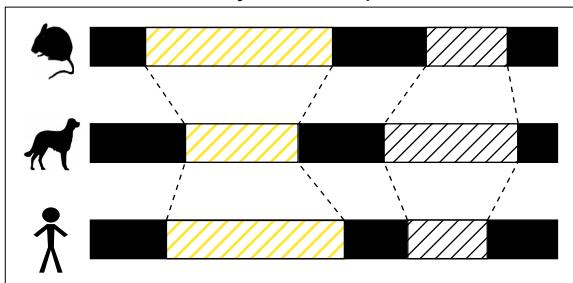
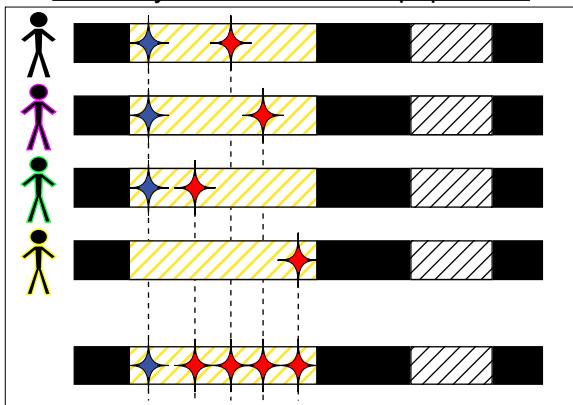


a

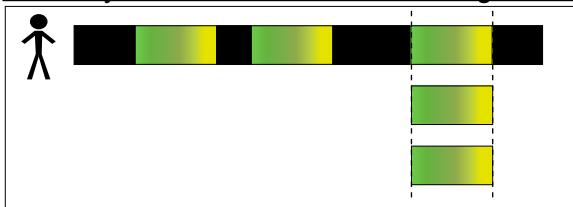
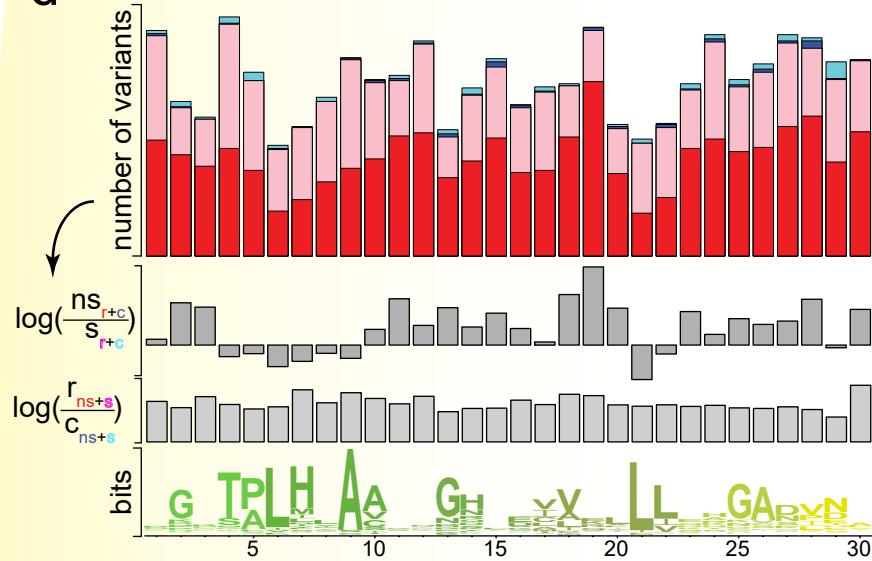
Similarity across species

**b**

Similarity within the human population

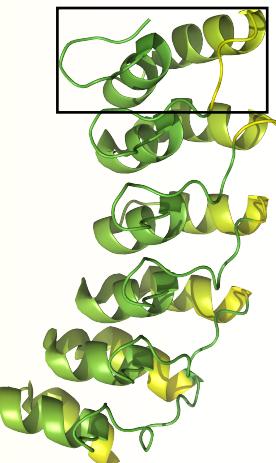
**C**

Similarity within the human reference genome

**d**

common/rare
non-synonymous

(c/r)	(ns/s)
c	s
c	ns
r	s
r	ns
*	variant



ankyrin
domains