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# Report of a *Lactuca altaica* Fisch. & C.A. Mey and *L. serriola* L. collecting expedition in Uzbekistan

Itinerary, collected material and data

Kik C, A Makhmudov & FO Khassanov



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Centre for Genetic Resources, the Netherlands (CGN), Wageningen University & Research  
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CGN is ISO 9001 certified.

Picture front page: *Lactuca altaica* in a farmer's field (KMK 58).

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# Foreword

The mission of the Centre for Genetic Resources, the Netherlands (CGN) is to contribute to the conservation, development and sustainable use of plant, animal and forest genetic resources, and hence to global food security, a more sustainable production, rural development, and the conservation of cultural heritage.

To that end, CGN currently holds collections of over 20 crops and a total number of accessions of more than 22,500 of interest to breeders, researchers and other users. Annually around 5000 seed samples are distributed.

To contribute to an effective global system of *ex situ* collections, for each of its collections CGN has analysed the coverage of the crop genepool by the germplasm in its own collection and those of others. In a number of cases, CGN has been able to identify gaps in the total set of collections of a specific crop. Some genetic diversity that is known or can be assumed to exist, appeared poorly represented or even absent from the genebank collections. Such cases warrant new collecting missions, if we wish to conserve as wide a diversity for the crop genepool as possible.

The species *Lactuca altaica* forms such a case. Therefore in 2019 CGN carried out a collecting mission in Uzbekistan, in close collaboration with its local counterparts. The present report provides details of the results of this collecting mission. Seven plant breeding companies co-financed the mission, a fact that is duly recognized and appreciated.

During the mission in total 2493 seed samples were collected from 21 populations of *L. altaica*, 28 populations of *L. serriola* and 13 mixed populations. Upon regeneration, the samples will be made available under the terms and conditions of the Standard Material Transfer Agreement.

This collecting mission formed an activity jointly undertaken by partners in Uzbekistan and the Netherlands. The support from the national authorities in Uzbekistan is duly recognized.

# 1 Introduction

In the global plant genetic database GENESYS ([www.genesys-pgr.org](http://www.genesys-pgr.org)), 16519 *Lactuca* accessions are present amongst which 12,980 *Lactuca sativa* L. and 3539 crop wild relatives (CWR). *Lactuca altaica* Fisch. & C.A. Mey. is present in this database with only 34 accessions, of which 10 originate from Uzbekistan. *Lactuca serriola* L. is ubiquitous present in GENESYS with 2141 accessions of which 61 originate from Uzbekistan. In total CGN maintains 5 and 803 accessions of *L. altaica* and *L. serriola* respectively and from Uzbekistan CGN holds 1 and 53 accessions of *L. altaica* and *L. serriola* respectively.

According to Web of Science *L. serriola* has often been a subject of study (n= 531). However this is not the case for *L. altaica* as only 12 records were found. Possibly this is due to the fact that many taxonomists view *L. altaica* as a synonym of *L. serriola* (Cichoriae Portal; [www.cichoriae.e-taxonomy.net/portal](http://www.cichoriae.e-taxonomy.net/portal)) and therefore report findings for this species under the species name of *L. serriola*. Both species belong to the primary genepool of lettuce (Zohary 1991) and *L. serriola* has been (and is) used to a large extent in the breeding of new lettuce cultivars (e.g. Lebeda et al 2014). Furthermore it is known that both species can be intercrossed successfully (Lindquist 1960).

The distribution area of both species is quite different, whereas *L. serriola* is a cosmopolitan species, *L. altaica* can be found in the region in between East Anatolia to West China (Lebeda et al 2004; Fig.1).



**Figure 1** Probable distribution area of *Lactuca altaica* (red circle).

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Given the fact that only a few accessions of *Lactuca altaica* are present in genebanks worldwide and that the species could harbour agronomically interesting traits, a collecting mission to Uzbekistan, a country that is located within the centre of biodiversity of the species, is clearly warranted.

Therefore in the context of an already existing joint plant genetic resources cooperation between the Uzbek Academy of Sciences (Uzbekistan) and CGN (the Netherlands), a collecting project was set-up to sample *Lactuca altaica*. For *L. serriola* the need to collect in Uzbekistan is less compared to *L. altaica*. However, large areas in this country were not yet explored as previous collecting only took place in the eastern part of the country. Therefore, collecting in this expedition took place primarily in the central and southern part of the country.

In order to make the collecting mission possible, a Memorandum of Understanding (MoU) was signed using the Standard Material Transfer Agreement (SMTA) for the exchange of the material collected. The MoU was signed in January 2019 and the SMTA in August 2019 between the appropriate authorities of both countries and these document formed the legal basis of the expedition (Appendix 1).

## 2 Objectives of the expedition

The major aim of this single crop expedition is to broaden the *Lactuca altaica* and *L. serriola* collections maintained at CGN by collecting wild populations which can subsequently be used after regeneration for breeding and research purposes, which consequently contributes to the international need for the conservation of plant genetic resources (PGR).

## 3 Members of the collecting team

- Prof Dr Furkat Khasanov and Dr Aziz Makhmudov, Institute of Botany, Academy Sciences of Uzbekistan, 100125, Durmon yuli str., 32 Tashkent, Uzbekistan; E-mail: fkhasanov1@mail.ru
- Dr Chris Kik, Centre for Genetic Resources, the Netherlands (CGN), Droevedaalsesteeg 1, 6708 PB Wageningen, the Netherlands; E-mail: chris.kik@wur.nl

## 4 Exploration area and expedition period

The exploration area was situated primarily in the central and southern part of Uzbekistan (Fig. 2). The area covered during the mission was ca. 550 km from North to South and ca. 400 km from East to West. In total around 4000 km was travelled during a period of four weeks (July 18 – August 15).



**Figure 2** The exploration area in Uzbekistan with the locations of the various *Lactuca* accessions indicated; green: *L. serriola*, yellow: *L. altaica* and blue: *Lactuca* populations of mixed or unclear identity (see for details Appendix 3).

Temperatures during daytime in Uzbekistan were in between 40 and 50 °C. For transport a Chevrolet Nexia LTZ was used. Overnight stays were in different places: B&Bs, farms, houses and outside on higher elevations.

## 5 Data collecting, sampling procedure and seed cleaning

A field collecting form based upon a modified multi-crop passport descriptor list (MCPD; see: <http://eurisco.ipk-gatersleben.de>) was used to document the passport data of the accessions sampled (Appendix 3). All sampled material

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received a so-called collecting number, in this case KMKxxx. Latitude, longitude and altitude were determined via GPS (Garmin, eTrex 20) with an inaccuracy of 1-5 meters. Latitude and longitude were recorded using as map datum WGS84 and as position format hddd°ddd. Pictures were taken of the collecting site. Passport data can be found in Appendix 4.

Sampling of all *Lactuca* species took place on individual plants and the seeds of a minimum of 2-3 flowering heads were collected per individual plant. If not more than 2-3 flowering heads could be collected these seeds were transferred to a single glassine bag. If a minimum of 4-6 flowering heads with ripened seeds were present per individual, seeds were collected in two glassine bags (see Appendix 4). These two glassine bags were stapled to each other. Subsequently the single and double bags per accessions were put together in one cotton bag (18 x 30 cm<sup>2</sup>) which was tied together with a string. A plasticized paper tag with the collecting number was placed on the inside and on outside with a string which tied together the cotton bag. The glassine bags were individually marked during the expedition to be able later on to trace the origin of the seeds within a population. This was done by writing on the glassine bags the accession number, the number of the individual and an 'a' or a 'b' in case the seed of an individual was divided over two bags (example: KMK32-01-a and KMK32-01-b in case of two seed batches from plant 1 of population KMK032). During the expedition the cotton bags (with seeds) were kept under ambient room conditions. Cleaning of the seeds during the expedition was not necessary as the seeds were already reasonably clean when harvested. Upon arrival at CGN the seeds were transferred to a conditioned storage room with a temperature of 15 °C and 15% relative humidity and a receipt number (RNR) was assigned to each accession.

## 6 Results and discussion

Seeds were collected from 2493 plants from a total of 62 populations. Probably 21 populations consisted of *Lactuca altaica*, 28 of *Lactuca serriola* and 13 mixed or unclear identity (Appendix 4).

### 6.1 *Lactuca altaica* and *L. serriola* populations

*Lactuca altaica* could be recognized by its plant height, flower diameter and flowerhead size all of which are larger than in *L. serriola*. Also its leaf apex is more acute compared to *L. serriola* and its synflorescence more slender compared to the paniculiform synflorescence of *L. serriola*. However, when collecting in the field it is not always obvious if one or two species are present in a population. More detailed species identification will be carried out during the regeneration of the seed samples.

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Both species occur in disturbed habitats: road sides, fields & field margins, backyards and orchards (Photo 1a, b).



**Photo 1** *Lactuca* collecting sites: a. *Lactuca altaica* in orchard (KMK62), b. *Lactuca serriola* along a road (KMK06).

The mean elevation of the locations where the two *Lactuca* species occurred clearly differed from each other [ $F(1,47)=79.5^{***}$ ]. However, overlap of the altitudes of the collecting sites of the two species occurs (Table 1). Given the

clear difference in mean altitude of the collecting sites between the two species, it could be expected that *L. altaica* prefers a more cool and moist habitat compared to *L. serriola*. However *L. serriola* can also occur on higher elevations but also in steppe habitats, a habitat in which *L. altaica* was not observed throughout the expedition. When an alpine meadow on ca. 2500 m was visited during the expedition no *L. altaica* plants were observed. This may suggest that the maximum altitude where the species can survive is around ca. 2000 m.

**Table 1** The number of collected *Lactuca* populations, the total number of plants sampled per species, and the mean, its standard error and range of the altitude where the species were observed.

species	N populations	N plants	altitude	
			mean $\pm$ SE	range
<i>L. altaica</i>	21	793	1175 $\pm$ 54	731-1676
<i>L. serriola</i>	28	1210	591 $\pm$ 39	248-1066

The estimated number of plants occurring in populations varied from less than 100 to over 1000. Most populations consisted of 100-1000 plants. More *L. serriola* than *L. altaica* populations were present where over 1000 plants occurred (Table 2).

**Table 2** The number of *Lactuca altaica* and *L. serriola* populations per estimated size class (number of plants in a population).

category	< 100	100-1000	>1000
<i>L. altaica</i>	1	13	7
<i>L. serriola</i>	0	15	13

The total number of sampled plants was 2493 of which 2003 were sampled from *L. serriola* (n= 1140) and *L. altaica* (n= 863). The ranges, means and their standard errors of the number of plants, the number of single bags and the number of double bags per *L. altaica* and *L. serriola* population from which seed was sampled is presented in Table 3.

From Table 3 it can be concluded that both species did not differ for the mean number of plants from which seed was collected, and the mean number of single and double bags. Furthermore, from the plants from which seed was harvested only half of them had enough seed for two bags.

**Table 3** The ranges, means and their standard errors of the a. number of plants from which seed was harvested, b. sampled single glassine bags and c. sampled double glassine bags from *L. altaica* and *L. serriola* populations.

	<i>L. altaica</i>		<i>L. serriola</i>	
	range	mean $\pm$ SE	range	mean $\pm$ SE
number of plants	7-70	41 $\pm$ 3	16-69	41 $\pm$ 2
single bags	2-40	18 $\pm$ 3	2-65	21 $\pm$ 3
double bags	0-50	23 $\pm$ 4	0-57	19 $\pm$ 3

## 6.2 Mixed populations

Thirteen populations were sampled in which *L. altaica* and *L. serriola* most probably co-occurred. However this observation was not based on a detailed investigation of the sampled plants, but rather on a general impression when collecting seeds. Therefore the total number of such populations could be different after careful examination. This of course is also true for the populations that were classified as *L. altaica* or *L. serriola*. More detailed examination during regeneration is expected to reveal the correct taxonomic status of the populations.

The mixed populations, like the single species populations, were also observed in ruderal habitats like road sides, fields & field margins, backyards and orchards (Photo 2).



**Photo 2** *Lactuca* collecting sites: mixed population in vineyard (KMK30).

The altitude where the mixed populations were found was in between those of *L. altaica* and *L. serriola* and differed significantly from these species [ $F(2,61)=37.5^{***}$ ]. However the range in elevation of the mixed populations overlapped with those of the two *Lactuca* species (Table 4).

**Table 4** The number of collected mixed *Lactuca* populations, the total number of plants sampled and the mean, its standard error and the range of the altitude where the mixed populations were observed.

species	N populations	N plants	altitude	
			mean $\pm$ SE	range
mixed	13	490	918 $\pm$ 74	604-1469
<i>L. altaica</i>	21	793	1175 $\pm$ 54	731-1676
<i>L. serriola</i>	28	1210	591 $\pm$ 39	248-1066

The estimated number of plants occurring in mixed populations varied from less than 100 to over 1000. Most populations consisted of 100-1000 plants, which was also true for the single-species populations (Table 5).

**Table 5** The number of mixed populations per estimated size class (number of plants in a population). From a single mixed population (KMK46) the number of plants was not recorded.

category	< 100	100-1000	>1000
mixed	3	6	3
<i>L. altaica</i>	1	13	7
<i>L. serriola</i>	0	15	13

From Table 6 it can be concluded that the mixed populations did not differ from the *L. altaica* and *L. serriola* populations with respect to the number of plants, single and double bags.

**Table 6** The ranges, means and their standard errors of the a. number of plants from which seed was harvested, b. sampled single glassine bags and c. sampled double glassine bags from mixed *Lactuca* populations.

	mix		<i>L. altaica</i>		<i>L. serriola</i>	
	range	mean $\pm$ SE	range	mean $\pm$ SE	range	mean $\pm$ SE
number of plants	2-68	38 $\pm$ 5	7-70	41 $\pm$ 3	16-69	41 $\pm$ 2
single bags	2-44	20 $\pm$ 4	2-40	18 $\pm$ 3	2-65	21 $\pm$ 3
double bags	0-45	17 $\pm$ 4	0-50	23 $\pm$ 4	0-57	19 $\pm$ 3

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## 7 Conclusions

- a. This collecting mission was carried out in the context of a joint PGR project between Institute of Botany, Academy Sciences of Uzbekistan (Uzbekistan) and the Centre for Genetic Resources, the Netherlands (CGN), Wageningen University & Research (the Netherlands).
- b. A SMTA was signed between the competent national authorities of Uzbekistan and the Netherlands. This formed the legal basis of the expedition.
- c. From sixty-two *Lactuca* populations seeds were collected during the mission which can be subdivided in 21 *Lactuca altaica*, 28 *Lactuca serriola* and 13 populations of mixed or unclear identity.
- d. Seed samples were taken from individual plants; from every plant a minimum of two flowerheads were sampled.
- e. In total from 2493 plants seeds were harvested in glassine bags; from *L. altaica* 793 plants were sampled, from *L. serriola* 1210 and from mixed populations 490.
- f. On average seeds were harvested from ca. 40 plants per population.
- g. A clear difference in altitude between the collecting sites was found for the *Lactuca* species: *L. altaica* was found on higher elevations, *L. serriola* on lower elevations and the mixed populations in between. However also overlap in altitude between the three groups was observed. This difference in altitude occurrence might point at a habitat preference of *L. altaica* for more moist and cool conditions in contrast to *L. serriola*.

## 8 References

Lebeda A, Dolezalova I, Ferakova V et al (2004). Geographical distribution of wild *Lactuca* species (Asteraceae, Lactuceae). *Bot Rev* 70(3): 328-356

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Lindqvist K (1960). Cytogenetic studies in the serriola group of *Lactuca*. *Hereditas* 46 (1-2): 75-151

Zohary D., 1991. The wild genetic resources of cultivated lettuce (*Lactuca sativa* L.). *Euphytica* 53, 31-35

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## 9 Acknowledgements

This expedition was made possible by the financial support from breeding companies affiliated to PLANTUM-NL and the Dutch Ministry of Agriculture, Nature and Food Quality (LVN). The support and hospitality of local people in Uzbekistan was of great value. Furthermore CK would like to express his sincere gratitude to the Furkat Khassanov and Aziz Makhudov who made this collecting mission a success.

# Appendix 1 SMTA (art 10 and annex I)

## ARTICLE 10 — SIGNATURE/ACCEPTANCE

The **Provider** and the **Recipient** may choose the method of acceptance unless either party requires this **Agreement** to be signed.

### Option 1 –Signature\*

I, (*Full Name of Authorized Official*), represent and warrant that I have the authority to execute this **Agreement** on behalf of the **Provider** and acknowledge my institution's responsibility and obligation to abide by the provisions of this **Agreement**, both by letter and in principle, in order to promote the conservation and sustainable use of **Plant Genetic Resources for Food and Agriculture**.

Signature.....  
Name of the **Provider**.....

Date.....13.08.2019.....

I, (*Full Name of Authorized Official*), represent and warrant that I have the authority to execute this **Agreement** on behalf of the **Recipient** and acknowledge my institution's responsibility and obligation to abide by the provisions of this **Agreement**, both by letter and in principle, in order to promote the conservation and sustainable use of **Plant Genetic Resources for Food and Agriculture**.

Signature.....  
Name of the **Recipient**.....

Date.....4.09.2019.....

### Option 2 – Shrink-wrap Standard Material Transfer Agreements\*

The **Material** is provided conditional on acceptance of the terms of this **Agreement**. The provision of the **Material** by the **Provider** and the **Recipient**'s acceptance and use of the **Material** constitutes acceptance of the terms of this **Agreement**.

### Option 3 – Click-wrap Standard Material Transfer Agreement\*

I hereby agree to the above conditions.

\* Where the **Provider** chooses signature, only the wording in Option 1 will appear in the Standard Material Transfer Agreement. Similarly where the **Provider** chooses either shrink-wrap or click-wrap, only the wording in Option 2 or Option 3, as appropriate, will appear in the Standard Material Transfer Agreement. Where the "click-wrap" form is chosen, the **Material** should also be accompanied by a written copy of the Standard Material Transfer Agreement.

#### LIST OF MATERIALS PROVIDED

This Annex contains a list of the Material provided under this Agreement, including the associated information referred to in Article 5b.

This information is either provided below or can be obtained at the following website: (*URL*).

The following information is included for each **Material** listed: all available passport data and, subject to applicable law, any other associated, available, non-confidential descriptive information.

(List)

The *Lactuca serriola* and *L. altaica* material collected encompassed CGN collecting numbers KMK01-KMK62.

More information can be gained from:

<https://missions.cgn.wur.nl/> or [www.wur.nl/cgn](http://www.wur.nl/cgn)



KTone

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## Appendix 2 Expedition collecting form

### Uzbekistan - the Netherlands *Lactuca* expedition 2019

*Team/collector(s)* ..... *Collecting number* .....

*Date* ..... *Photo number* .....

*Crop name* ..... *Cultivar name* .....

*Latin species name* .....

*Locality* .....

.....  
*Latitude* ..... *Longitude* ..... *Altitude* .....

*Number of plants sampled*:

*Estimated size of population sampled*:

*Topography*...swamp...flood plain...level...undulating...hilly...steep...mountainous

**Biological status of accession**

100) Wild

20) Farm or cultivated habitat

21) Field

22) Orchard

23) Backyard, kitchen or home garden

24) Fallow land

25) Pasture

200) Weedy

30) Market or shop

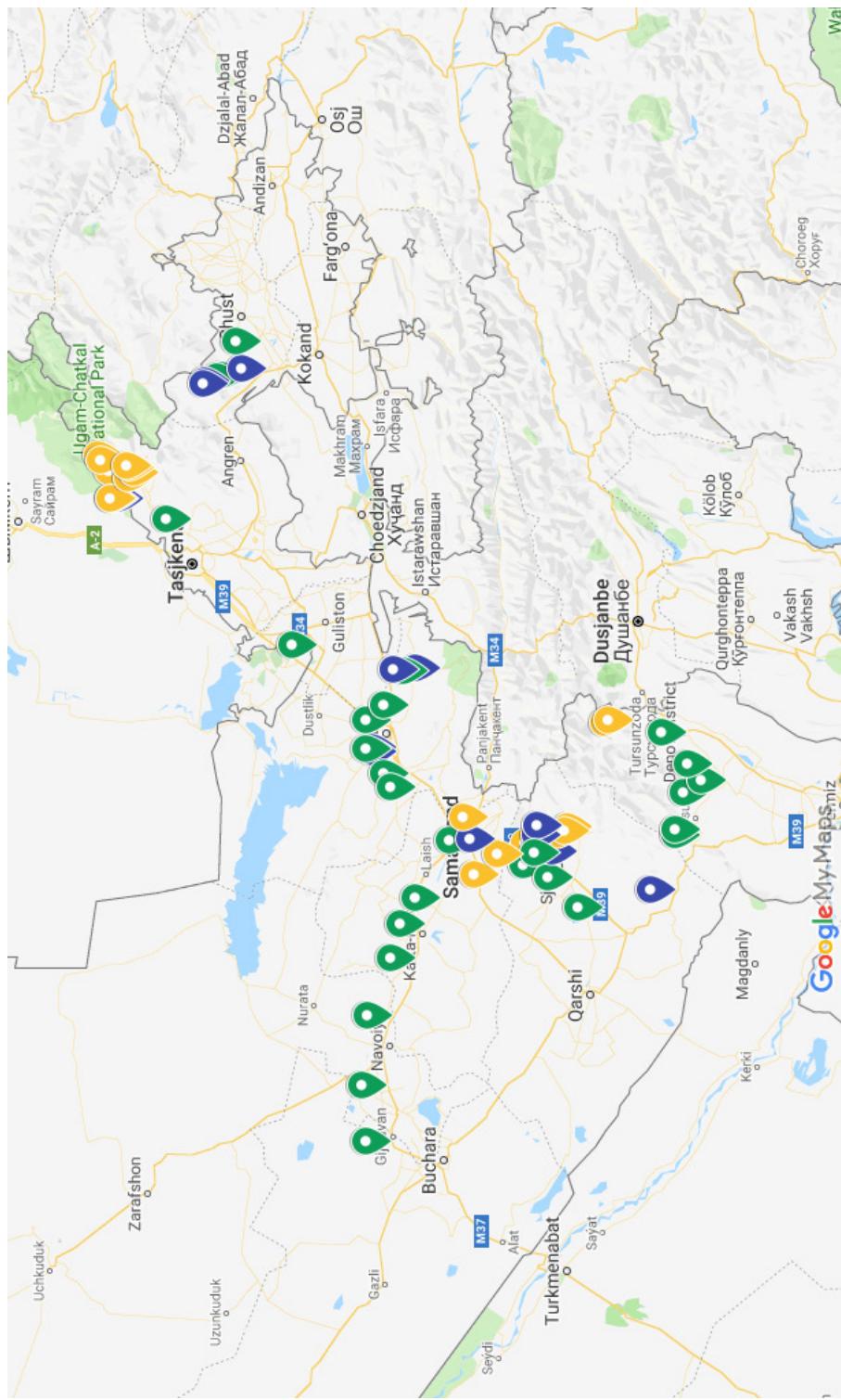
300) Traditional cultivar/landrace

60) Weedy, disturbed or ruderal habitat

&lt;/div

## Appendix 3 Map of locations where collecting took place in Uzbekistan

Collecting numbers (KMxxx) are indicated for the *Lactuca* material collected; yellow marker: *Lactuca serriola*, green: *Lactuca alticaria*, blue: mixed *Lactuca* population.



## Appendix 4 Passport data of *Lactuca* accessions (KMK xxx) collected in Uzbekistan

Date: day-month-year, latitude and longitude determined via GPS: mapdatum WGS84, position format for longitude and latitude: hddd.ddddd', altitude in meters; population type: W: wild material.

RNR	Country	Collection number	Photo map	Photo nr	Date	Name of species	Name of genus during collection	Name of species during collection	Photo to double	Latitude	Longitude	Population type	Topography	Elevation	Number of single bags	Number of double bags	Number of double bags	Number of single bags	Population size	Remarks
190206	KM/K01	U2B	109	536-533	20-7-2019	<i>L serriola</i>	<i>Beiderge</i>	<i>Gulistan</i>	41.5231	69.92512	1319	W	hilly	road side, field	29	29	34	ca 200	ca 10000 salt, heat tolerant?	
190207	KM/K02	U2B	109	534-536	21-7-2019	<i>L serriola</i>			40.55818	69.90912	267	W	level	road side	2	2	0	2	5-10	
190208	KM/K03	U2B	109	539-543	21-7-2019	L. alticaria ?			40.05078	67.75627	604	W	hilly	road side	5	5	10	ca. 100		
190209	KM/K04	U2B	109	544-546	21-7-2019	<i>L serriola</i> - <i>L alticaria</i>			40.04962	67.75080	641	W	hilly	road side, field margin	5	5	5	5	5-10	
190210	KM/K05	U2B	109	565-566	22-7-2019	<i>L serriola</i>	<i>Kushkudau</i>		39.78075	66.55570	606	W	level	road side, field margin	65	4	69	>10000		
190211	KM/K06	U2B	109	567-569	22-7-2019	<i>L serriola</i>	<i>Kattakungan</i>		39.87548	66.54475	555	W	undulating-level	road side, field margin	32	0	32	100-1000		
190212	KM/K07	U2B	109	570-572	22-7-2019	<i>L serriola</i>	<i>Kattakungan</i>		39.87548	66.54475	448	W	level	road side, field margin	26	36	0	ca. 100		
190213	KM/K08	U2B	109	573-574	22-7-2019	<i>L serriola</i>	<i>Nurjichon</i>		40.07793	65.61207	407	W	level	road side, field margin	32	16	48	100-1000		
190214	KM/K09	U2B	109	569-591	23-7-2019	<i>L serriola</i>	<i>Yobkent</i>		40.08362	64.80390	248	W	level	road side, field margin	21	11	32	100-1000		
190215	KM/K10	U2B	109	592-594	23-7-2019	<i>L serriola</i>	<i>Yul'kapa</i>		40.10812	65.04762	327	W	level	road side, field margin	26	6	32	100-1000	<i>L tatarica</i> , photo 109-595-596, salt stressed roadside burning. Sampled at 109-590-801	
190216	KM/K11	U2B	109	597-601	23-7-2019	<i>L serriola</i>	<i>Sana-kond</i>		39.57502	67.01601	261	W	level	road side, field margin	37	2	39	>10000		
190217	KM/K12	U2B	109	603-604	24-7-2019	<i>L serriola</i>	<i>Kitab</i>		39.11243	66.81901	612	W	level	road side, field margin	40	1	41	>10000	near to orchard	
190218	KM/K13	U2B	109	605-606	24-7-2019	<i>L serriola</i>	<i>Yakabog</i>		38.96155	66.72084	614	W	level	orchard	20	24	44	>10000	sampled in apple orchard	
190219	KM/K14	U2B	109	607-609	24-7-2019	<i>L serriola</i>	<i>Guar</i>		38.77446	66.47889	526	W	level	road side, field margin	10	23	33	>10000	very dry and hot habitat	
190220	KM/K15	U2B	109	610-612	24-7-2019	<i>L serriola</i>	<i>Derband</i>		38.16055	67.06093	971	W	undulating-hilly	road side, field margin	38	4	42	100-1000	hot and dry habitat	
190221	KM/K16	U2B	109	613-616	25-7-2019	<i>L serriola</i>	<i>Rudina</i>		38.10861	67.40065	781	W	level	backyard	13	33	46	>10000		
190222	KM/K17	U2B	109	620-621	25-7-2019	<i>L serriola</i>	<i>Shurchay</i>		37.99886	67.50285	532	W	level	backyard	11	20	31	ca. 100	along farmhouse	
190223	KM/K18	U2B	109	622-623	25-7-2019	<i>L serriola</i>	<i>Deynsay</i>		38.07970	67.62884	537	W	level	field margin	9	33	42	>10000	good population	
190224	KM/K19	U2B	109	624-626	25-7-2019	L. alticaria	<i>Sharquni</i>		38.60423	67.9656	1451	W	undulating	backyard	40	6	46	ca. 100	good population of <i>L alticaria</i>	
190225	KM/K20	U2B	109	637-640	25-7-2019	<i>L serriola</i>	<i>Shargun</i>		38.58491	67.98461	1206	W	hilly	pasture	18	6	24	ca. 100		
190226	KM/K21	U2B	109	641-643	26-7-2019	<i>L serriola</i>	<i>Denov</i>		38.24735	67.8802	457	W	undulating-level	field margin	11	19	30	100-1000		
190227	KM/K22	U2B	109	644-646	26-7-2019	<i>L serriola</i>	<i>Baysun</i>		38.15773	67.10224	540	W	level	field margin	24	12	36	>10000	+ 2 movies	
190228	KM/K23	U2B	109	651-652	26-7-2019	<i>L serriola</i>	<i>Dzhambanabad</i>		38.63183	67.51973	1043	W	level	roadside	13	0	13	ca. 100	dry, gypsum soil, some plants <i>altica</i> ?	
190229	KM/K24	U2B	109	652-653	26-7-2019	L. alticaria	<i>Tarkaray</i>		39.41948	67.74065	1145	W	undulating	field margin	35	1	36	100-1000	on mudwall, stones under leaves	
190230	KM/K25	U2B	109	670-672	28-7-2019	<i>L serriola</i> (+ <i>L alticaria</i> )?	<i>Ozbegim</i>		38.91504	66.23242	856	W	undulating-level	backyard	13	32	45	ca. 100	95% <i>ser</i> + 5% <i>alt</i> ?	
190231	KM/K26	U2B	109	673-675	28-7-2019	L. alticaria	<i>Minjur</i>		38.91058	67.05617	1016	W	steep	field margin	10	30	40	ca. 1000	larger flowers	
190232	KM/K27	U2B	109	676-679	28-7-2019	L. alticaria	<i>Katakul</i>		38.83842	67.12817	1515	W	undulating	roadside	23	14	37	100-1000		
190233	KM/K28	U2B	109	694-696	29-7-2019	L. alticaria	<i>Tatar</i>		38.85300	67.99807	1126	W	undulating	undulating	11	10	21	100-500		
190234	KM/K29	U2B	109	697-700	30-7-2019	L. alticaria	<i>Sevaz</i>		39.10779	66.59923	701	W	undulating	orchard	3	41	44	>10000	good population	
190235	KM/K30	U2B	109	701-703	30-7-2019	L. alticaria	<i>Iskon</i>		39.03981	67.06405	900	W	undulating	orchard	19	30	49	100-200	mixed population?	
190236	KM/K31	U2B	109	704-705	30-7-2019	L. alticaria	<i>Miraki</i>		39.02879	67.13934	992	W	hilly	road side, field margin	7	37	44	100-1000		

Number of species	Number of individuals during expedition	Photo nr	Date	Name	Habitat	Erosion	Expedition during	Photo map	Country	Photo nr	Date	Name	Habitat	Erosion	Expedition during	Photo map	Country	Photo nr	Date	Name	Habitat	Erosion	Expedition during	Photo map	Country																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																																							
1902337	KM132	U2B	109	706-707	L	1902337	20-21-2019	L	U2B	109	708-712	30-7-2019	L	1902338	U2B	109	714-715	31-7-2019	L	1902339	U2B	109	716-717	31-7-2019	L	190240	U2B	109	718-723	31-7-2019	L	190241	U2B	109	724-725	1-8-2019	L	190242	U2B	109	726-727	1-8-2019	L	190243	U2B	109	728-729	1-8-2019	L	190244	U2B	109	730-732	1-8-2019	L	190245	U2B	109	733-735	2-8-2019	L	190246	U2B	109	736-738	2-8-2019	L	190247	U2B	109	739-740	2-8-2019	L	190248	U2B	109	741-743	4-8-2019	L	190249	U2B	109	744-745	4-8-2019	L	190250	U2B	109	746-749	5-8-2019	L	190251	U2B	109	750-751	5-8-2019	L	190252	U2B	109	752-753	5-8-2019	L	190254	U2B	109	756-758	5-8-2019	L	190255	U2B	109	759-761	5-8-2019	L	190256	U2B	109	762-763	6-8-2019	L	190257	U2B	109	764-766	6-8-2019	L	190258	U2B	109	767-768	6-8-2019	L	190259	U2B	109	769-770	6-8-2019	L	190260	U2B	109	771-772	7-8-2019	L	190261	U2B	109	773-775	7-8-2019	L	190262	U2B	109	776-777	7-8-2019	L	190263	U2B	109	778-784	7-8-2019	L	190264	U2B	109	785-791	10-8-2019	L	190265	U2B	109	792-794	10-8-2019	L	190266	U2B	109	795-798	10-8-2019	L	190267	U2B	109	799-802	10-8-2019	L	190268	U2B	109	803-806	10-8-2019	L	190269	U2B	109	807-810	10-8-2019	L	190270	U2B	109	811-814	10-8-2019	L	190271	U2B	109	815-818	10-8-2019	L	190272	U2B	109	819-822	10-8-2019	L	190273	U2B	109	823-826	10-8-2019	L	190274	U2B	109	827-830	10-8-2019	L	190275	U2B	109	831-834	10-8-2019	L	190276	U2B	109	835-838	10-8-2019	L	190277	U2B	109	839-842	10-8-2019	L	190278	U2B	109	843-846	10-8-2019	L	190279	U2B	109	847-850	10-8-2019	L	190280	U2B	109	851-854	10-8-2019	L	190281	U2B	109	855-858	10-8-2019	L	190282	U2B	109	859-862	10-8-2019	L	190283	U2B	109	863-866	10-8-2019	L	190284	U2B	109	867-870	10-8-2019	L	190285	U2B	109	871-874	10-8-2019	L	190286	U2B	109	875-878	10-8-2019	L	190287	U2B	109	879-882	10-8-2019	L	190288	U2B	109	883-886	10-8-2019	L	190289	U2B	109	887-890	10-8-2019	L	190290	U2B	109	891-894	10-8-2019	L	190291	U2B	109	895-898	10-8-2019	L	190292	U2B	109	899-902	10-8-2019	L	190293	U2B	109	903-906	10-8-2019	L	190294	U2B	109	907-910	10-8-2019	L	190295	U2B	109	911-914	10-8-2019	L	190296	U2B	109	915-918	10-8-2019	L	190297	U2B	109	919-922	10-8-2019	L	190298	U2B	109	923-926	10-8-2019	L	190299	U2B	109	927-930	10-8-2019	L	190300	U2B	109	931-934	10-8-2019	L	190301	U2B	109	935-938	10-8-2019	L	190302	U2B	109	939-942	10-8-2019	L	190303	U2B	109	943-946	10-8-2019	L	190304	U2B	109	947-950	10-8-2019	L	190305	U2B	109	951-954	10-8-2019	L	190306	U2B	109	955-958	10-8-2019	L	190307	U2B	109	959-962	10-8-2019	L	190308	U2B	109	963-966	10-8-2019	L	190309	U2B	109	967-970	10-8-2019	L	190310	U2B	109	971-974	10-8-2019	L	190311	U2B	109	975-978	10-8-2019	L	190312	U2B	109	979-982	10-8-2019	L	190313	U2B	109	983-986	10-8-2019	L	190314	U2B	109	987-990	10-8-2019	L	190315	U2B	109	991-994	10-8-2019	L	190316	U2B	109	995-998	10-8-2019	L	190317	U2B	109	999-1002	10-8-2019	L	190318	U2B	109	1003-1006	10-8-2019	L	190319	U2B	109	1007-1010	10-8-2019	L	190320	U2B	109	1011-1014	10-8-2019	L	190321	U2B	109	1015-1018	10-8-2019	L	190322	U2B	109	1019-1022	10-8-2019	L	190323	U2B	109	1023-1026	10-8-2019	L	190324	U2B	109	1027-1030	10-8-2019	L	190325	U2B	109	1031-1034	10-8-2019	L	190326	U2B	109	1035-1038	10-8-2019	L	190327	U2B	109	1039-1042	10-8-2019	L	190328	U2B	109	1043-1046	10-8-2019	L	190329	U2B	109	1047-1050	10-8-2019	L	190330	U2B	109	1051-1054	10-8-2019	L	190331	U2B	109	1055-1058	10-8-2019	L	190332	U2B	109	1059-1062	10-8-2019	L	190333	U2B	109	1063-1066	10-8-2019	L	190334	U2B	109	1067-1070	10-8-2019	L	190335	U2B	109	1071-1074	10-8-2019	L	190336	U2B	109	1075-1078	10-8-2019	L	190337	U2B	109	1079-1082	10-8-2019	L	190338	U2B	109	1083-1086	10-8-2019	L	190339	U2B	109	1087-1090	10-8-2019	L	190340	U2B	109	1091-1094	10-8-2019	L	190341	U2B	109	1095-1098	10-8-2019	L	190342	U2B	109	1099-1102	10-8-2019	L	190343	U2B	109	1103-1106	10-8-2019	L	190344	U2B	109	1107-1110	10-8-2019	L	190345	U2B	109	1111-1114	10-8-2019	L	190346	U2B	109	1115-1118	10-8-2019	L	190347	U2B	109	1119-1122	10-8-2019	L	190348	U2B	109	1123-1126	10-8-2019	L	190349	U2B	109	1127-1130	10-8-2019	L	190350	U2B	109	1131-1134	10-8-2019	L	190351	U2B	109	1135-1138	10-8-2019	L	190352	U2B	109	1139-1142	10-8-2019	L	190353	U2B	109	1143-1146	10-8-2019	L	190354	U2B	109	1147-1150	10-8-2019	L	190355	U2B	109	1151-1154	10-8-2019	L	190356	U2B	109	1155-1158	10-8-2019	L	190357	U2B	109	1159-1162	10-8-2019	L	190358	U2B	109	1163-1166	10-8-2019	L	190359	U2B	109	1167-1170	10-8-2019	L	190360	U2B	109	1171-1174	10-8-2019	L	190361	U2B	109	1175-1178	10-8-2019	L	190362	U2B	109	1179-1182	10-8-2019	L	190363	U2B	109	1183-1186	10-8-2019	L	190364	U2B	109	1187-1190	10-8-2019	L	190365	U2B	109	1191-1194	10-8-2019	L	190366	U2B	109	1195-1198	10-8-2019	L	190367	U2B	109	1199-1202	10-8-2019	L	190368	U2B	109	1203-1206	10-8-2019	L	190369	U2B	109	1207-1210	10-8-2019	L	190370	U2B	109	1211-1214	10-8-2019	L	190371	U2B	109	1215-1218	10-8-2019	L	190372	U2B	109	1219-1222	10-8-2019	L	190373	U2B	109	1223-1226	10-8-2019	L	190374	U2B	109	1227-1230	10-8-2019	L	190375	U2B	109	1231-1234	10-8-2019	L	190376	U2B	109	1235-1238	10-8-2019	L	190377	U2B	109	1239-1242	10-8-2019	L	190378	U2B	109	1243-1246	10-8-2019	L	190379	U2B	109	1247-1250	10-8-2019	L	190380	U2B	109	1251-1254	10-8-2019	L	190381	U2B	109	1255-1258	10-8-2019	L	190382	U2B	109	1259-1262	10-8-2019	L	190383	U2B	109	1263-1266	10-8-2019	L	190384	U2B	109	1267-1270	10-8-2019	L	190385	U2B	109	1271-1274	10-8-2019	L	190386	U2B	109	1275-1278	10-8-2019	L	190387	U2B	109	1279-1282	10-8-2019	L	190388	U2B	109	1283-1286	10-8-2019	L	190389	U2B	109	1287-1290	10-8-2019	L	190390	U2B	109	1291-1294	10-8-2019	L	190391	U2B	109	1295-1298	10-8-2019	L	190392	U2B	109	1299-1302	10-8-2019	L	190393	U2B	109	1303-1306	10-8-2019	L	190394	U2B	109	1307-1310	10-8-2019	L	190395	U2B	109	1311-1314	10-8-2019	L	190396	U2B	109	1315-1318	10-8-2019	L	190397	U2B	109	1319-1322	10-8-2019	L	190398	U2B	109	1323-1326	10-8-2019	L	190399	U2B	109	1327-1330	10-8-2019	L	190400	U2B	109	1331-1334	10-8-2019	L	190401	U2B	109	1335-1338	10-8-2019	L	190402	U2B	109	1339-1342	10-8-2019	L	190403	U2B	109	1343-1346	10-8-2019	L	190404	U2B	109	1347-1350	10-8-2019	L	190405	U2B	109	1351-1354	10-8-2019	L	190406	U2B	109	1355-1358	10-8-2019	L	190407	U2B	109	1359-1362	10-8-2019	L	190408	U2B	109	1363-1366	10-8-2019	L	190409	U2B	109	1367-1370	10-8-2019	L	190410	U2B	109	1371-1374	10-8-2019	L	190411	U2B	109	1375-1378	10-8-2019	L	190412	U2B	109	1379-1382	10-8-2019	L	190413	U2B	109	1383-1386	10-8-2019	L	190414	U2B	109	1387-1390	10-8-2019	L	190415	U2B	109	1391-1394	10-8-2019	L	190416	U2B	109	1395-1398	10-8-2019	L	190417	U2B	109	1399-1402	10-8-2019	L	190418	U2B	109	1403-1406	10-8-2019	L	190419	U2B	109	1407-1410	10-8-2019	L	190420	U2B	109	1411-1414	10-8-2019	L	190421	U2B	109	1415-1418	10-8-2019	L	190422	U2B	109	1419-1422	10-8-2019	L	190423	U2B	109

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Wageningen University & Research  
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The mission of Wageningen University & Research is "To explore the potential of nature to improve the quality of life". Under the banner Wageningen University & Research, Wageningen University and the specialised research institutes of the Wageningen Research Foundation have joined forces in contributing to finding solutions to important questions in the domain of healthy food and living environment. With its roughly 30 branches, 5,000 employees and 10,000 students, Wageningen University & Research is one of the leading organisations in its domain. The unique Wageningen approach lies in its integrated approach to issues and the collaboration between different disciplines.

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