, 1957	1	1	1	1	1	1	0.00000000
	<i>Microcotyle pomacanthae</i>	Hargis, 1957	1	1	1	1	1	1	0.00000000
	<i>Microcotyle pomatome</i>	Hargis, 1957	1	1	1	1	1	1	0.00000000
	<i>Microcotyle pseudoheteracantha</i>	Hargis, 1957	1	1	1	1	1	1	0.00000000
	<i>Microcotyle pseudomugilis</i>	Hargis, 1957	1	1	1	1	1	1	0.00000000
	<i>Pyragraphorophorus hippus</i>	Hargis, 1957	1	1	1	1	1	1	0.00000000
GASTROCOTYLIDAE	<i>Amphipolycoyte clorascombrus</i>	Hargis, 1957	1	1	1	1	1	1	0.00000000
n (genera) = 8	<i>Lithidocotyle acanthophallus</i>	Hargis, 1957	2	1	1	1	1	2	0.30103000
n (species) = 8	<i>Protomicrocotyle nitrobilis</i>	Hargis, 1957	1	1	1	1	1	1	0.00000000
	<i>Pseudoxazine mexicana</i>	Hargis, 1957	2	1	1	1	1	2	0.30103000
	<i>Pseudomazocraes selene</i>	Hargis, 1957	1	1	1	1	1	1	0.00000000
	<i>Scomberocotyle scomberomore</i>	Hargis, 1957	2	1	1	1	1	2	0.30103000
	<i>Thoracocotyle crocea</i>	Hargis, 1957	2	1	1	1	1	2	0.30103000
	<i>Vallezia oligoplitis</i>	Hargis, 1957	1	1	1	1	1	1	0.00000000
ONCHOBOTHRIIDAE	<i>Acanthobothrium bajaense</i>	Appy & Dailey, 1973	1	1	1	1	1	1	0.00000000
n (genera) = 8	<i>Acanthobothrium bullardi</i>	Ghoshroy & Cairn, 2001	1	1	1	1	1	1	0.00000000
n (species) = 36	<i>Acanthobothrium dasi</i>	Ghoshroy & Cairn, 2001	1	1	1	1	1	1	0.00000000
	<i>Acanthobothrium dollyae</i>	Caira & Burge, 2001	1	1	1	1	1	1	0.00000000
	<i>Acanthobothrium heterodonti</i>	Caira & Zahner, 2001	1	1	1	1	1	1	0.00000000
	<i>Acanthobothrium maryanskii</i>	Caira & Burge, 2001	1	1	1	1	1	1	0.00000000
	<i>Acanthobothrium puertecitense</i>	Caira & Zahner, 2001	1	1	1	1	1	1	0.00000000
	<i>Acanthobothrium rajvi</i>	Ghoshroy & Cairn, 2001	1	1	1	1	1	1	0.00000000
	<i>Acanthobothrium royi</i>	Caira & Burge, 2001	1	1	1	1	1	1	0.00000000
	<i>Acanthobothrium santarosense</i>	Caira & Zahner, 2001	1	1	1	1	1	1	0.00000000
	<i>Acanthobothrium soberoni</i>	Ghoshroy & Cairn, 2001	1	1	1	1	1	1	0.00000000
	<i>Bilocutaneus pritchardae</i>	Caira & Ruhnke, 1990	1	1	1	1	1	1	0.00000000
	<i>Calliobothrium eschrichtii</i>	Nasin, Caira & Euzet, 1997	1	1	1	1	1	1	0.00000000
	<i>Calliobothrium evani</i>	Nasin, Caira & Euzet, 1997	1	1	1	1	1	1	0.00000000
	<i>Calliobothrium hayhowi</i>	Nasin, Caira & Euzet, 1997	1	1	1	1	1	1	0.00000000
	<i>Calliobothrium pellucidum</i>	Nasin, Caira & Euzet, 1997	1	1	1	1	1	1	0.00000000
	<i>Calliobothrium riseri</i>	Nasin, Caira & Euzet, 1997	1	1	1	1	1	1	0.00000000

C E S T O D A	LITOBOTHRIIDAE	<i>Calliobothrium violae</i>	Nasin, Caira & Euzet, 1997	1	1	1	1	1	1	0.00000000	
		<i>Erudituncus musteli</i>	Healy, Scholz & Caira, 2001	1	1	1	1	1	1	0.00000000	
		<i>Pachybothrium hudsoni</i>	Caira & Pritchard, 1986	1	1	1	1	1	1	0.00000000	
		<i>Pedibothrium kerkhami</i>	Caira, 1992	1	1	1	1	1	1	0.00000000	
		<i>Pedibothrium longispine</i>	Caira, 1992	1	1	1	1	1	1	0.00000000	
		<i>Pedibothrium brevispine</i>	Caira, 1992	1	1	1	1	1	1	0.00000000	
		<i>Pedibothrium globicephalum</i>	Caira, 1992	1	1	1	1	1	1	0.00000000	
		<i>Pedibothrium maccallumi</i>	Caira, 1992	1	1	1	1	1	1	0.00000000	
		<i>Pedibothrium manteri</i>	Caira, 1992	1	1	1	1	1	1	0.00000000	
		<i>Pedibothrium servatorium</i>	Caira, 1992	1	1	1	1	1	1	0.00000000	
		<i>Pedibothrium tollorense</i>	Caira & Rasolofonirina, 1998	1	1	1	1	1	1	0.00000000	
		<i>Pedibothrium veravalensis</i>	Caira, 1992	1	1	1	1	1	1	0.00000000	
		<i>Pinguicollum pinguicollum</i>	Caira & Keeling, 1996	4	1	1	1	1	4	0.60205999	
		<i>Platybothrium harpago</i>	Healy, 2002	1	1	1	1	1	1	0.00000000	
		<i>Platybothrium hypoprioni</i>	Healy, 2002	1	1	1	1	1	1	0.00000000	
		<i>Platybothrium auriculatum</i>	Healy, 2002	1	1	1	1	1	1	0.00000000	
		<i>Platybothrium cervinum</i>	Healy, 2002	1	1	1	1	1	1	0.00000000	
		<i>Platybothrium parvum</i>	Healy, 2002	1	1	1	1	1	1	0.00000000	
		<i>Platybothrium spinulifera</i>	Healy, 2002	1	1	1	1	1	1	0.00000000	
A C A N T H O C E P H A L A	PSEUDOPHYLLIDEA	<i>Litobothrium alopias</i>	Olson & Caira, 2001	1	1	1	1	1	1	0.00000000	
		<i>Litobothrium amplifica</i>	Olson & Caira, 2001	1	1	1	1	1	1	0.00000000	
		<i>Litobothrium amschensis</i>	Olson & Caira, 2001	1	1	1	1	1	1	0.00000000	
		<i>Litobothrium conformis</i>	Olson & Caira, 2001	1	1	1	1	1	1	0.00000000	
		<i>Litobothrium daileyi</i>	Olson & Caira, 2001	1	1	1	1	1	1	0.00000000	
		<i>Litobothrium gracile</i>	Olson & Caira, 2001	1	1	1	1	1	1	0.00000000	
		<i>Litobothrium janovyi</i>	Olson & Caira, 2001	1	1	1	1	1	1	0.00000000	
		<i>Litobothrium nickoli</i>	Olson & Caira, 2001	1	1	1	1	1	1	0.00000000	
		<i>Cephalochlamys compactus</i>	Jackson & Tinsley, 2001	5	2	2	1	1	375254	5.57432533	
		<i>Cephalochlamys namaquensis</i>	Jackson & Tinsley, 2001	2	2	2	1	1	375251	5.57432186	
		<i>Paracephalochlamys papillonis</i>	Jackson & Tinsley, 2001	1	1	1	1	1	1	0.00000000	
NEOECHINORHYNCHIDAE	<i>Floridosentus elongatus</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000		
	<i>Gracilisentis gracilisentis</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000		
	<i>Micrasentis wardae</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000		
	<i>Neoechinorhynchus agilis</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000		
	<i>Neoechinorhynchus cariopodi</i>	Arai, 1989; Margolis & Arthur, 1979	1	1	1	1	1	1	0.00000000		
	<i>Neoechinorhynchus chilkaense</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000		
	<i>Neoechinorhynchus crassus</i>	Arai, 1989; Margolis & Arthur, 1979	7	5	3	3	1	90954348	7.95882346		
	<i>Neoechinorhynchus cristatus</i>	Arai, 1989; Margolis & Arthur, 1979	9	6	4	3	1	91327598	7.96060204		
	<i>Neoechinorhynchus cylindratus</i>	Arai, 1989; Margolis & Arthur, 1979	16	13	7	5	1	181155168	8.25805073		
	<i>Neoechinorhynchus notemigoni</i>	Arai, 1989; Margolis & Arthur, 1979	1	1	1	1	1	1	0.00000000		
	<i>Neoechinorhynchus pungitius</i>	Arai, 1989; Margolis & Arthur, 1979	5	5	4	4	1	135867648	8.13311606		
	<i>Neoechinorhynchus rutili</i>	Arai, 1989	40	26	14	10	1	399020069	8.60099474		
	<i>Neoechinorhynchus saginatus</i>	Arai, 1989	12	8	4	3	1	91329588	7.96061150		
	<i>Neoechinorhynchus salmonis</i>	Arai, 1989	14	10	5	4	1	136243878	8.13431700		
	<i>Neoechinorhynchus strigosus</i>	Arai, 1989; Margolis & Arthur, 1979	6	5	5	4	1	136238905	8.13430114		
	<i>Neoechinorhynchus tenellus</i>	Arai, 1989; Margolis & Arthur, 1979	5	3	3	2	1	46038053	7.66311695		

		<i>Neoechinorhynchus tumidus</i>	Arai, 1989; Margolis & Arthur, 1979	11	8	4	4	1	135870636	8.13312561
		<i>Octospinifer macilentus</i>	Arai, 1989; Margolis & Arthur, 1979	3	1	1	1	1	3	0.47712125
		<i>Tanaorhamphus longirostris</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	CYSTIDICOLIDAE	<i>Ascarophis curvicauda</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
n (genera) = 9		<i>Ascarophis filiformis</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
n (species) = 26		<i>Ascarophis morrhuae</i>	Margolis & Arthur, 1979; Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Ascarophis pacificus</i>	Love & Moser, 1983	6	6	5	5	1	180408696	8.25625747
		<i>Ascarophis sebastodis</i>	Margolis & Arthur, 1979; Love & Moser, 1983	21	10	8	4	1	137351681	8.13783398
		<i>Ascarophis skrjabini</i>	Love & Moser, 1983	4	3	2	2	1	45664801	7.65958157
		<i>Caballeronema wardlei</i>	Margolis & Arthur, 1979; Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Ctenascarophis gastricus</i>	Love & Moser, 1983	2	2	1	1	1	1001	3.00043408
		<i>Cystidicola cristomermi</i>	Margolis & Arthur, 1979	1	1	1	1	1	1	0.00000000
		<i>Cystidicola farionis</i>	Margolis & Arthur, 1979	23	12	6	5	1	180784922	8.25716221
		<i>Cystidicola stigmatura</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Cystidicola tenuissima</i>	Love & Moser, 1983	6	4	2	2	1	45665800	7.65959107
		<i>Cystidicoloides prevosti</i>	Margolis & Arthur, 1979	1	1	1	1	1	1	0.00000000
		<i>Cystidicoloides tenuissima</i>	Margolis & Arthur, 1979	15	8	4	4	1	135870640	8.13312562
		<i>Metabronema laticauda</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Parascarophis galeata</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Salvelinema ishii</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Salvelinema salmincola</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Salvelinema walkeri</i>	Love & Moser, 1983	2	1	1	1	1	2	0.30103000
		<i>Spinictectus beaveri</i>	Love & Moser, 1983	4	2	1	1	1	1003	3.00130093
		<i>Spinictectus carolin</i>	Margolis & Arthur, 1979	1	1	1	1	1	1	0.00000000
		<i>Spinictectus echenei</i>	Love & Moser, 1983	6	5	3	2	1	46040047	7.66313576
		<i>Spinictectus gordani</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Spinictectus gracilis</i>	Margolis & Arthur, 1979	1	1	1	1	1	1	0.00000000
		<i>Spinictectus mollis</i>	Love & Moser, 1983	18	16	9	6	1	225691217	8.35351466
		<i>Spinictectus pacificus</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	ALLANTONEMATIDAE	<i>Heterotylenchus autumnalis</i>	Poinar, 1979	1	1	1	1	1	1	0.00000000
	DIPLOGASTERIDAE	<i>Mesodiplogaster theritteri</i>	Poinar, 1979	4	3	1	1	1	2001	3.30124709
	HETERORHABDITIDAE	<i>Heterorhabditis bacteriophora</i>	Poinar, 1979	4	4	3	1	1	750499	5.87535012
n (genera) = 1; n (species) = 2		<i>Heterorhabditis heliothidis</i>	Poinar, 1979	1	1	1	1	1	1	0.00000000
MERMITHIDAE		<i>Agamermis decudata</i>	Poinar, 1979	1	1	1	1	1	1	0.00000000
n (genera) = 10		<i>Filpjivimermis leipsandra</i>	Poinar, 1979	10	6	3	2	1	46041046	7.66314518
n (species) = 12		<i>Gastromermis viridis</i>	Poinar, 1979	2	2	1	1	1	1001	3.00043408
		<i>Hydromermis churchillensis</i>	Poinar, 1979	4	2	1	1	1	1003	3.00130093
		<i>Hydromermis contorta</i>	Poinar, 1979	5	3	2	1	1	376252	5.57547882
		<i>Isomermis lairdi</i>	Poinar, 1979	6	4	1	1	1	3000	3.47712125
		<i>Isomermis wisconsensis</i>	Poinar, 1979	6	1	1	1	1	6	0.77815125
		<i>Limnomeritis rosea</i>	Poinar, 1979	3	2	1	1	1	2002	3.30146407
		<i>Mermis nigrescens</i>	Poinar, 1979	2	1	1	1	1	2	0.30103000
		<i>Mesomeritis flumenalis</i>	Poinar, 1979	58	37	12	5	1	183010057	8.26247496
			Poinar, 1979	9	2	1	1	1	1008	3.00346053

		<i>Octomyomermis muspratti</i>	Poinar, 1979	10	2	1	1	1	1009	3.00389117
		<i>Strelkovimermis peterseni</i>	Poinar, 1979	3	1	1	1	1	3	0.47712125
	NEOTYLENCHIDAE	<i>Deladenus siricidicola</i>	Poinar, 1979	6	3	2	2	1	45664803	7.65958159
	STEINERNEMATIDAE	<i>Neoaplectana bibionis</i>	Poinar, 1979	3	2	1	1	1	1002	3.00086772
n (genera) = 1		<i>Neoaplectana carpocapsae</i>	Poinar, 1979	7	6	3	3	1	92069112	7.96411395
n (species) = 4		<i>Neoaplectana glaseri</i>	Poinar, 1979	5	5	4	2	1	46412299	7.66663308
		<i>Neoaplectana leucaniae</i>	Poinar, 1979	2	2	2	1	1	375251	5.57432186
	CALIGIDAE	<i>Anuretes heckelli</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
n (genera) = 10		<i>Anuretes quadrilaterus</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
n (species) = 80		<i>Caligodes laciniatus</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Caligus asymmetricus</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Caligus auxisi</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Caligus belones</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Caligus bonito</i>	Love & Moser, 1983	8	7	5	2	1	46785547	7.67011171
		<i>Caligus chelifer</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Caligus chorinemi</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Caligus clemensi</i>	Margolis & Arthur, 1979; Love & Moser, 1983	13	8	7	6	2	2629059755	9.41980046
		<i>Caligus confusus</i>	Love & Moser, 1983	2	2	2	1	1	375251	5.57432186
		<i>Caligus constrictus</i>	Love & Moser, 1983	2	2	2	1	1	375251	5.57432186
		<i>Caligus coryphaenae</i>	Love & Moser, 1983	12	8	7	3	2	2496552369	9.39734068
		<i>Caligus curtus</i>	Margolis & Arthur, 1979	7	7	4	3	2	2495440590	9.39714723
		<i>Caligus diaphanus</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Caligus elongatus</i>	Love & Moser, 1984	23	23	17	12	2	2887757255	9.46056068
		<i>Caligus epidemicus</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Caligus germani</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Caligus glandifer</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Caligus hobsoni</i>	Love & Moser, 1983	15	12	7	3	1	92444334	7.96588030
		<i>Caligus irritans</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Caligus klavrei</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Caligus lacustris</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Caligus lunatus</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Caligus macarovi</i>	Margolis & Arthur, 1979; Love & Moser, 1983	2	2	2	2	1	45663801	7.65957206
		<i>Caligus minimus</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Caligus mutabilis</i>	Love & Moser, 1983	9	7	5	1	1	1496998	6.17522122
		<i>Caligus olsoni</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Caligus orientalis</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Caligus patulus</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Caligus pectinatus</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Caligus pelamydus</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Caligus platytarsis</i>	Love & Moser, 1983	3	3	2	1	1	376250	5.57547651
		<i>Caligus praetextus</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Caligus productus</i>	Love & Moser, 1983	4	4	4	2	2	2450523302	9.38925884
		<i>Caligus quadatus</i>	Love & Moser, 1983	17	12	8	2	1	47898307	7.68032016
		<i>Caligus rectus</i>	Love & Moser, 1983	5	5	4	1	1	1123749	6.05066932
		<i>Caligus robustus</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
				1	1	1	1	1	1	0.00000000

C O P E P O D A	<i>Caligus rufimaculatus</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	<i>Caligus serratus</i>	Love & Moser, 1983	3	3	2	2	1	45664800	7.65958156
	<i>Caligus spinosurculus</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	<i>Caligus tenax</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	<i>Caligus tenuicaudatus</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	<i>Caligus tenuifrucatus</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	<i>Dentigrypus curtus</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	<i>Lepeophtheirus abdominalis</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	<i>Lepeophtheirus appendiculatus</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	<i>Lepeophtheirus bifidus</i>	Love & Moser, 1983	4	4	2	1	1	377248	5.57662695
	<i>Lepeophtheirus bifurcatus</i>	Love & Moser, 1983	3	3	2	2	1	45664800	7.65958156
	<i>Lepeophtheirus brachyurus</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	<i>Lepeophtheirus breviventris</i>	Margolis & Arthur, 1979; Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	<i>Lepeophtheirus constrictus</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	<i>Lepeophtheirus crassus</i>	Love & Moser, 1983	3	1	1	1	1	3	0.47712125
	<i>Lepeophtheirus cuneifer</i>	Love & Moser, 1983	2	1	1	1	1	2	0.30103000
	<i>Lepeophtheirus dissimilatus</i>	Love & Moser, 1983	10	10	7	6	2	2629061735	9.41980078
	<i>Lepeophtheirus edwardsi</i>	Love & Moser, 1983	8	8	7	4	1	136981414	8.13666164
	<i>Lepeophtheirus emiens</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	<i>Lepeophtheirus goniostii</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	<i>Lepeophtheirus hastatus</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	<i>Lepeophtheirus hexagrammi</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	<i>Lepeophtheirus hospitalis</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	<i>Lepeophtheirus longipes</i>	Margolis & Arthur, 1979; Love & Moser, 1983	6	6	3	3	1	90955342	7.95882821
	<i>Lepeophtheirus longispinosus</i>	Love & Moser, 1983	5	4	4	2	1	46411303	7.66662376
	<i>Lepeophtheirus nanaimoensis</i>	Margolis & Arthur, 1979	1	1	1	1	1	1	0.00000000
	<i>Lepeophtheirus nordmanni</i>	Margolis & Arthur, 1979; Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	<i>Lepeophtheirus oblitus</i>	Margolis & Arthur, 1979; Love & Moser, 1983	7	2	2	1	1	375256	5.57432764
	<i>Lepeophtheirus parvicirrus</i>	Margolis & Arthur, 1979; Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	<i>Lepeophtheirus parviventris</i>	Margolis & Arthur, 1979; Love & Moser, 1983	21	18	11	5	2	2586739228	9.41275265
	<i>Lepeophtheirus parvus</i>	Love & Moser, 1983	3	3	2	1	1	376250	5.57547651
	<i>Lepeophtheirus paulus</i>	Margolis & Arthur, 1979; Love & Moser, 1983	6	1	1	1	1	6	0.77815125
	<i>Lepeophtheirus pollachioides</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	<i>Lepeophtheirus pravipes</i>	Margolis & Arthur, 1979; Love & Moser, 1983	4	4	4	3	2	2495437602	9.39714671
	<i>Lepeophtheirus salmonis</i>	Love & Moser, 1983	14	7	5	4	1	136240902	8.13430751
	<i>Lepeophtheirus thompsoni</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	<i>Pseudocaligus apodus</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	<i>Pseudolepeophtheirus longicauda</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	<i>Pupulina brevicauda</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	<i>Pupulina minor</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	<i>Sciaenophilus bennetti</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	<i>Tuxophorus caligodes</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
	<i>Demoleus heptapus</i>	Love & Moser, 1983	2	2	2	2	1	45663801	7.65957206
	<i>Demoleus latus</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
<b>PANDARIDAE</b>									
n (genera) = 12									

## ARTHROPODA

A R T H R O P O D A	n (species) = 28	<i>Dinemoura discrepans</i>	Love & Moser, 1983	2	1	1	1	1	2	0.30103000
		<i>Dinemoura latifolia</i>	Love & Moser, 1983	5	5	4	2	1	46412299	7.66663308
		<i>Dinemoura producta</i>	Love & Moser, 1983	6	6	4	2	1	46413295	7.66664240
		<i>Echthrogaleus coleoptratus</i>	Love & Moser, 1983	6	6	5	4	2	2540351000	9.40489373
		<i>Echthrogaleus denticulatus</i>	Love & Moser, 1983	4	4	4	2	1	46411302	7.66662375
		<i>Gangliopus pyriformis</i>	Love & Moser, 1983	3	3	3	2	1	46038051	7.66311693
		<i>Nesippus borealis</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Nesippus crypturus</i>	Love & Moser, 1983	5	3	2	1	1	376252	5.57547882
		<i>Nesippus gracilis</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Nesippus orientalis</i>	Love & Moser, 1983	6	3	2	1	1	376253	5.57547997
		<i>Nesippus tigris</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Nogagus ambiguus</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Pagina tunica</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Pandarus bicolor</i>	Love & Moser, 1983	8	8	6	5	2	2584892945	9.41244256
		<i>Pandarus carcharini</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Pandarus cranchii</i>	Love & Moser, 1983	10	7	4	2	1	46414293	7.66665174
		<i>Pandarus floridanus</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Pandarus katoi</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Pandarus satyrus</i>	Love & Moser, 1983	4	4	3	2	1	46039049	7.66312634
		<i>Pandarus sinuatus</i>	Love & Moser, 1983	5	4	3	2	2	2450151050	9.38919286
		<i>Pandarus spinaciatacanthiae</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Pannosus japonicus</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
		<i>Perissopus dentatus</i>	Love & Moser, 1983	4	2	2	1	1	375253	5.57432417
		<i>Perissopus oblongatus</i>	Love & Moser, 1983	11	9	8	6	2	2629429016	9.41986145
		<i>Phyllosthyreus cornutus</i>	Love & Moser, 1983	3	3	3	2	1	46038051	7.66311693
		<i>Prasaeetes rhinodontis</i>	Love & Moser, 1983	1	1	1	1	1	1	0.00000000
IXODIDAE		<i>Dermacentor albipictus</i>	Kennedy & Newman, 1986	8	7	3	2	1	46042038	7.66315454
n (genera) = 3		<i>Dermacentor andersoni</i>	Kennedy & Newman, 1986	25	18	10	4	1	138093141	8.14017211
n (species) = 19		<i>Dermacentor variabilis</i>	Kennedy & Newman, 1986	19	16	9	4	1	137724882	8.13901241
		<i>Haemophysalis leporis-palustris</i>	Kennedy & Newman, 1986	9	7	4	3	1	91328592	7.96060676
		<i>Ixodes angustus</i>	Kennedy & Newman, 1986	23	13	6	3	1	90076065	7.95460941
		<i>Ixodes cookei</i>	Kennedy & Newman, 1986	11	10	7	3	1	92442351	7.96587098
		<i>Ixodes echotonae</i>	Kennedy & Newman, 1986	3	3	2	2	1	45664800	7.65958156
		<i>Ixodes hearlei</i>	Kennedy & Newman, 1986	1	1	1	1	1	1	0.00000000
		<i>Ixodes hexagononus</i>	Kennedy & Newman, 1986	1	1	1	1	1	1	0.00000000
		<i>Ixodes kingi</i>	Kennedy & Newman, 1986	5	4	2	2	1	45665799	7.65959106
		<i>Ixodes marmotae</i>	Kennedy & Newman, 1986	1	1	1	1	1	1	0.00000000
		<i>Ixodes marxi</i>	Kennedy & Newman, 1986	5	5	3	3	1	90954346	7.95882346
		<i>Ixodes muris</i>	Kennedy & Newman, 1986	12	10	4	2	1	46417271	7.66667960
		<i>Ixodes pacificus</i>	Kennedy & Newman, 1986	8	8	5	3	1	91700840	7.96237331
		<i>Ixodes rugosus</i>	Kennedy & Newman, 1986	1	1	1	1	1	1	0.00000000
		<i>Ixodes sculptus</i>	Kennedy & Newman, 1986	5	3	3	2	1	46038053	7.66311693
		<i>Ixodes soricis</i>	Kennedy & Newman, 1986	2	1	1	1	1	2	0.30103000
		<i>Ixodes spinipalpus</i>	Kennedy & Newman, 1986	6	6	5	3	1	91698851	7.96236389
		<i>Ixodes texanus</i>	Kennedy & Newman, 1986	8	8	4	2	1	46415284	7.66666101

HARPYRHYNCHIDAE n (genera) = 2 n (species) = 114	<i>Harpyrhynchus</i> sp. 1 - 91	Moss, 1979	1	1	1	1	1	1	0.0000000
	<i>Harpyrhynchus brevis</i>	Moss, 1979	29	27	7	1	1	2256213	6.35338010
	<i>Harpyrhynchus coturnix</i>	Moss, 1979	2	2	2	1	1	375251	5.57432186
	<i>Harpyrhynchus cylindripalpis</i>	Moss, 1979	6	6	4	1	1	1124745	6.05105407
	<i>Harpyrhynchus monstrosis</i>	Moss, 1979	9	8	3	2	1	46043032	7.66316391
	<i>Harpyrhynchus</i> n.sp.1	Moss, 1979	2	1	1	1	1	2	0.3010300
	<i>Harpyrhynchus nidulens</i>	Moss, 1979	7	7	5	1	1	1496996	6.17522064
	<i>Harpyrhynchus novoplumeris</i>	Moss, 1979	7	7	5	2	1	46785546	7.67011170
	<i>Harpyrhynchus pilirostris</i>	Moss, 1979	2	2	2	1	1	375251	5.57432186
	<i>Harpyrhynchus plumeris</i>	Moss, 1979	7	5	4	2	1	46412301	7.66663310
	<i>Harpypalpus</i> sp. 1 - 11	Moss, 1979	1	1	1	1	1	1	0.0000000
	<i>Harpypalpus holopus</i>	Moss, 1979	3	3	3	1	1	749501	5.87477222
	<i>Harpypalpus longipes</i>	Moss, 1979	7	6	4	1	1	1124746	6.05105446
	<i>Harpypalpus</i> n.sp. 1	Moss, 1979	2	2	2	1	1	375251	5.57432186
CERATOPHYLLIDAE n (genera) = 16 n (species) = 53	<i>Amalaraeus athabascae</i>	Kennedy & Newman, 1986	7	5	2	2	1	45666797	7.65960055
	<i>Amalaraeus dissimilis</i>	Kennedy & Newman, 1986	26	15	6	3	1	92078041	7.96415607
	<i>Amphalius runatus necopinus</i>	Kennedy & Newman, 1986	2	1	1	1	1	2	0.30103000
	<i>Ceratophyllus adustus</i>	Kennedy & Newman, 1986	1	1	1	1	1	1	0.0000000
	<i>Ceratophyllus diphinis</i>	Kennedy & Newman, 1986	1	1	1	1	1	1	0.0000000
	<i>Ceratophyllus gallinae</i>	Kennedy & Newman, 1986	3	3	2	1	1	376250	5.57547651
	<i>Ceratophyllus garei</i>	Kennedy & Newman, 1986	8	7	5	3	1	91699847	7.96236861
	<i>Ceratophyllus lunatus tundrensis</i>	Kennedy & Newman, 1986	10	7	3	2	1	46042040	7.66315456
	<i>Ceratophyllus niger</i>	Kennedy & Newman, 1986	1	1	1	1	1	1	0.0000000
	<i>Ceratophyllus petrochelidoni</i>	Kennedy & Newman, 1986	1	1	1	1	1	1	0.0000000
	<i>Dactylophysylla comis</i>	Kennedy & Newman, 1986	1	1	1	1	1	1	0.0000000
	<i>Dactylophysylla rara</i>	Kennedy & Newman, 1986	2	2	2	2	1	45663801	7.65957206
	<i>Dasyphylax gallinulae perpinnatus</i>	Kennedy & Newman, 1986	3	3	2	1	1	376250	5.57547651
	<i>Foxellia ignota</i>	Kennedy & Newman, 1986	11	7	5	2	1	46785550	7.67011174
	<i>Malaraeus bitterrootensis</i>	Kennedy & Newman, 1986	4	3	2	2	1	45664801	7.65958151
	<i>Malaraeus euphorbi</i>	Kennedy & Newman, 1986	5	4	3	2	1	46039050	7.66312635
	<i>Malaraeus telchinus</i>	Kennedy & Newman, 1986	15	9	5	3	1	91701839	7.96237805
	<i>Megabothris abanitis</i>	Kennedy & Newman, 1986	31	17	9	4	1	137725878	8.13901555
	<i>Megabothris acerbus</i>	Kennedy & Newman, 1986	5	4	2	1	1	377249	5.57662810
	<i>Megabothris asio</i>	Kennedy & Newman, 1986	35	19	6	3	1	92081984	7.96417467
	<i>Megabothris atrox</i>	Kennedy & Newman, 1986	5	3	2	1	1	376252	5.57547882
	<i>Megabothris calcariifer gregsoni</i>	Kennedy & Newman, 1986	17	8	3	3	1	90957340	7.95883775
	<i>Megabothris clantonii</i>	Kennedy & Newman, 1986	1	1	1	1	1	1	0.0000000
	<i>Megabothris greenlandicus</i>	Kennedy & Newman, 1986	15	8	3	2	1	46043038	7.66316397
	<i>Megabothris lucifer</i>	Kennedy & Newman, 1986	11	7	3	2	1	46042041	7.66315457
	<i>Megabothris quirini</i>	Kennedy & Newman, 1986	40	21	7	4	1	136994264	8.13670238
	<i>Monopsyllus ciliatus protinus</i>	Kennedy & Newman, 1986	20	14	6	3	1	92077049	7.96415138
	<i>Monopsyllus eumolpi</i>	Kennedy & Newman, 1986	18	12	5	3	1	91704812	7.96239212
	<i>Monopsyllus thambus</i>	Kennedy & Newman, 1986	7	5	2	2	1	45666797	7.65960055
	<i>Monopsyllus tolli</i>	Kennedy & Newman, 1986	2	2	2	2	1	456663801	7.65957206

R  
A

R	<i>Monopsyllus vison</i>	Kennedy & Newman, 1986	29	16	7	3	1	92448294	7.9658989	
A	<i>Monopsyllus wagneri</i>	Kennedy & Newman, 1986	25	16	8	4	1	137357610	8.1378527	
	<i>Nosopsyllus fasciatus</i>	Kennedy & Newman, 1986	11	8	3	2	1	46043034	7.66316393	
	<i>Opisocrotis bruneri</i>	Kennedy & Newman, 1986	10	7	4	3	1	91328593	7.96060677	
	<i>Opisocrotis labis</i>	Kennedy & Newman, 1986	5	4	2	2	1	44665799	7.6499751	
	<i>Opisocrotis tuberculatus</i>	Kennedy & Newman, 1986	10	6	4	2	1	46413299	7.6666424	
	<i>Opisodasy keeni keeni</i>	Kennedy & Newman, 1986	12	7	4	2	1	46414295	7.6666517	
	<i>Opisodasy pseudoaureotomys</i>	Kennedy & Newman, 1986	8	6	4	2	1	46413297	7.6666424	
	<i>Opisodasy vesperalis</i>	Kennedy & Newman, 1986	5	5	4	3	1	91326599	7.9605972	
	<i>Orchopeas caedens</i>	Kennedy & Newman, 1986	23	13	5	2	1	46791505	7.6701670	
	<i>Orchopeas howardi howardi</i>	Kennedy & Newman, 1986	6	4	1	1	1	3000	3.4771212	
	<i>Orchopeas leucopus</i>	Kennedy & Newman, 1986	23	13	4	2	1	46420249	7.6667074	
	<i>Orchopeas nepos</i>	Kennedy & Newman, 1986	7	5	2	2	1	45666797	7.6596005	
	<i>Orchopeas sextentatus agilis</i>	Kennedy & Newman, 1986	17	14	6	3	1	92077046	7.9641513	
	<i>Oropsylla alaskenensis</i>	Kennedy & Newman, 1986	6	5	5	2	1	46783556	7.6700932	
	<i>Oropsylla arctomyia</i>	Kennedy & Newman, 1986	18	13	6	3	1	92076060	7.9641467	
	<i>Oropsylla idahoensis</i>	Kennedy & Newman, 1986	18	7	5	3	1	91698957	7.9623686	
	<i>Oropsylla rupestris</i>	Kennedy & Newman, 1986	13	9	6	3	1	92070297	7.9641195	
	<i>Tarsopsylla coloradensis</i>	Kennedy & Newman, 1986	6	5	3	2	1	46040047	7.6631357	
	<i>Thrassis acamantis</i>	Kennedy & Newman, 1986	11	9	3	2	1	46044026	7.6631732	
	<i>Thrassis bacchi bacchi</i>	Kennedy & Newman, 1986	5	3	2	2	1	456664802	7.6595815	
	<i>Thrassis peteolatus</i>	Kennedy & Newman, 1986	7	4	3	1	1	750502	5.8753518	
	<i>Thrassis spenceri</i>	Kennedy & Newman, 1986	5	2	2	2	1	45663804	7.6595720	
A	PHILOPTERIDAE	<i>Cotingocola dimorpha</i>	Clayton & Price, 1998	1	1	1	1	1	0.0000000	
L	n (genera) = 1	<i>Cotingocola fizpatricki</i>	Clayton & Price, 1998	1	1	1	1	1	0.0000000	
L	n (species) = 8	<i>Cotingocola gracilis</i>	Clayton & Price, 1998	1	1	1	1	1	0.0000000	
O		<i>Cotingocola meridae</i>	Clayton & Price, 1998	2	1	1	1	1	0.3010300	
P		<i>Cotingocola parnipapillae</i>	Clayton & Price, 1998	2	2	1	1	1	1001	3.0004340
H		<i>Cotingocola rupicolae</i>	Clayton & Price, 1998	3	3	1	1	1	2000	3.3010300
A		<i>Cotingocola stotzi</i>	Clayton & Price, 1998	1	1	1	1	1	0.0000000	
G		<i>Cotingocola tergalis</i>	Clayton & Price, 1998	3	1	1	1	1	3	0.4771212
	n (genera) = 25	<i>Antrodiella liebmanni</i>	Lindblad, 2000	1	1	1	1	1	0.0000000	
	n (species) = 47	<i>Antrodiella semispina</i>	Lindblad, 2000	1	1	1	1	1	0.0000000	
		<i>Antrodiella sp.</i>	Lindblad, 2000	2	2	2	2	1	45663801	7.6595720
		<i>Aurifaria luteoumbrina</i>	Lindblad, 2000	4	4	4	4	1	135866651	8.1331128
		<i>Ceriporia alachuana</i>	Lindblad, 2000	5	5	4	3	1	91326599	7.9605972
		<i>Ceriporia ferruginicincta</i>	Lindblad, 2000	1	1	1	1	1	0.0000000	
		<i>Ceriporia microspora</i>	Lindblad, 2000	2	2	2	2	1	45663801	7.6595720
		<i>Ceriporia reticulata</i>	Lindblad, 2000	1	1	1	1	1	0.0000000	
		<i>Ceriporia xylostromatoides</i>	Lindblad, 2000	11	10	10	7	1	269478851	8.4305246
		<i>Ceriporiopsis latemarginata</i>	Lindblad, 2000	1	1	1	1	1	0.0000000	
		<i>Ceriporiopsis lowei</i>	Lindblad, 2000	1	1	1	1	1	0.0000000	

A P H Y L L O P H O R A L E S	<i>Ceriporiopsis mucida</i>	Lindblad, 2000	8	8	7	7	1	268375030	8.4287421
	<i>Ceriporiopsis sp.</i>	Lindblad, 2000	1	1	1	1	1	1	0.0000000
	<i>Coriolopsis byrsina</i>	Lindblad, 2000	12	12	10	7	1	269480831	8.4305278
	<i>Coriolopsis polyzona</i>	Lindblad, 2000	1	1	1	1	1	1	0.0000000
	<i>Coriolopsis rigidula</i>	Lindblad, 2000	13	11	7	6	1	224950728	8.3520874
	<i>Cotylidia aurantiaca</i>	Lindblad, 2000	2	2	2	2	1	45663801	7.6595720
	<i>Cristelloporia dimittita</i>	Lindblad, 2000	2	2	2	2	1	45663801	7.6595720
	<i>Datronia caperata</i>	Lindblad, 2000	10	10	9	7	1	269112564	8.4299339
	<i>Datronia scutellata</i>	Lindblad, 2000	2	2	2	2	1	45663801	7.6595720
	<i>Datronia steroidea</i>	Lindblad, 2000	1	1	1	1	1	1	0.0000000
	<i>Diplomitoporus costaricensis</i>	Lindblad, 2000	3	3	3	3	1	90952351	7.9588139
	<i>Earliella scabrosa</i>	Lindblad, 2000	7	7	6	4	1	136611155	8.1354861
	<i>Filoboletus gracilis</i>	Lindblad, 2000	2	2	2	2	1	45663801	7.6595720
	<i>Fomes fasciatus</i>	Lindblad, 2000	1	1	1	1	1	1	0.0000000
	<i>Fomitopsis feei</i>	Lindblad, 2000	2	2	2	2	1	45663801	7.6595720
	<i>Fuscocerrerena portoricensis</i>	Lindblad, 2000	1	1	1	1	1	1	0.0000000
	<i>Ganoderma amazonense</i>	Lindblad, 2000	1	1	1	1	1	1	0.0000000
	<i>Ganoderma australe</i>	Lindblad, 2000	2	2	2	2	1	45663801	7.6595720
	<i>Gleophyllum striatum</i>	Lindblad, 2000	3	3	2	2	1	45664800	7.6595815
	<i>Gleoporus thelephoroides</i>	Lindblad, 2000	4	4	4	3	1	91325602	7.9605925
	<i>Hexagonia hydnoides</i>	Lindblad, 2000	2	2	2	2	1	45663801	7.6595720
	<i>Hexagonia papryacea</i>	Lindblad, 2000	3	3	3	3	1	90952351	7.9588139
	<i>Hymenochaete berteroii</i>	Lindblad, 2000	1	1	1	1	1	1	0.0000000
	<i>Hymenochaete dura</i>	Lindblad, 2000	1	1	1	1	1	1	0.0000000
	<i>Hymenochaete epichlora</i>	Lindblad, 2000	2	2	1	1	1	2	0.3010300
	<i>Hymenochaete sp.</i>	Lindblad, 2000	1	1	1	1	1	1	0.0000000
	<i>Inonotus dryophilus</i>	Lindblad, 2000	1	1	1	1	1	1	0.0000000
	<i>Melanoporella carbonacea</i>	Lindblad, 2000	1	1	1	1	1	1	0.0000000
	<i>Microporellus obovatus</i>	Lindblad, 2000	2	2	2	2	1	45663801	7.6595720
	<i>Nigroporus vinosus</i>	Lindblad, 2000	2	2	2	2	1	45663801	7.6595720
	<i>Perenniporia ochroleuca</i>	Lindblad, 2000	3	3	3	2	1	46038051	7.66311692
	<i>Perenniporia roseo-isabellina</i>	Lindblad, 2000	1	1	1	1	1	1	0.0000000
	<i>Perenniporia tepeitensis</i>	Lindblad, 2000	1	1	1	1	1	1	0.0000000
	<i>Perenniporia tephropora</i>	Lindblad, 2000	5	5	4	4	1	135867648	8.13311606
	<i>Phellinus chrysaeus</i>	Lindblad, 2000	2	2	2	2	1	45663801	7.6595720
	<i>Phellinus contiguus</i>	Lindblad, 2000	4	4	3	3	1	90953349	7.95881865

#### Appendix 1. Host specificity of selected parasite species.

## APPENDIX II.

Appendix 2. The C++ code for algorithm to compute host specificity rank and corresponding index value.

```

typedef double IndexValue;

IndexValue host_index(const int host_species, const int host_genera,
                      const int host_families, const int host_orders,
                      const int host_classes)
{
    IndexValue value = 0;
    if (host_classes > 1) {
        value = host_index(n_species, n_genera, n_families, n_orders,
                             host_classes-1);
        IndexValuef_index = host_index(host_species, host_genera, host_families,
                                   host_orders, 1);
        IndexValueo_index = host_index(host_classes, host_classes, host_classes,
                                   host_classes, 1);
        value += f_index - o_index + 1;
    } else if (host_orders > 1) {
        value = host_index(n_species, n_genera, n_families, host_orders-1, 1);
        IndexValuef_index = host_index(host_species, host_genera, host_families,
                                   1, 1);
        IndexValueo_index = host_index(host_orders, host_orders, host_orders, 1, 1);
        value += f_index - o_index + 1;
    } else if (host_families > 1) {
        value = host_index(n_species, n_genera, host_families-1, 1, 1);
        IndexValuef_index = host_index(host_species, host_genera, 1, 1, 1);
        IndexValueo_index = host_index(host_families, host_families, 1, 1, 1);
        value += f_index - o_index + 1;
    } else if (host_genera > 1) {
        for (int g = 1; g < host_genera; g++) {
            value += n_species - g + 1;
        }
        value += host_species - host_genera + 1;
        // The approachon the followinglines follows the same recursive
        // logicas above, but is less computationallyefficient than
        // the code actuallyused. It producesidenticalresults.
        //
        // value= host_index(n_species, host_genera-1, 1, 1, 1);
        // IndexValuef_index = host_index(host_species, 1, 1, 1, 1);
        // IndexValueo_index = host_index(host_genera, 1, 1, 1, 1);
        // value += f_index - o_index + 1;
    } else {
        value = host_species;
    }
    return value;
}

```

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