

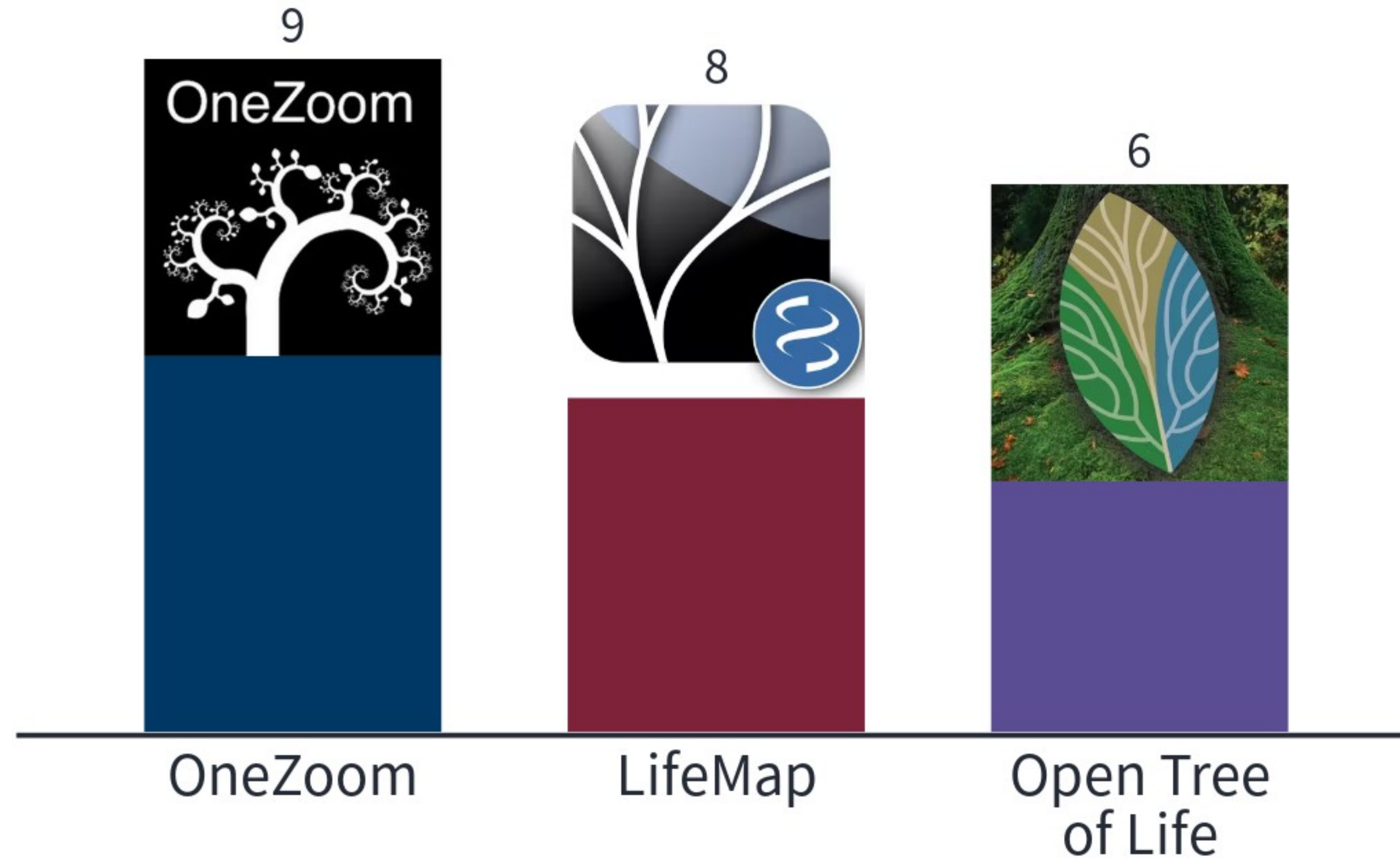


# Criteria for evaluating

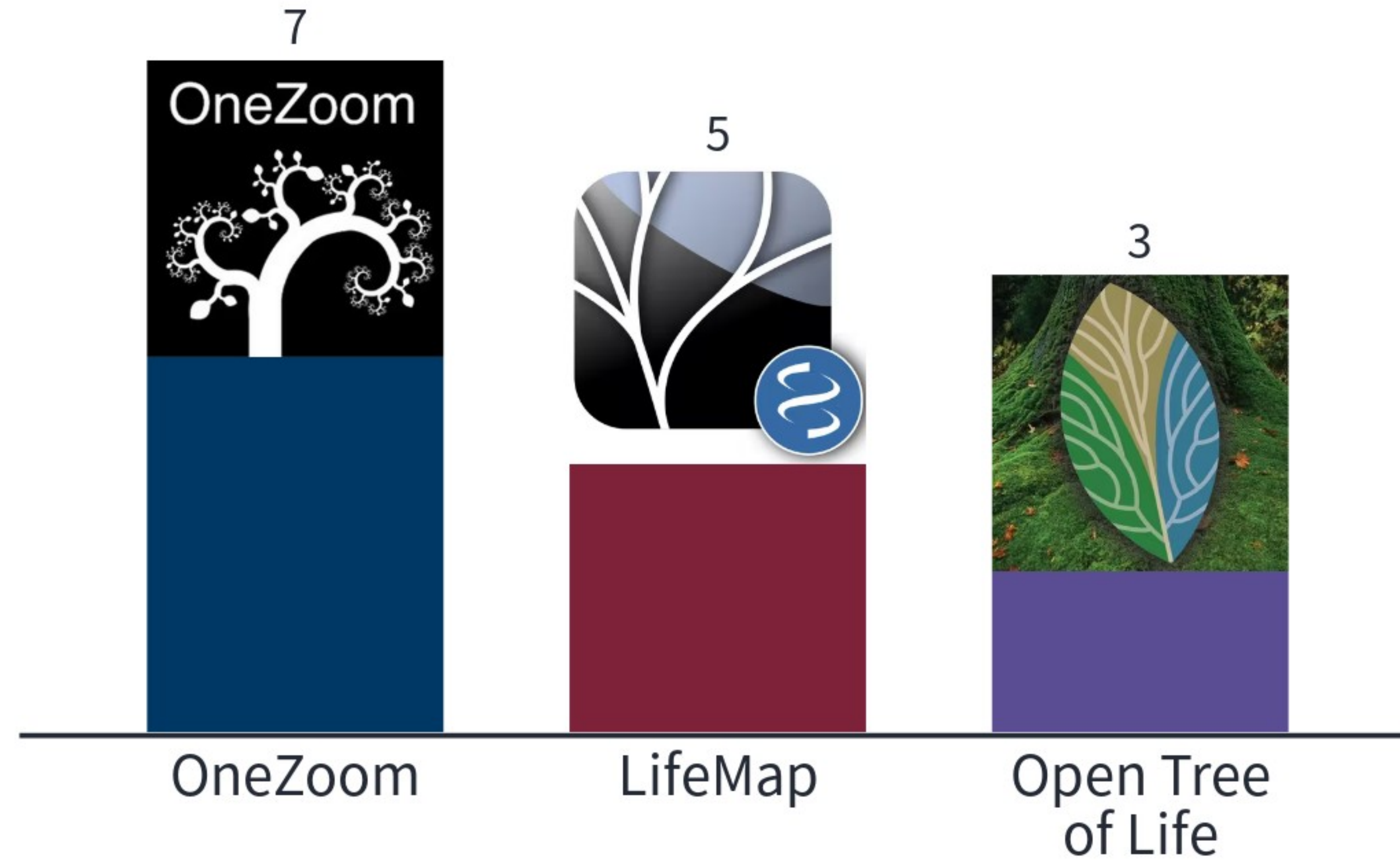
- Bullet 1
- Bullet 2
- Bullet 3



# Ease of finding a species or taxonomic group

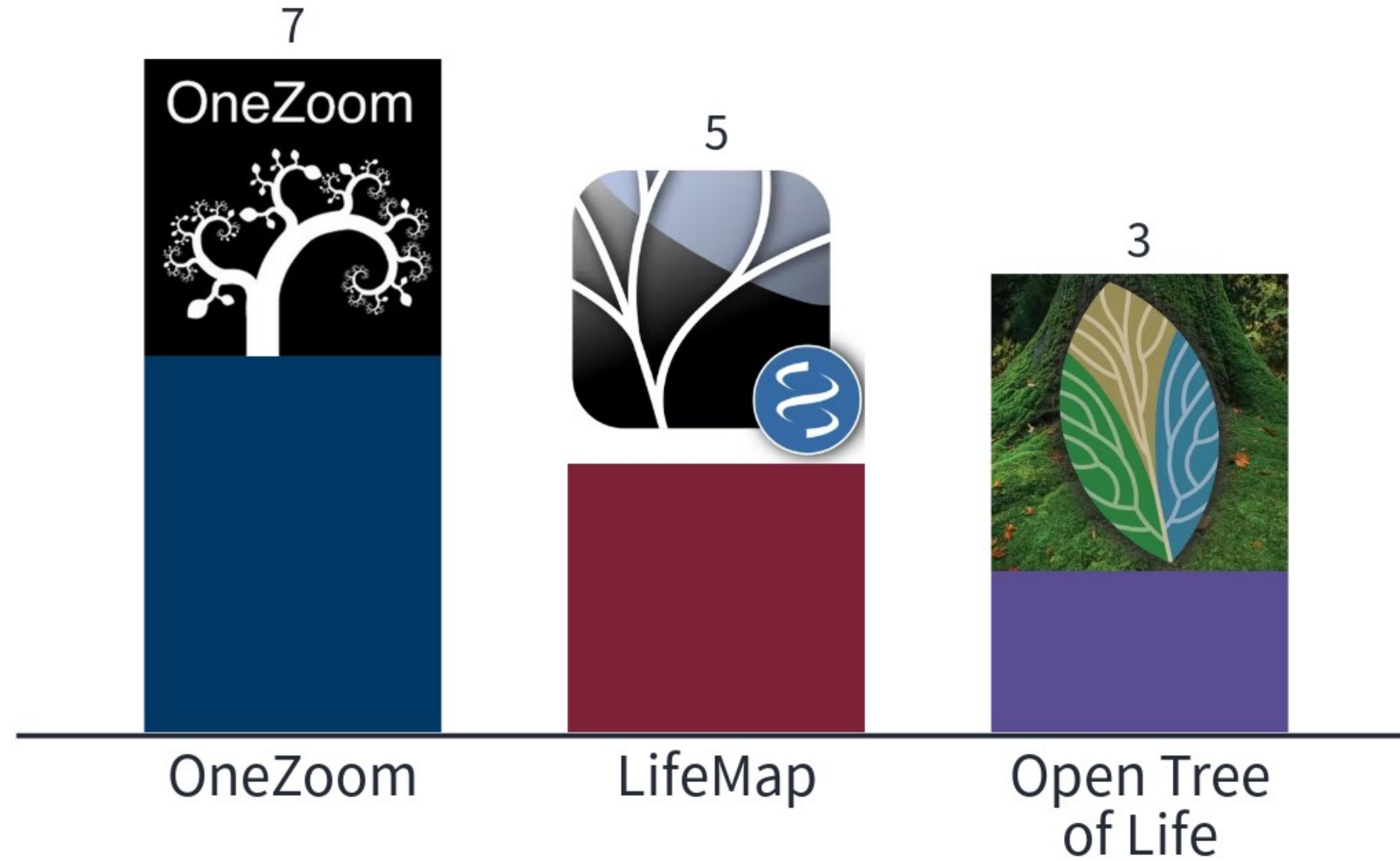


# Ease of finding common ancestor of two species

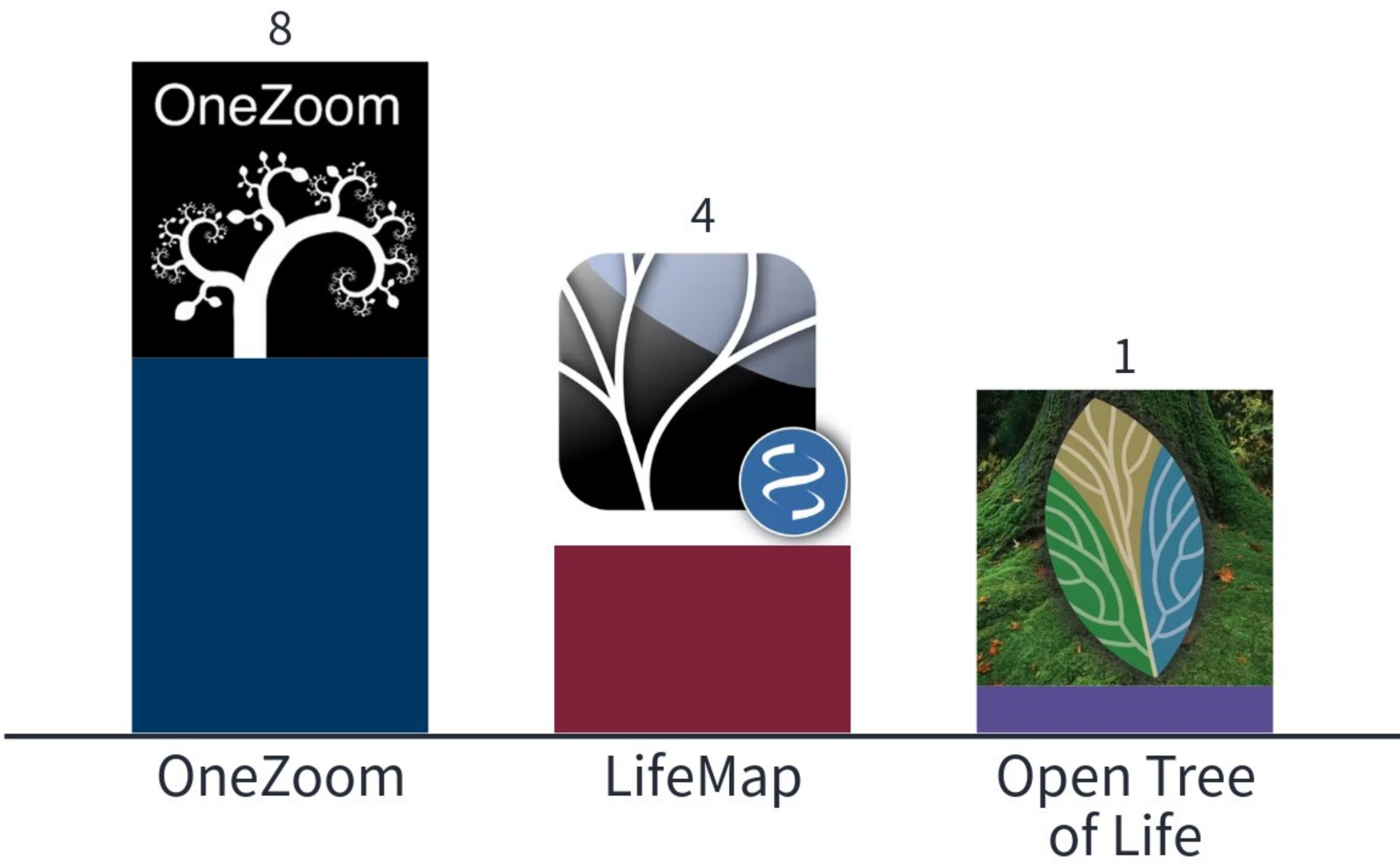




# Can you discover the size of a taxonomic group



# Wow factor





# Any other thoughts or comments?

How easy it is to navigate around the tree eg. zoom function & whether you can interpret the timeline. Links to other data (eg. sequences)

I noticed a difference between onezoom and the other 2. The latter tended to have more "species" in the family I was looking at - but to my knowledge these aren't widely recognised as species (more subs/ hybrids) which I think is a little misleading

The ability to use common names highly varied between different species in all the trees.