

| Movement type | Method | movement phases | parasite | host | outcome | possible transient phase processes | possible transient phase factors | reference |
|---------------|--------------------------------------|-----------------|--|--|--------------------------------|--|------------------------------------|---|
| dispersal | genetics (gene flow) | D - A | freshwater trematodes | long-finned eel (<i>A. dieffenbachii</i>) | overlapping genetic structures | none cited | none cited | Blasco-costa et al. 2012 |
| | genetics (gene flow) | D - A | marine trematode (<i>Gymnophallus choledochus</i>) | birds (e.g. ducks, gulls, wader) | overlapping genetic structures | none cited | none cited | Feis et al. 2015 |
| | genetics (assignment tests) | D - A | black-tailed prairie dog <i>Cynomys ludovicianus</i> | fleas <i>Oropsylla hirsuta</i> | unrelated genetic structures | none cited | none cited | Jones and Britten 2010 |
| | genetics (viral gene flow) | D - A | vampire bats <i>Desmodus rotundus</i> | rabies virus <i>Lyssavirus, Rabodyviridae</i> | overlapping genetic structures | none cited | landscape structure | Streiker et al. 2016 |
| local | mark-recapture | D - A | Swallow bugs (<i>Oeciacus vicarius</i>) | cliff swallows (<i>Petrochelidon pyrrhonota</i>) | increased prevalence | none cited | none cited | Brown and Brown 2004 |
| | GPS tracking | D - T - A | canine distemper virus (<i>Morbillivirus</i>) | serengethi lions (<i>Panthera leo</i>) | no effect by nomads | recovery | host grouping, movement behaviour | Craft et al. 2011 |
| | mark-recapture | D - A | nematodes (<i>Pharyngodon wandillahensis</i>) | pygmy blue-tongued lizard (<i>Tiliqua adelaidensis</i>) | increased transmission | infection forcing (environmental transmission) | habitat | Fenner et al. 2011 |
| migration | Citizen Science surveys | D - T - A | protozoa (<i>Ophryocystis elektroscirrha</i>) | monarch butterfly (<i>Danaeus plexippus</i>) | decreased prevalence | infection forcing | host grouping, habitat | Altizer et al. 2000, Bartel et al. 2011 |
| | remote tracking (radio telemetry) | D - T | chytrid fungus (<i>Batrachochytrium dendrobatidis</i>) | common toad (<i>Bufo bufo</i>) | decreased prevalence | recovery | habitat | Daversa et al. unpublished |
| | mark-recapture | D - A | avian influenza virus | mallard ducks (<i>Anas platyrhynchos</i>) | increased prevalence | none cited | none cited | van Dijk et al. 2014 |
| | Citizen Science surveys | D - T - A | bacteria (<i>Mycoplasma gallisepticum</i>) | house finches (<i>Carpodacus mexicanus</i>) | increased prevalence | none cited | none cited | Hosseini et al. 2011 |
| | mark-recapture | D - T - A | avian influenza virus | pink-footed geese (<i>Anser brachyrhynchus</i>) | decreased prevalence | recovery | host grouping (allopatry), habitat | Hoye et al. 2011 |
| | Field surveys | T | sea lice (<i>Lepeophtheirus salmonis, Caligus clemensi</i>) | pink salmon (<i>Oncorhynchus gorbuscha</i>) | decreased prevalence | no infections present | host grouping (allopatry) | Krkosek et al. 2007 |
| | remote tracking (GPS) | D - A | ticks (<i>Ixodes ricinus</i>) | red deer (<i>Cervus elaphus</i>) | decreased prevalence | recovery | habitat (elevation) | Mysterud et al. 2016 |
| | mark-recapture | D - A | trematodes | galaaxid fish (<i>Galaxia spp.</i>) | decreased prevalence | recovery | habitat (water salinity) | Poulin. et al. 2012 |