

# Fokken van gezonde rashonden

<https://www.wur.nl/nl/project/fokken-van-gezonde-rashonden.htm>

Wetenschapswinkelproject, Tweede bijeenkomst begeleidingscommissie

12/10/2023 Jack Windig, Talita Kuijpers, Rita Hoving



# Agenda tweede bijeenkomst begeleidingscie “Fokken van gezonde rashonden”

- Welkom - Lèneke Pfeiffer, Wetenschapswinkel
- N.a.v. verslag 12/4
- Resultaten tot nu toe – Jack Windig, Talita Kuipers
  - ACT groep
  - Wetterhoun
  - Bouvier
  - Verdere plannen
- Reacties
- Afronding project: hoe opgedane kennis breder delen?

# Onderzoeksvragen

- Hoe kan fokkerij voor gezonde honden het beste vorm worden gegeven?
- Specifiek per ras
  - Welke gezondheidsproblemen met welke frequentie?
  - Huidige populatiegrootte / structuur en inteelt(toename)?
  - Inteelttoename in toekomst bij verschillend beleid

# Onderzoeksvragen

- Hoe kan fokkerij voor gezonde honden het beste vorm worden gegeven?
- Algemeen rashondfokkerij
  - Welke rol kunnen DNA typeringen spelen?
  - Wanneer en hoe kan fokwaardeschatting worden opgezet, en hoeveel zal dit bijdragen tot een betere gezondheid?
  - Wanneer en hoe kan outcross worden opgezet, en hoeveel zal dit bijdragen tot een betere gezondheid?

# ACT Group



**Breeding  
Better  
Buddies**

ACT GROUP 3.079  
SWEN, TALITA, ELLEN, CATO, MAX, KAREL, MIA

- 7 Students 8 weeks
  - Literature review
  - Interviews
- Report
  - Genetics
  - Perspective owners and breeders
  - Ethics
  - Recommendations

# Genetics

## DNA USAGE

- Use of DNA / SNP chips
  - DNA tests for diseases
    - How breed specific?
    - What to do with carriers?
  - Determine inbreeding and relatedness
    - Gain relative to pedigree?
  - High cost
  - Ownership data?

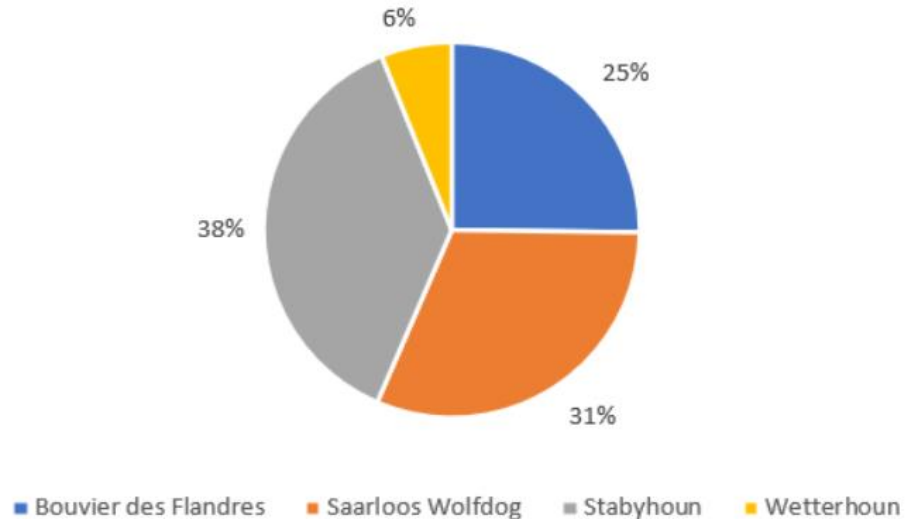
# Genetics

- Outcross
  - Trade off – genetic diversity / breed purity
- Use look a likes
  - SNP chip to identify breed purity and relatedness
- Estimate breeding values
  - Polygenic traits and multiple traits
- Genetic management
  - Mean kinships and Optimal contributions

# Perspective of owners and breeders

- Interviews via facebook
- Massive respons
  - Bouvier 62
  - Saarloos 77
  - Staby 92
  - Wetterhoun 15

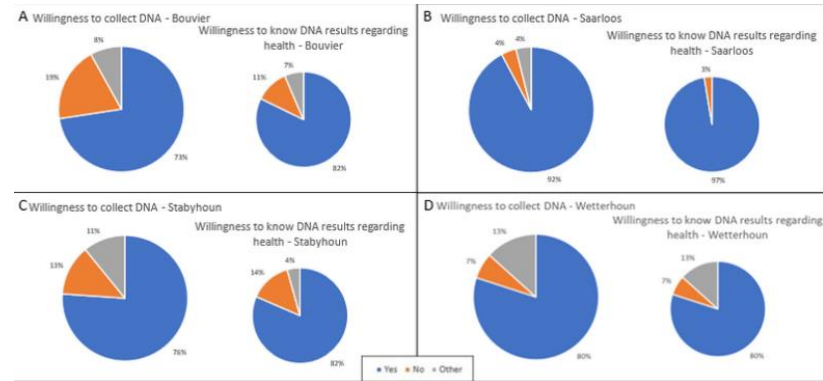
Distribution of the 246 responses





# Main findings 1

- Few pups used for further breeding
  - Selection on appearance
  - Difficulty finding breeders
  - >90% of pups get a pedigree
- Willingness to collect DNA
  - 73 – 92% positive
  - 80% - 97% Want to know results of health tests



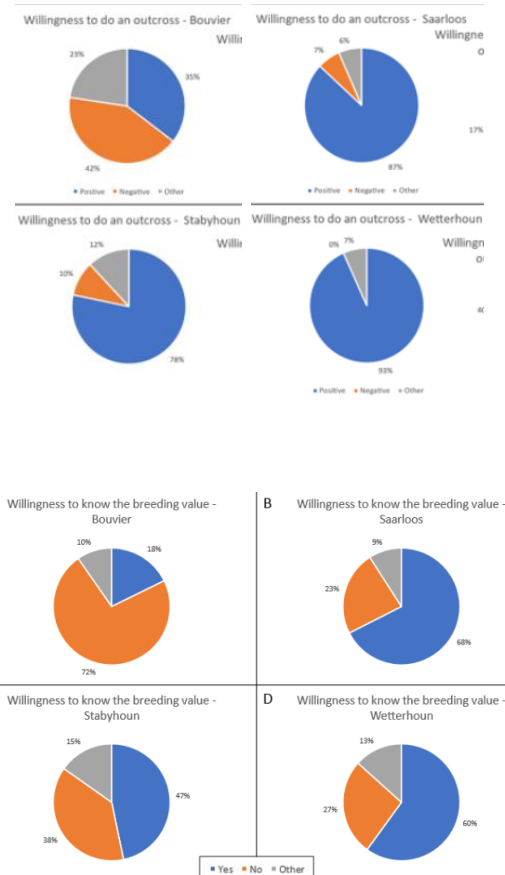
# Main findings 2

## ■ Outcross

- Divided opinions
- Bouvier more negative
- Saarloos and Wetterhoun most positive

## ■ Breeding values often mistrusted

- “Too materialistic, capitalistic, scientific”
- “Should accept dogs as they are”
- Confused with economic value



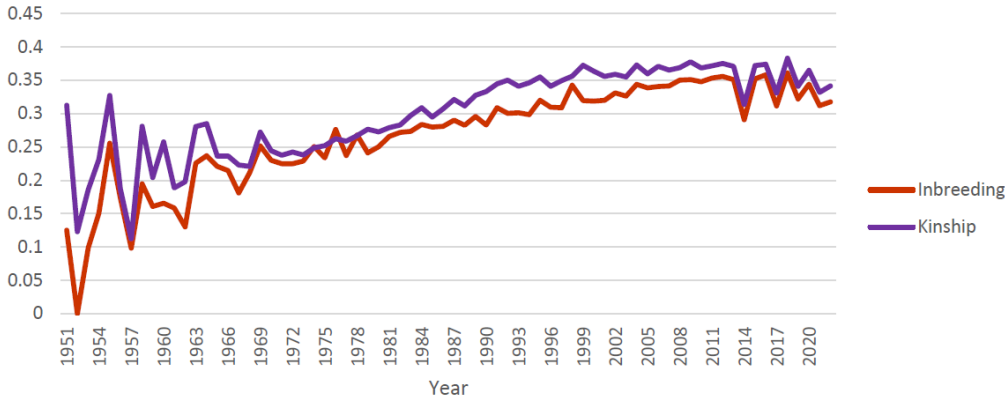
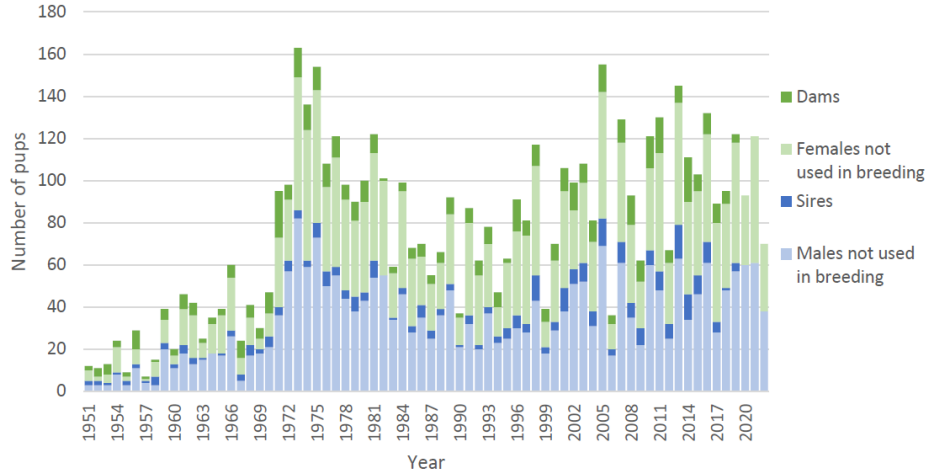


## Recommendations

- Keep updating the databases regularly
  - Look into the gEBV
  - Use Sweden & UK for reference
- Consider look-alike population as a reserve
- Promote communication between breeders
- Organise breeders' info markets
- Weighted disease index

# Wetterhoun: Population structure and inbreeding

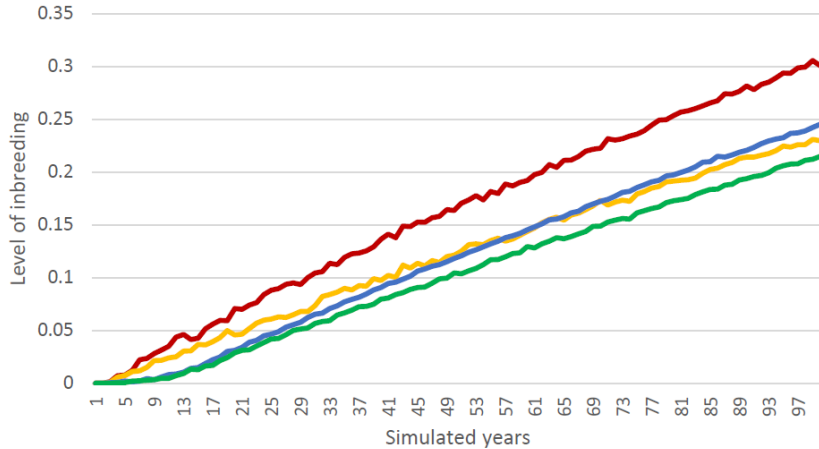
BSc thesis  
Aisha Boering



- Population structure
  - Rather small breed
  - Currently more males used for breeding
- Inbreeding
  - Has been way too high
  - Decreased in final decade

Period	Inbreeding rate	Kinship rate
<b>1951-1970</b>	3.66	0.12
<b>1970-1985</b>	2.12	2.62
<b>1985-1995</b>	2.00	3.68
<b>1995-2004</b>	1.48	1.16
<b>2004-2012</b>	1.32	0.57
<b>2012-2022</b>	-1.51	-1.41

# Wetterhoun: Genetic management



Method	Inbreeding rate
None	2.34
NVSW regulations	1.32
Minimise kinship parents	1.17
Use Mean Kinships	1.01
Kinship parents + MK breed	0.99

- Current policy reduces rate considerably
  - But not below 1%
- Mean kinship most effective
  - At 1%

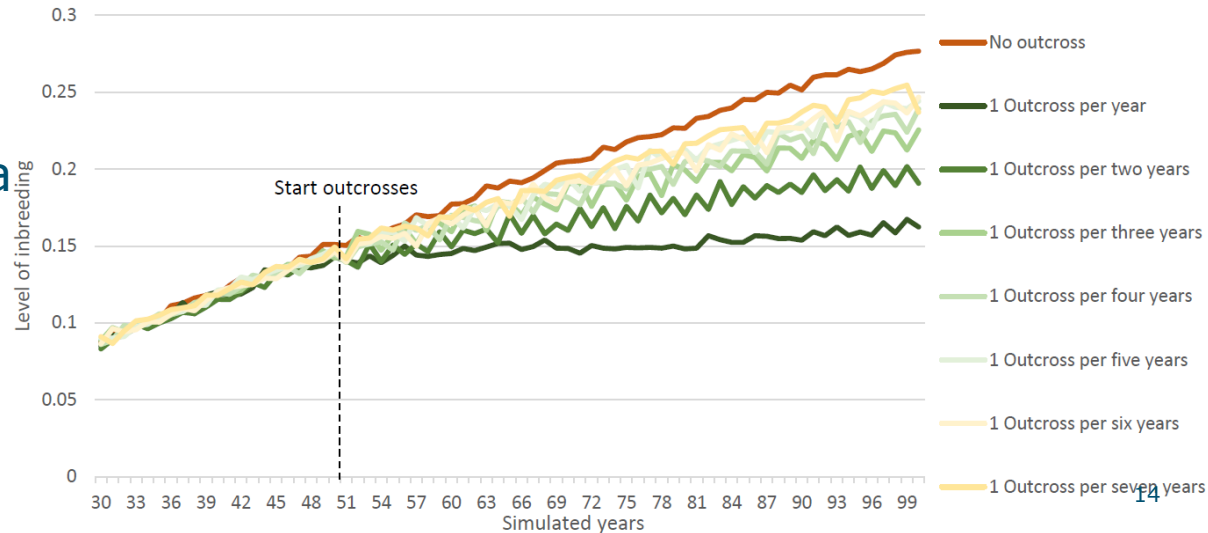
# Wetterhoun: Outcross



- Outcrosses
- Should be repeated for a lasting effect on inbreeding
- Useful for a reset

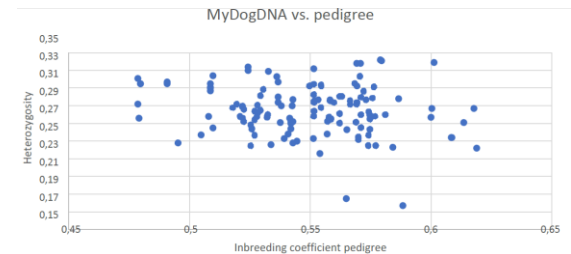
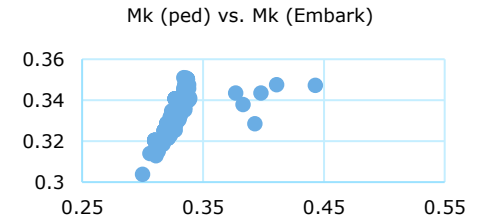
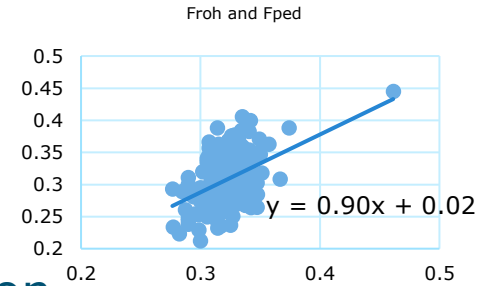
## ■ Increased litter size

- WH x WH: 5.9 pups
- WH x other: 7.8 pups
- WH x look alike: 4.0 pups



# Stabij + Saarloos (2019)

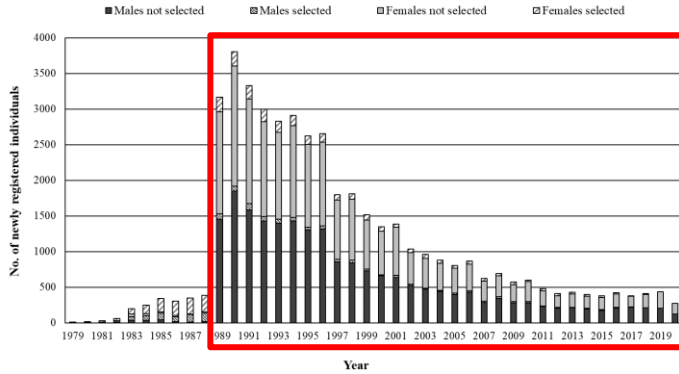
- Both: MK best method
- Stabij
  - DNA (Embark) vs. pedigree: high correlation
  - Depends on method
- Saarloos
  - DNA (My Dog) vs. pedigree: no match
  - Outcross
    - Backcrosses reduce effect
    - Repeats needed



# Summary Bouvier Pedigree analysis

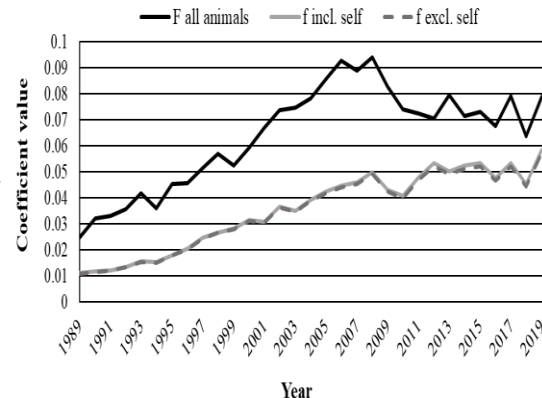
Research practice  
Talita Kuipers

Number of newly registrations per year (1979 – 2020)



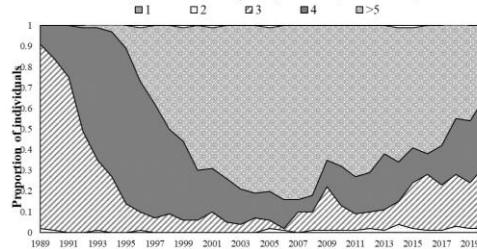
- Pups born per year decreased
  - >3000 before 1993
  - Around 400 since 2011

Inbreeding (1989 – 2020)



Period	deltaF	deltaf
1981-1990	0.93%	0.30%
1991-2000	1.29%	0.95%
2001-2010	0.77%	0.63%
2011-2020	0.45%	0.42%

Ancestral generations completely known (1989 – 2020)

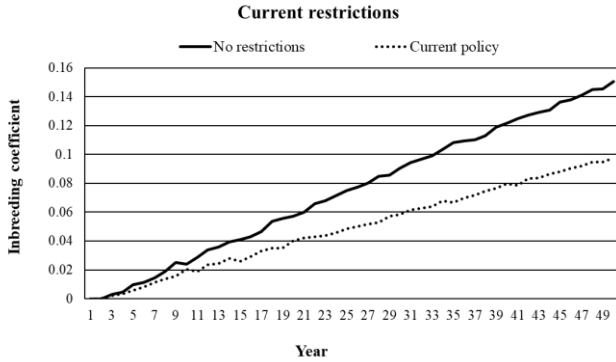


- Inbreeding rate has been too high
- And possibly still is on the high side

Pedigree not complete after 2001!



# Summary Bouvier simulations



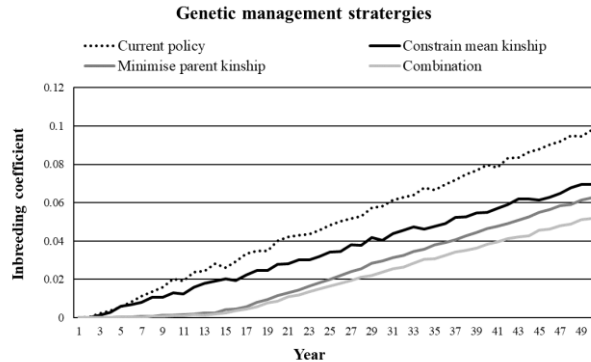
## Inbreeding rate:

1.34% No restrictions

0.83% Current restrictions

- Current restrictions are important
- Further reduction possible

- Recommend a combination of Mean kinship and minimising parent kinship



0.83% Current restrictions

0.56% Use Mean Kinship

0.58% Minimise Kinship parents

0.43% MK + Kinship parents

# Genetic (?) problems in the Bouvier

Disease	Age of onset	Prevalence	Possible mode of Inheritance
HD	Varying	now low	Polygenic
ED	Varying	now low	Polygenic
PPM	>6 weeks	17.1 %	Recessive, dominant or polygenic?
PHTVL/ PHPV	Undetermined	10.6 %	Autosomal incomplete dominant
Hereditary cataracts	Few weeks to months	Congenital: 9.5 % Non-congenital: 27.6 %	Autosomal recessive/dominant ?
RD	2 – 3 months	8.2 %	Autosomal recessive (X-linked?)
Distichiasis	0.3 – 8.9 years	11.6 %	Autosomal dominant ?
Corneal dystrophy	Varying	9.5 %	Sex linked recessive?
PRA	Early: 2 – 6 weeks Late: 2 – 5 years	0 %	Autosomal recessive
Entropion	4 – 7 months	4.8 %	Polygenic?
Microphthalmia	Undetermined	-	Autosomal recessive
Primary Glaucoma	± 7 years	“not free of ICAA”: 78.4%	Polygenic (?)

- More than 80% of the dogs have some sort of genetic disease
- Genetic background often unclear

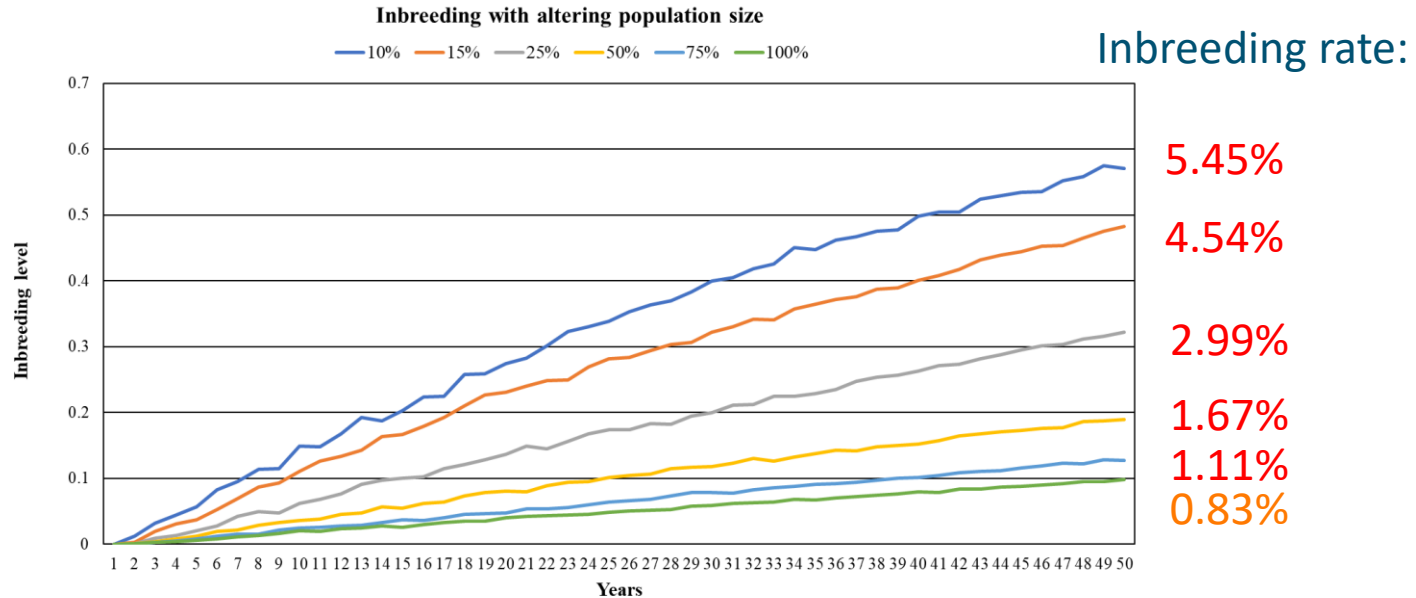
# Priority for reducing Genetic problems

Disease	Age of onset	Prevalence	Severity
Primary Glaucoma	± 7 years	“not free of ICAA”: 78.4%	Severe, but late
Hereditary cataracts	Few weeks to months	Congenital: 9.5 % Non-congenital: 27.6 %	Severe
PPM	>6 weeks	17.1 %	Potentially severe
Distichiasis	0.3 – 8.9 years	11.6 %	Moderately severe
PHTVL/ PHPV	Undetermined	10.6 %	Moderate?
Corneal dystrophy	Varying	9.5 %	Less severe
RD	2 – 3 months	8.2 %	Less severe
Entropion	4 – 7 months	4.8 %	Not severe
PRA	Early: 2 – 6 weeks Late: 2 – 5 years	0 %	Severe
Microphthalmia	Undetermined	-	-
ED	Varying	now low	Moderately severe
HD	Varying	now low	Moderately severe

- Table based on prevalence
- Severity gives similar priority
- Onset changes priority
- Discussion over first two

- Theoretical formula: Priority = severity \* healthy days lost
- Hard to determine severity

# Reduce breeding population?



There is **no scope** to reduce # of individuals in the breeding population!

# Recommended steps

## Short term

- Include Mean Kinship and Minimise parental kinship
- Prioritise diseases

## Long term

- Create disease inventory
- Unravel genetic background
- Reassess priority list
- Calculate EBVs
- Create health index
- Select (and Mate) based on EBVs, Inbreeding coefficients, and Health index

# How to select against multiple genetic diseases

- Animal breeders' approach: construct an (economic) index

- Define breeding goal (H)
- Determine economic weight of traits
  - $H = a_1X + a_2Y + a_3Z + \dots$
- Measure traits influencing performance
- Determine heritabilities and genetic correlations
- Estimate breeding value of individuals

- Weighing information of relatives, predictor traits etc.

- Example: cattle breeding – Total merit index, production index, udder health index, fertility index, etc.

CRV Zwartbont

InSire-stieren	naam	ID	afstamming	A2A2	CRV Genomisch	CRV Fokwaarde	GEWENDE					EINDWEE											
							Levensduur	Productie	Udder	Fert	Levensduur	Productie	Udder	Fert									
542024	ENCOURAGE	243185	Pregrants' Trait	✓	+8%	122%	412	104	102	106	106	104	1791	134	128	112	92	638	112	706	99	106	
542072	FLAGSTONE	422561	Teddy's + Robert	✓	+8%	15%	356	98	102	106	106	102	1584	440	114	118	70	557	110	511	105	106	
542041	FASTLANE	242610	Stampten's Leden	✓	+7%	14%	312	102	106	107	102	104	216	523	124	81	48	376	107	549	104	104	
542034	NOBANNER	243160	Pregrants' Robert	✓	+7%	+8%	303	102	102	106	107	106	105	756	143	116	91	42	375	108	523	104	104
542108	SAMMIE P BP	234156	Larsch PP's Marie	✓	+8%	+13%	301	98	99	111	105	106	99	868	131	131	86	60	443	105	620	105	106
542088	LIBERATE	342516	Gigetta's Ruth Fleur	✓	+5%	+13%	298	104	100	108	102	105	96	1400	137	128	97	77	538	105	573	101	105
542102	ARTEMIS PP	243156	Mara P's Cedric	✓	+5%	+15%	281	105	102	107	104	103	100	990	148	117	89	51	413	106	533	101	106
542075	BORESO	321465	Rosema's Frith	✓	+8%	+14%	281	101	102	109	103	100	107	1408	120	110	89	59	493	105	693	98	109
542023	THERAPY	242610	Larsen's Nobby	✓	+8%	+13%	279	100	104	110	105	109	102	928	121	117	38	49	294	107	584	105	106

# Work needed for a health index

- Better knowledge of genetics needed
  - Mode of inheritance for monogenic traits
  - Heritabilities and genetic correlations for polygenic traits needed
- Create index
  - How to weigh Severity?
- Set up estimation of breeding values
  - Overcome mistrust -> communication needed
  - Look at GB and Sweden
- Advice breeders
  - Include MK in advice

# What is next

- 2 BSc projects ongoing (Schapendoes + Saarloos?)
  - Outcross
  - DNA vs. pedigree
- Final Report
  - Overview results ACT group and theses
  - General advice for breed societies
    - When and how to use DNA typing?
    - When and how to do an outcross?
    - How to deal with multiple genetic diseases
  - Breeding value estimation



# Tijd voor discussie

Priority with mutiple diseases

Outcross

Use of DNA

Breeding values

And more...

