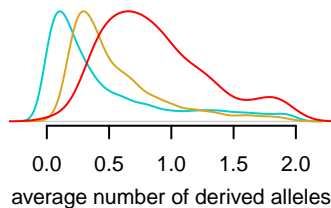
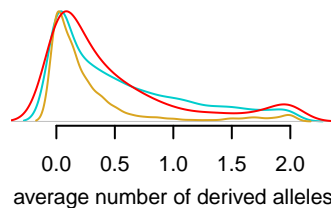


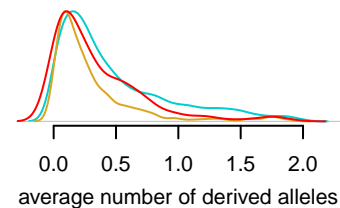
**red : is1**



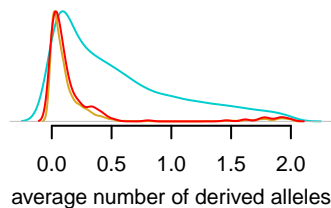
**red : is2**



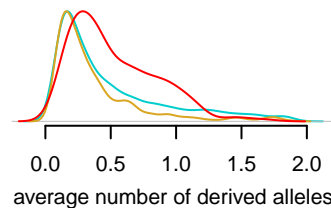
**red : is3**



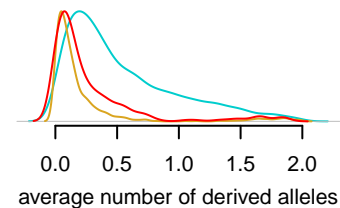
**red : is4**



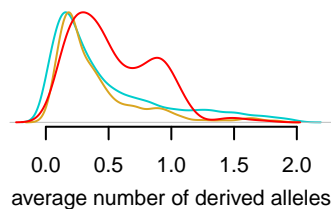
**red : isA**



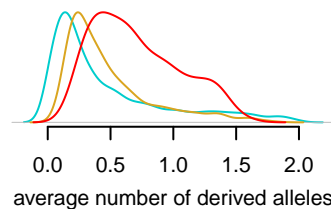
**red : isD**



**red : isO**



**red : isN**



— top 25% kME genes  
— all genes in module  
— genes not in module