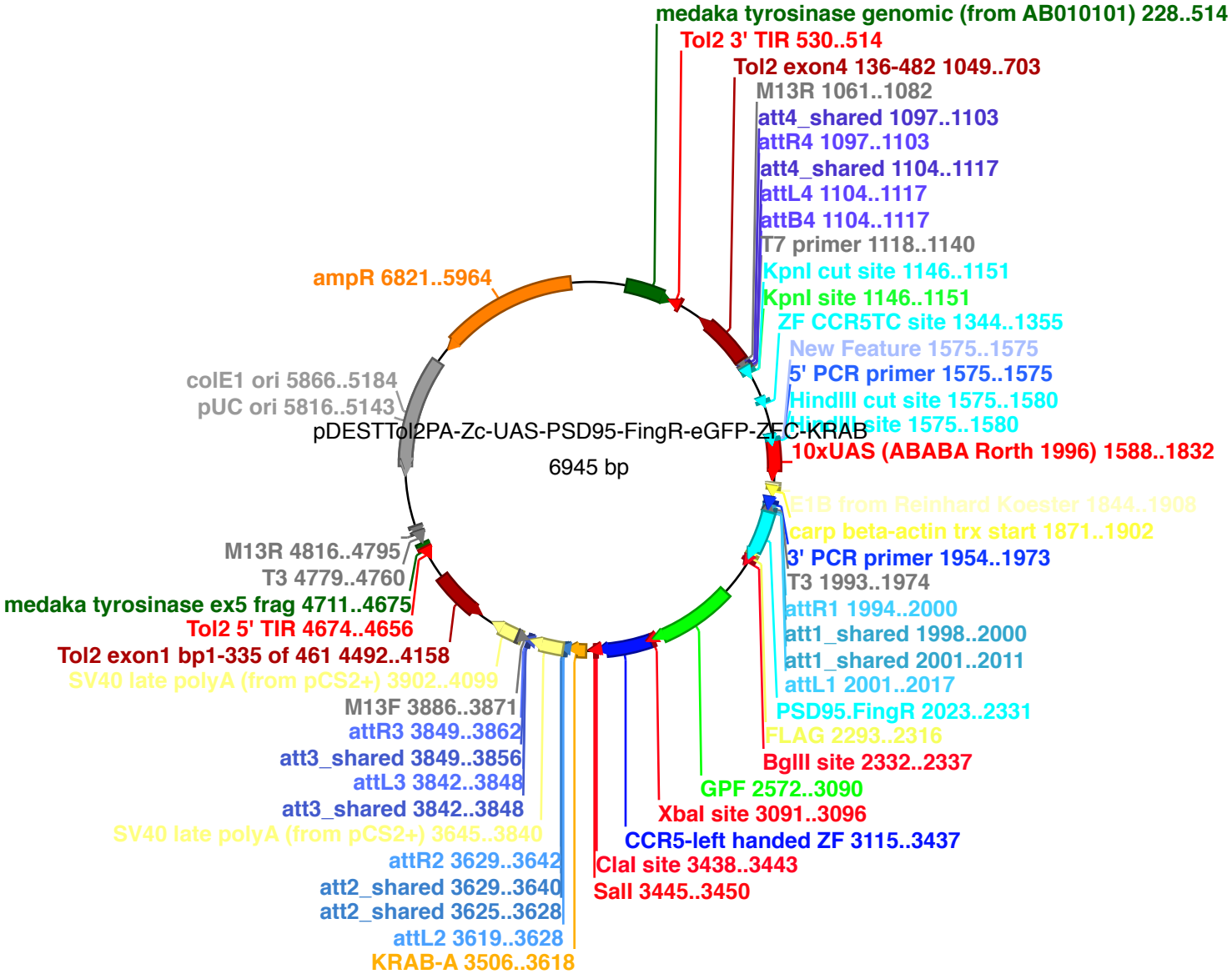
medaka tyrosinase genomic (from AB010101) 228..514
Tol2 3' TIR 530..514
Tol2 exon4 136-482 1049..703
M13R 1061..1082
att4_shared 1097..1103
attR4 1097..1103
att4_shared 1104..1117
attL4 1104..1117
attB4 1104..1117
T7 primer 1118..1140
KpnI cut site 1146..1151
KpnI site 1146..1151
ZF CCR5TC site 1344..1355
New Feature 1575..1575
5' PCR primer 1575..1575
HindIII cut site 1575..1580
HindIII site 1575..1580
10xUAS (ABABA Rorth 1996) 1588..1832
E1B from Reinhard Koester 1844..1908
carp beta-actin trx start 1871..1902
3' PCR primer 1954..1973
T3 1993..1974
attR1 1994..2000
att1_shared 1998..2000
att1_shared 2001..2011
attL1 2001..2017
PSD95.FingR 2023..2331
FLAG 2293..2316
BglII site 2332..2337
GPF 2572..3090
XbaI site 3091..3096
CCR5-left handed ZF 3115..3437
ClaI site 3438..3443
SalI 3445..3450
attR2 3629..3642
att2_shared 3629..3640
att2_shared 3625..3628
attL2 3619..3628
KRAB-A 3506..3618
SV40 late polyA (from pCS2+) 3645..3840
att3_shared 3842..3848
attL3 3842..3848
attR3 3849..3862
att3_shared 3849..3856
M13F 3886..3871
SV40 late polyA (from pCS2+) 3902..4099
Tol2 exon1 bp1-335 of 461 4492..4158
Tol2 5' TIR 4674..4656
medaka tyrosinase ex5 frag 4711..4675
T3 4779..4760
M13R 4816..4795
pUC ori 5816..5143
colE1 ori 5866..5184
ampR 6821..5964
pDESTTol2PA-Zc-UAS-PSD95-FingR-eGFP-ZFC-KRAB
6945 bp