



# Mapping out African genomics research with 'sf'

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 EXEGETIC



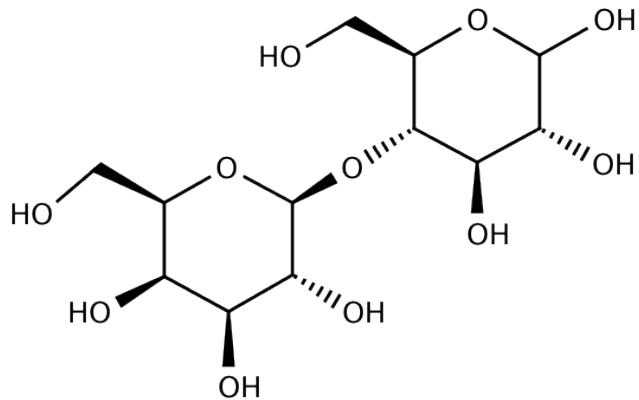
# What is the Human Genome?



The collection of all of our genes,  
the biological material  
which shapes much of who we are

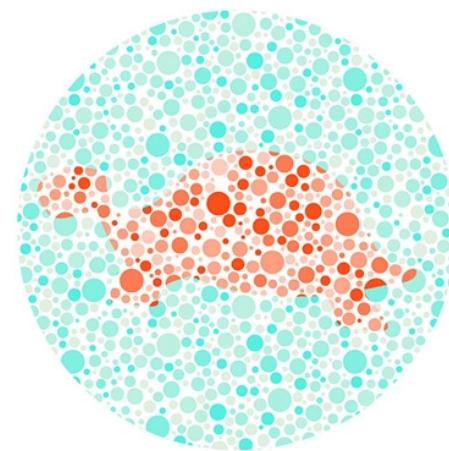
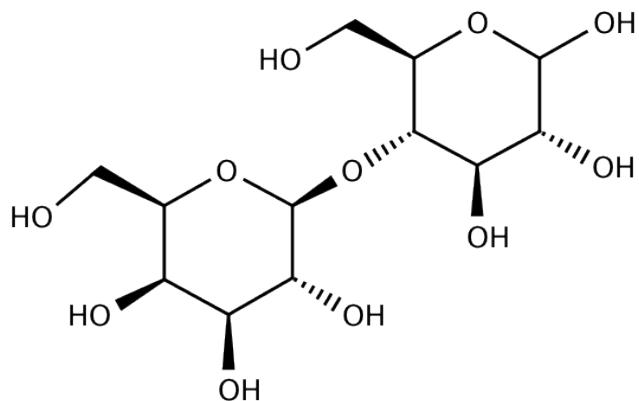
## Our genes influence traits as diverse as:

- whether we can digest lactose



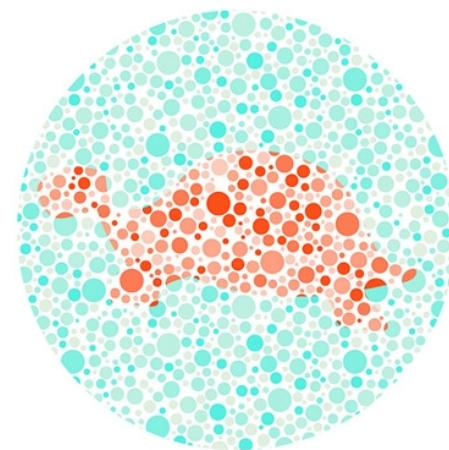
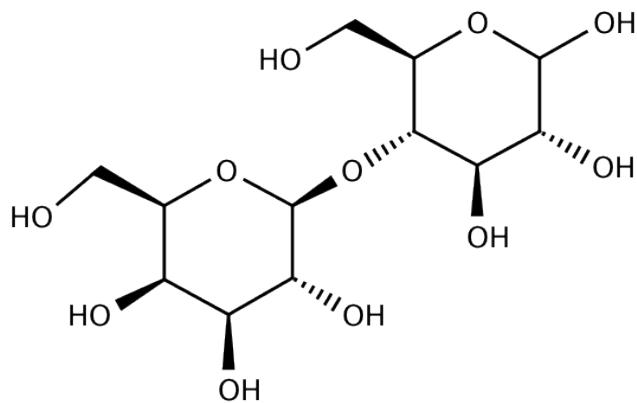
# Our genes influence traits as diverse as:

- whether we can digest lactose
- our ability to distinguish pink from blue



# Our genes influence traits as diverse as:

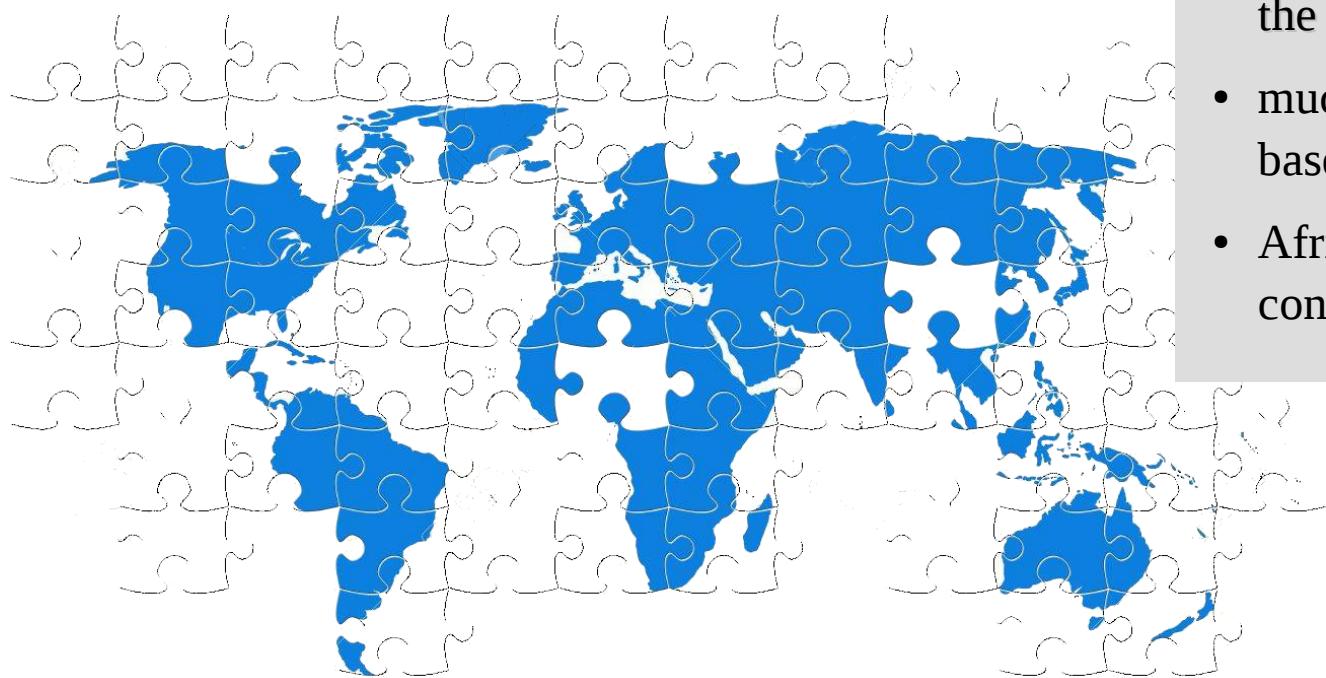
- whether we can digest lactose
- our ability to distinguish pink from blue



- our risk of developing heart disease



# What makes African genome data so valuable?



- we've only just scratched the surface
- much of what we know is based on European populations
- African populations harbour considerable genetic diversity

## Recent studies exploring the human genome in African populations

VISUALIZING  
PROGRESS USING 'SF'

```
# load multiple packages in a single command using 'easypackages'
library(easypackages)
packages("sf", "dplyr", "ggplot2")

# read in world borders data set (http://thematicmapping.org/downloads/world\_borders.php)
map_data <- st_read(inDir, 'TM_WORLD_BORDERS-0.3.shp')
```

```
# subset african data with a bit of help from dplyr
map_data <- map_data %>%
  filter(REGION==2)
```

Simple feature collection with 6 features and 11 fields

geometry type: MULTIPOLYGON

dimension: XY

bbox: xmin: -8.667223 ymin: -18.01639 xmax: 31.30278 ymax: 37.09139

epsg (SRID): 4326

proj4string: +proj=longlat +datum=WGS84 +no\_defs

	FIPS	ISO2	ISO3	UN	NAME	AREA	POP2005	REGION	SUBREGION	LONG	LAT	geometry
1	AG	DZ	DZA	12	Algeria	238174	32854159	2	15	2.632	28.163	MULTIPOLYGON (((2.96361 36....
2	AO	AO	AGO	24	Angola	124670	16095214	2	17	17.544	-12.296	MULTIPOLYGON (((11.75083 -1....
3	BN	BJ	BEN	204	Benin	11062	8490301	2	11	2.469	10.541	MULTIPOLYGON (((2.484418 6....
4	CF	CG	COG	178	Congo	34150	3609851	2	17	15.986	-0.055	MULTIPOLYGON (((12.77905 -4....
5	CG	CD	COD	180	Democratic Republic of the Congo	226705	58740547	2	17	23.654	-2.876	MULTIPOLYGON (((12.95305 -5....
6	BY	BI	BDI	108	Burundi	2568	7858791	2	14	29.887	-3.356	MULTIPOLYGON (((29.2299 -3....

```

# read in sampling info
sampling_info <- (read.csv(paste0(inDir, "/sampling_info.csv")))

# add coordinates derived from map_data
sampling_data <- full_join(sampling_info, sampling_coordinates, by = 'NAME')

```

		NAME	MAG	LON	LAT
1		Uganda	100	32.3860	1.28000
2		South Africa	100	23.1210	-30.55800
3		Ethiopia	120	39.6160	8.62600
4		Nigeria	99	8.1050	9.59400
5		Gambia	113	-15.3860	13.45300
6		Kenya	101	37.8580	0.53000
7		Sierra Leone	85	-11.7920	8.56000
8		Nigeria	109	8.1050	9.59400
9		South Africa	8	23.1210	-30.55800
10		South Africa	8	23.1210	-30.55800
11		Burkina Faso	34	-1.7400	12.27800
12		Mali	50	-3.5240	17.35000
13		Nigeria	49	8.1050	9.59400
14		Ghana	26	-1.2070	7.96000
15		Benin	50	2.4690	10.54100
16		Botswana	48	23.8150	-22.18200
17		Zambia	41	26.3200	-14.61400
18		Cameroon	50	12.2770	5.13300
19		Other	400	102.3446	7.36965
20		Guinea	46	-10.9420	10.43900
21		Cote d'Ivoire	40	-5.5560	7.63200
22	Democratic Republic of the Congo		23	23.6540	-2.87600
23		Uganda	50	32.3860	1.28000

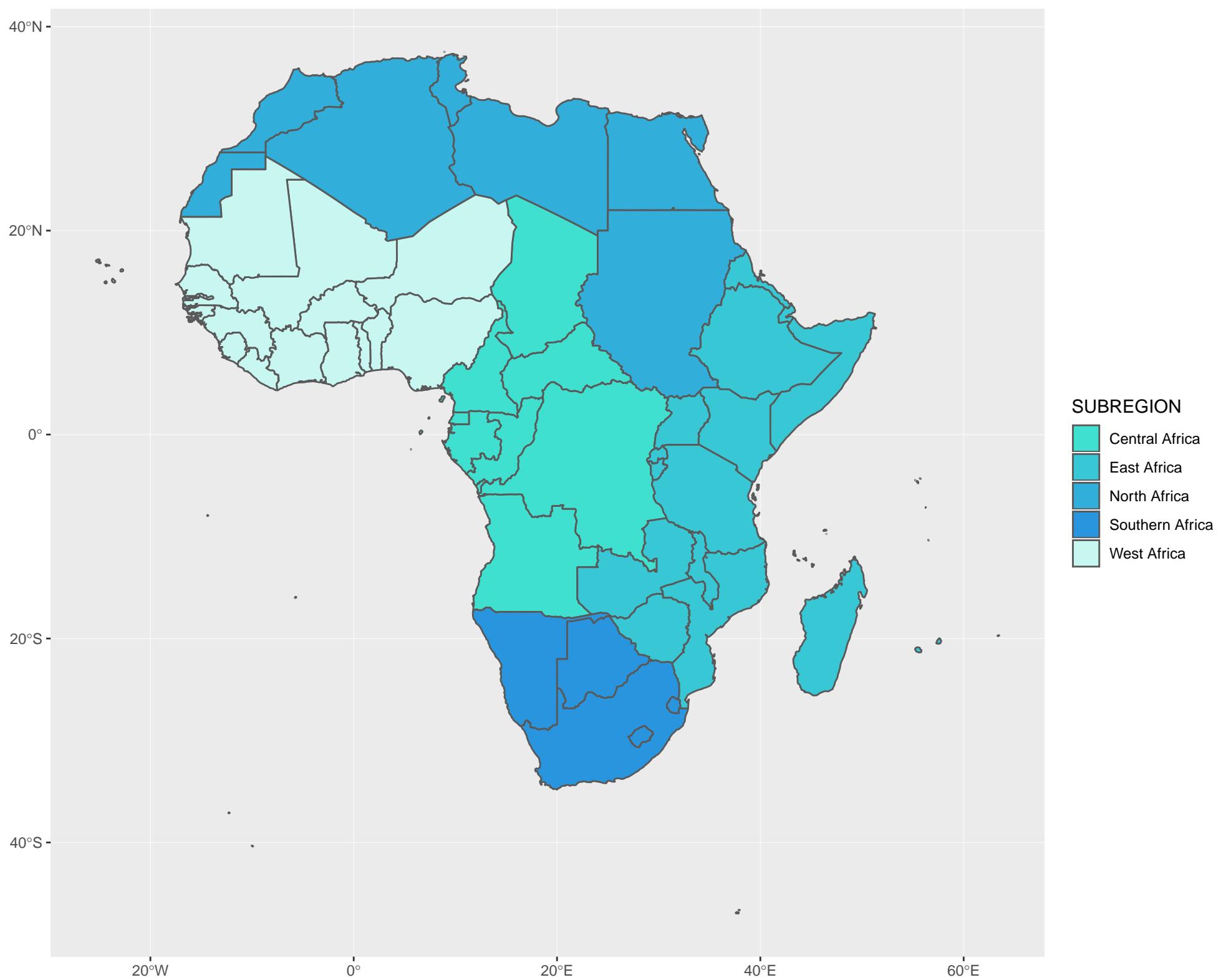
```
# quickly find colours in between
```

```
colorRampPalette(colors=c("Turquoise", "Blue"))(10)
```

```
[1] "#40E0D0" "#38C7D5" "#31AEDA" "#2A95DF" "#237CE4" "#1C63EA" "#154AEF" "#0E31F4" "#0718F9" "#0000FF"
```

```
# plot map_data
```

```
ggplot(map_data) + geom_sf(aes(fill = SUBREGION)) +  
scale_fill_manual("SUBREGION", values = c("#40E0D0", "#38C7D5",  
"#31AEDA", "#2A95DF", "#C8F6F1"))
```



```
# add country polygons, with labels above or below country midpoints  
p <- ggplot() +  
  
  geom_sf(data = map_data, aes(fill = SUBREGION), color = 'transparent', lwd = 0.5) +  
  
  geom_label(data = map_data, aes(map_data$LON, map_data$LAT, label = map_data$NAME),  
             alpha = 0, size = 2.8, label.size = 0) +  
  
  scale_fill_manual("Region", values = c("#40E0D0", "#38C7D5", "#31AEDA",  
                         "#2A95DF", "#C8F6F1")) +
```

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  scale_fill_manual("Region", values = c("#40E0D0", "#38C7D5", "#31AEDA",  
                     "#2A95DF", "#C8F6F1")) +  
  
# specify map variables  
  
q <- p +  
  
theme(legend.justification = c(0.6, -0.1), legend.position = c(0.28, 0.2), axis.title.x = element_blank(),  
      axis.title.y = element_blank(), panel.background = element_rect(fill = "#DADADA")
```

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  geom_label(data = map_data, aes(map_data$LON, map_data$LAT, label = map_data$NAME),
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        axis.title.y = element_blank(), panel.background = element_rect(fill = "#DADADA"))
# add sampling data, scaled by number of participants
r <- q +
  geom_point(data = sampling_data, aes(x = LON, y = LAT), stroke = 0,
             shape = 21, fill = "#6D6D6D", cex = (sqrt(sampling_data$MAG)/2))
```

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p <- ggplot() +
  geom_sf(data = map_data, aes(fill = SUBREGION), color = 'transparent', lwd = 0.5) +
  geom_label(data = map_data, aes(map_data$LON, map_data$LAT, label = map_data$NAME),
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r <- q +
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             shape = 21, fill = "#6D6D6D", cex = (sqrt(sampling_data$MAG)/2))
# export figure
r %>% ggsave(filename=paste0(outDir, "/figure.pdf"), width = 28, height = 21, unit = "cm")
```

