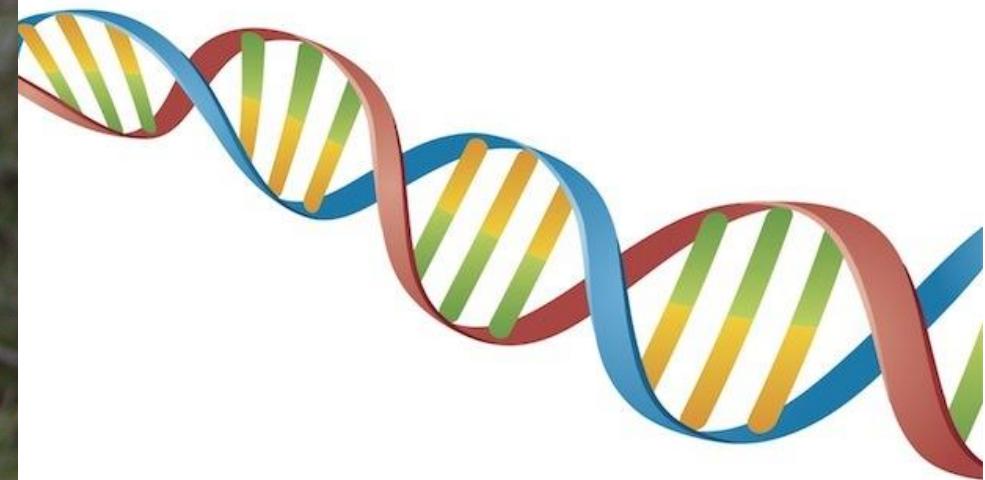


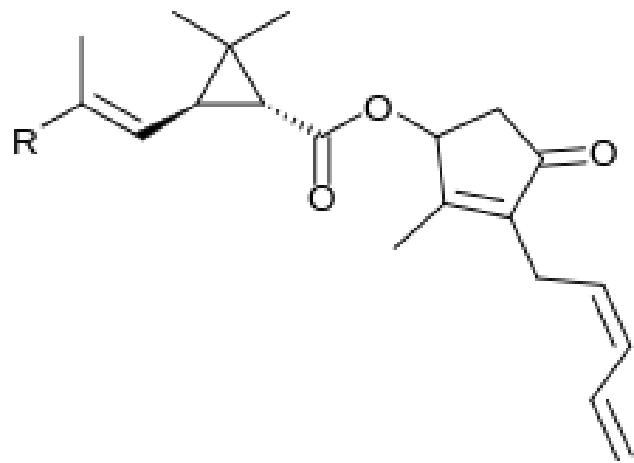
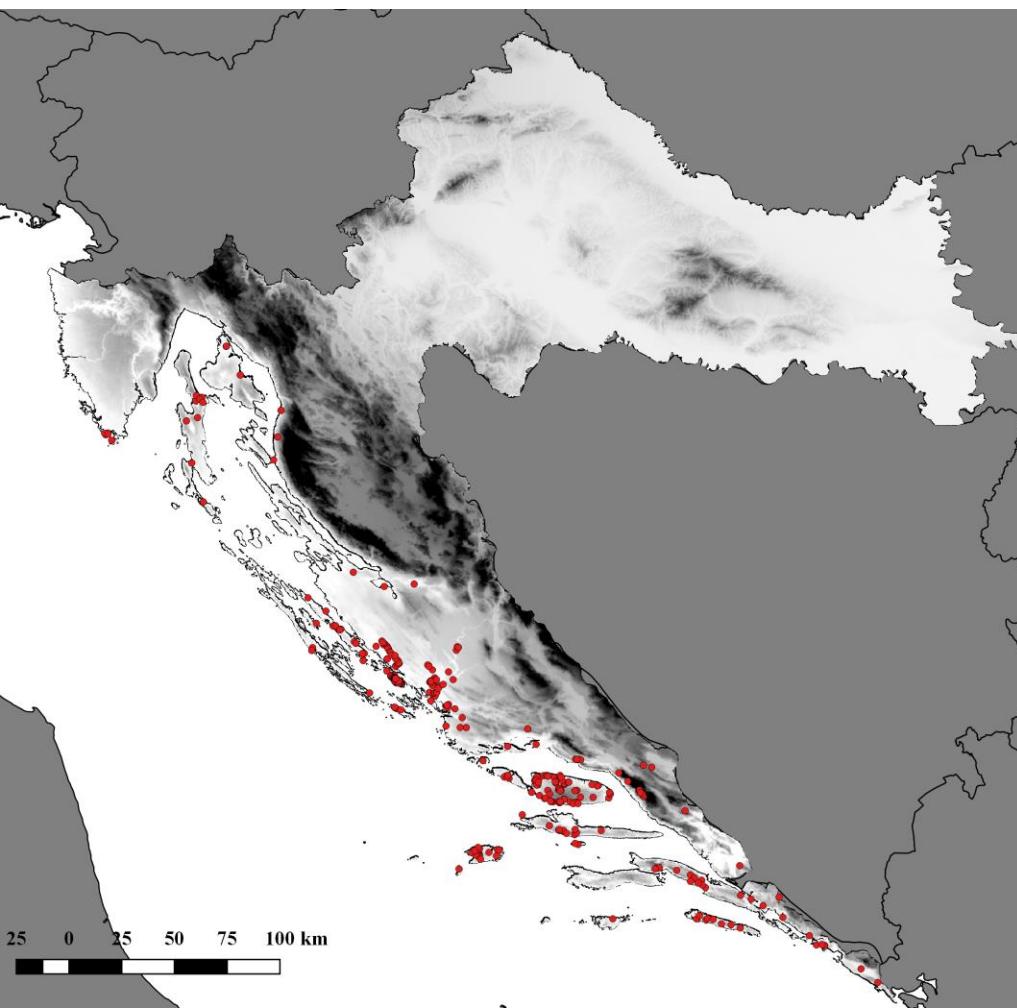
RAZVITAK MIKROSATELITNIH BILJEGA DALMATINSKOG BUHAĆA POMOĆU SEKVENCIRANJA VISOKE PROSUPNOSTI



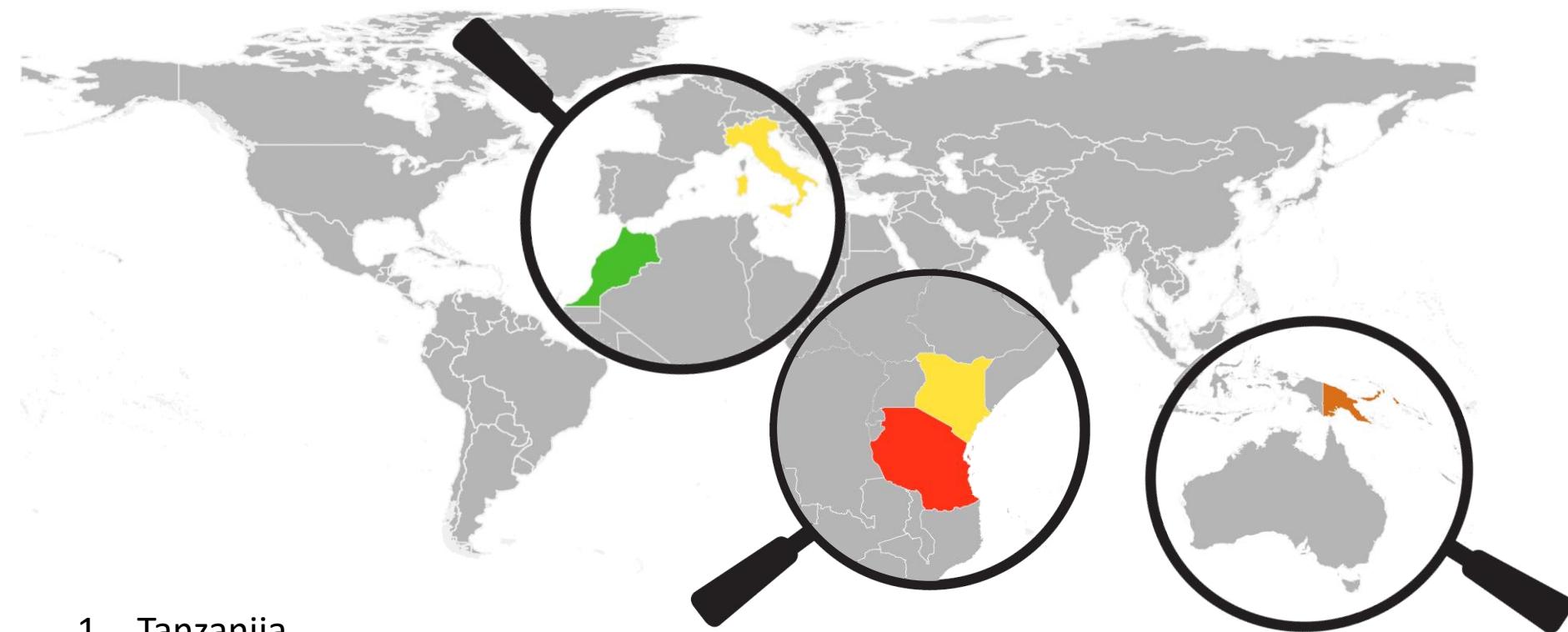
Filip Varga, mag. biol. exp.

Sveučilište u Zagrebu, Agronomski fakultet, Zagreb
Znanstveni centar izvrsnosti za bioraznolikost i molekularno oplemenjivanje bilja, Zagreb
E-mail: fvarga@agr.hr

Tanacetum cinerariifolium (Asteraceae)



PROIZVODNJA U SVIJETU (2015. godina)



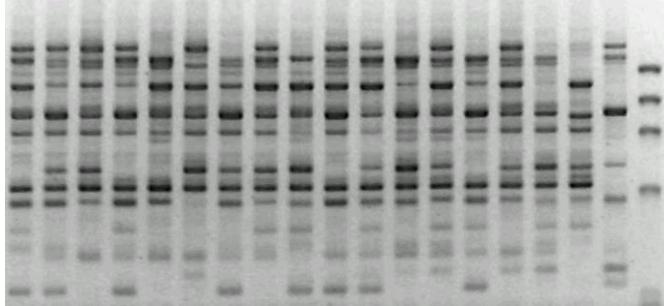
1. Tanzanija
2. Papua Nova Gvineja
3. Kenija
4. Italija
5. Maroko

VELIČINA GENOMA DALMATINSKOG BUHAČA ($2n$)

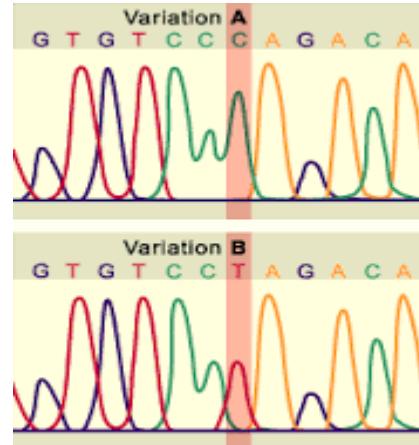


0 2 4 6 8 10 12 14 16 /Gbp

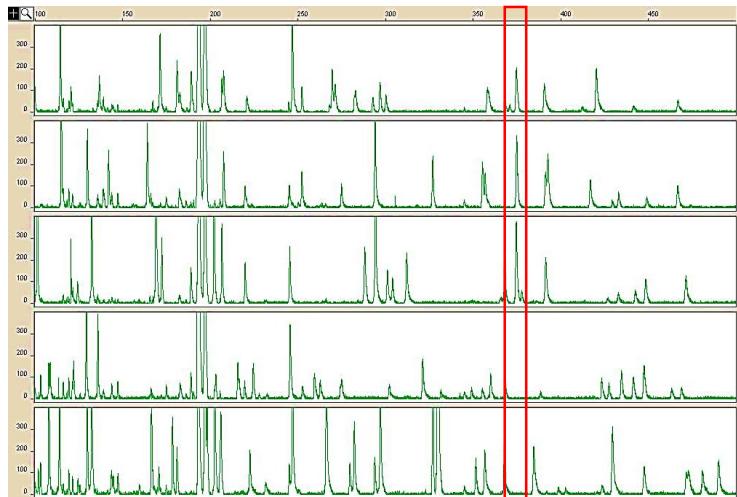
MOLEKULARNI BILJEZI



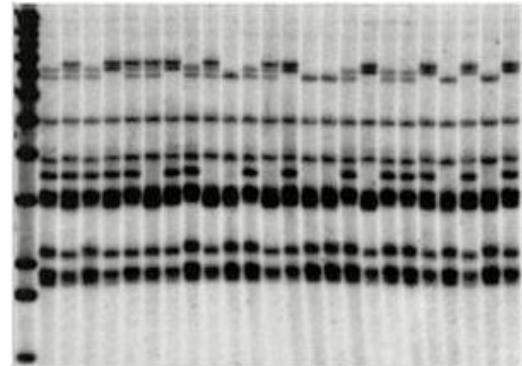
RAPD



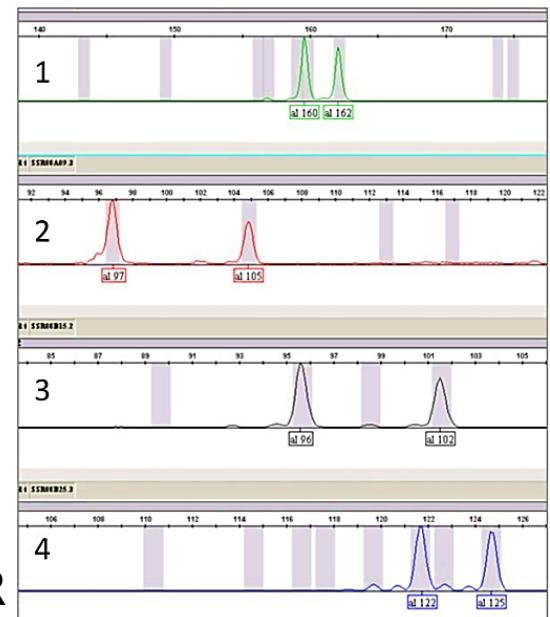
SNP



AFLP



RFLP



SSR

MIKROSATELITNI BILJEZI

- temelje se na umnažanju kratkih ulomaka DNA u kojima se svojstveni motiv (motif) sastavljen od 1 do 6 nukleotidnih baza ponavlja

$(AC)_7$

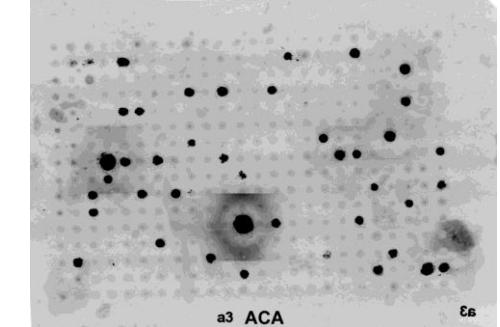
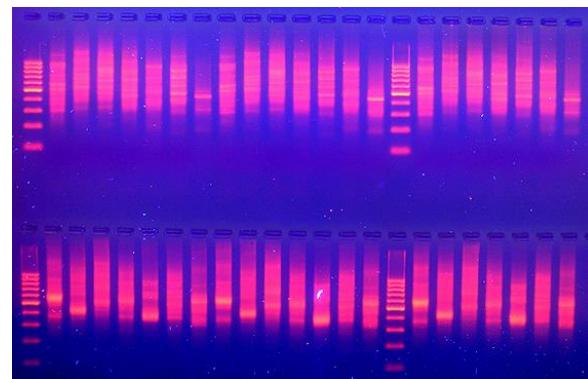
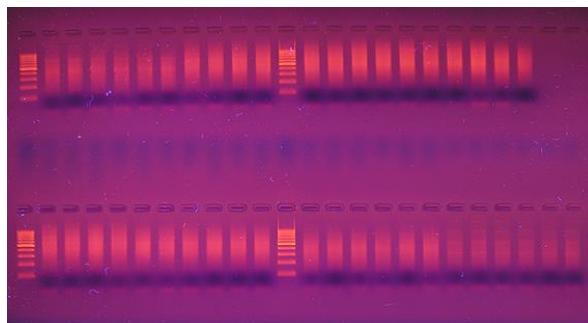
```
GAATCGAACCCACGACCTGCAC $\textcolor{red}{ACACACACACACAC}$ CCCTCTTCCATTATGC  
TAGAACCATTTATCTGTTATGTTCTTTCAATACTTATTAAAGTCGATTTT  
TTTATTTAGTGTATAAGCTGTTCAGCTTATTCGGCTTATTGATAAGCTA  
AATTTTGTTCTTATTTCTAAATACTAGATTAAATTCACACTTCTGTA  
TTTTTGGTAATTTACTATTCTACTATTATTAAATTAAATTCTGTT  
TTAATTAAAAAATTCTATAATAAAAATTAAATCAAAAAAATATAAACTTT  
TTTTAAAAATCTTATTAAATTGTAGTTCTGTGAAAATT
```

1. jednostavni (simple): $(AT)_n$
2. sastavljeni (compound): $(AT)_n(CA)_m$
3. prekinuti (interrupted): $(AT)_n-CCA-(AT)_m$

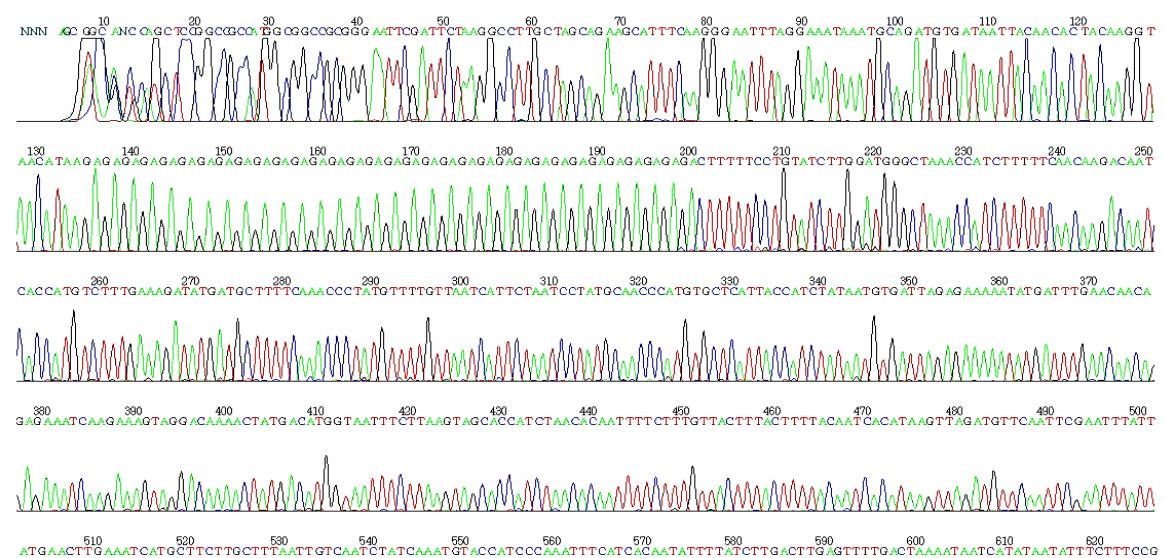
- česti mikrosatelitni motivi kod biljaka:
(GA), (GT), (AGA), (ACA), (ACC), (AGG)

RAZVOJ MIKROSATELITNIH BILJEGA

- (1) Izolacija DNA
 - (2) Fragmentiranje genomske DNA "koktelom" endonukleaza
 - (3) Hibridizacija
 - (4) PCR umnožavanje fragmenata
 - (5) Ligacija u vektor, selekcija
 - (6) Sekvenciranje



File: Zlatko_01-G20-T7.ab1 Run Ended: 2008/7/2 12:41:2 Signal G:703 A:857 C:548 T:704
Sample: Zlatko_01-G20_T7 Lane: 50 Base spacing: 14.42 959 bases in 11351 scans Page 1 of 1



SEKVENCIRANJE NOVE GENERACIJE



(1) Izolacija DNA (Qiagen Dneasy Plant Mini Kit)

(2) Sekvenciranje (Illumina NExtSeq sustav, UK)

- rezultat 6.619.805 očitanja visoke kvalitete (1,98 Gbp genoma)
- daljnje pročišćavanje podataka i sastavljanje 35.758 kontiga (N50=360 bp, Avg=348 bp, max=37.391 bp)

(3) Identifikacija SSR biljega (MicroSATellite identification tool)

- identificirano 1169 mikrosatelitnih sekvenci



DI-NUKLEOTIDNI MOTIVI
(904)

TRI-NUKLEOTIDNI MOTIVI
(170)

TETRA-NUKLEOTIDNI MOTIVI
(73)

OSTALI MOTIVI
(4)

HEKSA-NUKLEOTIDNI MOTIVI
(10)

PENTA-NUKLEOTIDNI MOTIVI
(8)

(4) Dizajn početnica (PRIMER3)

No	SSR Marker	Repeat Motif	Product Size	Left Primer	Right Primer	OriginalSeqID
1	TcUz001	(TC)8	127	GTTGTTTGCTGCGTGT	CGATGGAAGAAGAAGATGGTGG	Tan_OPT1_contig_487
2	TcUz002	(GA)6	127	ACGCAGACAATGACACGGTA	GGGCTGCGTTTGTATATT	Tan_OPT1_contig_4268
3	TcUz003	(AC)6	143	ACCAGAGAAAGTGTCAAAGTCA	TTTTGGCATGCAAGTTAGGC	Tan_OPT1_contig_1893
4	TcUz004	(AG)6	143	GAAAACGTTGAGATTTGTTCA	TCAACAAGCACCTTCTTAGCTT	Tan_OPT1_contig_1970
5	TcUz005	(CA)7	150	TGCTAGCCAACTATTACCCCT	TGTTTACGTGCATCATT CGC	Tan_OPT1_contig_75
6	TcUz006	(AT)8	151	CGGGGATTGCCTTCTCTTCT	AAAGCCTTCAGAGCCTCAGT	Tan_OPT1_contig_46
7	TcUz007	(AT)6	152	GCGTCTTCTCCTTGGCAAAG	GCCCAGCACTGCACAAATAT	Tan_OPT1_contig_57_2
8	TcUz008	(CA)7	152	TGAAAATGTTCCCCAATGC	AAAGTCCAAGGGCAGAGAC	Tan_OPT1_contig_9369
9	TcUz009	(AT)6	153	AAACCGTCTGCATCAATAAACAA	ACAAGTTCAAGATCAGGTTAGC	Tan_OPT1_contig_1612
10	TcUz010	(CA)7	160	TGGTCGTGTGTAGGGAGAG	TGGTCCCCTATTCCTGTATCC	Tan_OPT1_contig_1269
11	TcUz011	(GT)8	180	CAGAAAGTGAATAAAGCATGA	TGACCTCTACCCACCTCGAC	Tan_OPT1_contig_7243
12	TcUz012	(AT)7	188	TGACTTCATTTGGCGGAT	GCCCAGCACTGCACAAATAT	Tan_OPT1_contig_57_3
13	TcUz013	(AT)7	194	ATGGCCCATGATGATTAAAAA	TGAGTTGGTTTTGGGAAG	Tan_OPT1_contig_34036
14	TcUz014	(GT)6	195	CAAGGGCGCGAGTGTAAAG	TTTGGGCTTAATTCCTGCTT	Tan_OPT1_contig_1622
15	TcUz015	(AC)7	216	TTCAAAACATTCTCCACCACA	GGATTGATGAGATGGCATGAT	Tan_OPT1_contig_20748
16	TcUz016	(TA)11	217	GCACATGTATTTCTGTATCG	TCATAAGCGATATGAAGAGTGC	Tan_OPT1_contig_5862
17	TcUz017	(TA)44	220	AGTGTGGATCCATGACAAACA	TGGGTAAAATGCCGAACTGC	Tan_OPT1_contig_709
18	TcUz018	(TG)6	241	TGGTGGTTACAGCTAGTGTGTG	AAAAAGCGATCTTAGCCCTCT	Tan_OPT1_contig_30112
19	TcUz019	(GT)6	245	AAAGTCCAAGGGCGAGAT	CATGTTGATTGCTCTGTCTGA	Tan_OPT1_contig_8820
20	TcUz020	(TA)7	246	ACAAGAGCATTTATTCGGC	AGTTATCTTGTTCATGCCGA	Tan_OPT1_contig_991
21	TcUz021	(GTT)6	112	CCTGAGCATTCCAGTCCCA	GCTAGTTGTTATGAGTGC GGG	Tan_OPT1_contig_711
22	TcUz022	(TGG)6	114	GGAGGAGGTTCTTGAGCAGA	TTCCCTAAACTCCACCGG	Tan_OPT1_contig_547
23	TcUz023	(TTC)5	118	ACGAAGATACTGAACCA	TCTGTATGAGTGTGGAGAGA	Tan_OPT1_contig_649
24	TcUz024	(AAT)6	122	TGTTTCTTGTCTTGAGCCGG	AGGCTGGGATTACACTTCC	Tan_OPT1_contig_193
25	TcUz025	(GAG)7	127	TGAGATGGATGTAGAGGCCG	CTAAATGGCTCCGTCTCCT	Tan_OPT1_contig_634
26	TcUz026	(TTA)5	131	TCCGAAACAAC TCAACTACGA	TGATGTTGCA GTTGTAAAGGA	Tan_OPT1_contig_177
27	TcUz027	(CTC)5	133	GTAGGTACCACAGGAGAGTCG	ATGGTGATGATGAGGCCCTCC	Tan_OPT1_contig_210
28	TcUz028	(GTT)7	141	CTCCGCACTAAAACAACCA	GTCCTTGCTACCTAAGTGC	Tan_OPT1_contig_528
29	TcUz029	(ACA)7	147	GAAACCAAGCTGCTAACCC	AATGACCCCTGAGCTCCACAC	Tan_OPT1_contig_441
30	TcUz030	(GGT)5	157	ACCTTACGGCTGCTTGAGA	GTGGCAGCAGACCGAAC	Tan_OPT1_contig_34
31	TcUz031	(CAA)6	180	TCAACATGGCCCGTGAAC TA	CCTGCCTCCTGTTTGTGT	Tan_OPT1_contig_1557
32	TcUz032	(ATG)5	187	CGAAGCCGTCCTATCGTCTA	ACTCACTTGGGTTGGGACA	Tan_OPT1_contig_1909
33	TcUz033	(AAT)5	195	GGTGTAGCACGAAGCTTCC	CGACAATTGGGTATCGTT	Tan_OPT1_contig_2158
34	TcUz034	(GTT)5	202	CCTGAGCATTCCAGTCCCA	TGCAACAAGATTGCCACTT	Tan_OPT1_contig_712
35	TcUz035	(TTC)5	208	GCCTCAACTACCTTTTGATT	GGTGAGTTGAGCTTCTGG	Tan_OPT1_contig_11937
36	TcUz036	(GGT)5	208	GGACCAAAAGGGATATGGTG	CCCTTTTACCCACACACAC	Tan_OPT1_contig_17371
37	TcUz037	(ATG)5	209	CGAAGTCCGCGTTAGTGA	GCCCTACGATCCGAACTAA	Tan_OPT1_contig_2666
38	TcUz038	(ACC)5	230	TTTCGCTCTCATCAGTTCA	ACCACGTACGCCACCAAAA	Tan_OPT1_contig_24814
39	TcUz039	(GAT)5	230	GCGAGTTTGTTCGCGATAG	TTTCGATTGGATCAATCAC	Tan_OPT1_contig_31285
40	TcUz040	(GAA)6	247	TGGATCCGTCAAAATCGATCT	CCTGATCCCCGTACATATAATCA	Tan_OPT1_contig_39

(5) Testiranje novorazvijenih biljega na polimorfizam

(6) Selekcija najprikladnijih biljega za korištenje u analizi populacijske genetike dalmatinskog buhača



Istraživanje provedeno u sklopu projekta:

Genetska osnova insekticidnog potencijala dalmatinskog buhača (*Tanacetum cinerariifolium* /Trevir./ Sch. Bip.)



Projekt je u cjelovitosti financiran od strane Hrvatske zaklade za znanost

