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# **Update on Human Granulocytic Anaplasmosis**

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**ABSTRACT:** Granulocytic anaplasmosis (GA) is a disease of humans, domestic animals, and wildlife caused by *Anaplasma phagocytophilum*, formerly *Ehrlichia phagocytophila*, *E. equi*, and the unnamed agents of "human granulocytic ehrlichiosis" (HGE). This pathogen is inoculated into host skin by the bite of *Ixodes* spp. ticks, including *I. pacificus* in California and *I. scapularis* in the eastern U.S. After inoculation, *A. phagocytophilum* disseminates to the blood and is phagocytosed into host neutrophils. The clinical characteristics of GA in people vary from no symptoms to fever, headache, neurological symptoms, and occasionally death. Horses with GA may experience high fever, depression, reduced ability to eat, limb edema, jaundice, and ataxia. GA is an emerging disease in the eastern U.S. but only a handful of human cases have been reported in the western U.S., despite relatively common reports of disease in horses and dogs. Wildlife and dog-sentinel studies have clarified that infection is common in the coast range mountains and Sierra Nevada foothills, with ongoing research focusing on ecological determinants that can modify the prevalence of infection in any particular area. One known reservoir is the dusky-footed woodrat. However, important, poorly understood ecological determinants serve to modify the probability of rodent as well as human infections, including climate, vegetation, and possibly the presence of other reservoir-competent rodents and nidicolous woodrat-specialist ticks such as *I. spinipalpis*.

**KEY WORDS:** Anaplasma phagocytophilum, disease, dusky-footed woodrat, Ixodes pacificus, Ixodes spinipalpis, Neotoma fuscipes, tick-borne disease

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## THE ORGANISM AND DISEASE Phylogeny and Biological Characteristics

Granulocytic anaplasmosis (GA) is a disease of humans, domestic animals, and wildlife caused by the newly renamed pathogen, Anaplasma phagocytophilum (Dumler et al. 2001). This new species combines Ehrlichia phagocytophila, E. equi, and the unnamed agents of "human granulocytic ehrlichiosis" (HGE), which cross-react serologically (Walker and Dumler 1995) and are indistinguishable morphologically (Popov et al. 1998). There is microvariability in the 16S rRNA gene of strains from the upper midwestern, eastern, and western U.S. among different host species (Chen et al. 1994, Reubel et al. 1998), but analysis of the 16S, 444 ep-Ank, and groESL heat shock protein genetic regions from equine and human-origin isolates from northern California suggested that the HGE agent and E. equi in the U.S. are synonymous, with multiple strains that may vary geographically but do not appear to be hostdifferentiated (Chae et al. 2000).

A. phagocytophilum shares common biological characteristics and immunopathological mechanisms in hosts with other ehrlichial and rickettsial bacteria. This organism is inoculated into host skin by tick bite, in context with tick saliva, which contains an array of anti-inflammatory and immunomodulatory chemicals during feeding to prevent host blood clotting, platelet aggregation, NK cell activity, and T<sub>H</sub>1 cytokine responses (Wikel 1999). After tick inoculation and local host immune responses, A. phagocytophilum disseminates to the blood, where this obligate intra-cellular pathogen must persist within host neutrophils. The bacterium enters the neutrophil after attaching via an outer surface protein to a host cell receptor and being phagocytosed into a small host membrane-bound site called a

phagosome. *A. phagocytophilum*-infected neutrophils secrete chemokines that recruit macrophages, which then secrete IFN-γ and IL-10 and amplify local and systemic inflammation (Dumler *et al.* 2000, Klein *et al.* 2000). Within the phagosome, *A. phagocytophilum* can inhibit fusion of this protected environment with the lysosome, overcoming initial host immune attacks against the bacteria. Ultimately, however, when host cells are primed with IFN-γ, they are able to kill the intracellular *A. phagocytophilum* (Webster *et al.* 1998, Mott *et al.* 1999).

### **Clinical Outcomes of Infection**

The clinical characteristics of GA in people are variable including pyrexia, headache, myalgia, nausea, and ataxia (Goodman *et al.* 1996). More severe cases may result in organ failure, susceptibility to opportunistic infections, neuritis, or respiratory complications, with a case fatality rate up to 5% (Walker and Dumler 1995, Foley 2000). GA in horses is characterized by high fever, depression, reduced ability to eat, limb edema, petechiation, icterus, and ataxia (Madigan 1993). Counts of blood cells may document anemia, low platelet counts, and low white blood cell counts. Because much of the pathogenesis of GA is immune-mediated, there may be downstream immune-mediated sequelae, including possible polyarthritis and vasculitis.

# Laboratory Findings in GA and Diagnosis

Although most cases of GA occur in areas where tick biting is common, not all human patients report a tick bite and not all dog or horse owners recognize that their animal has been infected. The symptoms and laboratory diagnostic results of GA are non-specific and may include elevated liver enzymes, thrombocytopenia, and neutro-

penia. In some cases, the symptoms (e.g., headache and fever) are severe, and yet there are only mild or no laboratory abnormalities.

Probably the most rapid method for diagnosis of GA is to find the characteristic membrane-bound inclusions (morulae) in neutrophils on blood smears, but the proportion of neutrophils with visible morulae may be as low as 1-2%. The Centers for Disease Control (CDC) criteria for a confirmed diagnosis require visualization of morulae be accompanied by positive titers (≥64), probably because some cellular inclusions may appear similar to morulae. Although a single titer ≥64 may be part of a presumptive diagnosis, this does not document active infection. However, a 4-fold rise in specific titer over 4 weeks is considered confirmatory for GA. Titers become positive approximately 2 weeks after onset of clinical signs and remain elevated after exposure for months to years, so there is a brief window early in the infection when the patient may be ill but seronegative.

Culture of *A. phagocytophilum* is time-consuming, specialized, and somewhat expensive. Polymerase chain reaction (PCR) for specific DNA is extremely valuable for confirming a suspected diagnoses but generally not available rapidly enough in practice for use in primary diagnosis and to guide treatment. There are multiple published protocols for PCR of *A. phagocytophilum*. PCR may be most useful early in acute infection before the patient seroconverts, late in infection to rule-out chronic infection, or when serological titers to several agents are similar.

#### ECOLOGY Emergence

Human granulocytic anaplasmosis (HGA) emerged dramatically in the eastern and midwestern U.S. over the past 25 years. Since first reported in Wisconsin in 1994, over 400 cases of HGA have been diagnosed in the upper midwestern and northeastern United States (CDC 2000). In contrast, only 8 human cases have been documented in California as of April 2002, despite being reportable to the California Department of Health Services (CDHS), availability of diagnostic laboratory support, and dissemination of information to physicians by CDHS personnel and others. Yet, GA is common in dogs and horses (Madigan and Gribble 1987, Foley et al. 2001). However, following targeting of the north coast of California by state and local health officials, 4 cases of HGE were documented in one county within 12 months where no case had previously been confirmed (Foley et al. Thus, cases within the state are no doubt 1999*a*). somewhat under-reported.

#### **Tick Vectors**

Worldwide, Anaplasma phagocytophilum is vectored by at least 4 tick species and hosted by dozens or hundreds of mammals, reptiles, and birds. Ixodes scapularis is the vector in the eastern U.S., I. pacificus in California, I. ricinus in Europe, and I. persulcatus in Asia (MacLeod and Gordon 1933, Pancholi et al. 1995, Richter et al. 1996, Telford et al. 1996, Des Vignes and Fish 1997, Petrovec et al. 1997). Locally important or enzootic vectors may include I. hexagonus in England

and *I. spinipalpis* in the western U.S.

The spatial distribution of human and equine cases of GA in the western U.S. corresponds with the range of Ixodes pacificus, the western black-legged tick, a common tick throughout much of low mountain California (Furman and Loomis 1984, Madigan and Gribble 1987, Richter et al. 1996). I. pacificus has been recorded from 55 of 58 counties in California, and adult I. pacificus containing A. phagocytophilum DNA have been detected in Alameda, El Dorado, Humboldt, Napa, Orange, Sacramento, Santa Cruz, Sonoma, and Yolo Counties (Barlough et al. 1997, Kramer et al. 1999; Nicholson et al. 1999; Lane et al. 2001), although tick surveillance has been limited. The vector competence of I. pacificus for A. phagocytophilum has been evaluated. Infected I. pacificus individuals can be found in nature (Richter et al. 1996; Barlough et al. 1997, Kramer et al. 1999, Lane et al. 2001); horses can be experimentally infected using field-caught, naturally-infected ticks (Reubel et al. 1998); and the agent can be transmitted from experimentally-infected horses to naïve horses by feeding of *I. pacificus* (Richter *et al.* 1996).

The prevalence of A. phagocytophilum in ticks ranges from 0-51% in the eastern U.S. (Magnarelli et al. 1995, Pancholi et al. 1995, Daniels et al. 1997, Varde et al. 1998), but typically is much lower in the western U.S., from 0.8-11% (Barlough et al. 1997, Kramer et al. 1999, Nicholson et al. 1999, Lane et al. 2001). The prevalence in *I. pacificus* nymphs is also low, with 0/465 nymphs positive even where the adult prevalence was 11% (Lane et al. 2001) and only 1/47 pools positive from Sonoma County (Barlough et al. 1997). Yet, nymphs may cause most human, deer, and mountain lion cases, because these peak typically in June and July, corresponding to nymphal tick abundance (Lane 1990; Foley et al. 1998, 1999b, 2001; CDC 2000). In contrast, cases in horses and dogs seem to peak in the winter, corresponding to peak activity of I. pacificus adults (Madigan and Gribble 1987, Vredevoe et al. 1999, Foley et al. 2001), although dog cases also occur in late spring, at the time of the emergence of nymphs (Foley et al. 2001). phagocytophilum is not transovarially transmitted among ticks (Lewis 1979, Munderloh and Kurtti 1995).

## **Mammalian Hosts and Reservoirs**

Knowledge of natural reservoirs of A. phagocytophilum in California is limited. Immature I. pacificus commonly feed on lizards, birds, and small mammals, and occasionally on deer and carnivores (Furman and Loomis 1984). Adult I. pacificus occur frequently on dogs, coyotes, bears, bobcats, horses, deer, cattle, humans, and other large animals (Furman and Loomis 1984, Westrom et al. 1985, Lane 1994). The PCRprevalence of A. phagocytophilum in cervids ranges from 9-64% across Connecticut, Wisconsin, and Maryland (Bakken et al. 1996, Belongia et al. 1997, Massung et al. 1998, Magnarelli et al. 1999a). In California, the A. phagocytophilum PCR-prevalence was 5% in Columbia black-tailed deer (Odocoileus hemionus columbianus) and 31% in tule elk (Cervus elaphus nannodes) (Foley et al. 1998). DNA sequencing from the blood of deer and elk confirmed that California cervid isolates were homologous with horse isolates from northern California (Foley *et al.* 1998). Infected cervids have been observed from Sonoma, Marin, and Mendocino Counties. Because experimental studies have not been performed in deer, it is not known how they contribute to the maintenance of infection in nature, although their role as reservoirs would be limited to the number of nymphs they support.

Infections with A. phagocytophilum are common in several rodent species. Only a few incidents of infection with A. phagocytophilum have been reported in house mice (Mus musculus), chipmunks (Tamias spp.), and voles (Microtus pennsylvanicus), all in the eastern U.S. (Tyzzer 1938, Telford et al. 1996, Walls et al. 1997). The primary reservoir for A. phagocytophilum in the eastern U.S. is the white-footed mouse (*Peromyscus leucopus*), with a seroprevalence from 1-50% (Telford et al. 1996, Walls et al. 1997, Nicholson et al. 1998, Stafford et al. However, even though the PCR-prevalence ranged from 10.5-36%, infections in these rodents were transient (Walls et al. 1997, Nicholson et al. 1998). Reservoir competence was documented by experimental transmission of A. phagocytophilum by feeding larval I. scapularis on infected P. leucopus, allowing these larvae to molt, and transmitting the infection to naïve P. leucopus individuals via nymphs (Telford et al. 1996). There is not much evidence of anaplasmosis in other Peromyscus spp., with 11% seropositive in P. boylei, 6% in P. maniculatus, and 8% in P. gossypinus (Nicholson et al. 1998, 1999). PCR-positivity in P. maniculatus ranged from 0-21% (Nicholson et al. 1998, 1999; Zeidner et al. 2000).

Woodrats are likely reservoirs for A. phagocytophilum in California. Serologic evidence of exposure was found in Neotoma fuscipes (17.4-26.0%), N. lepida (25%), N. albigula (1%), and N. mexicana (4.7%) (Nicholson et al. 1998, 1999), and A. phagocytophilum DNA was recovered from seropositive woodrats (Nicholson et al. 1999). Titers in woodrats averaged 194 (geometric mean titer) compared with 410 in all seropositive *Peromyscus* (Nicholson 1998). Persistent infection (1-8 months) with A. phagocytophilum was documented in wild-caught N. fuscipes with no clinical signs (Nicholson et al. 1999, Foley et al. 2002). The agent in woodrat blood was detectable by PCR and was transmissible via needle inoculation to susceptible woodrats. Neotoma fuscipes is particularly common in oak woodlands, which is also optimal habitat for I. pacificus.

Infection with *A. phagocytophilum* has been described also in carnivores, including coyotes (Pusterla *et al.* 2000), mountain lions (Foley *et al.* 199*b*), black bears (Walls *et al.*, unpubl. data), dogs (Madewell and Gribble 1982, Greig *et al.* 1996, Foley *et al.* 2001), and raccoons, cats, skunks, and bobcats (J. Foley, unpubl. data). The seroprevalence in coyotes, 46%, is surprisingly high, and a PCR-positive coyote has been described (Pusterla *et al.* 2000). The titers in coyotes were relatively high as well, with two coyotes having titers ≥1,280. Seropositive coyotes were reported from Los Angeles, Santa Clara, and Tehama Counties, and a PCR-positive coyote was detected in Santa Clara County (Pusterla *et al.* 2000). Mountain lions had relatively high serologic (17%) and PCR-prevalence (16%) of *A. phagocytophilum*, although

with low titers (Foley *et al.* 1999*b*). If infections in mountain lions are clinically similar to infections in cats, the disease may be mild (Lewis *et al.* 1975). The prevalence of *A. phagocytophilum* in mountain lions varied from 19% in north coastal California to 25% in the Sierra Nevada and 0% from Monterey. Both coyotes and mountain lions have extensive home ranges or migration patterns and are likely to encounter hundreds of individual ticks from several independent foci, from which they might acquire *A. phagocytophilum*.

Dogs in California develop relatively mild clinical signs of anaplasmosis and have been used as sentinels of human risk, because of the frequency with which they are bitten by I. pacificus and their close association with people (Foley et al. 2001). The California statewide seroprevalence of A. phagocytophilum in dogs was 8.7% (Foley et al. 2001). Dogs in the eastern U.S. also have been reported to experience infection with A. phagocytophilum. Clinical, serological, and molecular descriptions were provided for 17 dogs with anaplasmosis from Minnesota and Wisconsin (Greig et al. 1996) and 3 dogs from North Carolina and Virginia (Goldman et al. 1998). A serosurvey of tick-borne disease in Oklahoma documented 33% prevalence of antibodies to A. phagocytophilum in dogs, but 0/100 in horses (Rodgers et al. 1989). In Connecticut and New York, 9.4% of ill dogs had antibodies to A. phagocytophilum (Magnarelli et al. 1997). In California, canine GA has been reported from Marin County (northern coast range) and the Sierra Nevada foothills (Madewell and Gribble 1982). A large serosurveillance study of 1,082 healthy dogs across California using Geographic Information Systems technology indicated a Bayes' adjusted prevalence (BAP) 0.01 to 0.47 (Humboldt County) (Foley et al. 2001). Other north coast counties with moderately high seroprevalence were Mendocino, Sonoma, Lake, and Marin (from 0.15-0.20). Counties in the Sierra Nevada foothills had lower seroprevalence, ranging from 0.09-0.13. Santa Cruz County had a BAP of 0.10.

In contrast to dogs, horses (particularly older animals) experience relatively severe clinical disease (Madigan and Gribble 1987). Equine granulocytic anaplasmosis is common in California, with early reports from horses in northern coastal ranges (Sonoma and Mendocino Counties) and the foothills of the Sierra Nevada (Gribble 1969, Madigan and Gribble 1987, Madigan et al. 1990). Case reports were described for 5 clinically affected horses from the northeastern U.S. (Madigan et al. 1996). In horses from Connecticut and New York with clinical signs of anaplasmosis, the seroprevalence was 29% (Magnarelli et al. 1999b), and in well horses from Minnesota and Wisconsin, the seroprevalence was 17.6% from tick-endemic vs. 3.8% from non tick-endemic areas (Bullock et al. 2000). Infection with human origin A. phagocytophilum protected horses from challenge with E. equi (Barlough et al. 1995). Equine cases in California are reported regularly from counties in the northern coast ranges and Sierra Nevada foothills, with seroprevalence in horses from coast range and Sierra foothills of 10% (Madigan et al. 1990). A questionnaire-based study evaluated the spatial distribution of equine anaplasmosis (Vredevoe et al. 1999). Of 62 practices in northern California, cases were diagnosed in the foothills of the Sierra Nevada and coast range mountains with spatial clustering based on nearest neighbor statistical analysis.

In summary, GA is a threat to humans, domestic animals, and wildlife in a range of habitats typical of *I. pacificus* ticks, although the spatial ecology is characterized by distinct patchiness. There appears to be a low disease incidence in humans but not other animals, suggesting key features of the ecology remain unidentified.

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