

Disentangling arthropod and plant resources consumed by *Orius* spp. in peach and alfalfa crops by metagenomic analysis

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Table S1. Arthropod (ZBJ-ArtF1c/ZBJ-ArtR2c and mICOLintF/HC02198) and plant (ITS-S2F/ITS4R and cA49325/trnL110R) primer pairs used in this study. Also indicated the sequence of each forward (F) and reverse (R) primer and the length of the amplified fragment.

	Sequence 5'- 3'(F)	Sequence 5'- 3'(R)	Reference	Fragment (bp)
ZBJ-ArtF1c/ZBJ-ArtR2c	AGATATTGGAACWTTATATTTATTTTGG	WACTAATCAATTWCCAAATCCTCC	Zeale et al. 2011	157
mICOLintF/HC02198	GGWACWGGWTGAACWGTWTAYCCYCC	TAAACTTCAGGGTGACCAAAAAATCA	Leray et al. 2013/Folmer et al. 1994	313
ITS-S2F/ITS4R	ATGCGATACTTGGTGTGAAT	TCCTCCGCTTATTGATATGC	Chen et al. 2010/White et al. 1990	350
cA49325/trnL110R	CGAAATCGGTAGACGCTACG	GATTTGGCTCAGGATTGCC	Taberlet et al. 2007/Borsch et al. 2003	80

Table S2. Accession numbers of the sequences from the databases used as reference to obtain the percentages of similarity.

Accession number	species	Databases used
MG007856.1	<i>Orius laevigatus</i>	GenBank
MG007857.1		
KM021482.1	<i>Orius niger</i>	
KM021745.1		
KM021890.1		
KM022197.1		
KM022760.1		
KM022882.1		
KM022999.1		
SONE138-11	<i>Orius majusculus</i>	Bold
SONE141-11		

Table S3. Percentages of relative read abundance (RRA%) of the reads amplified from resources consumed by each arthropod and plant primer pairs in each library (L) in the trials included in the study: **(1)** Arthropod taxa detected in the analyzed *Orius*; **(2)** Plant taxa detected in the analyzed *Orius*. Those percentages eliminated from the analysis for exceeding the 1% threshold are shown in bold and grey. Art1= ZBJ-ArtF1c/ZBJ-ArtR2c; Art2= mICOLintF/HC02198; PI1= ITS-S2F/ITS4R; PI2= cA49325/trnL110R; NA= Not amplified.

(1)

Primers used	L1		L3		L5		L7		L9		L11	
Taxa detected	Art1	Art2	Art1	Art2	Art1	Art2	Art1	Art2	Art1	Art2	Art1	Art2
<i>Orius</i>	-	-	-	NA	-	-	-	-	-	NA	-	-
<i>Orius laevigatus</i>	-	-	100	NA	8,88	-	-	-	-	NA	87,23	-
<i>Orius majusculus</i>	-	-	-	NA	-	-	-	-	-	NA	-	-
<i>Orius niger</i>	-	-	-	NA	-	-	-	-	-	NA	-	-
<i>Aeolothrips intermedius</i>	-	-	-	NA	-	-	73,78	16,72	-	NA	0,21	-
<i>Frankliniella occidentalis</i>	-	100	-	NA	-	100	-	74,62	-	NA	-	73,11
Aphididae	100	-	-	NA	-	-	-	-	-	NA	-	-
<i>Therioaphis trifolii</i>	-	-	-	NA	-	-	-	-	-	NA	0,29	10,27
<i>Hypera</i>	-	-	-	NA	5,33	-	2,32	-	-	NA	-	-
<i>Liriomyza</i>	-	-	-	NA	3,55	-	21,58	8,66	-	NA	-	-
<i>Aedes caspius</i>	-	-	-	NA	-	-	-	-	-	NA	0,38	8,07
Ceratopogonidae	-	-	-	NA	-	-	-	-	-	NA	0,21	4,89
<i>Tanytarsus volgensis</i>	-	-	-	NA	68,64	-	-	-	-	NA	-	-
<i>Orthocladinae</i>	-	-	-	NA	13,61	-	2,32	-	-	NA	-	-
<i>Cicadellidae</i>	-	-	-	NA	-	-	-	-	100	NA	11,68	-

(2)

Primers used	L2		L4		L6		L8		L10		L12	
Taxa detected	PI1	PI2	PI1	PI2	PI1	PI2	PI1	PI2	PI1	PI2	PI1	PI2
Streptophyta	-	-	NA	-	-	-	NA	-	-	2,17	100	31,69
Asparagales	-	-	NA	-	-	7,07	NA	-	-	-	-	-
Caryophyllales	-	-	NA	-	-	-	NA	-	-	-	-	8,69
Asteraceae	-	-	NA	12,95	-	2,18	NA	-	-	42,04	-	-
Poaceae	-	-	NA	-	-	1,25	NA	-	-	-	-	-
Solanaceae	-	2,67	NA	-	-	-	NA	-	-	-	-	-
<i>Poa annua</i>	-	-	NA	-	-	-	NA	-	-	1,78	-	-
<i>Pinus</i>	-	9,45	NA	15,83	-	2,18	NA	-	-	-	-	1,36
Fabaceae	-	40,86	NA	41,73	-	87,32	NA	100	-	42,93	-	58,26
Rosaceae	-	47,02	NA	29,50	-	-	NA	-	-	11,08	-	-
<i>Medicago sativa</i>	-	-	NA	-	100	-	NA	-	59,65	-	-	-
<i>Prunus persica</i>	100	-	NA	-	-	-	NA	-	40,35	-	-	-

Table S4. Taxa obtained from each library and with each primer pair. NT= not tested. NAMP = Not amplified.

species/sample	Crop	Date	# of individuals/ Sample size	Library #	Arthropod primer pairs		Plant primer pairs	
					ZBJ-ArtF1c/ZBJ-ArtR2c	mICOLintF/HC02198	ITS-S2F/ITS4R	CA49325/trnL110R
<i>Orius laevigatus</i>	Peach	June-2016	7	L1	<i>Orius laevigatus</i> Aphididae	<i>Orius</i> <i>Orius laevigatus</i> <i>Frankliniella occidentalis</i>	NT	NT
				L2	NT	NT	<i>Prunus persica</i>	Solanaceae <i>Pinus</i> Rosaceae Fabaceae
<i>Orius majusculus</i>			21	L3	<i>Orius laevigatus</i> <i>Orius majusculus</i>	<i>Orius</i> <i>Orius majusculus</i>	NT	NT
				L4	NT	NT	NAMP	Asteraceae <i>Pinus</i> Fabaceae Rosaceae
<i>Orius niger</i>	Alfalfa	June-2016	9	L5	<i>Hypera</i> <i>Tanytarsus volgensis</i> Orthocladinae <i>Liriomyza</i> <i>Orius laevigatus</i> <i>Orius niger</i>	<i>Orius niger</i> <i>Frankliniella occidentalis</i>	NT	NT
				L6	NT	NT	Streptophyta	Streptophyta Caryophyllales Fabaceae <i>Pinus</i>
		June-2016	25	L7	<i>Hypera</i> Orthocladinae <i>Liriomyza</i> <i>Orius niger</i> <i>Aeolothrips intermedius</i>	<i>Liriomyza</i> <i>Orius niger</i> <i>Frankliniella occidentalis</i> <i>Aeolothrips intermedius</i>	NT	NT

				L8	NT	NT	<i>Medicago sativa</i>	Asparagales Asteraceae Poaceae <i>Pinus</i> Fabaceae
		July-2017	13	L9	<i>Orius</i> <i>Orius niger</i> Cicadellidae	<i>Orius</i> <i>Orius niger</i>	NT	NT
				L10	NT	NT	NAmp	Fabaceae
		September-2017	22	L11	<i>Orius</i> <i>Orius laevigatus</i> <i>Orius niger</i> Cicadellidae	<i>Aedes caspius</i> Ceratopogonidae <i>Orius</i> <i>Orius niger</i> <i>Therioaphis trifolii</i> Cicadellidae <i>Frankliniella occidentalis</i>	NT	NT
				L12	NT	NT	<i>Medicago sativa</i> <i>Prunus persica</i>	Streptophyta Asteraceae Fabaceae Rosaceae <i>Poa annua</i>
<i>Prunus persica</i>	Peach		1 cm ²	L13	NT	NT	<i>Prunus persica</i>	Rosaceae
<i>Medicago sativa</i>	Alfalfa		1 cm ²	L14	NT	NT	<i>Medicago sativa</i>	Fabaceae

Table S5. Percentages of similarity means by pairs from the region amplified by each arthropod primer pair used calculated from the sequences obtained in the databases of three amplified *Orius* species.

Species comparison	ZBJ-ArtF1c/ZBJ-ArtR2c (%)	mICOLintF/HC02198 (%)
<i>O. majusculus</i> - <i>O. laevigatus</i>	92,04	87,29
<i>O. majusculus</i> - <i>O. niger</i>	92,42	90,85
<i>O. laevigatus</i> - <i>O. niger</i>	94,34	91,53

Table S6. Mean of the calculated percentages of similarity between the obtained number of OTUs for the predator (*O. majusculus* or *O. niger*) and for the prey (other *Orius* species) using the ZBJ-ArtF1c/ZBJ-ArtR2c pair of primers.

Predator species	Prey species	Library	# Predator OTUs	# Prey OTUs	Mean % similarity between OTUs
<i>O. majusculus</i>	<i>O. laevigatus</i>	L3	5	5	91,90
<i>O. niger</i>	<i>O. laevigatus</i>	L5	6	1	92,73
<i>O. niger</i>	<i>O. laevigatus</i>	L11	2	1	93,94

Table S7. Number of reads and OTUs obtained per library regarding the amplified *Orius* species, and the taxon finally assigned with each pair of arthropod primer used. The species with the highest number of reads and OTUs within a library and pair of primer used is indicated in grey.

Predator	Library	Primer pair	<i>Orius</i> species		
			# Reads	# OTUs	Taxon assigned
<i>Orius laevigatus</i>	L1	ZBJ-ArtF1c/ZBJ-ArtR2c	56868	23	<i>O. laevigatus</i>
		mICOLintF/HC02198	2015	3	<i>O. laevigatus</i>
<i>Orius majusculus</i>	L3	ZBJ-ArtF1c/ZBJ-ArtR2c	24906	5	<i>O. majusculus</i>
			6668	5	<i>O. laevigatus</i>
		mICOLintF/HC02198	1635	2	<i>O. majusculus</i>
<i>Orius niger</i>	L5	ZBJ-ArtF1c/ZBJ-ArtR2c	109935	6	<i>O. niger</i>
			15	1	<i>O. laevigatus</i>
		mICOLintF/HC02198	10679	2	<i>O. niger</i>
	L7	ZBJ-ArtF1c/ZBJ-ArtR2c	79012	3	<i>O. niger</i>
		mICOLintF/HC02198	21255	2	<i>O. niger</i>
	L9	ZBJ-ArtF1c/ZBJ-ArtR2c	25677	1	<i>O. niger</i>
		mICOLintF/HC02198	19202	1	<i>O. niger</i>
	L11	ZBJ-ArtF1c/ZBJ-ArtR2c	38716	2	<i>O. niger</i>
			3668	1	<i>O. laevigatus</i>
		mICOLintF/HC02198	23446	38	<i>O. niger</i>
		Total	423697	95	

Table S8. Number of template-mismatches of each arthropod pair of primers with each amplified taxon. NS = no sequence present in the databases.

Taxon	mICOLintT/HC02198		ZBJ-ArtF1c/ ZBJ-ArtR2c	
	Forward (mICOLintT)	Reverse (HC02198)	Forward (ZBJ-ArtF1c)	Reverse (ZBJ-ArtR2c)
<i>Hypera</i>	1	NS	0	2
<i>Liriomyza</i>	1	3	1	1
<i>Tanytarsus volgensis</i>	1	NS	3	8
Orthocladinae	0	NS	0	6
Ceratopogonidae	1	NS	3	4
<i>Aedes caspius</i>	1	1	1	5
<i>Orius</i>	2	4	3	3
<i>Orius laevigatus</i>	0	NS	1	0
<i>Orius majusculus</i>	1	NS	1	0
<i>Orius niger</i>	0	NS	0	2
Aphididae	1	NS	3	2
<i>Therioaphis trifolii</i>	2	NS	3	8
Cicadellidae	1	NS	4	2
<i>Aeolothrips intermedius</i>	0	NS	3	6
<i>Frankliniella occidentalis</i>	1	NS	4	5

Figure S1. Percentage of predator and prey taxa amplified by each arthropod primer pairs from each library: **(A)** Taxa amplified by ZBJ-ArtF1c/ZBJ-ArtR2c primer pairs; **(B)** Taxa amplified by mICOLintF/HC02198 primer pairs. Predator (green); Prey (orange).

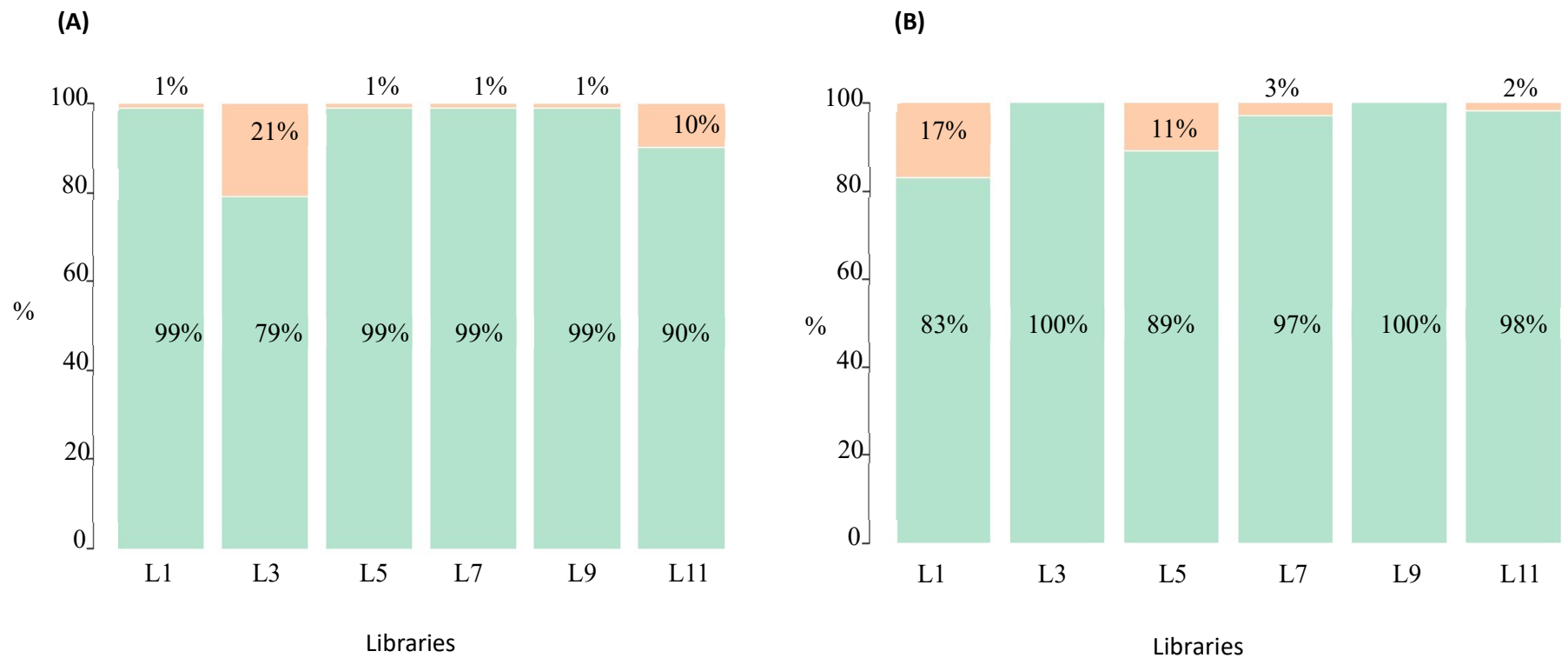
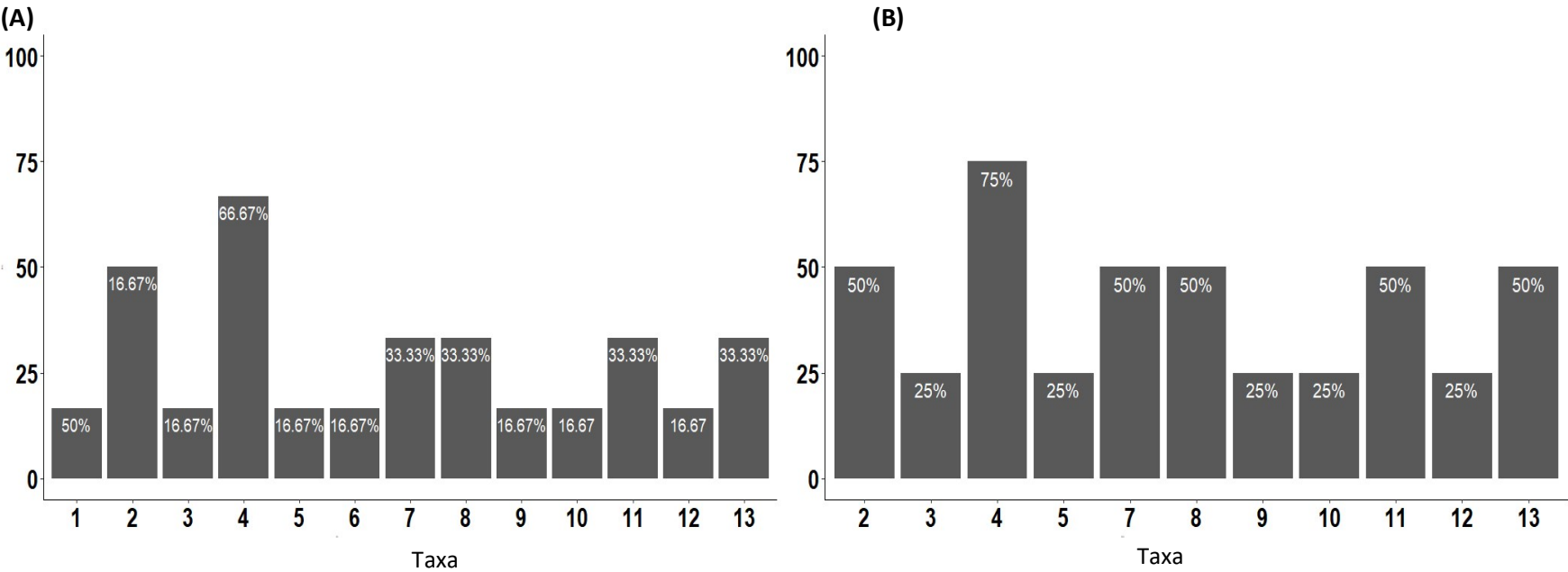


Figure S2. Percentage of frequency of occurrence (FOO%) of the **(A)** arthropod taxa resources items consumed by all analysed *Orius*; **(B)** arthropod taxa resources items consumed by *Orius niger*; **(C)** plant taxa resources items consumed by all *Orius*; **(D)** of the plant taxa resources items consumed by *Orius niger*.



- 1-*Orius*

2-*Orius laevigatus*

3-*Aeolothrips intermedius*

4-*Frankliniella occidentalis*
- 5-Aphididae

6- *Therioaphis trifolii*

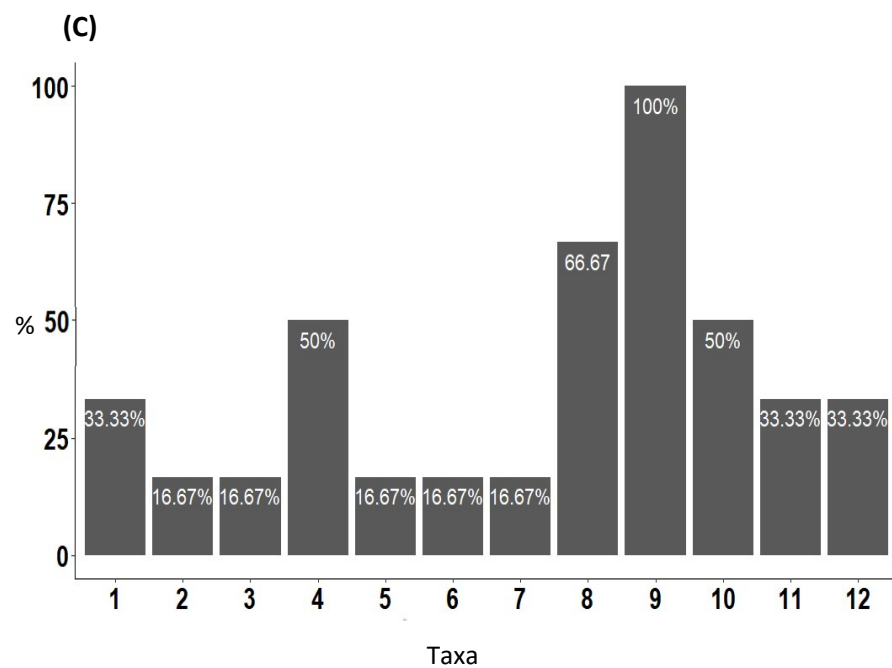
7-*Hypera*

8-*Liriomyza*
- 9- *Aedes caspius*

10- *Tanytarsus volgensis*

11- Orthocladiinae

12- *Ceratopogonidae*
- 13-Cicadellidae



1-Streptophyta

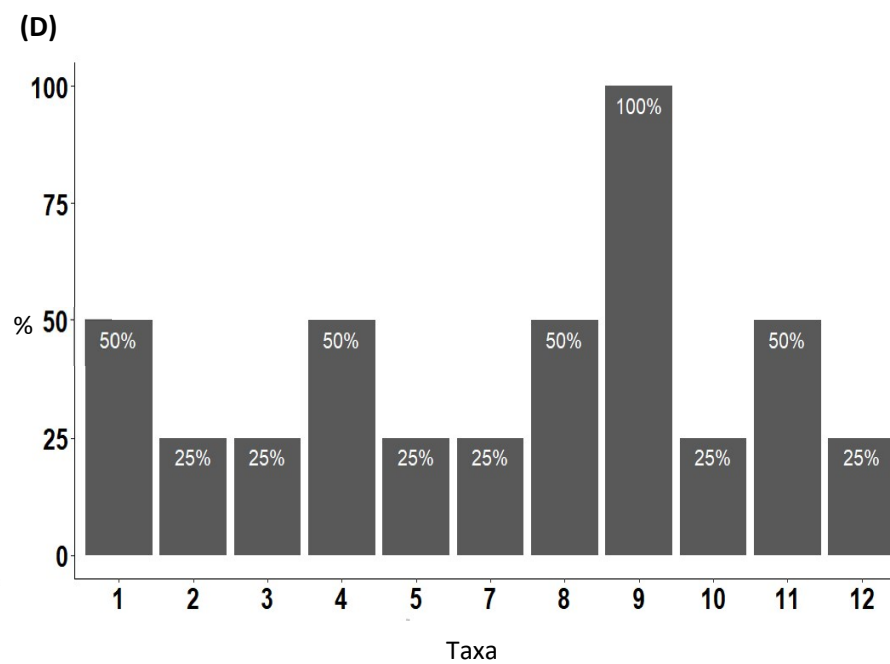
2-Asparagales

3-Caryophyllales

4-Asteraceae

5-Poaceae

6-Solanaceae



7-*Poa annua*

8-*Pinus*

9-Fabaceae

10-Rosaceae

11-*Medicago sativa*

12-*Prunus persica*