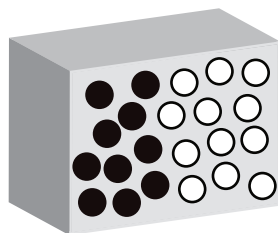


RDA or dbRDA?

	RDA	dbRDA
Missing	Sensitive	Not sensitive
Noise	Less	More



What is my sampling scheme?



Individual-based



GENOTYPES

p

n		SNP1 SNP2 SNP3...		
		AA	TT	GC
	IndA	AA	TT	GC
	IndB	AA	TT	GG
	IndC	AC	TT	GG
	IndD	AC	TA	GG

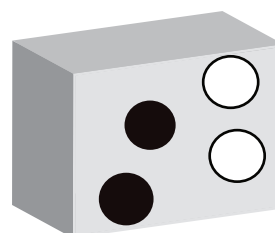
Can missing data bias my results?

No:
Few 0s or
0 = absence

distance-based
RDA

Yes:
0 different from
absence

RDA on genotypes



Population-based



ALLELE FREQUENCIES

p

n		SNP1 SNP2 SNP3...		
		0.12	0.05	0.41
	PopA	0.12	0.05	0.41
	PopB	0.26	0.01	0.01

*Do I have enough samples
(n observations $>$ p variables)?*

No: $n < p$

distance-based
RDA at the individual level

Yes : $n > p$

RDA on allele frequencies