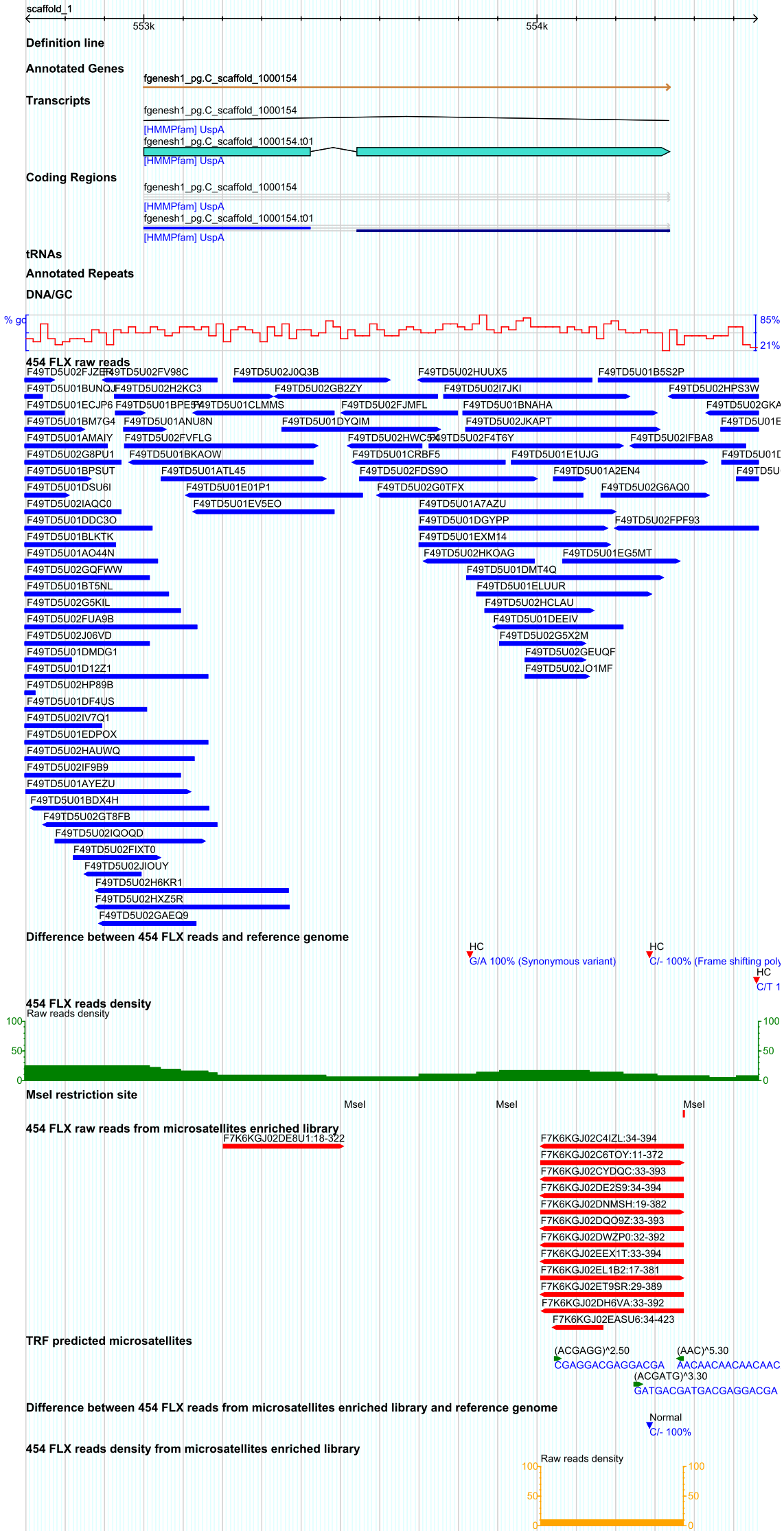
scaffold_1
553k
554k
Definition line
Annotated Genes
fgenesh1_pg.C_scaffold_1000154
Transcripts
fgenesh1_pg.C_scaffold_1000154
[HMMPfam] UspA
fgenesh1_pg.C_scaffold_1000154.t01
[HMMPfam] UspA
Coding Regions
fgenesh1_pg.C_scaffold_1000154
[HMMPfam] UspA
fgenesh1_pg.C_scaffold_1000154.t01
[HMMPfam] UspA
tRNAs
Annotated Repeats
DNA/GC
85%
21%
454 FLX raw reads
Difference between 454 FLX reads and reference genome
HC
G/A 100% (Synonymous variant)
HC
C/- 100% (Frame shifting poly
HC
C/T 1
454 FLX reads density
Raw reads density
100
50
0
MseI restriction site
MseI
MseI
MseI
454 FLX raw reads from microsatellites enriched library
F7K6KGJ02DE8U1:18-322
F7K6KGJ02C4IZL:34-394
F7K6KGJ02C6TOY:11-372
F7K6KGJ02CYDQC:33-393
F7K6KGJ02DE2S9:34-394
F7K6KGJ02DNMSH:19-382
F7K6KGJ02DQO9Z:33-393
F7K6KGJ02DWZP0:32-392
F7K6KGJ02EEX1T:33-394
F7K6KGJ02EL1B2:17-381
F7K6KGJ02ET9SR:29-389
F7K6KGJ02DH6VA:33-392
F7K6KGJ02EASU6:34-423
TRF predicted microsatellites
(ACGAGG)^2.50
CGAGGACGAGGACGA
(AAC)^5.30
AACAACAACAACAAC
(ACGATG)^3.30
GATGACGATGACGAGGACGA
Difference between 454 FLX reads from microsatellites enriched library and reference genome
Normal
C/- 100%
454 FLX reads density from microsatellites enriched library
Raw reads density
100
50
0