

# Hypertrophic Cardiomyopathy

## In brief

Hypertrophic cardiomyopathy is the most common cardiac disease in cats worldwide. It is characterized by the increase in wall thickness of the left ventricle and the intraventricular septum, eventually leading to congestive heart failure. In the Maine Coon, a point mutation (A31P) in the MYBPC3 gene has been found causative for the disease. The relative risk for developing HCM is around 1.8 in heterozygous cats and 18 in homozygous cats compared to Maine Coons negative for A31P. HCM has also been found in about 5.4% of cats negative for A31P, thus this mutation is not sole cause of HCM in Maine Coons.

## Clinical overview

Hypertrophic cardiomyopathy is a cardiac disease that is characterized by increased wall thickness and mass of the left ventricle and/or the intraventricular septum. This causes turbulence in blood flow and increased venous pressure in the left atrium and lungs. The clinical signs most commonly noted with the disease are respiratory signs associated with congestive heart failure such as tachypnea (increased breathing speed), exercise intolerance and panting, difficulty breathing, and coughing. Thromboembolism may also occur and affected cats have an increased risk of sudden cardiac death.

In Maine Coon cats and the majority of other cat breeds, clinical and echocardiographic signs appear after the cat has reached breeding age. The most common age of diagnosis is 5-7 years. In Ragdoll cats however, the disease has an early onset with an average age of diagnosis being only 15 months. Both in Maine Coons and Ragdolls, cats homozygous for the breed-specific causative mutation have earlier onset of the disease and increased severity of the disease than those with only one copy.

## Disease severity

Considerable


## Clinical signs

- respiratory signs associated with congestive heart failure
- thromboembolism
- increased left ventricle wall thickness

## Mode of Inheritance

Autosomal                      Dominant  
(Incomplete Penetrance)

## Results of the genetic test are reported as follows:

Clear  
 At risk

## Mutation(s) found in:

Maine Coon  
Maine Coon Polydactyl

## References

### Online Database

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### Scientific Articles

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