

MHC associations of myositis in Hungarian Vizsla dog

Lorna Kennedy



Centre for Integrated Genomic Medical Research, Manchester, UK

Why dogs?!?!

1 in 10 people in the UK own a dog (that's about 6 million dogs!!) How many worldwide?

Large investment in working dogs

- Guide dogs, hearing dogs
- Farm dogs, drug sniffers
- Locating earthquake victims
- Detecting cancer, epilepsy
- heart attacks
 - Health care, insurance
 - -Pet food





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Advantages of the dog as a model

Medical knowledge about dogs: second only to humans! Dog genome sequenced in 2005 Many genes very similar to human genes

Causal gene mutations for many monogenic canine diseases have been identified. can be traced by synteny to human genome



Dogs as a model for complex disease Many different breeds (>400) subject to intense selection for particular traits e.g. short legs, some bottlenecks: eg 2nd world war Each breed is genetically distinct Within breed very homogeneous <u>Complex diseases: may be subtypes</u>, but within breed, disease likely to be same 100 cases/100 controls for GWAS, not 1000s study same diseas/different genetic backgrounds



Canine myositis in Hungarian Vizslas

- A breed specific polymyositis is frequently observed in the Hungarian Vizsla.
- Affected dogs present with difficulty eating and drinking, regurgitation, and sialorrhea.
- Possible masticatory muscle atrophy and exercise intolerance
- Clinical response to immunosuppressive therapies points to an immune-mediated aetiology.
- Clinical and histological similarities with the immune-mediated myopathies observed in humans.
- MHC class II associations reported in the human conditions.













Affected dogs with megaesophagus – in "Bailey chair" to aid swallowing







Affected dog with sialorrhea: drooling!







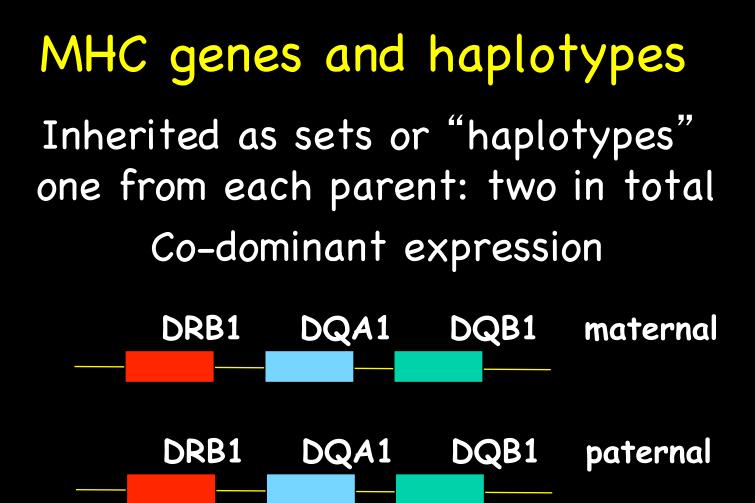






Canine Major Histocompatibility Complex





N.B. No DP expressed in dogs



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Canine MHC – DLA: very variable Alleles and haplotypes in the domestic dog: 15,500 dogs from <200 breeds

GeneNo of allelesDRB1302DQA139DQB11553 locus haplotypes>300

Variable breed distribution



Average per breed: 5-7 haplotypes 35% pedigree dogs are homozygous for MHC

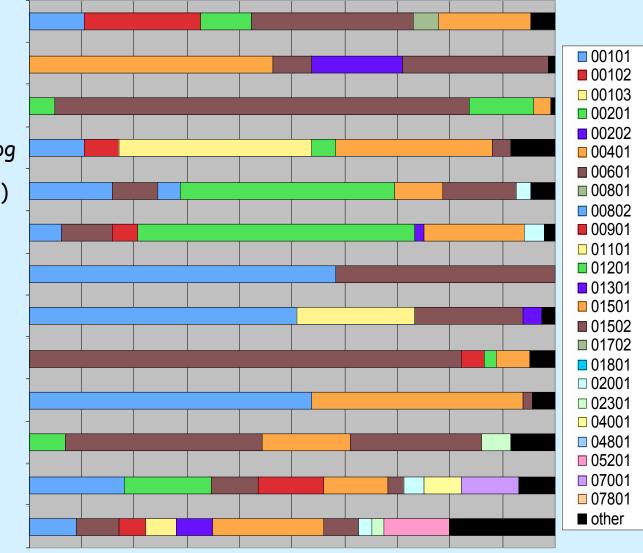


DLA-DRB1 alleles in some breeds

- n
- 115 Beagle
- 170 Boxer
- 321 Doberman
- 327 German shepherd dog
- 816 Labrador (Retriever)
- 381 Retriever (Golden)
- 43 Rottweiler
- 73 Setter (English)
- 151 Spaniel (Cocker)
- 228 Terrier (Westie)
- 291 Terrier (Yorkshire)
- 177 Husky

CIGMR

109 Mongrel (Brazilian)



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MHC diversity in breeds

Intrabreed

- some breeds have few alleles and haplotypes
- most breeds have 4-5 frequent haplotypes
- no breed has all known alleles/haplotypes

Interbreed

- some alleles and haplotypes are only found in a few breeds - "restricted"
- other alleles and haplotypes in many breeds



MHC associations with autoimmune diseases

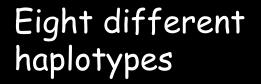
Identified for:

Diabetes, Hypothyroid disease, IMHA, SLO, SLE, Addison's Disease, Meningiocephalitis, Anal furunculosis, Chronic hepatitis, Pancreatitis

Different breeds can have different associations Same haplotype can be associated with several diseases

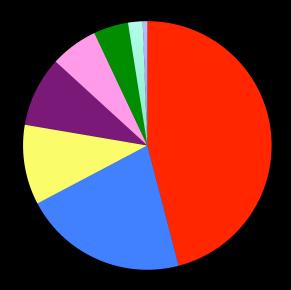


MHC haplotypes in Hungarian Vizslas



One very frequent (46%) One frequent (21%) Two less frequent (10%) Four others (<6%)

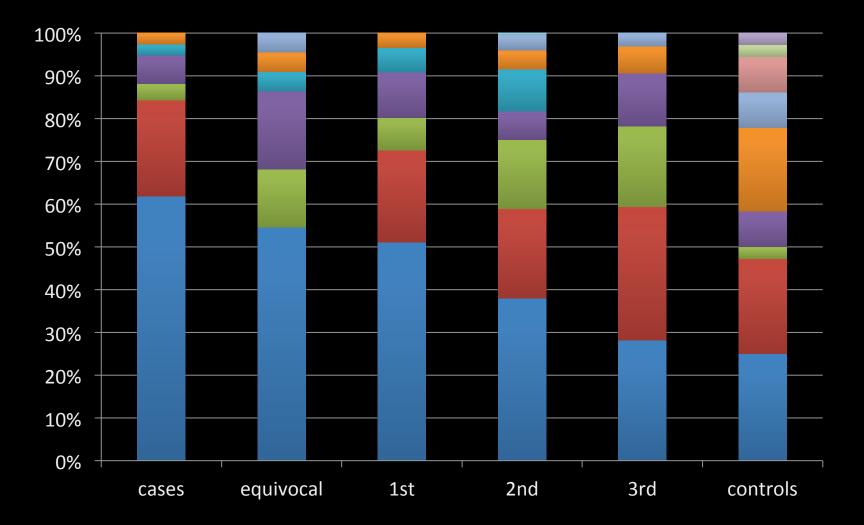






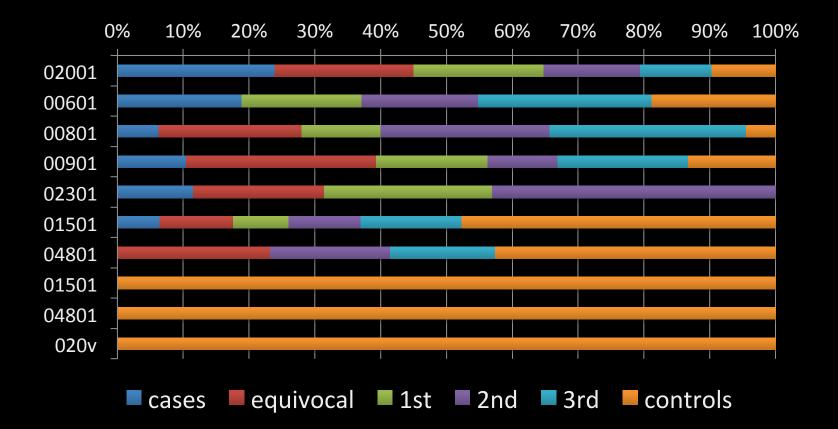


Haplotype frequencies by group





Distribution of groups by haplotype





DLA haplotypes found in Hungarian Vizsla n=341

| DLA | DRB1 | DQA1 | DQB1 | % | Disease risk |
|-----|-------|--------|--------|------|-------------------------------------|
| | | | | | Canine myositis in Vizslas |
| 1 | 02001 | 00401 | 01303 | 45.9 | Also with immune diseases in Saluki |
| | | | | | Protective SLO in Gordon setters |
| 2 | 00601 | 005011 | 00701 | 21.3 | (IMHA; in other breeds) |
| 3 | 00801 | 00301 | 00401 | 10.4 | |
| 4 | 00901 | 00101 | 008011 | 9.1 | (Diabetes, RA; in other breeds) |
| 5 | 02301 | 00301 | 00501 | 6.2 | |
| 6 | 01501 | 00601 | 02301 | 4.5 | (Addisons; in NSDTR) |
| 7 | 04801 | 00101 | 008011 | 1.9 | |
| 8 | 01501 | nt | 059v | 0.6 | Only found in vizslas from USA |

SLO = Symmetrical Lupoid Onychodystrophy

An aside, for breeders!!! How should we use MHC data?

Some evidence that homozygous individuals are slightly less able to respond to infection

"heterozygous advantage"

- Some evidence that (human) couples who are both homozygous for the same MHC haplotype are at increased risk of spontaneous abortion......
- So maybe we should avoid mating identically MHC homozygous dogs
- MUCH MORE important to not mate affected dogs, or those with affected 1st degree relatives



Take home messages

Strong MHC association with myositis in the Vizsla

Future work: GWAS

For breeders:



- DLA results should not be used as the only tool for mate selection
- A genetic test will need

many more dogs of known disease status to be tested to identify relative contribution of different genes



13th May 2014

With thanks to Bill Ollier, Jonathan Massey, Simon Rothwell, Dimitra Zante and



Clare Rusbridge (Veterinarian) Di Addicott for sample collection

Funding: UK Hungarian Vizsla Breed Clubs

Pictures: Di Addicott





13th May 2014

Canine myositis in Hungarian Vizslas

- Masticatory muscle myositis (MMM)
- Mean age of onset 2.3 years (range 0.2-8.8 years)
- Slightly more male dogs
- The most consistent clinical signs are dysphagia and regurgitation (due to tongue, pharyngeal and oesophageal dysfunction; and masticatory muscle atrophy)
- A marked elevation in muscle enzymes was an indicator of disease, but Vizsla Polymyositis could not be ruled out if muscle enzymes were normal

