# Genome-Wide Association & Whole Genome Sequencing of an Inherited Progressive Retinal Atrophy in the Bengal Cat

<u>AR Personett</u>, B Gandolfi, JW Pearce, WM Young, LJ Castaner, LA Lyons College of Veterinary Medicine, University of Missouri, Columbia, MO

# **1. Introduction**

- \* Inherited retinal diseases of humans are a leading cause of blindness throughout the world. These include retinitis pigmentosa, Leber's congenital amaurosis and some forms of macular degeneration (1).
- Progressive retinal atrophy (PRA) is a term describing a group of inherited retinal diseases in animals, similar to those of humans (1).
- \* PRA is characterized by degeneration of the retinal photoreceptors of the eye, which eventually leads to blindness. It is an age-dependent disease with varying rates of progression (2).







- \* Animal models of human inherited retinal degenerations are useful for development of genetic testing methods and therapies to benefit both animals and humans (2).
- \* Four types of PRA have been found in cats—two in Abyssinian cats and one in Persians (1). The fourth is a novel, autosomal recessive PRA that has been observed in the Bengal cat breed.

## **2. Objectives**

- Characterize the phenotype of a novel feline PRA using state-of-the-art imaging methods
- Confirm the chromosomal location of the PRA with GWAS and haplotype analysis
- Determine the causal mutation of the PRA in Bengal cats



Figure 3: A trio of cats with the known PRA genotype from the Bengal colony was whole genome sequenced. These included 1 carrier cat (left) and 2 affected cats (center & right).

### 4. Results

- Complete ophthalmic examinations of affected kittens showed the earliest signs of retinal photoreceptor degeneration to be around 15 weeks of age. Degeneration was determined to be complete by 8 months of age.
- \* The GWAS suggested a 6 Mb region between base pairs 20,000,000 and 26,000,000 on chromosome A3 to be strongly associated with PRA (Figure 4).
- \* Whole genome sequencing revealed 569 polymorphisms in 90 genes located in the associated region provided by the GWAS.
- Of the SNPs that were Sanger Sequenced, only 1 was still concordant after population screening.



Figure 1: Optical Coherence Tomography scans. Cross-sectional images were taken of the area of the retina outlined in green (a). Retinal thickness is decreased in the 6 month old affected cat (c) compared to the non-affected cat (**b**).



Figure 2: Fundus exam images. a. Non-affected cat with normal retinal vessels. b. Affected cat at 4 months of age with early stage PRA. c. Affected cat at 6 months of age with mid-end stage PRA.

### 3. Methods

- Complete ophthalmic exams including biomicroscopy and indirect ophthalmoscopy were performed every 4 weeks on 4 affected Bengal kittens in an established breeding colony.
- Optical Coherence Tomography (OCT) scans were performed on all kittens every 4 weeks and within 1 week of complete ophthalmic examination (Figures 1 & 2).
- \* A genome-wide association study (GWAS) was previously performed using the Illumina

Figure 4: Manhattan plot summarizing the case–control GWAS for Bengal cats with PRA. A significant association with the disease was obtained with SNPs making up a 6 Mb region on chromosome A3 (chromosome 3 here).

# **5.** Conclusion

- Complete ophthalmic examinations confirmed the phenotype of a mid-onset form of PRA in the Bengal cat breed.
- OCT scans further characterized the phenotype and retinal thickness changes associated with PRA in affected cats. Retinal thickness data will be quantified at a later time.
- \* The concordant SNP is currently being evaluated with additional Sanger Sequencing.
- \* If the mutation is confirmed, a genetic test can be produced to eliminate the disease in the breed.
- \* A feline model could identify a new gene associated with retinal degeneration in humans and other species. Animal models can be used further to develop gene and drug therapies.

# 6. Acknowledgements

- Stipend was provided by Merial, a Sanofi Company
- Study was funded in part previously by the National Center for Research Resources R24

Infinium Feline 63K iSelect DNA array data from 45 affected cases and 53 unaffected controls.

#### \* A trio of cats with the known PRA genotype from the Bengal colony (Figure 3) was whole genome sequenced. Sequencing data was screened and polymorphisms in the associated region were evaluated.

\* Five missense mutations in a gene known to be involved with vision in humans and one splice site variant in a gene involved in energy production were chosen to be verified with Sanger Sequencing.

RR016094 and is currently supported by the Office of Research Infrastructure Programs OD R24OD010928, the Winn Feline Foundation, and the Cat Health Network

References: 1 Rah, HC, Maggs, DJ, Blankenship, TN, Narfstrom, K, & LA Lyons. (2005) Early-Onset, Autosomal Recessive Progressive Retinal Atrophy in Persian Cats. Investigative Ophthalmology & Visual Science 46(5):1742-1747.

2 Alhaddad, H, Gandolfi, B, Grahn, RA, Rah, HC, Peterson, CB, Maggs, DJ, Good, KL, Pedersen, NC, & LA Lyons. (2014) Genome-wide Association and Linkage Analyses Localize a Progressive Retinal Atrophy Locus in Persian Cats. Mammalian Genome 25:354-362.