cross accuracy

--rows--
parms_species
- Drosophila_melanogaster
- none

parms_typeSimulations
- arroundHomRegions
- arroundRepeats
- fast
- realSVs
- uniform

parms_numeric_sample
- 0
- 1
- 2

parms_numeric_run
- 0
- 1
- 2

--cols--

test_species
- Drosophila_melanogaster

test_typeSimulations
- arroundHomRegions
- arroundRepeats
- realSVs
- uniform

test_numeric_sample
- 0
- 1
- 2

test_numeric_run
- 0
- 1
- 2

test_simName
- sim1
- sim2

precision

1.0
0.8
0.6
0.4
0.2
0.0

test_species
test_typeSimulations
test_numeric_sample
test_numeric_run
test_simName

parms_idx

parms_species
parms_typeSimulations
parms_numeric_sample
parms_numeric_run

test_idx