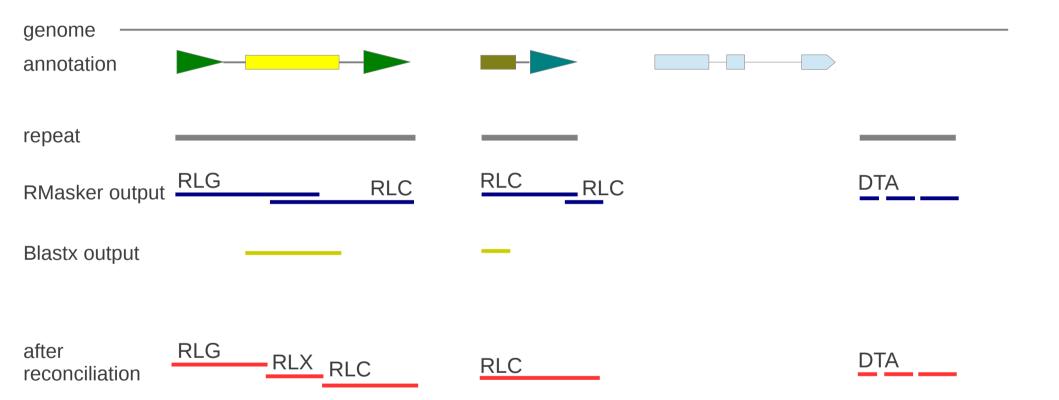
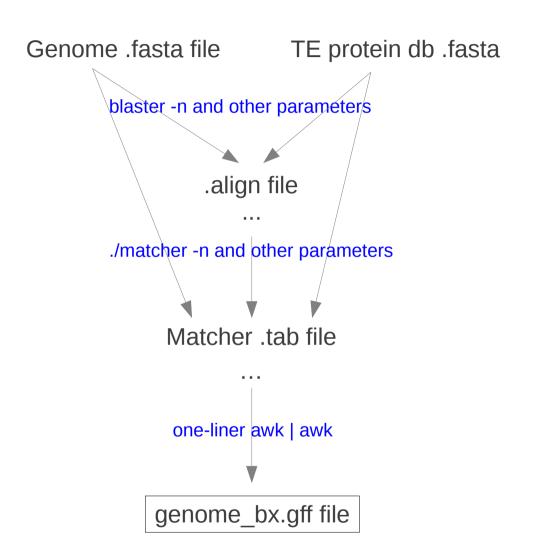
Repeat detection and reconciliation



Blaster pipeline for Blastx

RepeatMasker



Genome .fasta file
Nt repeat db .fasta

RepeatMasker -gff ... -engine -lib genome.fasta

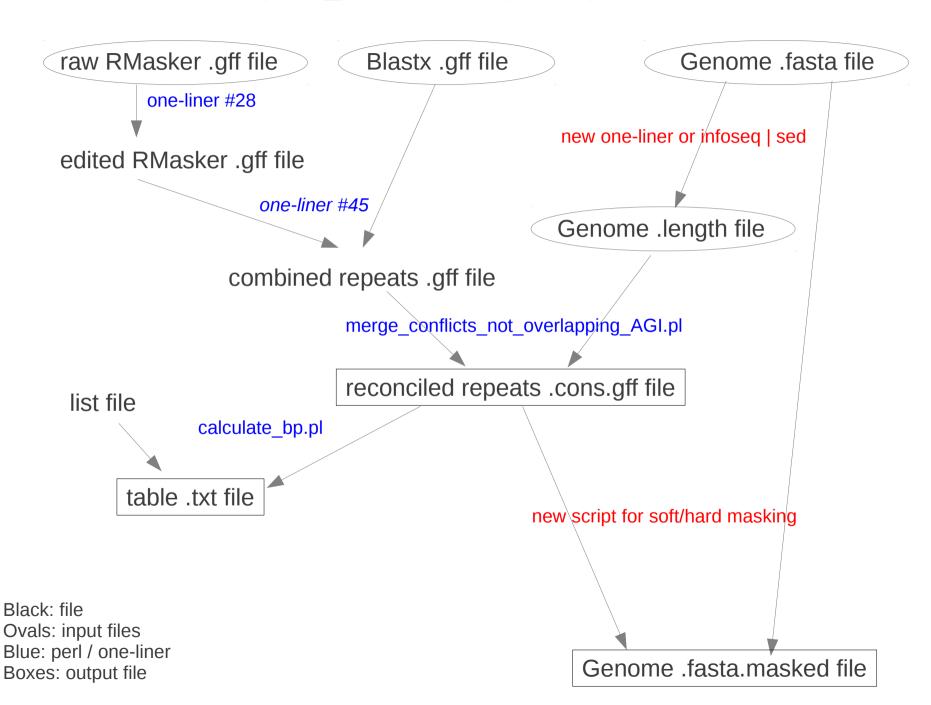
Genome.out.gff file

Genome.out
Genome.ori.out
Genome.cat
Genome.tbl
Genome.masked
Genome.ref

Black: file

Blue: perl / one-liner Boxes: output file

Repeat_Annotation.pl script



Automated Repeat Annotation Pipeline - Output

BRN	91964	482
BTN	44386	620
BXX	228	4
DHH	5375702	15679
DTA	1623797	7171
DTB	61	1
DTC	6057630	16390
DTH	1075348	5857
DTI	24208069	112938
DTM	2861511	9027
DTR	1344	32
DTT	360884	1785
DTX	1929242	13990
DXX	301879	4060
RIJ	12285	94
RIL	3242076	6670
RIT	288	6
RIX	146190	307
RLB	190	4
RLC	9150797	13560
RLE	15063	96
RLG	34398152	34349
RLX	258441	2323
RPP	27	1
RSU	1642616	11314
RXX	41432	818
RYD	230	5
SRC	17435	98
SRR	27722	182
SRS	7052	274
SRT	104059	261
SRX	887	6
SSS	1646444	30066
SSL	987034	33456
SXX	30	1
XXX	839220	12392

Obart_chr1	repeat	SSL	137	186				Name=SSL 0
Obart chr1	repeat	SSL	724	753	•	•	•	Name=SSL 1
Obart chr1	repeat	DTI	2554	2730	•	•	•	Name=DTI 2
Obart_chr1	repeat	DTH	3011	3032		•		Name=DTH 3
Obart_chr1	repeat	XXX	3033	3125		•		Name=XXX 4
Obart_chr1	repeat	RIX	3126	3167		•		Name=RIX 5
Obart chr1	repeat	SSS	3544	3574				Name=SSS 6
Obart chr1	repeat	DTI	7174	7219				Name=DTI 7
Obart chr1	repeat	DTI	9101	9363				Name=DTI 8
Obart chr1	repeat	DTI	11210	11308				Name=DTI 9
Obart_chr1	repeat	DTX	11309	11395				Name=DTX 10

>chr12