

Population Genetics for Living Documents

learns from population genetics how to accelerate the improvement of complicated text/code bases while also improving their stability via smart combinations of the 5 fundamental factors of evolution

0. Reproduction

computers track genealogy via version control systems

1. Mutation

changes by users, reviewers, authors, new todos, data ...

2. Selection

recrushing review by multitudes of views and people with extended real-world testing by beginners and experts

3. Population size

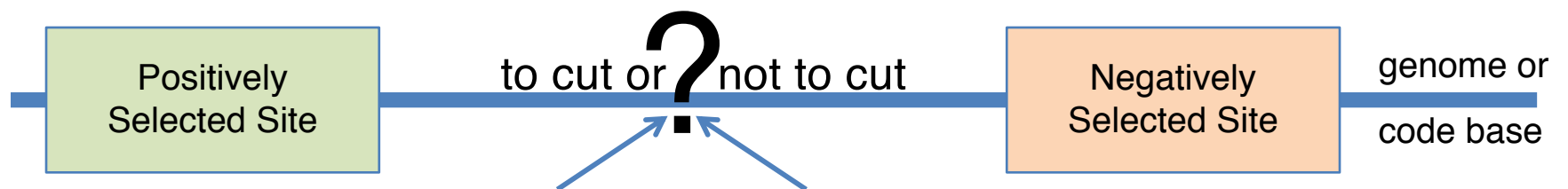
more feedback is often better but small groups can be key

4. Recombination

with a user-friendly quality annotation algebra for telling computers & editors which cut/replace options work well

5. Migration

support for traveling lightly to new systems for finding new views to test for improving overall capability integration



Recombination enables selection of excellence by **breaking up mixed bags** ... if mix is unintended

by **keeping mixed bags** combos are tested via **Linkage** but its inability to allow for sorting pros/cons costs quality

Asexual modes of inheritance are so loyal to all copy errors linked to their DNA that they collapse in echo-chambers of self-similarity after losing all relevance and connection to their environment.

... not even the Coronavirus does that as it keeps recombining ...