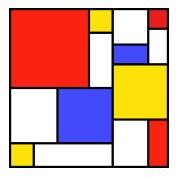
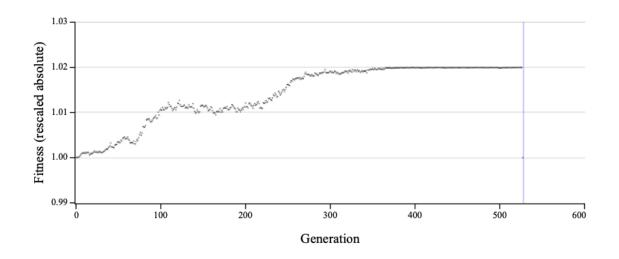
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#9: Selective Sweeps

- a selective sweep model involves:
 - a sweep mutation, generally beneficial
 - followed until establishment / completion
 - often conditional on success



- sweeps can be:
 - "hard" introduced in a single ancestor
 - "soft" from multiple ancestral lineages

- sweeps can be:
 - "hard" introduced in a single ancestor
 - "soft" from multiple ancestral lineages
- sweeps can be:
 - from a new, introduced mutation
 - from identical recurrent mutations
 - from standing genetic variation

- a sweep has to start somewhere!
- select candidate subpopulation(s)
- get the genomes from the candidates
 - p1.genomes
 - c(p1,p2) genomes
 - sim_subpopulations_genomes
- draw a genome or genomes to modify:
 - sample()

- a sweep involves a mutation
 - might be introduced
 - might be pre-existing
- introduce a sweep mutation:
 - addNewMutation()
 - addNewDrawnMutation()
- transmute into a sweep mutation:
 - setMutationType()
 - setSelectionCoefficient()

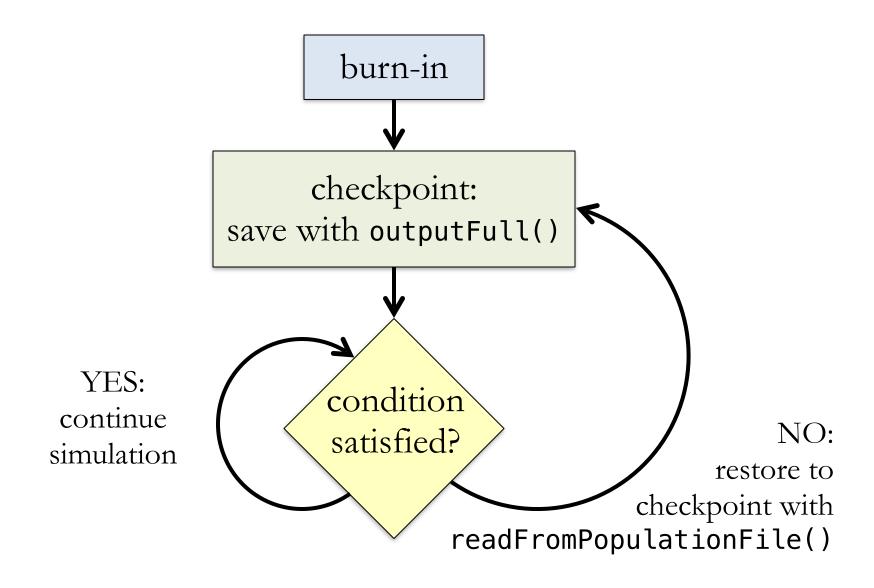
- you might want to know
 - when it's lost
 - when it's established
 - when it's fixed

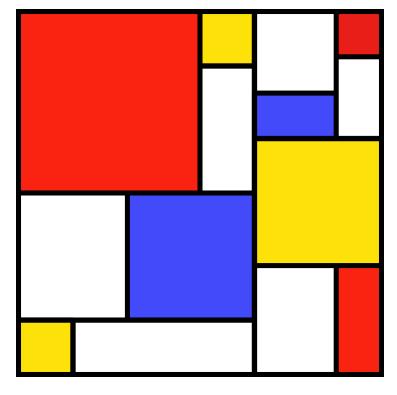
- you might want to know
 - when it's lost
 - when it's established
 - when it's fixed
- monitor frequency:
 - countOfMutationsOfType()
 - mutationsOfType()
 - mutationFrequencies()
 - mutationCounts()

- often desirable to be "conditional"
 - guarantees a particular outcome
 - doesn't bias the process leading there

- often desirable to be "conditional"
 - guarantees a particular outcome
 - doesn't bias the process leading there
- a conditional simulation should:
 - save at a key point when the condition begins
 - watch for deviation from the requirement
 - restore to the key point with a new seed

- to make a conditional simulation...
- save / restore population state:
 - outputFull()
 - readFromPopulationFile()
- manage the random number seed:
 - setSeed() (this is optional)
- conditionality can be on anything!





SLiM Workshop Exercise #9