BASICS

The organism responsible is the fungus *Ophidiomyces ophiidiicola*, within the family Onygenacea. Snake Fungal Disease (SFD) was first definitively identified in a population of Timber Rattlesnakes residing in New Hampshire in 2006.

**KNOWN AFFECTED** species include milk snakes, black rat snakes, garter snakes, timber rattlesnakes, eastern massasauga, cotton mouth snakes, and black racer snakes.

The characteristic **CLINICAL SIGN** of SFD is facial swelling. The disease can progress from the nasal cavity internally via the eyes, throat, and lungs causing eye infections and pneumonia. The fungus additionally **SPREADS EXTERNALLY** along the neck, body, and tail forming scattered nodules (lumps) or ulcerations.

The fungus can be shed into the environment by infected animals and **SPREAD** from the environment to other snakes, particularly in animals that share dens. There is no definitive evidence of snake to snake transmission. Spread of the fungus to new locations may occur when people track contaminated **SOIL** imbedded in clothing or shoes.

SFD is **DIAGNOSED** by identification of the typical skin lesions as well as laboratory identification of the **FUNGUS** by culture or DNA detection and microscopic examination of tissues.

**TREATMENT** with antifungal medications has not been successful in colubrid snakes.
Several studies have indicated that **TEMPERATURE** is a significant factor affecting the growth of *O. ophiodiicola*. This suggests that populations hibernating in the lower thermal range of 0 °C - 10 °C should have a reduced infection during the spring and summer than snakes that **HBERNATE** in the upper thermal range of 0 °C - 10 °C. In addition, data suggest that with increasing global temperatures, snake populations will be more vulnerable to *O. ophiodiicola*.

**DIAGNOSIS** Methods to identify the fungus include histopathological examination via skin biopsy, fungal culture and real-time or quantitative polymerase chain reaction (rtPCR and qPCR).

Due to its close relationship to *Chrysosporium anamorph Nannizziopsis vriessii* (CANV) complex, *O. ophiodiicola* infections may have been **MISDIAGNOSED**.

**DETAILS**

*Ophidiomyces ophiodiicola* is characterized as an **ENVIRONMENTAL SAPROBE**, meaning that it normally feeds on decaying organic matter in the environment. This is evident due to its highly tolerant nature; it can thrive in a wide pH range (5 - 11), it is drought tolerant, and can utilize a number of complex carbon, nitrogen and sulfur compounds. Evidence that this organism is a saprobe makes it likely that infection of snakes is opportunistic.

*Ophidiomyces ophiodiicola* is closely related to **CHRYSPORIUM ANAMORPH NANNIZZIOPSIS VRIESSII (CANV)** complex. Many early reports of snake fungal disease identify CANV as the causative organism but the fungus has since been reclassified.

Both **COLUBRIDS AND PIT VIPERS** in Eastern and Midwestern United States have been identified with SFD. Recent advances in molecular diagnostics have allowed identification of cases dating back as far as 1986.

**CLINICAL SIGNS** Incubation period is between 30 to 37 days with some showing clinical signs as early as day 12 of inoculation. In rare cases where there are wounds secondary to the infection, the fungus can penetrate the body and cause a **SYSTEMIC FUNGAL INFECTION** resulting in nodules on the coelomic fat pad, kidneys, liver and air sac. Experimental data shows snakes surviving an average of 90 days with SFD and having a 40% **MORTALITY RATE**.

**TRANSMISSION** Since *O. ophiodiicola* is an environmental saprobe, it is likely that the fungus resides in the soil. There have been a number of cases of captive snake populations becoming infected as well.

Snake fungal disease nodule.
Photo by Brad M. Glorioso, USGS