

# Establishing and maintaining a novel domestic cat breed through genomics and pedigree analysis

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### **BACKGROUND INFORMATION**

- Breeding cats for a desired phenotype carries the risk of inadvertent inbreeding depression
- Closely monitoring pedigrees and genotypes of cats can guide breeding practices to:
  - Maintain and refine desired breed attributes
  - Minimize or prevent adverse health conditions
- Population Management or Species Survival Plans (PMPs/SSPs) are used to manage captive populations in zoos. Similar genetic analyses can be used to develop Breed Management Plans (BMPs)
- A cat colony in Qatar is being used to establish a new breed of cat, the Katari

# RESEARCH QUESTION

How can genetic data be used to understand and maintain a cat colony and establish a new breed of cat?

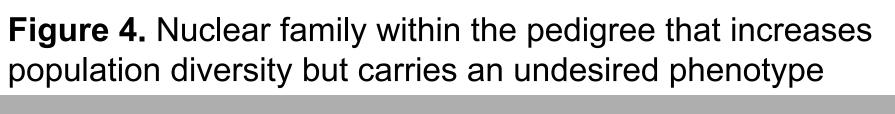
### **OBJECTIVES**

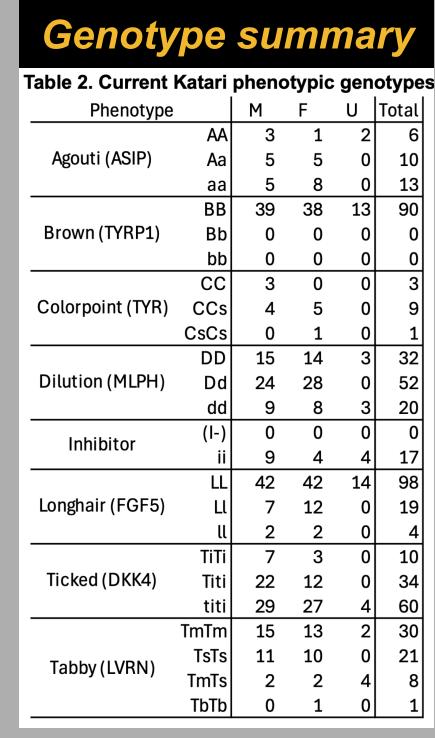
- Parentage verify the colony
- Determine known variants of phenotypic traits like coat color, pattern, length, etc.
- Determine genetic diversity of founders
- Determine presence of known disease variants
- Develop recommendations for further breeding pairs to:
- Refine the desired Katari phenotype
- Eliminate or minimize undesired allele variants
- Reduce inbreeding coefficients
- Recommend general husbandry and wellness protocols to maintain a healthy colony

### MATERIALS AND METHODS

- Constructed colony pedigree using Pedigree-Draw (PedDraw)
- Mitotyped founders and key cats:
- DNA isolation from buccal swabs
- Mitochondrial control region (mtDNA CR) PCR
- Sanger sequencing of mtDNA CR
- Determined mitotype of mtDNA CR DNA using Sequencher software
  Conctyped cate for disease and trait variants
- Genotyped cats for disease and trait variants
- Conducted parentage analyses using whole genome sequencing (WGS), microsatellite (STR), and single nucleotide polymorphism (SNP) data
- ☑ Genotyped cats for parentage & ancestry verification (~121-138 SNPs)

# 24143 24469 1137 24469 24402 225 226 248 243 244 24503 24499 24503 24499 Figure 4. Nuclear family within the pedigree that increases

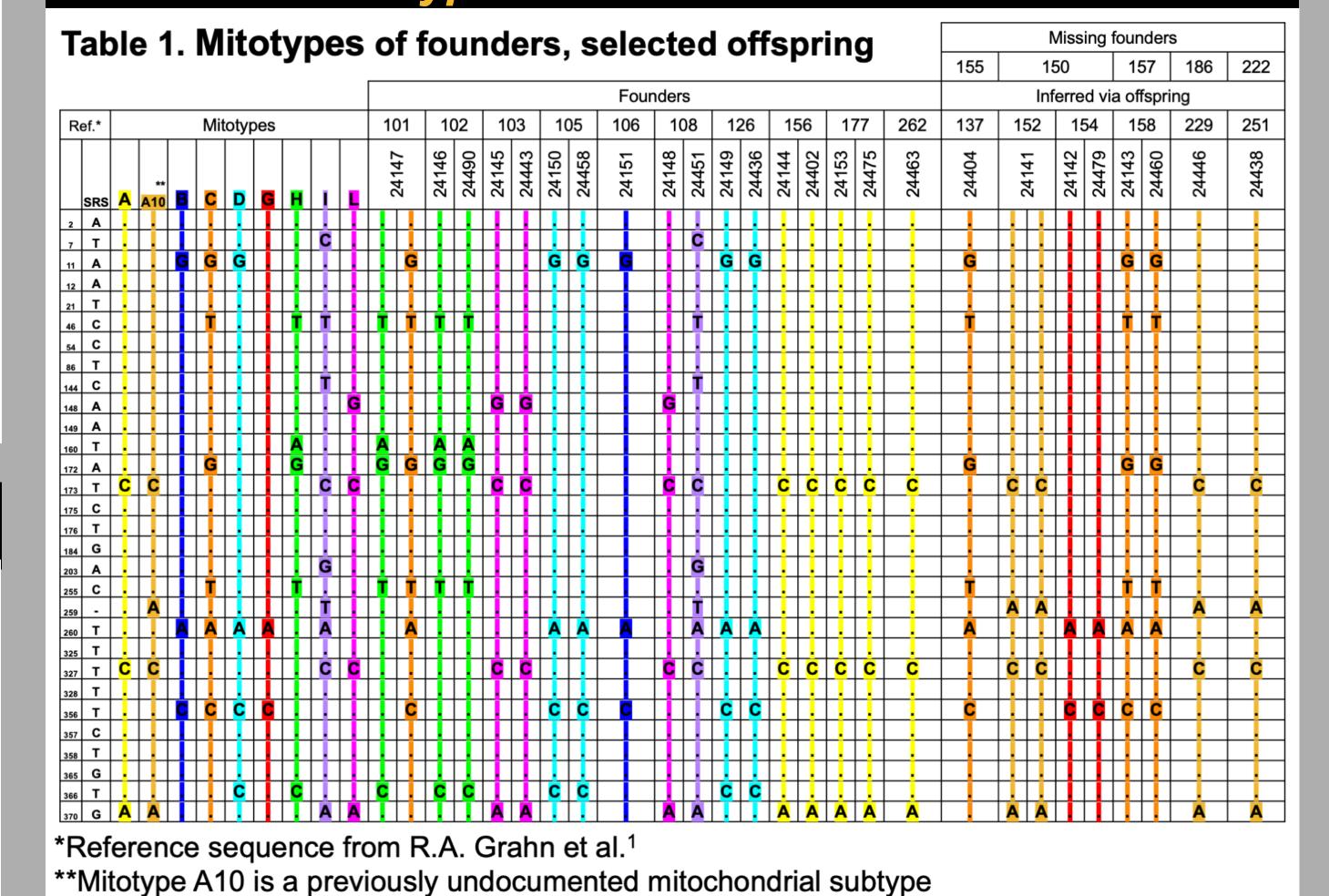




### **RESULTS**

- 171 cats currently included in pedigree (Fig. 1)
- ★ 16 founders (8 male, 8 female)
- 9 different mitotypes; 5 maternal origins (Table 1)
- ★ A novel mitochondrial subtype (A10) was discovered in three cats
- Representative phenotypes illustrated in Figure 2
- Current phenotypic genotypes summarized in Table 2
- Novel variants may explain unknown phenotypic variance (Fig. 3)
- All putative disease variants were likely benign polymorphisms
- The Katari breed should resemble cats in early stages of domestication, with colors conducive to desert survival
- Phenotypic traits of extant domesticated cats (long hair, orange color, etc.) should be minimized
- Maintaining genetic diversity may require retention of some undesirable traits until further generations (Fig. 4)
- Population modeling will help determine breeding pairs, colony size, and founder importation
- The pedigree currently has minimal inbreeding; the colony is only in its second and third generations

# Katari cat mitotypes



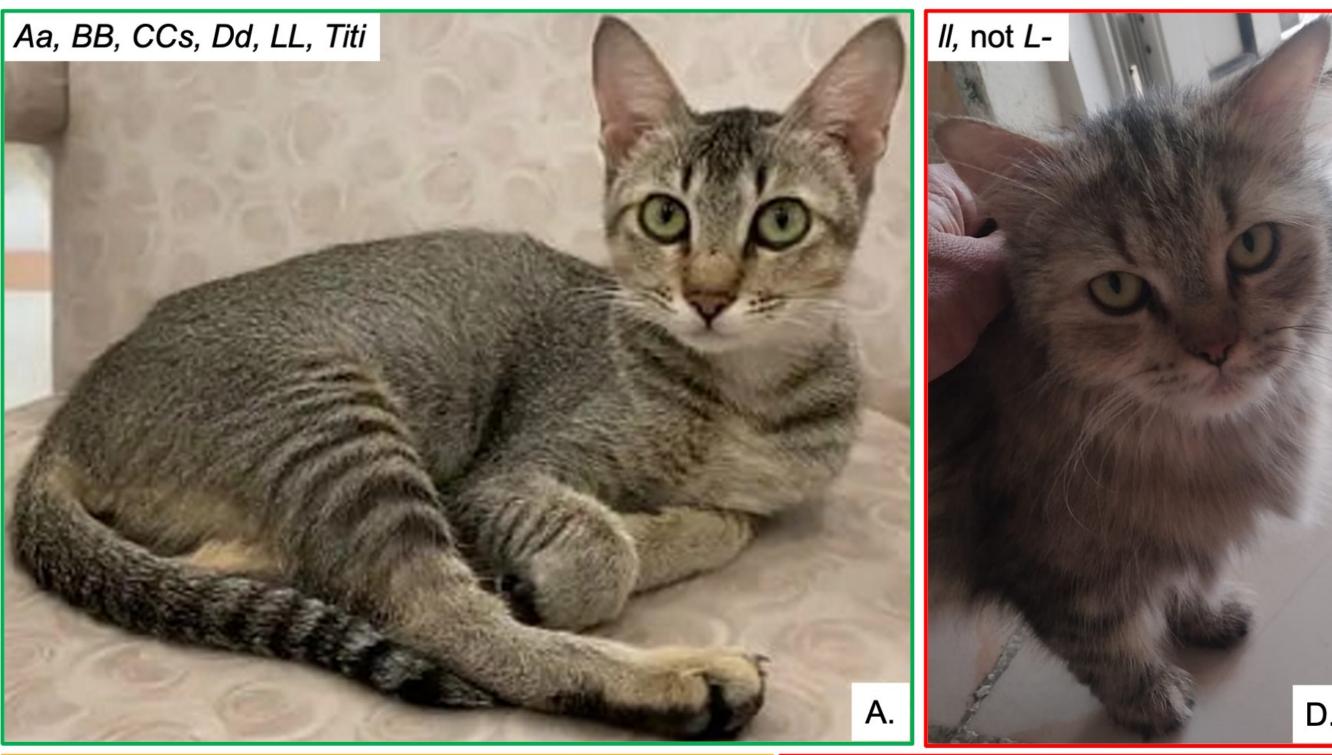
## Mystery Katari phenotypes





data. A) Cat 213; lighter brown coloration than DNA indicates. B) Cat 108; appears dilute, yet genotype indicates dense coloration. C) Cat 150; coloration implies SS (high white) genotype, yet no kittens descended from that cat has Ss (bicolor) fur. D) Cat 218; looks like it should be homozygous recessive for ASIP (aa), yet inherited at least one (A-) from a parent.

### Katari cat phenotype





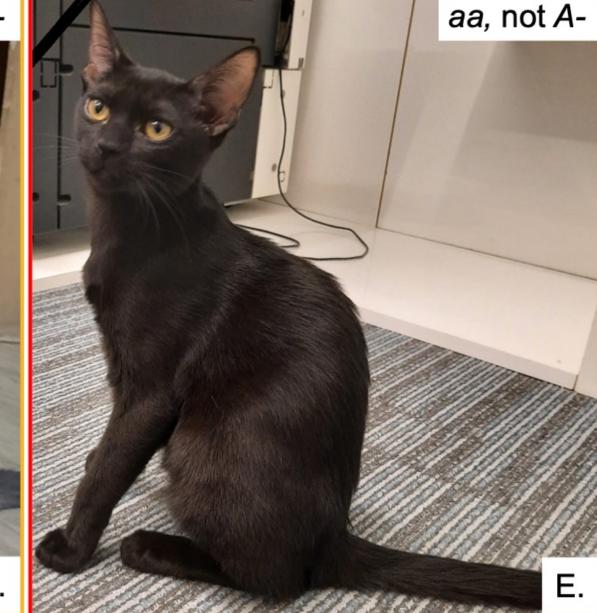




Figure 2. Current Katari
phenotype overview. A) ideal
Katari phenotype, cat 101.
Agouti, black, full color, dense,
shorthaired, ticked.\* B) alternate
Katari phenotype under
consideration, Cat 111. Blue, not
black. C-E) undesired Katari

phenotypes. C) Cat 126; mackerel tabby, not ticked. D) Cat 157; longhaired, not shorthaired. E) Cat 116; solid, not agouti.

\*Information on causative variants for coat phenotype gathered from OMIA<sup>2</sup>

### **FUTURE DIRECTIONS**

- Population management software will be used to recommend breeding practices that minimize inbreeding and maximize genetic diversity
- Communication and collaboration will continue with the Katari colony to:
  - Sample and sequence more cats to complete population genomic data
  - Improve healthcare and husbandry practices

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- Sanger & whole genome sequencing Genomics Technology Core, MU
- Genotyping Veterinary Genetics Laboratory, UC Davis

Katari colony pedigree

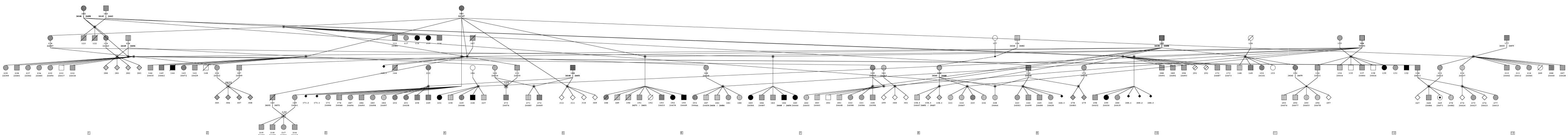


Figure 1. Katari cat colony pedigree as of 8/5/24. 171 cats are currently included on the extended pedigree. 18 founders are present. 162 kittens were born across 39 matings in 35 different mating pairs; approx. 4 kittens born per litter on average. 62 female kittens (~38.3%), 68 male kittens (~42%), and 32 kittens of unknown sex (~19.8%) were born; 7 were stillborn, indicating approx. 4.32% loss.