

# Genome-wide Association Study of Type A Pulmonic Stenosis in Bulldogs



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## Pulmonic Stenosis

- Pulmonic stenosis (PS) is the most common congenital heart defect in dogs.<sup>1</sup>
- Clinical signs include: exercise intolerance, collapse, and congestive heart failure.<sup>2</sup>
- Type A PS (fused pulmonic valve leaflets) is the most frequent form observed (Fig 1).<sup>2</sup>
- Bulldogs are overrepresented for PS suggesting a genetic cause.<sup>2</sup>
- Hypothesis:** PS in Bulldogs is associated with genetic variants that disrupt cardiac development.

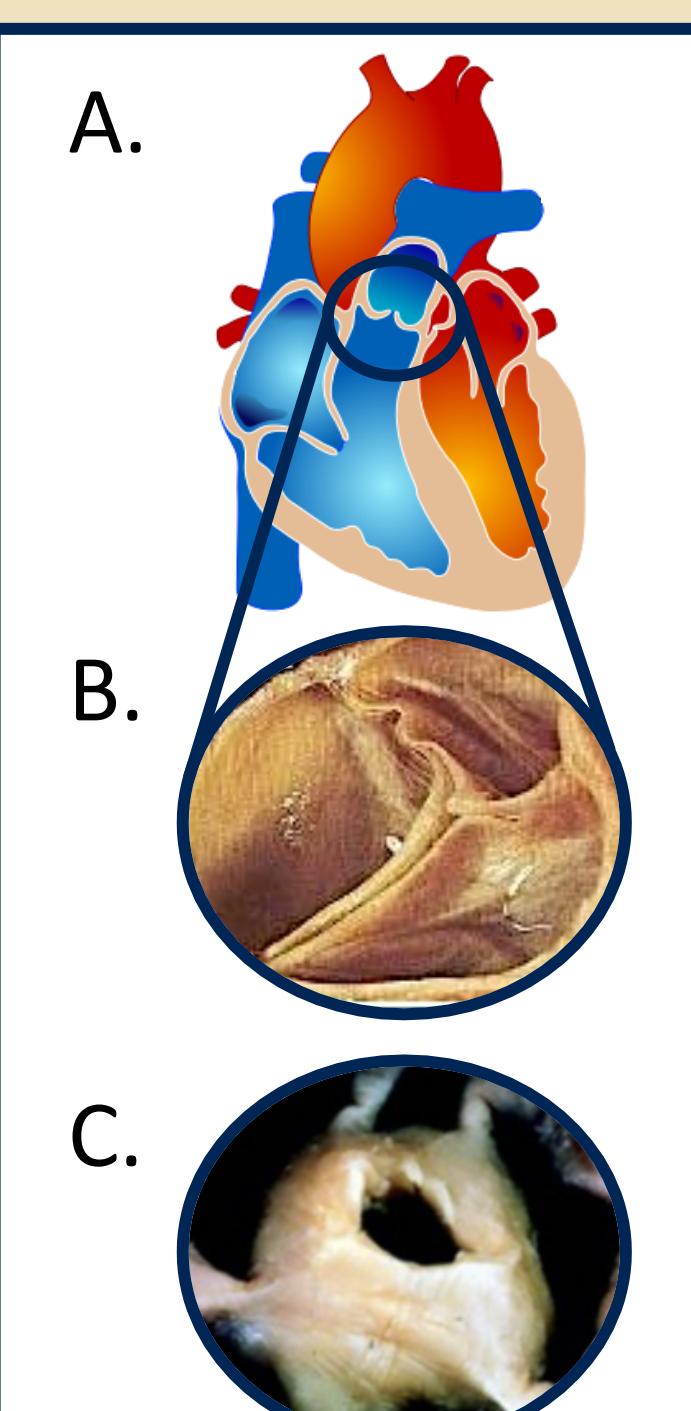


Figure 1. A. Location of pulmonary valve  
B. Normal valve anatomy  
C. Type A valve anatomy with dome shape and commissural fusion

## Result: Pulmonic Stenosis Incidence

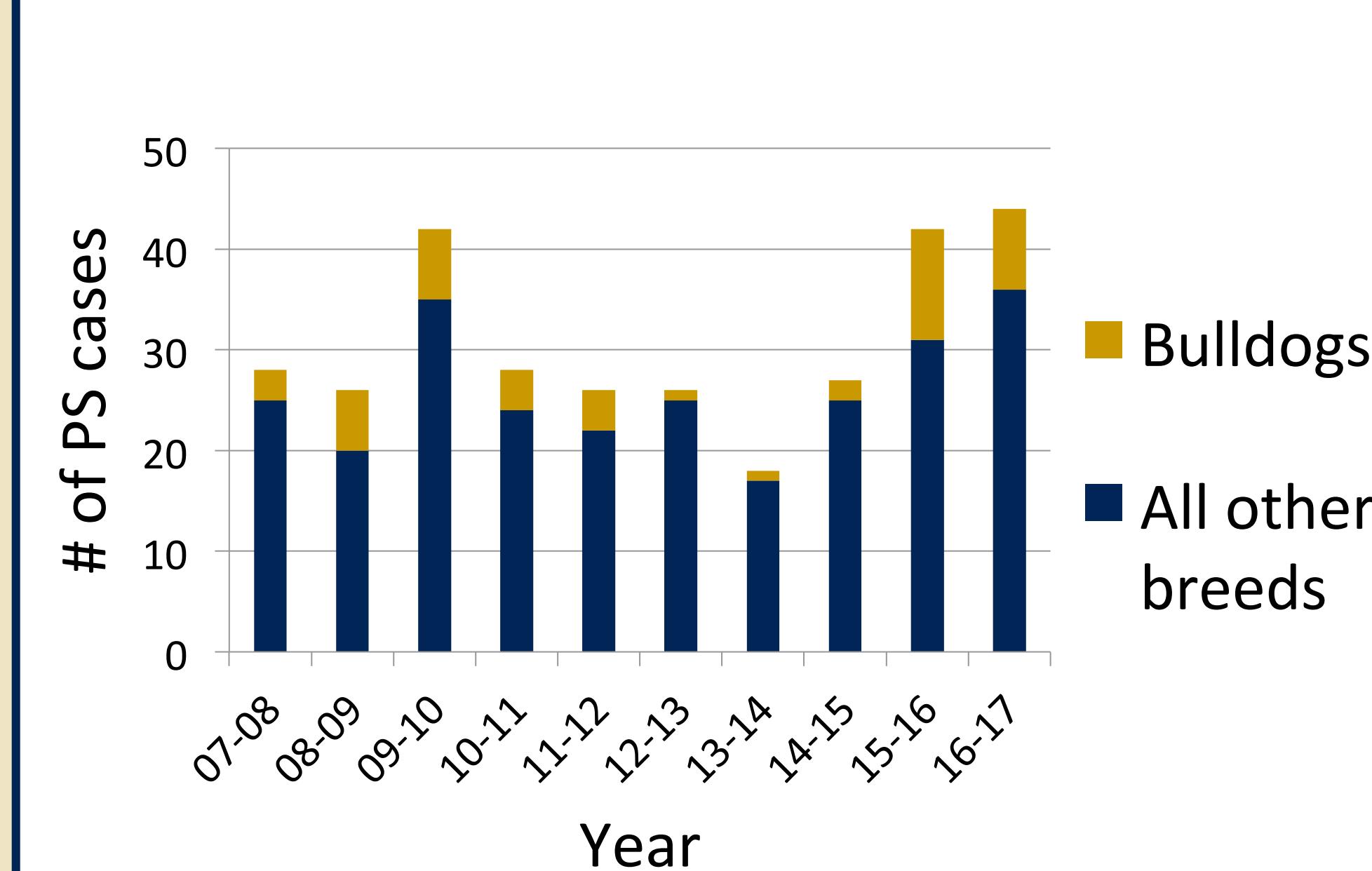


Figure 2. Incidence of PS each year for 10 years at the UC Davis Veterinary Medical Teaching Hospital (VMTH).

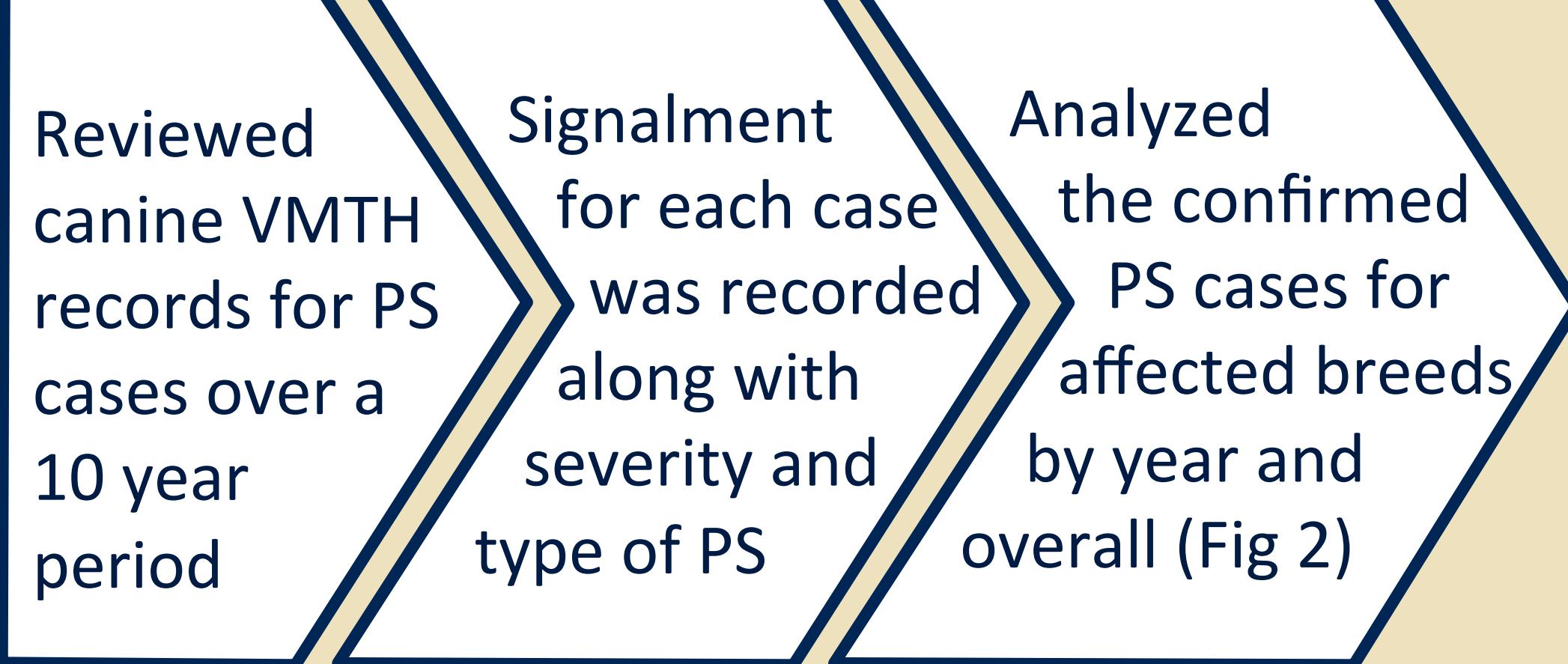
The top five breeds affected with PS as a percent of total PS cases were:

- Bulldogs – 15%
- Pitbull terrier – 11%
- French Bulldog – 6%
- Chihuahua – 4%
- German Shepherd – 3%

4% of all Bulldogs that were seen at VMTH between 2007-2017 were diagnosed with PS.



## Methods: Pulmonic Stenosis Incidence



## Result: GWAS

- 121,832 markers remained after quality control and filtering (call rate >0.90; MAF <0.05).
- 12 cases and 45 controls were included in the analysis after PCA outliers were removed.
- There was mild population stratification present with an inflation factor ( $\lambda$ ) of 1.18.
- A Bonferroni significant association was present on chromosome 7 after both EMMA (p<sub>adjusted</sub> = 9.89x10<sup>-3</sup>) and 100,000 permutation testing (p<sub>adjusted</sub> = 6.1x10<sup>-3</sup>) (Fig 3).
- The extent of population stratification after mixed linear model analysis (EMMA) correction (Fig 4).
- There is one gene located over the three significant SNPs, PTPN2.

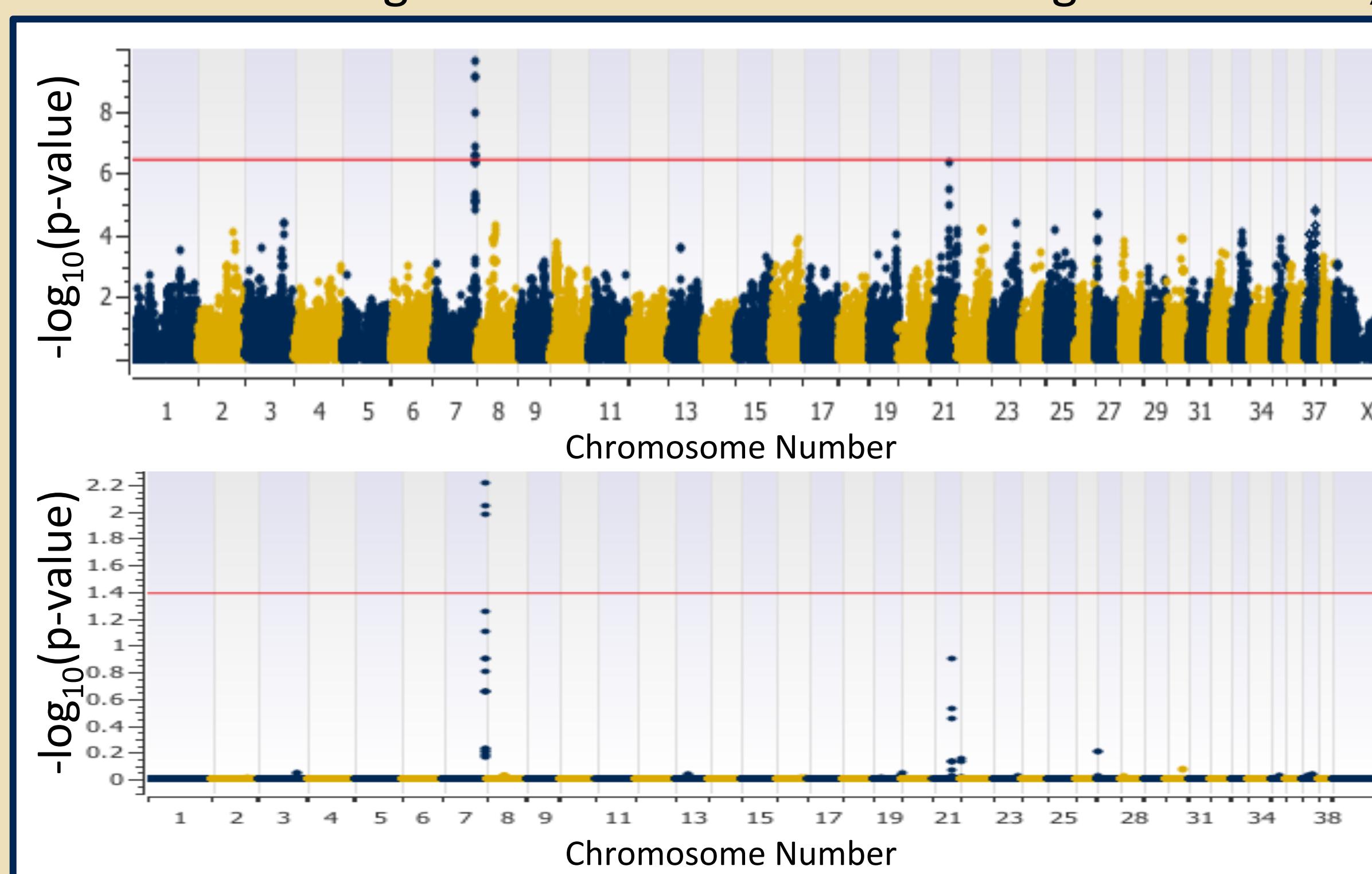


Figure 3. A. EMMA Manhattan Plot of Type A PS B. 100,000 permutation testing. SNPs are color-coded by chromosome. Red line indicates Bonferroni genome-wide significance.

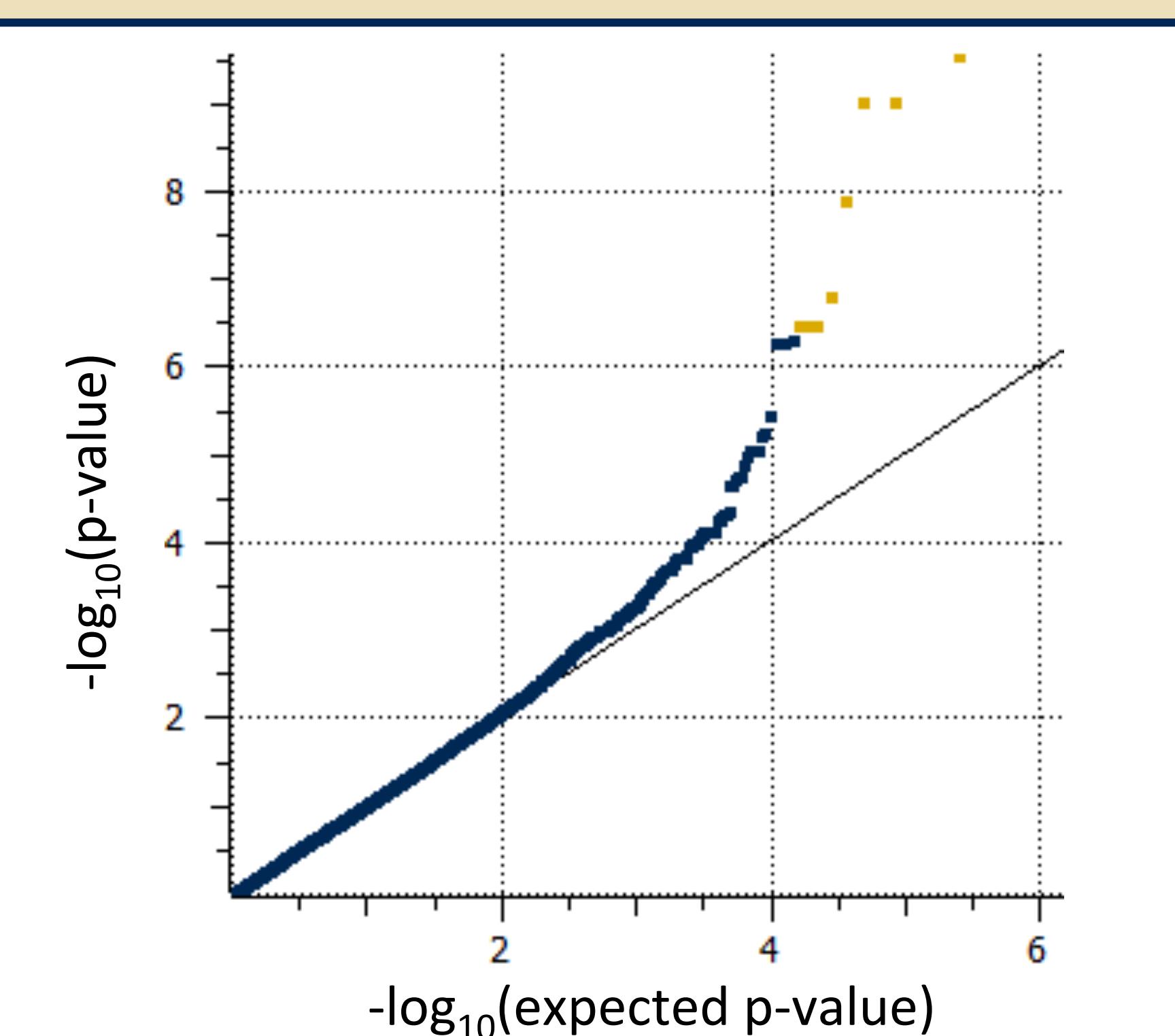


Figure 4. EMMA QQ plot demonstrating the remaining population stratification. Gold dots are Bonferroni significant SNPs.

## Methods: GWAS

Phenotyped and collected DNA on 13 cases and 62 controls

Genotyped samples on Illumina 230K Canine SNP Array

Golden Helix Software:

- Quality control and filtering of SNPs
- Removal of sample outliers on a PCA plot
- Additive Case-Control X<sup>2</sup> association test
- EMMA to correct population stratification
- Bonferroni correction for multiple testing
- 100,000 permutations testing

Identification of candidate genes

## Discussion

Pulmonic stenosis is a significant disease in Bulldogs. At the UC Davis Veterinary Medical Teaching Hospital 4% of all Bulldog patients were diagnosed with PS. They were the most affected breed accounting for 15% of all PS cases. A genetic investigation into the cause of PS in Bulldogs revealed a statistically significant region on chromosome 7 (CFA 7) and a suggestive region on CFA 21. On CFA 7 a likely candidate gene is PTPN2. PTPN2 is in the same family as PTPN11, which causes Noonan syndrome in humans.<sup>3</sup> It has been hypothesized previously that Bulldogs may be afflicted with a canine form of Noonan syndrome, but the genetic evidence has yet to be found.<sup>4</sup> Further sequencing in the region is necessary to identify a possible causal variant.

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