Dataset Statistics
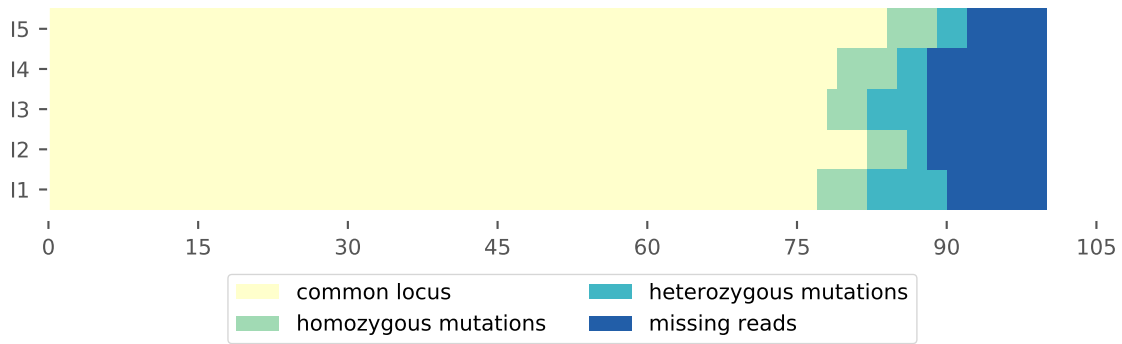
Distribution of Locus Types per Individual

Legend: common locus, homozygous mutations, heterozygous mutations, missing reads

Global Distribution of Locus Event Types

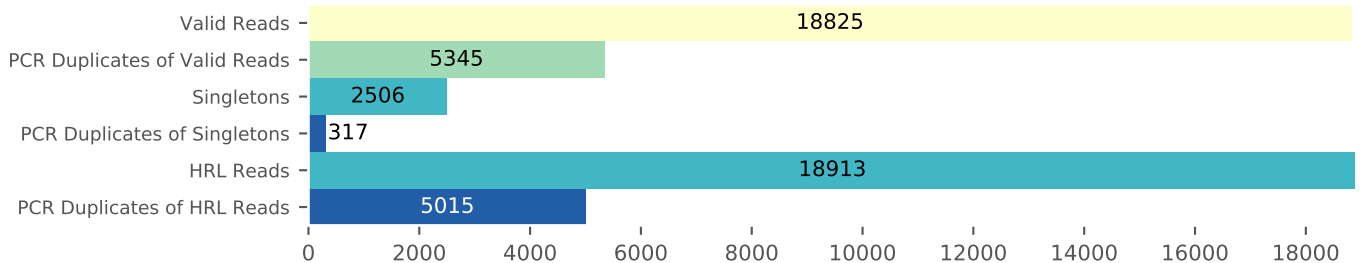| Event Type	Count
common	400
mutation homozygous	24
mutation heterozygous	22
dropout	54

PCR Ratio

Category	PCR Duplicates Ratio
All Reads	0.2653
Valid Reads	0.2839
HRL Reads	0.2652
Singletons	0.1265

Legend: Actual Reads, PCR Duplicates

Read Origin Breakdown

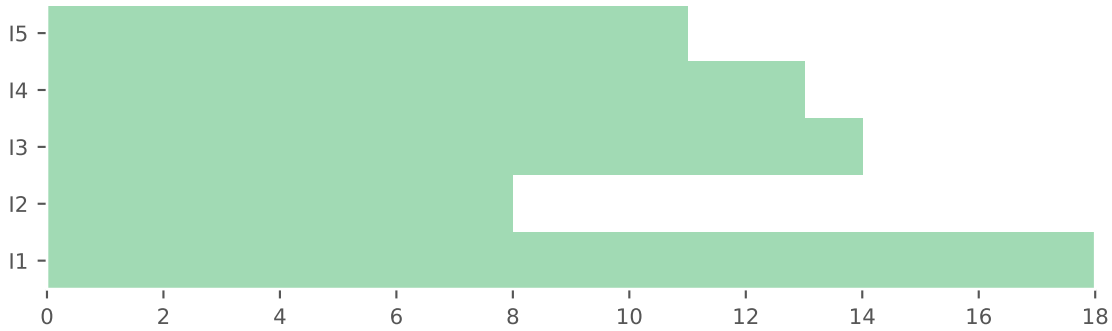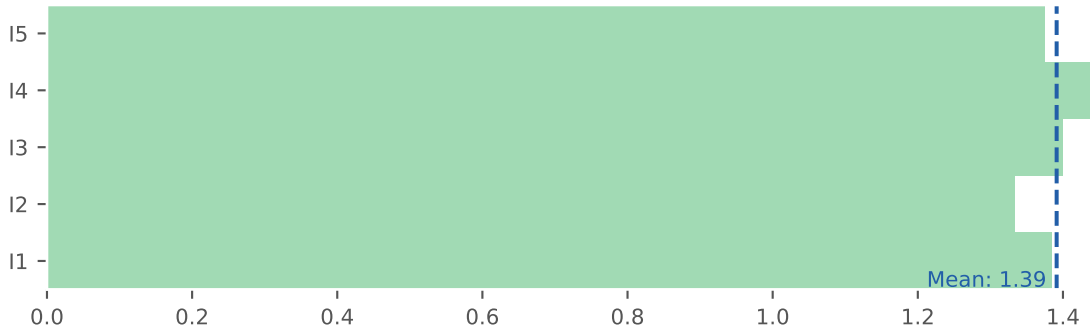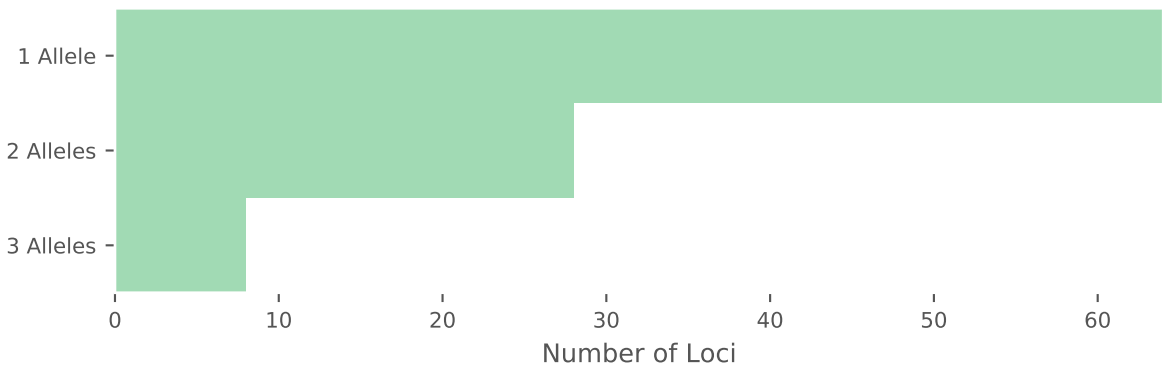| Category	Count
Valid Reads	18825
PCR Duplicates of Valid Reads	5345
Singletons	2506
PCR Duplicates of Singletons	317
HRL Reads	18913
PCR Duplicates of HRL Reads	5015
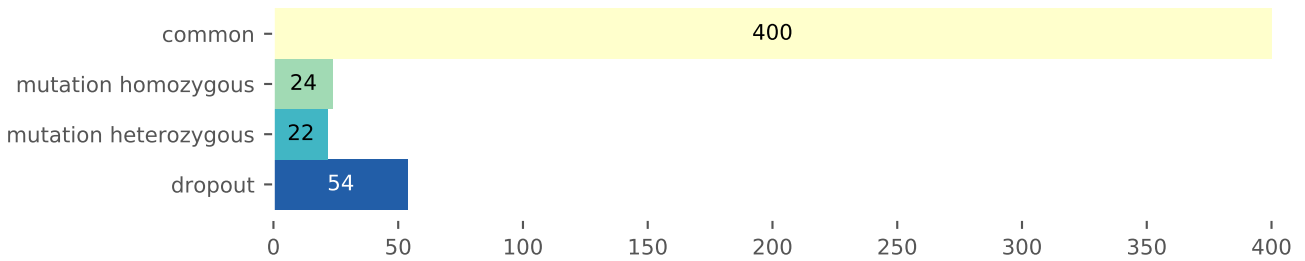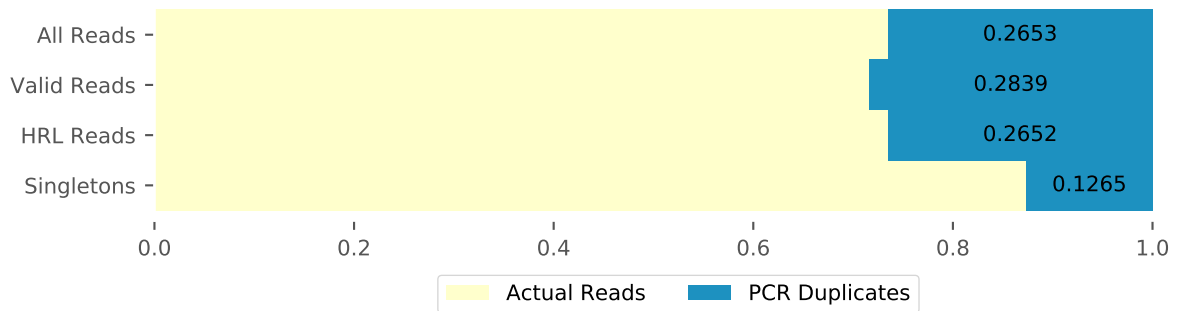
Dataset Statistics

Total number of mutations per individual

Mean number of mutations per mutation event

Mean: 1.39

Distribution of #Different Alleles per Locus

Number of Loci

Distribution of Individual Coverage Values

Nr of Loci

Coverage