

Chromosomal and DNA barcode analysis of the *Polyommatus (Agrodiaetus) damone* (Eversmann, 1841) species complex (Lepidoptera, Lycaenidae)

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Abstract

The *Polyommatus (Agrodiaetus) damone* (Eversmann, 1841) species complex comprises from 5 to 8 species distributed in southeastern Europe and southern Siberia. Here we used chromosomal and DNA-barcode markers in order to test the taxonomic hypotheses previously suggested for this complex. We revealed that all taxa within this group demonstrate chromosomal stasis and share the same or very similar haploid chromosome number ($n = 66$ or $n = 67$). This finding is unexpected since the karyotypes are known to be very diverse and species-specific within the other taxa of the subgenus *Agrodiaetus* Hübner, 1822. Analysis of the mitochondrial gene *COI* revealed six diverged clusters of individuals within the complex. Each cluster has a specific geographic distribution and is characterized by distinct morphological features in the wing pattern. The clusters mostly (but not always) correlate with traditionally recognized species. As a result of our study, we describe a new subspecies *P. (A.) iphigenides zarmitanus* **subsp. nov.** from Uzbekistan and Tajikistan and show that the taxon originally described as *Lycaena kindermannii* var. *melania* Staudinger, 1886 represents a subspecies *P. (A.) iphigenides melanius* (Staudinger, 1886). *Polyommatus (A.) samusi* Korb, 2017 (**syn. nov.**) and *P. (A.) melanius komarovi* Korb, 2017 (**syn. nov.**) are considered here as junior subjective synonyms of *P. (A.) iphigenides iphigenides* (Staudinger, 1886).

Keywords

Agrodiaetus, chromosomal stasis, chromosome, *COI*, DNA barcoding, karyosystematics, taxonomy

Introduction

The *Polyommatus* (*Agrodiaetus*) *damone* (Eversmann, 1841) species complex is a monophyletic group (Vershina and Lukhtanov 2017) that comprises from 5 to 8 species distributed in SE Europe, Central Asia and S Siberia (Eckweiler and Bozano 2016). The taxa of the complex were previously revised by Staudinger (1899), Forster (1956, 1960), Dantchenko and Lukhtanov (1993) and Dantchenko (1997). There are also limited molecular (Wiemers 2003; Kandul et al. 2004, 2007; Lukhtanov et al. 2005, 2009; Vodolazhsky et al. 2011; Vodolazhsky and Stradomsky 2012) and chromosomal (Lukhtanov 1989; Kandul 1997; Lukhtanov et al. 1997; Lukhtanov and Dantchenko 2002a; Lukhtanov et al. 2005) data for a few taxa of the complex. However, the complex has never been systematically studied by using chromosomal and molecular markers, although such an approach is considered as an essential requirement for revealing taxonomic structure in the subgenus *Agrodiaetus* (Lukhtanov and Dantchenko 2002b; Kandul et al. 2004).

Here we analyzed karyotypes and mitochondrial DNA-barcodes of all species of the *P. (A.) damone* complex in order to test the taxonomic hypotheses previously suggested for this group (see the references above).

The taxa *P. (A.) damone walteri* Dantchenko et Lukhtanov, 1993, *P. (A.) damone fabiani* Bálint, 1997 and *P. (A.) damone bogdoolensis* Dantchenko et Lukhtanov, 1997 are not considered in this paper since neither chromosomal nor molecular data are available. This also applies to *P. (A.) carmon altaiensis* (Forster, 1956), recently treated by Eckweiler and Bozano (2016) as a separate species. All these taxa represent the most eastern populations of the *P. (A.) damone* complex distributed in Mongolia, Altai and southwestern Siberia. Morphologically they are close to other populations of *P. damone* or to *P. mediator* Dantchenko et Churkin, 2003. Their study will become possible in the future as soon as the material suitable for molecular and chromosomal analyses becomes available.

Material and methods

Molecular methods and DNA barcode analysis

Standard *COI* barcodes (658-bp 5' segment of mitochondrial *cytochrome oxidase subunit I*) were studied. *COI* sequences were obtained from 44 specimens representing the *P. damone* species group and from two samples [*P. damon* (Denis et Schiffermüller, 1775) and *P. icarus* (Rottemburg, 1975)] which were selected as outgroup (Table 1). Legs were sampled from these specimens, and sequence data from the DNA barcode region of *COI* were obtained at the Canadian Centre for DNA Barcoding (CCDB, Biodiversity Institute of Ontario, University of Guelph) using protocols described in Hajibabaei et al. (2005), Ivanova et al. (2006) and deWaard et al. (2008). Specimens examined are deposited in the Zoological Institute of the Russian Academy of Sciences, St. Petersburg, Russia and in the McGuire Center for Lepidoptera and Biodiversity

Table 1. Specimens of the *Polyommatus (Agrodiaetus) damone* complex used in the DNA-barcode analysis.

Species and subspecies	Sequence code	Field code	GenBank number	Country	Locality	Reference
<i>P. damon</i>	FJ663230	n/a	FJ663230	Kazakhstan	Altai	Lukhtanov et al. 2009
<i>P. damone altaicus</i>	FJ663229	LOWA298-06	FJ663229	Kazakhstan	Saur-Tarbagatai Mts	Lukhtanov et al. 2009
<i>P. damone altaicus</i>	FJ663228	LOWA299-06	FJ663228	Kazakhstan	Saur-Tarbagatai Mts	Lukhtanov et al. 2009
<i>P. damone altaicus</i>	BPAL3395-16	CCDB-25452_F10	MW186990	Russia	Altai, Jarbalyk	This study
<i>P. damone altaicus</i>	BPAL838-11	CCDB-05724_G06	MW186700	Kazakhstan	Saur Mts, Saikan	This study
<i>P. damone altaicus</i>	BPAL839-11	CCDB-05724_G07	MW186701	Kazakhstan	Saur Mts, Saikan	This study
<i>P. damone altaicus</i>	AY496734	n/a	AY496734	Russia	Altai, Aktrash	Kandul et al. 2004
<i>P. damone altaicus</i>	BPAL3394-16	CCDB-25452_F09	MW186989	Russia	Altai, Chermal	This study
<i>P. damone</i>	BPAL836-11	CCDB-05724_G04	MW186988	Russia	Volga, Volsk	This study
<i>P. damone</i>	BPAL837-11	CCDB-05724_G05	MW186992	Russia	Volga, Volsk	This study
<i>P. damone</i>	BPAL835-11	CCDB-05724_G03	MW186999	Russia	Volga, Akulovka	This study
<i>P. damone</i>	AY496735	n/a	AY496735	Russia	South Urals, Guberli Mts, Aadaevo	Kandul et al. 2004
<i>P. damone irinae</i>	BPAL833-11	CCDB-05724_G01	MW186997	Russia	Volgograd Region, Olkhovka	This study
<i>P. damone irinae</i>	BPAL834-11	CCDB-05724_G02	MW186998	Russia	Volgograd Region, Olkhovka	This study
<i>P. damone irinae</i>	AY496736	n/a	AY496736	Russia	Volgograd Region, Olkhovka	Kandul et al. 2004
<i>P. damone pljushchii</i>	AY496774	n/a	AY496774	Russia	Crimea, Ai Petri	Kandul et al. 2004
<i>P. damone ssp.</i>	BPAL524-11	n/a	MW186991	Kazakhstan	Karaganda Region, Akchatau	This study
<i>P. damone tanais</i>	BPAL825-11	CCDB-05724_F05	MW186993	Ukraine	Amvrosievka	This study
<i>P. damone tanais</i>	BPAL826-11	CCDB-05724_F06	MW186994	Ukraine	Amvrosievka	This study
<i>P. damone tanais</i>	BPAL827-11	CCDB-05724_F07	MW186995	Ukraine	Amvrosievka	This study
<i>P. damone tanais</i>	BPAL828-11	CCDB-05724_F08	MW186996	Ukraine	Amvrosievka	This study
<i>P. damone tanais</i>	KC692328	n/a	KC692328	Russia	Rostov Region, Belaya Kalitva	Vodolazhsky and Stradomsky 2012
<i>P. icarus</i>	HM913968	n/a	HM913968	Italy	39.9919°N, 15.7931°E	GenBank
<i>P. iphigenides</i>	n/a	LOWA422-06	FJ663238	Kyrgyzstan	Transalai Mts, Nura	Lukhtanov et al. 2009
<i>P. iphigenides</i>	n/a	LOWA423-06	FJ663237	Kyrgyzstan	Transalai Mts, Nura	Lukhtanov et al. 2009
<i>P. iphigenides iphigenides</i>	n/a	LOWA424-06	FJ663236	Kyrgyzstan	Transalai Mts, Nura	Lukhtanov et al. 2009
<i>P. iphigenides iphigenides</i>	n/a	LOWA514-06	FJ663235	Kyrgyzstan	Alai, Tengizbai Pass	Lukhtanov et al. 2009
<i>P. iphigenides iphigenides</i>	n/a	LOWA515-06	FJ663234	Kyrgyzstan	Alai, Tengizbai Pass	Lukhtanov et al. 2009
<i>P. iphigenides iphigenides</i>	BPAL1586-12	CCDB-03032_F06	MW194007	Tajikistan	Iskanderkul	This study
<i>P. iphigenides iphigenides</i>	BPAL1587-12	CCDB-03032_F07	MW194008	Tajikistan	Iskanderkul	This study
<i>P. iphigenides iphigenides</i>	AY496758	n/a	AY496758	Kazakhstan	Shymkent Region, Ugamski Mts	Kandul et al. 2004
<i>P. iphigenides iphigenides</i>	AY557155	WE98001	AY557155	Kyrgyzstan	25 km S Song Kul Lake	Wiemers 2003
<i>P. iphigenides melanius</i>	BPALB479-18	CCDB-23848_A04	MW186954	Tajikistan	Alai Mts, Jirgatal	This study
<i>iphigenides melanius</i>	BPALB480-18	CCDB-23848_A05	MW186955	Tajikistan	Alai Mts, Jirgatal	This study
<i>P. iphigenides melanius</i>	BPALB481-18	CCDB-23848_A06	MW186956	Tajikistan	Alai Mts, Jirgatal	This study
<i>P. iphigenides melanius</i>	BPALB482-18	CCDB-23848_A07	MW186957	Tajikistan	Alai Mts, Jirgatal	This study
<i>P. iphigenides melanius</i>	BPALB483-18	CCDB-23848_A08	MW186958	Tajikistan	Alai Mts, Jirgatal	This study
<i>P. iphigenides melanius</i>	BPALB484-18	CCDB-23848_A09	MW186959	Tajikistan	Alai Mts, Jirgatal	This study
<i>P. iphigenides melanius</i>	BPALB556-18	CCDB-23848_G09	MW186960	Tajikistan	Peter I Mts, Khorakul Lake	This study
<i>P. iphigenides melanius</i>	BPALB558-18	CCDB-23848_G11	MW186961	Tajikistan	Peter I Mts, Mingbulak	This study
<i>P. iphigenides melanius</i>	BPALB559-18	CCDB-23848_G12	MW186962	Tajikistan	Peter I Mts, Mingbulak	This study
<i>P. iphigenides zarmitanus</i>	BPAL1390-12	CCDB-03030_E12	MW186963	Uzbekistan	Nuratau Mts, Zarmitan	This study
<i>P. iphigenides zarmitanus</i>	BPAL1391-12	CCDB-03030_F01	MW186964	Uzbekistan	Nuratau Mts, Zarmitan	This study
<i>P. iphigenides zarmitanus</i>	BPAL1392-12	CCDB-03030_F02	MW186965	Uzbekistan	Nuratau Mts, Zarmitan	This study
<i>P. iphigenides zarmitanus</i>	BPAL1394-12	CCDB-03030_F04	MW186967	Uzbekistan	Nuratau Mts, Zarmitan	This study
<i>P. iphigenides zarmitanus</i>	BPAL1514-12	CCDB-03031_H05	MW186968	Uzbekistan	Hissar Range, Tamshush	This study
<i>P. iphigenides zarmitanus</i>	BPAL1515-12	CCDB-03031_H06	MW186969	Uzbekistan	Hissar Range, Tamshush	This study
<i>P. iphigenides zarmitanus</i>	BPAL1533-12	CCDB-03032_B01	MW186970	Uzbekistan	Hissar Range, Sangardak	This study
<i>P. iphigenides zarmitanus</i>	BPAL1534-12	CCDB-03032_B02	MW186971	Uzbekistan	Hissar Range, Sangardak	This study
<i>P. iphigenides zarmitanus</i>	BPAL1535-12	CCDB-03032_B03	MW186972	Uzbekistan	Hissar Range, Sangardak	This study
<i>P. iphigenides zarmitanus</i>	BPAL1536-12	CCDB-03032_B04	MW186973	Uzbekistan	Hissar Range, Sangardak	This study

Species and subspecies	Sequence code	Field code	GenBank number	Country	Locality	Reference
<i>P. iphigenides zarmitanus</i>	BPAL1544-12	CCDB-03032_B12	MW186974	Uzbekistan	Hissar Range, Tamshush	This study
<i>P. iphigenides zarmitanus</i>	AY556853	DS01001	AY556853	Uzbekistan	Kitabsky reserve	Wiemers 2003
<i>P. iphigenides zarmitanus</i> (Holotype)	BPAL1393-12	CCDB-03030_F03	MW186966	Uzbekistan	Nuratau Mts, Zarmitan	This study
<i>P. juldusus</i>	BPAL852-11	CCDB-05724_H08	MW186985	Kazakhstan	Almaty Region, Kegen Pass	This study
<i>P. juldusus</i>	BPAL870-11	CCDB-05725_B03	MW186986	Kyrgyzstan	Issykkyl, Kadzhisai	This study
<i>P. juldusus kasachstanus</i>	AY496759	n/a	AY496759	Kazakhstan	Dzhungarian Alatau	Kandul et al. 2004
<i>P. juldusus kirgisorum</i>	BPAL1381-12	CCDB-03030_E03	MW186987	Kyrgyzstan	Shamsi	This study
<i>P. karatavicus</i>	BPAL040-10	RPVL-00040	MW186975	Kazakhstan	Karatau Mts, Minzhilgi	This study
<i>P. karatavicus</i>	BPAL041-10	RPVL-00041	MW186976	Kazakhstan	Karatau Mts, Minzhilgi	This study
<i>P. karatavicus</i>	BPAL042-10	RPVL-00042	MW186977	Kazakhstan	Karatau Mts, Minzhilgi	This study
<i>P. karatavicus</i>	BPAL1388-12	CCDB-03030_E10	MW186978	Kazakhstan	Karatau Mts	This study
<i>P. karatavicus</i>	AY496760	n/a	AY496760	Kazakhstan	Karatau Mts	Kandul et al. 2004
<i>P. mediator habievi</i>	JF343830	ILL087	JF343830	Mongolia	Arshantyn-Nuruu Mts	Vodolazhsky et al. 2011
<i>P. mediator habievi</i>	JF343829	ILL086	JF343829	Mongolia	Bayan Ulegei aimak, Elt Gol river	Vodolazhsky et al. 2011
<i>P. mediator mediator</i>	EF104602	n/a	EF104602	Mongolia	Altai Mts, Biger	Kandul et al. 2004
<i>P. phyllides askhabadicus</i>	BPAL864-11	CCDB-05725_A09	MW186983	Iran	Kuh e Sorkh Mts, Fariman	This study
<i>P. phyllides askhabadicus</i>	BPAL865-11	CCDB-05725_A10	MW186984	Iran	Kuh e Sorkh Mts, Fariman	This study
<i>P. phyllides askhabadicus</i>	AY954011	n/a	AY954011	Iran	Khorasan, Chakane	Lukhtanov et al. 2005
<i>P. phyllides kentauensis</i>	BPAL1382-12	CCDB-03030_E04	MW186980	Kazakhstan	Karatau Mts	This study
<i>P. phyllides kentauensis</i>	AY496769	n/a	AY496769	Kazakhstan	Karatau Mts	Kandul et al. 2004
<i>P. phyllides phyllides</i>	FJ663239	LOWA633-06	FJ663239	Tajikistan	Iskanderkul	Lukhtanov et al. 2009
<i>P. phyllides phyllides</i>	BPAL1328-12	CCDB-03029_H09	MW186979	Uzbekistan	Sairob	This study
<i>P. phyllides phyllides</i>	BPAL1578-12	CCDB-03032_E10	MW186981	Tajikistan	Iskanderkul	This study
<i>P. phyllides phyllides</i>	BPAL2660-14	CCDB-17967_H11	MW186982	Tajikistan	Sarsaryak	This study
<i>P. phyllides phyllides</i>	FJ663240	LOWA571-06	FJ663240	Uzbekistan	Nuratau Mts, Zarmitan	Lukhtanov et al. 2009
<i>P. phyllides phyllides</i>	AY496771	n/a	AY496771	Kazakhstan	Karzhantau Mts	Kandul et al. 2004
<i>P. phyllides phyllides</i>	AY496770	n/a	AY496770	Kazakhstan	Kirgizski Range	Kandul et al. 2004

(MGCL), Florida Museum of Natural History, University of Florida, Gainesville, Florida, USA. Photographs of these specimens, as well as collecting data are available in the Barcode of Life Data System (BOLD), projects Butterflies of Palearctic (BPAL) and Butterflies of Palearctic Part B (BPALB) at <http://www.boldsystems.org/>.

We also used 31 published *COI* sequences (Wiemers 2003; Kandul et al. 2004, 2007; Lukhtanov et al. 2005, 2009; Vodolazhsky et al. 2011; Vodolazhsky and Stradomsky 2012) which were downloaded from GenBank (Table 1).

Sequences were aligned using the BioEdit software (Hall 1999) and edited manually. Phylogenetic hypotheses were inferred using Bayesian inference as described previously (Vershina and Lukhtanov 2010; Przybyłowicz et al. 2014; Lukhtanov et al. 2016). Briefly, the Bayesian analysis was performed using the program MrBayes 3.2 (Ronquist et al. 2012) with default settings as suggested by Mesquite (Maddison and Maddison 2015): burn-in = 0.25, nst = 6 (GTR + I + G). Two runs of 10,000,000 generations with four chains (one cold and three heated) were performed. We checked runs for convergence and proper sampling of parameters [effective sample size (ESS) > 200] using the program tracer v1.7.1 (Rambaut et al. 2018). The first 25% of each run was discarded as burn-in. The consensus of the obtained trees was visualized using FigTree 1.3.1 (<http://tree.bio.ed.ac.uk/software/figtree/>).

Chromosomal analysis

Karyotypes were studied in 16 adult males representing four species (Table 2) and were processed as previously described (Lukhtanov et al. 2014; Vishnevskaya et al. 2016). Briefly, gonads were removed from the abdomen and placed into freshly prepared fixative (3:1; 96% ethanol and glacial acetic acid) directly after capturing the butterfly in the field. Testes were stored in the fixative for 3–36 months at +4 °C. Then the gonads were stained in 2% acetic orcein for 30–60 days at +18–20 °C. Different stages of male meiosis, including metaphase I (MI) and metaphase II (MII) were examined using an original two-phase method of chromosome analysis (Lukhtanov et al. 2006, 2008). Abbreviation *ca* (circa) means that the count was made with an approximation due to an insufficient quality of preparation or overlapping of some chromosomes or bivalents.

Leica DM2500 light microscope equipped with HC PL APO 100×/1.44 Oil CORR CS lens and S1/1.4 oil condenser head was used for bright-field microscopy analysis. Leica DM2500 light microscope equipped with HC PL APO 100×/1.40 OIL PH3 lens was used for phase-contrast microscopy analysis.

Results

DNA-barcode analysis

Phylogenetic analysis revealed six clusters of closely related individuals within the *P. (A.) damone* species complex (Fig. 1). Of these clusters, four groups were monophyletic and two groups were paraphyletic. The lineages of *P. (A.) damone* (I) and *P. (A.) karatavicus* Lukhtanov, 1990 (V) were highly supported. The lineage of *P. (A.) phyllides* (Staudinger, 1886) (VI) and the lineage [*P. (A.) mediator* Dantchenko et Churkin, 2003 + *P. (A.) juldusus kasachstanus* Lukhtanov et Dantchenko, 1994)] (II) had medium support. The clusters III [*P. (A.) iphigenides iphigenides* (Staudinger, 1886) + *P. (A.) iphigenides melanius* (Staudinger, 1886)] and VI (*P. (A.) iphigenides zarmitanus* subsp. nov.) appeared on the tree as two distinct, not closely related paraphyletic taxa.

Chromosomal analysis

Chromosomal analysis of three males of *P. (A.) damone damone*, of a single male of *P. (A.) damone tanais* Dantchenko et Pljushtch, 1993, of two males of *P. (A.) iphigenides iphigenides*, of three males of *P. (A.) iphigenides melanius*, of a single male of *P. (A.) phyllides phyllides*, of two males of *P. (A.) phyllides askhabadicus* (Forster, 1960) and four males of *P. (A.) iphigenides zarmitanus* subsp. nov. revealed the same (or almost the same) haploid chromosome number $n = 66$ or $n = 67$ in all studied taxa (Table 2). The karyotype structure was also found to be identical in all studied individuals, with three large bivalents in the center of metaphase plates (Fig. 2). Bivalent 1 was 1.2–1.5 times larger than bivalent 2, and the latter was 1.2–1.5 times larger than bivalent 3.

Table 2. Chromosome numbers of species of the *Polyommatus (Agrodiaetus) damone* complex collected by A. Dantchenko (AD), V. Lukhtanov (AV), and Yu. Budashkin and N. Kandul (B & K).

Species	ID number	Chromosome number	Country	Locality	date	Collector	Reference
<i>P. damone altaicus</i>	1987-445	n = ca65	Russia	Altai Mts, Tshulyshman River, 500 m	3–10 August 1987	VL	Lukhtanov 1989
<i>P. damone altaicus</i>	1997-1	n = ca65-67	Kazakhstan	near Zaisan city	23 June 1997	VL	Lukhtanov and Dantchenko 2002a
<i>P. damone altaicus</i>	1997-2	n = 67	Kazakhstan	Saur Mts, Saikan	2–3 July 1997	VL	Lukhtanov and Dantchenko 2002a
<i>P. damone damone</i>	94001	n = 66-67	Russia	Saratov Region, near Volsk	July 1994	AD	Lukhtanov et al. 1997
<i>P. damone damone</i>	94002	n = 67	Russia	Saratov Region, near Volsk	July 1994	AD	Lukhtanov et al. 1997
<i>P. damone damone</i>	94003	n = 66	Russia	Saratov Region, near Volsk	July 1994	AD	Lukhtanov et al. 1997
<i>P. damone damone</i>	94008	n = 67	Russia	Saratov Region, near Volsk	July 1994	AD	Lukhtanov et al. 1997
<i>P. damone damone</i>	94010	n = ca66-67	Russia	Saratov Region, near Radishevo	July 1994	AD	This study
<i>P. damone damone</i>	95DG5	n = ca67	Russia	South Ural, Kuvandyk	1995	AD	This study
<i>P. damone damone</i>	95DG6	n = ca66-67	Russia	South Ural, Kuvandyk	1995	AD	This study
<i>P. damone irinae</i>	AD00P077	n = ca67	Russia	Volgograd region	July 2000	AD	Kandul et al. 2007
<i>P. damone pljushchi</i>	95051	n = 65-67	Russia	Crimea, Ai-Petri, 1200 m	14 July 1995	B & K	Kandul 1997
<i>P. damone pljushchi</i>	95054	n = ca66-68	Russia	Crimea, Ai-Petri, 1200 m	14 July 1995	B & K	Kandul 1997
<i>P. damone pljushchi</i>	95055	n = ca65-67	Russia	Crimea, Ai-Petri, 1200 m	14 July 1995	B & K	Kandul 1997
<i>P. damone pljushchi</i>	96009	n = ca65-66	Russia	Crimea, Ai-Petri, 1200 m	10 July 1995	B & K	Kandul 1997
<i>P. damone pljushchi</i>	96010	n = 67	Russia	Crimea, Ai-Petri, 1200 m	10 July 1995	B & K	Kandul 1997
<i>P. damone pljushchi</i>	96011	n = 65	Russia	Crimea, Ai-Petri, 1200 m	10 July 1995	B & K	Kandul 1997
<i>P. damone pljushchi</i>	96012	n = 66-67	Russia	Crimea, Ai-Petri, 1200 m	10 July 1995	B & K	Kandul 1997
<i>P. damone pljushchi</i>	96017	n = ca66-68	Russia	Crimea, Ai-Petri, 1200 m	10 July 1995	B & K	Kandul 1997
<i>P. damone pljushchi</i>	95050	n = 66-67	Russia	Crimea, Ai-Petri, 1200 m	14 July 1995	B & K	Kandul 1997
<i>P. damone tanais</i>	95005	n = ca67	Ukraine	Don River basin, Shirokaya balka	26 May 1995	AD	This study
<i>P. iphigenides iphigenides</i>	irkeshitam	n = ca66-67	Kyrgyzstan	Transalai Mts (east), Irkeshitam	1996	VL	Lukhtanov and Dantchenko 2002a
<i>P. iphigenides iphigenides</i>	1996-4	n = ca66-67	Kyrgyzstan	Naryn Region, Chaek	4 July 1996	VL	This study
<i>P. iphigenides iphigenides</i>	1996-3	n = ca66	Kyrgyzstan	Moldatoo Mts, Teke-Uyuk	30 June 1996	VL	This study
<i>P. iphigenides iphigenides</i>	1995 – Chiitala	n = 67	Kyrgyzstan	Alai Mts, Chiitala village, 2300 m	1995	VL	Lukhtanov and Dantchenko 2002a
<i>P. iphigenides iphigenides</i>	1994-1	n = ca66-67	Tajikistan	Iskanderkul	July 1994	VL	This study
<i>P. iphigenides iphigenides</i>	95205	n = ca66-67	Kyrgyzstan	Alai Mts, Chiitala village, 2300 m	1995	VL	Lukhtanov and Dantchenko 2002a
<i>P. iphigenides iphigenides</i>	NK00P823 AY496758	n = ca65-67	Kazakhstan	Ugamski range	June 2000	VL	Lukhtanov et al. 2005
<i>P. iphigenides melanius</i>	068K18A	n = 66	Tajikaistan	Alai Mts, Jirgatol	July 2018	AD	This study
<i>P. iphigenides melanius</i>	077K18A	n = 67	Tajikaistan	Alai Mts, Jirgatol	July 2018	AD	This study
<i>P. iphigenides melanius</i>	Tj002	n = 66	Tajikaistan	Peter I Mts	July 20218	VL	This study
<i>P. iphigenides zarmitanus</i>	94L01	n = ca66-68	Uzbekistan	Nuratau Mts, Zarmitan, 1300 m	11–13 June 1994	VL	This study
<i>P. iphigenides zarmitanus</i>	94L03	n = ca68	Uzbekistan	Nuratau Mts, Zarmitan, 1300 m	11–13 June 1994	VL	This study
<i>P. iphigenides zarmitanus</i>	94L04	n = 67	Uzbekistan	Nuratau Mts, Zarmitan, 1300 m	11–13 June 1994	VL	This study
<i>P. iphigenides zarmitanus</i>	94L54	n = ca66-67	Uzbekistan	Hissar Range, Sangardak, 1600 n	2 July 1994	VL	Lukhtanov and Dantchenko 2002a
<i>P. iphigenides zarmitanus</i>	94L61	n = 67	Uzbekistan	Hissar Range, Tamshush, 1800 n	5–7 July 1994	VL	Lukhtanov and Dantchenko 2002a
<i>P. iphigenides zarmitanus</i>	94L64	n = 66	Uzbekistan	Hissar Range, Tamshush, 1800 n	5–7 July 1994	VL	Lukhtanov and Dantchenko 2002a
<i>P. iphigenides zarmitanus</i>	94L74	n = ca65-67	Uzbekistan	Samarkand Region, Aman-Kutan	7 July 1994	VL	Lukhtanov and Dantchenko 2002a
<i>P. iphigenides zarmitanus</i>	94L75	n = ca65-67	Uzbekistan	Samarkand Region, Aman-Kutan	7 July 1994	VL	Lukhtanov and Dantchenko 2002a

Species	ID number	Chromosome number	Country	Locality	date	Collector	Reference
<i>P. iphigenides zarmitanus</i>	1994-2	n = 67	Uzbekistan	Nuratau Mts, Zarmitan, 1300 m	11–13 June 1994	VL	This study
<i>P. iphigenides zarmitanus</i>	1994-3	n = 67	Uzbekistan	Tamshush	1994	VL	Lukhtanov and Dantchenko 2002a
<i>P. iphigenides zarmitanus</i>	1994-4	n = 67	Uzbekistan	Samarkand Region, Aman-Kutan	7 July 1994	VL	Lukhtanov and Dantchenko 2002a
<i>P. juldusus kasachstanus</i>	1997-3	n = 67	Kazakhstan	Dzhungarian Alatau, Kysylagash	June 2000	VL	Lukhtanov and Dantchenko 2002a
<i>P. karatavicus</i>	2000-K	n = 67	Kazakhstan	Karatau Mts, Minzhilgi	June 2000	VL	Lukhtanov et al. 2005
<i>P. phyllides askhabadicus</i>	F456	n = ca66-67	Iran	Kuh-e-Sorkh Mts., Torbat-e-Heydariyeh	7 July 2003	VL&D	This study
<i>P. phyllides askhabadicus</i>	VL03F523 AY954011	n = 67	Iran	Khorasan, Chakane	9 July 2003	VL&D	This study
<i>P. phyllides phyllides</i>	95204	n = ca65-67	Kyrgyzstan	Naryn Region, Moldatoo Mts, Chon-Konduk	26 July 1995	VL	This study
<i>P. phyllides phyllides</i>	NK00P672 AY496770	n = ca66-67	Kazakhstan	Kazakhstan, Kirgizski range,	June 2000	VL	Lukhtanov and Dantchenko 2002a
<i>P. phyllides phyllides</i>	NK00P808 AY496771	n = ca66-67	Kazakhstan	Kazakhstan, Karzhantau mts	June 2000	VL	Lukhtanov and Dantchenko 2002a

Discussion

Chromosomal stasis

It has been found that all taxa within *P. (A.) damone* species complex demonstrate chromosomal stasis and share the same or very similar haploid chromosomal number ($n = 66$ or $n = 67$). This result is unexpected since the karyotypes are known to be very diverse and species-specific in the subgenus *Agrodiaetus*.

It is believed that an unusual diversity of karyotypes is the most remarkable characteristic of *Agrodiaetus*. Species of this subgenus exhibit one of the highest ranges in chromosome numbers in the animal kingdom (Vershina and Lukhtanov 2017). In *Agrodiaetus* haploid chromosome numbers (n) range from $n = 10$ in *P. (A.) caeruleus* (Staudinger, 1871) to $n = 134$ in *P. (A.) shabrami* (Skala, 2001) (Lukhtanov et al. 2005). The genus *Polyommatus* as a whole shows numbers from $n = 10$ to $n = 226$ (Lukhtanov 2015). Additionally, the subgenus *Agrodiaetus* demonstrates a high level of karyotypic differentiation with respect to chromosome size (Lukhtanov and Dantchenko 2002b) and variation in number of chromosomes bearing ribosomal DNA clusters (Vershina et al. 2015). These differences provide reliable characters for species delimitation, description and identification (de Lesse 1960, 1963; Lukhtanov and Dantchenko 2002a, b).

The *P. (A.) damone* species complex represents an exception. In this group divergence in several phylogenetic lineages was not accompanied by changes in karyotypes, and the chromosome number $n = 66-67$ is the synapomorphic character for the species of the group.

DNA-barcode clusters

The DNA-barcode clusters revealed in our study correspond well to traditionally recognized species and certain specific geographic areas (Figs 3, 4). Cluster 1 includes

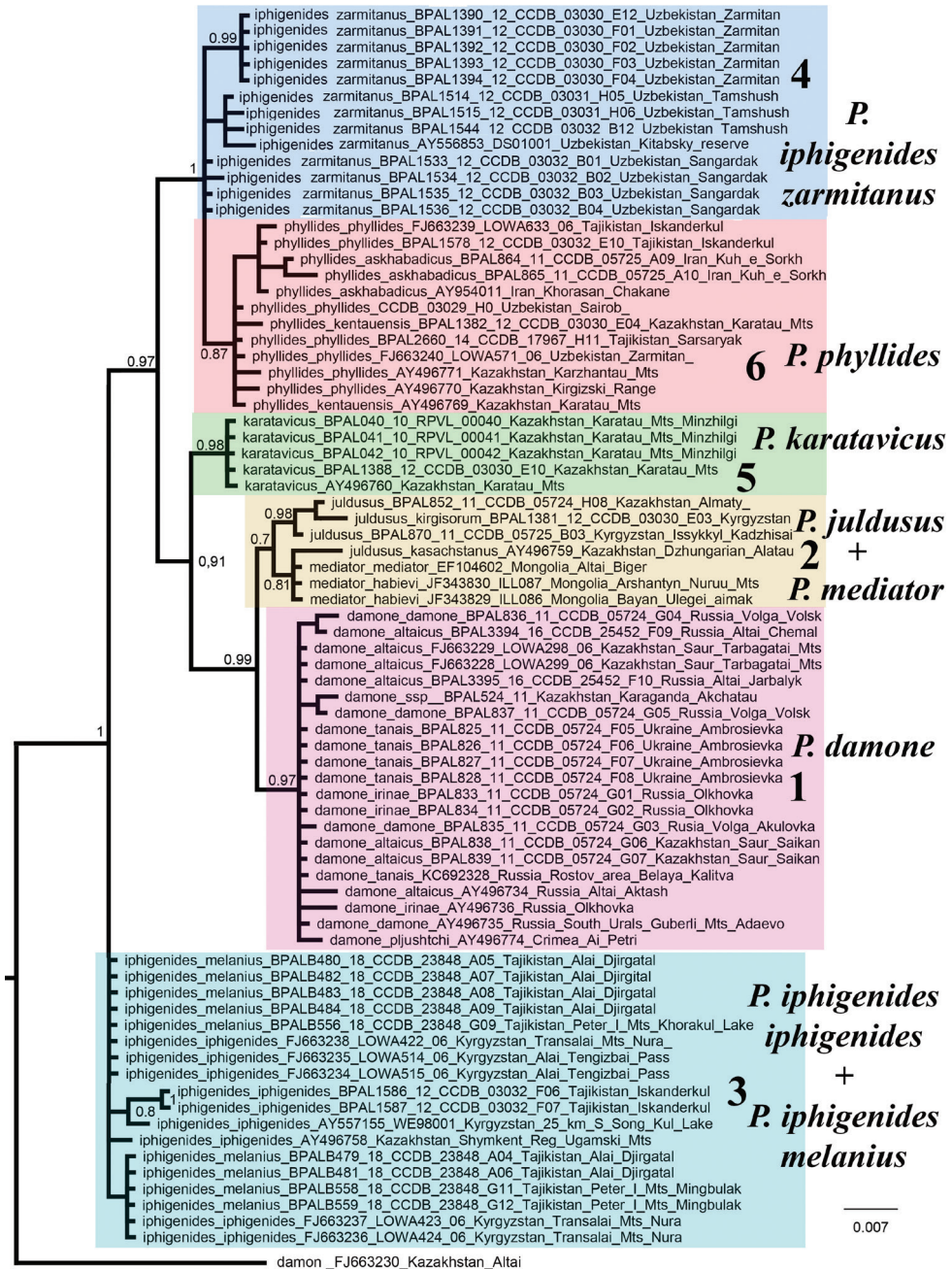


Figure 1. The Bayesian majority rule consensus tree of the analyzed samples of *Polyommatus* (*Agrodiaetus*) inferred from *COI* sequences. *Polyommatus icarus* is used to root the tree (not shown). Species and subspecies names, GenBank accession numbers, museum ID numbers and localities are shown to the right of the branches. Bayesian posterior probabilities higher than 0.5 are shown next to the recovered branches. 1–6 are clusters (see explanation in the text).

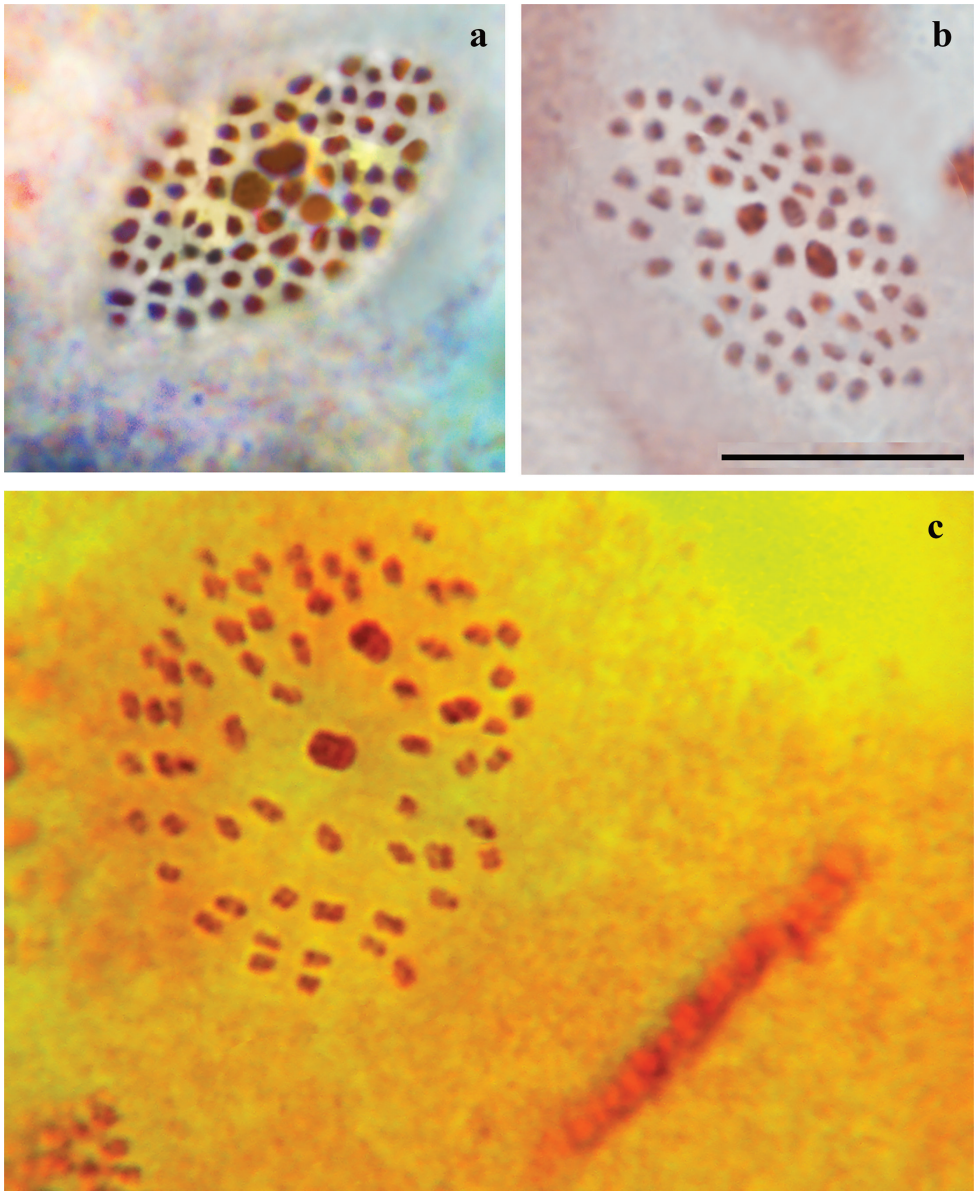


Figure 2. Karyotypes of *Polyommatus (Agrodiaetus) iphigenides melanius* and *P. (A.) phyllides askhabadicus*
a *P. (A.) iphigenides melanius*, sample 077K18A, MI, n = 67, phase-contrast **b** *P. (A.) iphigenides melanius*, sample 068K18A, MI, n = 66 **c** *P. (A.) phyllides askhabadicus*, sample F523, MI, n = 67. Scale Bar: 10 μ m.

specimens from the Crimea in the west to Altai and Saur-Tarbagatai Mts in the east and corresponds to *P. (A.) damone*. Cluster 2 includes specimens from NE Kyrgyzstan, SE Kazakhstan and SW Mongolia and corresponds to *P. (A.) juldusus* + *P. (A.) mediator*. Cluster 3 includes specimens from western and southern Kyrgyzstan, southern



Figure 3. Distribution areas of the COI clusters revealed in this study. Cluster 1 corresponds to *P. damone*. Cluster 2 corresponds to *P. juldusus* + *P. mediator*. Cluster 3 corresponds to *P. iphigenides* (including *P. iphigenides melanius*). Cluster 4 corresponds to *P. zarmitanus*. Cluster 5 corresponds to *P. karatavicus*.



Figure 4. Distribution area of *P. phyllides* (cluster 6).

Kazakhstan and Tajikistan and corresponds to *P. (A.) iphigenides iphigenides* + *P. (A.) iphigenides melanius*. Cluster 4 includes specimens from West Hissar in Uzbekistan and western Tajikistan and corresponds to *P. (A.) iphigenides zarmitanus* subsp. nov. which will be described below. Cluster 5 includes specimens from Karatau Mts in Kazakhstan and corresponds to *P. (A.) karatavicus*. Cluster 6 (Fig. 4) includes specimens from northeastern Iran to southeastern Kazakhstan and corresponds to *P. (A.) phyllides*.

Cluster 6 (= *P. phyllides*) is sympatric with cluster 2 (= *P. juldusus*) in northern Kyrgyzstan and southeastern Kazakhstan, with cluster 3 (= *P. iphigenides iphigenides* + *P. iphigenides melanius*) in Kyrgyzstan and Tajikistan, with cluster 4 (= *P. iphigenides zarmitanus*) in Uzbekistan and western Tajikistan, with cluster 5 (= *P. karatavicus*) in Karatau Mts in Kazakhstan (Eckweiler and Bozano 2016; our personal observations).

Taxonomic interpretations

Clusters 1 (*P. damone*), 2 (*P. juldusus* + *P. mediator*) and 5 (*P. karatavicus*)

We follow previous research (Dantchenko 2000; Dantchenko and Churkin 2003, Lukhtanov et al. 2005) in interpreting clusters 1 (*P. damone*), 2 (*P. juldusus* and *P. mediator*) and 5 (*P. karatavicus*) (see Taxonomic conclusions below). *P. (A.) mediator* was described as a species which is intermediate between *P. (A.) damone* and *P. (A.) juldusus*, but more similar to *P. (A.) juldusus* due to specific white pubescence of the costal area of the forewings (Dantchenko and Churkin 2003). This conclusion is now supported by molecular data: on the phylogenetic tree it appears as a clade, which also includes *P. (A.) juldusus kasachstanus*, and as a sister clade to *P. (A.) juldusus juldusus* + *P. (A.) juldusus kirgisorum*.

Up to our knowledge there are no data on sympatry of *P. (A.) mediator* and *P. (A.) damone* in Mongolia as it was reported or supposed earlier (Bálint and Johnson 1987; Bálint 1989).

Cluster 3 (*P. iphigenides iphigenides* + *P. iphigenides melanius*)

Polyommatus (Agrodiaetus) iphigenides is highly polymorphic with regard to the black suffusion on the wing upperside and the marginal and submarginal part of the wing underside in males as well as the white streak on hindwings in both sexes. In extreme cases, the suffusion can be practically absent resembling the upperside in *P. damone* or may extend almost to the discal spot which is observed as a fixed feature in two other taxa, *P. iphigenides melanius* and *P. juldusus kirgisorum*. The white streak is also very variable from clear visibility to complete absence. The taxa *P. (A.) samusi* Korb, 2017 (syn. nov.) and *P. (A.) melanius komarovi* Korb, 2017 (syn. nov.) are mainly described on the base of such extreme forms of the same population. Therefore, we consider these taxa as junior subjective synonyms of *P. (A.) iphigenides iphigenides*.

Cluster 3 also includes the taxon described as *Lycaena kindermanni* var. *Melania* Staudinger, 1886. For a long time, due to lack of material it had been considered to be a melanized form of *P. (A.) iphigenides iphigenides* (e.g. Forster 1960). But in recent years it has been treated as a separate species *P. (A.) melanius* with a local, nearly dot-like distribution in the border area between southwestern Kyrgyzstan and eastern Tajikistan in the Kyzylsu/Surkhob River basin (Dantchenko 2000; Eckweiler and Bozano 2016). We found that DNA barcodes of *P. (A.) iphigenides* and *P. (A.) melanius* are identical or differ by non-fixed 1–2 nucleotide substitutions. The main feature of *P. (A.) melanius*, a wide dark marginal border on the fore- and hindwings, is quite stable for the diagnosis of the taxon; however, the tendency towards such a wide border is expressed in different populations of *P. (A.) iphigenides*, too. Therefore, this trait can be hardly considered a species-specific character. Here we argue that *P. (A.) melanius* is rather a

subspecies *P. (A.) iphigenides* than a species. However, this is not a final conclusion. There is indirect evidence in favour of a possible species status of *P. (A.) melanius*, e.g. the distribution areas of *P. (A.) iphigenides iphigenides* and *P. (A.) iphigenides melanius* almost touch each other, and an intergradation zone would be expected between them. However, such a zone is still unknown, and specimens of *P. (A.) iphigenides iphigenides* and *P. (A.) iphigenides melanius* from very close localities are clearly differentiated. We suppose that genome-wide analysis may be useful to verify the taxonomic status of *P. (A.) iphigenides melanius*.

Cluster 4 (*P. iphigenides zarmitanus*)

Morphologically this group is close to *P. iphigenides iphigenides*, whereas with regard to mitochondrial DNA it is close to sympatric species *P. phyllides* which is morphologically very different. In our opinion, two alternative evolutionary scenarios can explain this pattern.

Scenario 1

The cluster 4 (*P. iphigenides zarmitanus*) and the lineage 6 (*P. phyllides*) are sister species which recently evolved from a common ancestor by means of sympatric speciation.

Scenario 2

Cluster 3 (*P. iphigenides*) and cluster 4 (*P. iphigenides zarmitanus*) are sister taxa evolved in allopatry; therefore, they share an ancestral type of the wing pattern and coloration, although differentiated with respect to DNA barcodes. The similarity between completely sympatric cluster 4 (*P. iphigenides zarmitanus*) and lineage 6 (*P. phyllides*) is a result of ancient mitochondrial introgression.

Analysis of multiple nuclear markers is required in order to distinguish between these two scenarios. Scenario 2 seems to be more probable since mitochondrial introgression is not a rare phenomenon in butterflies (e.g. Gompert 2008; Cong et al. 2017) and is also documented in the subgenus *Polyommatus* (*Agrodiaetus*) (Lukhtanov et al. 2015). Therefore, below we describe the new lineage discovered in West Hissar region as a subspecies of *P. iphigenides*.

Cluster 6 (*P. phyllides*)

There is no doubt that the cluster 6 (*P. phyllides*) is a distinct species, since it is a monophyletic lineage (Fig. 1), which is morphologically and ecologically differentiated (Dantchenko 2000, Eckweiler and Bozano 2016) and sympatric with *P. (A.) iphigenides iphigenides*, *P. (A.) iphigenides melanius*, *P. (A.) iphigenides zarmitanus*, *P. (A.) karatavicus* and *P. (A.) juldusus*.

New subspecies description

Polyommatus (Agrodiaetus) iphigenides zarmitanus subsp. nov.

<http://zoobank.org/092F10F6-B5E7-46C3-AA26-3A4D82F1F6D7>

Holotype. (Fig. 5a, b), male, BOLD process ID BPAL1393-12, field # CCDB-03030_F03, GenBank accession number MW186966; Uzbekistan, Samarqand Region, Nuratau Mts, near Zarmitan village, 40.40°N, 66.69°E, 1300 m, 11–13 June 1994, V. Lukhtanov leg., deposited in the Zoological Institute of the Russian Academy of Science (St. Petersburg).

COI barcode sequence of the holotype. ACATTATATTTTATTTTGTG-
GAATTTGAGCAGGAATAGTAGGGACATCCCTAAGAATTTTAATCCGTAT-
AGAATTGAGAACTCCTGGATCCTTAATTGGAGACGATCAAATTTATAA-
TACTATTGTTACAGCCCATGCATTTATTATAATTTTTTTTATAGTTATAC-
CTATTATAATTGGGGGATTTGGTAATTGATTAGTTCCTTTAATATTAG-
GAGCACCTGATATAGCCTTCCCCCGATTAAATAATATAAGATTCTGATT-
ATTACCGCCATCATTAATACTACTAATTTCCAGAAGAATTGTAGAAAATG-
GAGCAGGAACAGGATGAACAGTTTACCCCCCACTTTCATCTAATATT-
GCACATAGAGGATCATCTGTAGATTTAGCAATTTTCTCTCTTCATT-
TAGCAGGAATTTCTTCAATTTTAGGAGCAATTAATTTTATTACAAC-
ATTATTAACATACGGGTAAATAATTTATCATTTGATCAAATATCATTATT-
TATTTGAGCAGTAGGAATTACAGCATTATTATTACTTTTATCTTTACCTG-
TATTAGCTGGAGCAATTACCATATTATTAACAGATCGAAACCTTAATACCT-
CATTCTTTGACCCAGCTGGTGGGGGAGATCCAATTTTATATCAACATTTA.

Paratypes. 39 males, 14 females: Uzbekistan, Samarqand Region, Nuratau Mts, near Zarmitan village, 40.40°N, 66.69°E, 1300 m, 11–13 June 1994, V. Lukhtanov leg. 2 males: Uzbekistan, Qashqadaryo Region (old spelling: Kashkadarya Region), Hissar Range (west), near Tamshush village, 38.98°N, 67.35°E, 1800 m, 18–20 June 1994, V. Lukhtanov leg. 20 males: Uzbekistan, Qashqadaryo Region (old spelling: Kashkadarya Region), Hissar Range (west), near Tamshush village, 38.98°N, 67.35°E, 1800 m, 5–7 July 1994, V. Lukhtanov leg. 11 males, 2 females: Uzbekistan, Surxondaryo Region (old spelling: Surkhandarya Region), Hissar Range (west), Sangardak, 38.55°N, 67.50°E, 1600 m, 2 July 1994, V. Lukhtanov leg. 60 males, 21 females: Uzbekistan, Samarqand Region, Zeravshansky Range, Aman-Kutan, 1800 m, 39.27°N, 66.90°E, 7 July 1994, V. Lukhtanov leg. 13 males: Tajikistan, Sughd Region, Zeravshansky Range, Padzhrud village, 39.37°N, 68.03°E, 1300 m, 17 males, 13 males, 10 June 1994. All above paratypes are deposited in the Zoological Institute of the Russian Academy of Science (St. Petersburg). 5 males: Uzbekistan, [Jizzakh region], Usmat vic., 1700 m, 30.06.1988, V. Tshikolovets leg., in State Darwin Museum, Moscow. 15 males: [Uzbekistan], Aman-Kutan near Samarqand, 20 June 1938, A. Tsvetaev leg., in State Darwin Museum, Moscow. 26 males, 1 female: [Tajikistan], Hisar-Alai, Zeravshansky Range, Farob, 2000 m, 4 July 1998, G.D. Samodurov leg., in State Darwin Museum, Moscow. 1 female: Ta-

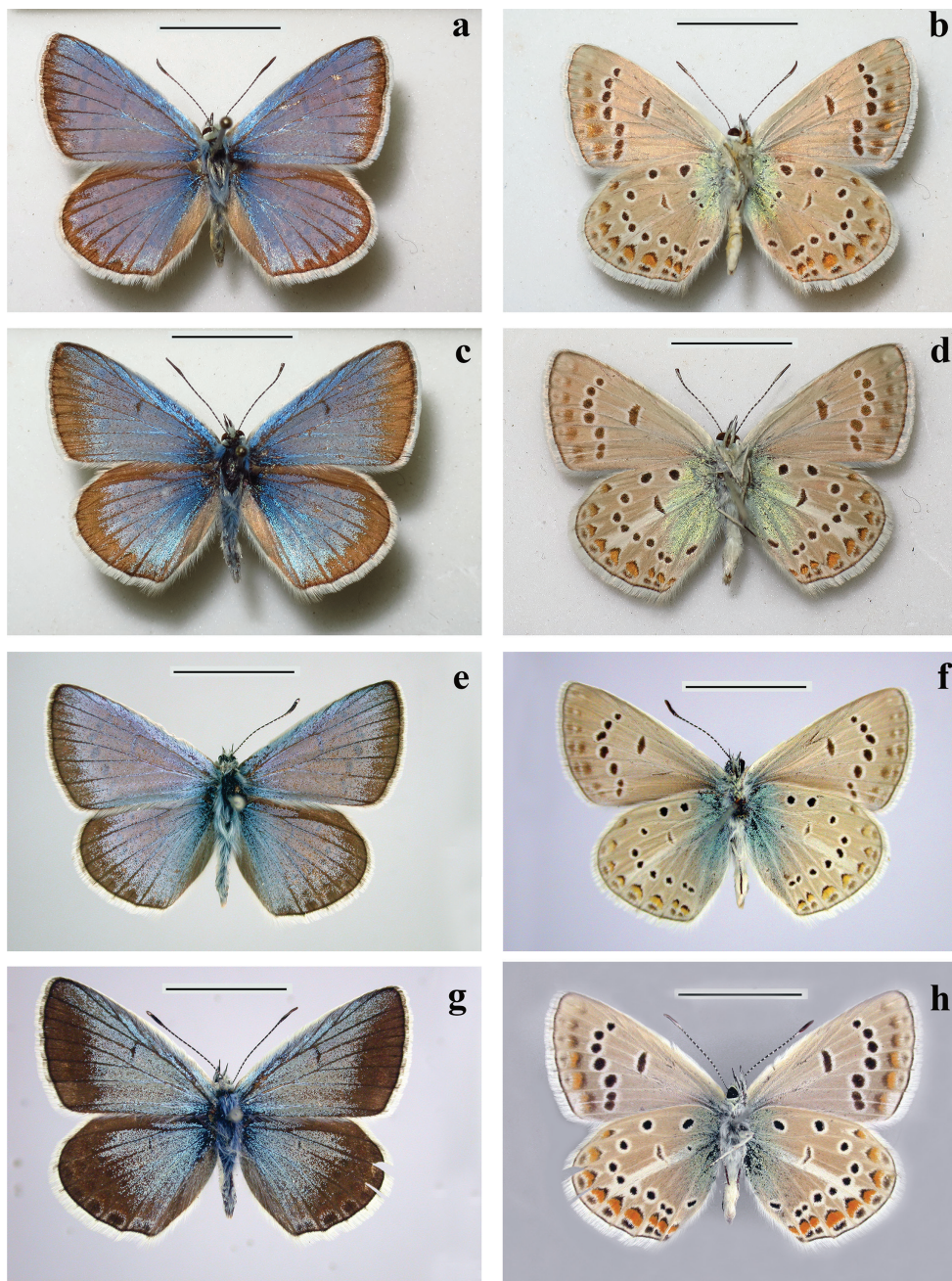


Figure 5. Males of *Polyommatus* (*Agrodiaetus*) *iphigenides* **a, b** upperside (**a**) and underside (**b**) of the holotype of *P. (A.) iphigenides zarmitanus* subsp. nov. **c, d** upperside (**c**) and underside (**d**) of *P. (A.) iphigenides iphigenides*, Tajikistan, Transalai Mts, Shibe village **e, f** upperside (**e**) and underside (**f**) of the Lectotype of *P. (A.) iphigenides iphigenides*, “Namangan”, in Museum für Naturkunde, Humboldt-Universität zu Berlin **g, h** upperside (**g**) and underside (**h**) of the Lectotype of *P. (A.) iphigenides melanius*, in Museum für Naturkunde, Humboldt-Universität zu Berlin. Scale Bars: 10 mm.

jikistan, West Hissar, Nofin lake, 2400, 17 July 1993, S. Churkin leg., in State Darwin Museum, Moscow. 32 males: [Uzbekistan], Aman-Kutan near Samarqand, 15–25 June 1938, A. Tsvetaev leg., in Zoological Museum Moscow University, Moscow (ZMMU). 7 males: [Uzbekistan], Aman-Kutan near Samarqand, 20–23 June 1938, G. Pashin leg., in ZMMU. 2 males: [Uzbekistan], Aman-Kutan near Samarqand, 27 July and 5 August 1937, A. G. Pashin leg., in ZMMU. 3 females: [Uzbekistan], Aman-Kutan near Samarqand, 15–26 June 1938, A. Tsvetaev leg., in ZMMU. 8 males: Tajikistan, West Hissar, Khazorchashma lake, 2800, 26 July 1993, S. Churkin leg.; 1 female: Tajikistan, West Hissar, Nofin lake, 2400, 17 July 1993, S. Churkin leg., in coll. Churkin (Reutov, Russia). 2 males, 1 female: Uzbekistan, West Hissar, Boysun Mts, Mochay, 1500 m, 26 June 1980, V. Tuzov leg., in coll. Tuzov (Moscow). 10 males, 1 female: [Uzbekistan], Aman-Kutan near Samarqand, 19–23 June 1938, A. Tsvetaev leg., in coll. Sochivko A. (Moscow). 10 males, 1 female: [Tajikistan], Hissar-Alai, Zeravshansky Range, Farob, 2000 m, 4 July 1998, L. Nikolaevsky leg., in coll. V. Kalinin, Moscow.

Description. Males. *Forewing* length 15–17 mm.

Upperside: Ground color bright glossy milky blue with narrow black marginal line, marginal part of forewings and hindwings dusted with black scales, discal strokes may be present or absent, veins darkened, costal area of the forewings white, hindwings with antemarginal spots, fringe white.

Underside: Forewing ground color light grey, submarginal row blurred, but clear visible; discoidal strokes black, bordered with white; postdiscal rows of black spots bordered with white, basal black spots absent; hindwing ground color light grey, basal area with strong greenish blue suffusion between wing root and basal spots; basal spots small, bordered with white, discal stroke less prominent than on forewings; postdiscal row of black spots bordered with white, submarginal and antemarginal marking strong and clear visible; submarginal row bordered distally with reddish lunules, more pronounced to anal end of row; white streak not contrasting, often hardly noticeable or absent at all, fringes pale grayish.

Genitalia. The male genitalia have a structure typical for other species of the subgenus *Agrodiaetus* (Coutsis 1986, Eckweiler and Bozano 2016).

Females. (Fig. 6a, b) *Forewing* length 15–17 mm.

Upperside: Ground color brown with slightly darker veins, discal strokes present, submarginal and antemarginal marking almost absent on fore wings and strong and clear visible on hindwings, antemarginal black spots on hindwings bordered with orange lunules, fringe whitish.

Underside: ground color and general design as in males but darker, brownish grey, greenish blue basal suffusion near invisible, white streak on hindwings clear visible, enlarged distally, fringe light greyish.

Diagnosis. The new subspecies is distinguished phenotypically from the most similar *P. iphigenides iphigenides* (Figs 5c–f, 6c, d) by the underside of the hind wing, which has a paler and less contrasting coloration. The white streak is also dim and weakly stands out against the background of the wing, is often reduced or absent. The same can be said about the basal greenish-blue suffusion: it is dim and weakly stands out

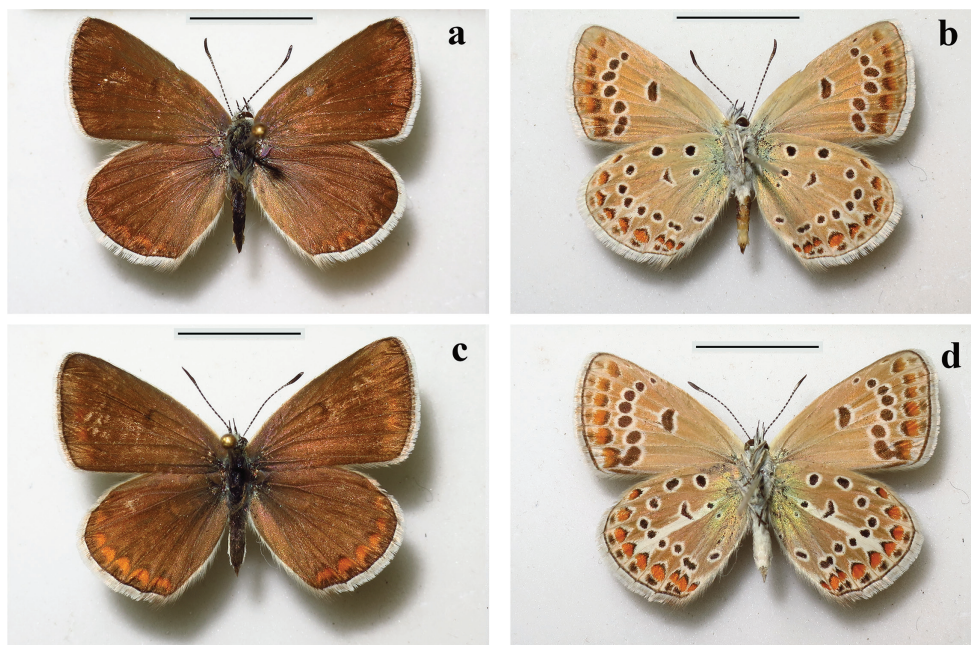


Figure 6. Females of *Polyommatus (Agrodiaetus) iphigenides* **a, b** upperside (**a**) and underside (**b**) of the paratype of *P. (A.) iphigenides zarmitanus* subsp. nov. **c, d** upperside (**c**) and underside (**d**) of *P. (A.) iphigenides iphigenides*, Tajikistan, Transalai Mts, Shibe village. Scale Bars: 10 mm.

against the background of the wing; its size, on average, is much smaller than that in *P. iphigenides iphigenides*. As a rule, it is limited by black dots of the basal row, while in *P. iphigenides iphigenides* it usually extends further in the distal direction, sometimes to spots of the discal row. This suffusion itself has a more greenish tint than that in *P. iphigenides iphigenides* (in the latter, it is more blue). The new species always has black dots of the basal row (although they are small), while in another species they are reduced.

The main differences between the species are still in the molecular characters. *Polyommatus iphigenides zarmitanus* can be distinguished from *P. iphigenides iphigenides* by using molecular markers from the *COI* gene. These mitochondrial diagnostic characters are in the following positions in the *COI* barcode region: adenine (A) in position 22, cytosine (C) in position 132, guanine (G) in position 180, cytosine (C) in position 286, guanine (G) in position 468, guanine (G) in position 468, and guanine (G) in position 627.

The new subspecies differs from sympatric (syntopic and synchronous) *P. phyllides* by milky blue (not greenish blue) wing upperside and white pubescence of the costal area of the forewings in males and by light grey color of the wing underside (*P. phyllides* has specific warm pinkish grey color of the wing underside). It also differs from *P. phyllides* by diagnostic nucleotide guanine (G) in position 627 of the *COI* barcode region.

Distribution area (Fig. 7). Uzbekistan: West part of the Hissar Range, Zeravshan Mts, Nuratau Mts, Boysun (= Baisuntau) Mts. Tajikistan: west part of the Zeravshan valley and Zeravshansky Range, West Hisar Range.



Figure 7. Distribution area of *P. (A.) iphigenides zarmitanus*. 1 is the type-locality, Zarmitan in Nuratau Mts. 2 is Padzhrud village in Zeravshansky Range. 3 is Aman-Kutan near Samarqand. 4 is Kitabsky Reserve in Hissar Range. 5 is Tamshush in Hissar Range. 6 is Sangardak in Hissar Range. 7 are Khazor-chashma and Nofin lakes in Hissar Range. 8 is Mochay in Boysun Mts.

Habitat and biology. Stony steppe and dry meadows from 1200 up to 2800 m alt. Flight period from late May to first decade of August in a single generation. The new subspecies flies syntopically and synchronously with the second generation of *P. (A.) phyllides*, but on average about one decade earlier. Host plant is preliminary determined as *Hedysarum* sp. (Fabaceae).

Etymology. The name *zarmitanus* is an adjective of the masculine gender. This name originates from Zarmitan, the village in Uzbekistan.

Taxonomic conclusion

The discovered topology (Fig. 1) can be considered as a signal to taxonomic rearrangement within the group. However, since the volume of the studied material of these taxa is small, we prefer to leave the existing taxonomic hypotheses. Additionally, we assume that the hypothesis of the existence of a species called *P. (A.) altaiensis* with subspecies

P. (A.) altaiensis altaiensis, *P. (A.) altaiensis bogdoolensis* Dantchenko et Lukhtanov, 1997, *P. (A.) altaiensis mediator* and *P. (A.) altaiensis habievi* Yakovlev, 2004 (Eckweiler and Bozano 2016) is speculative and not supported by significant morphological characters.

Based on the stated above, we propose the following taxonomic arrangement of the *P. damone* species complex:

***P. (A.) damone* (Eversmann, 1841)**

P. (A.) damone pljushtchi (Lukhtanov et Budashkin, 1993)

P. (A.) damone tanaïs Dantchenko et Pljushtch, 1993

P. (A.) damone irinae Dantchenko, 1997

P. (A.) damone damone (Eversmann, 1841)

P. (A.) damone altaicus (Elwes, 1899) (= *Lycaena damone* var. *sibirica* Staudinger, 1899; = *Agrodiaetus carmon altaiensis* Forster, 1956)

P. (A.) damone walteri Dantchenko et Lukhtanov, 1993

P. (A.) damone bogdoolensis Dantchenko et Lukhtanov, 1997

P. (A.) damone fabiani Bálint, 1997

***P. (A.) mediator* Dantchenko et Churkin, 2003** (= *Agrodiaetus mediator habievi* Yakovlev, 2004)

***P. (A.) juldusus* (Staudinger, 1886)** (= *Lycaena damone* var. *duplicata* A. Bang-Haas, 1910)

P. (A.) juldusus juldusus (Staudinger, 1886)

P. (A.) juldusus kirgisorum Lukhtanov et Dantchenko, 1994 (= *P. hyrsyz* Koçak et Kemal, 2001; = *P. kirgisorum khamul* Korb, 2009; = *P. kirgisorum gorthaur* Korb, 2009)

P. (A.) juldusus kasachstanus Lukhtanov et Dantchenko, 1994

P. (A.) juldusus rueckbeili Forster, 1960

P. (A.) juldusus tianchinensis Eckweiler, 2013

***P. (A.) iphigenides* (Staudinger, 1886)**

P. (A.) iphigenides iphigenides (Staudinger, 1886) (= *P. ishkashimicus alajanus* Korb, 1997; = *P. samusi* Korb, 2017, syn. nov.; = *P. melanius komarovi* Korb, 2017, syn. nov.)

P. (A.) iphigenides melanius (Staudinger, 1886)

P. (A.) iphigenides zarmitanus subsp. nov.

***P. (A.) karatavicus* Lukhtanov, 1990**

***P. (A.) phyllides* (Staudinger, 1886)**

P. (A.) phyllides phyllides (Staudinger, 1886)

P. (A.) phyllides askhabadicus (Forster, 1960)

P. (A.) phyllides kentauensis Lukhtanov, 1990

P. (A.) phyllides urumbash Churkin et Zhdanko, 2008

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References

- Bálint Z (1989) Recently collected lycaenid butterflies of Mongolia (V) (Lep., Lycaenidae). *Galathea* 5: 101–112.
- Bálint Z, Johnson K (1987) Reformation of the *Polyommatus* section with taxonomic and biogeographic overview (Lepidoptera, Lycaenidae, Polyommatini). *Neue Entomologische Nachrichten* 40: 1–68.
- Cong Q, Shen J, Borek D, Robbins RK, Opler PA, Otwinowski Z, Grishin NV (2017) When *COI* barcodes deceive: complete genomes reveal introgression in hairstreaks. *Proceedings of the Royal Society B – Biological Sciences* 284: 20161735. <https://doi.org/10.1098/rspb.2016.1735>
- Coutsis JG (1986) The blue butterflies of the genus *Agrodiaetus* Hübner (Lepidoptera, Lycaenidae): symptoms of taxonomic confusion. *Nota Lepidopterologica* 9: 159–169.
- Dantchenko AV (1997) Notes on the biology and distribution of the *damone* and *damocles* species-complexes of the subgenus *Polyommatus (Agrodiaetus)* (Lepidoptera: Lycaenidae). *Nachrichten des Entomologischen Vereins Apollo* 16: 23–42.
- Dantchenko AV (2000) Genus *Agrodiaetus*. In: Tuzov VK (Ed.) *Guide to the butterflies of Russia and adjacent territories*, Vol. 2. Sofia: Moscow, Pensoft, 196–214.
- Dantchenko A, Churkin S (2003) A new species of the genus *Polyommatus* Latreille, 1804 from Mongolia (Lepidoptera, Lycaenidae). *Neue Entomologische Nachrichten* 54: 5–13.
- Dantchenko A, Lukhtanov VA (1993) Zur Systematik und Verbreitung der Arten der *Polyommatus (Agrodiaetus) damone*-Gruppe (Lepidoptera, Lycaenidae) Südosteuropas und Südwestsibiriens. *Atalanta* 24(1/2): 75–83, 324–325.
- de Lesse H (1960) Speciation et variation chromosomique chez les Lepidopteres Rhopaloceras. *Annales des Sciences Naturelles Zoologie et Biologie Animale*, 12e série 2: 1–223.
- de Lesse H (1963) Variation chromosomique chez *Agrodiaetus carmon* H. S. et *A. cyanea* Stgr. *Revue Française d'Entomologie* 30: 177–181.

- deWaard JR, Ivanova NV, Hajibabaei M, Hebert PDN (2008) Assembling DNA barcodes: analytical protocols. In: Martin CC (Ed.) *Environmental Genomics, Methods in Molecular Biology*. Humana Press, Totowa, New Jersey, 410: 275–283. https://doi.org/10.1007/978-1-59745-548-0_15
- Eckweiler W, Bozano GC (2016) Guide to the butterflies of the Palearctic region. *Lycaenidae* part IV. Milano, Omnes artes, 132 pp.
- Forster W (1956) Bausteine zur Kenntnis der Gattung *Agrodiaetus* Scudd. (Lep. Lycaen) I. *Zeitschrift der Wiener Entomologischen Gesellschaft* 41: 42–61, 70–89, 118–127.
- Forster W (1960) Bausteine zur Kenntnis der Gattung *Agrodiaetus* Scudd. (Lep. Lycaen) II. *Zeitschrift der Wiener Entomologischen Gesellschaft* 45: 105–142.
- Gompert Z, Forister ML, Fordyce JA, Nice CC (2008) Widespread mito-nuclear discordance with evidence for introgressive hybridization and selective sweeps in *Lycaeides*. *Molecular Ecology* 17: 5231–5244. <https://doi.org/10.1111/j.1365-294X.2008.03988.x>
- Hajibabaei M, deWaard JR, Ivanova NV, Ratnasingham S, Dooph RT, Kirk SL, Mackie PM, Hebert PDN (2005) Critical factors for assembling a high volume of DNA barcodes. *Philosophical Transactions of the Royal Society of London – Series B, Biological Sciences* 360: 1959–1967. <https://doi.org/10.1098/rstb.2005.1727>
- Hall TA (1999) BioEdit: a user-friendly biological sequence alignment editor and analysis program for Windows 95/98/NT. *Nucleic Acids Symposium Series* 41: 95–98.
- Ivanova NV, deWaard JR, Hebert PDN (2006) An inexpensive, automation-friendly protocol for recovering high quality DNA. *Molecular Ecology Resources* 6: 998–1002. <https://doi.org/10.1111/j.1471-8286.2006.01428.x>
- Kandul NP (1997) The karyology and the taxonomy of the blue butterflies of the genus *Agrodiaetus* Hübner [1822] from the Crimea (Lepidoptera: Lycaenidae). *Atalanta* 28: 111–119.
- Kandul NP, Lukhtanov VA, Dantchenko AV, Coleman JWS, Sekercioglu CH, Haig D, Pierce NE (2004) Phylogeny of *Agrodiaetus* Hübner 1822 (Lepidoptera: Lycaenidae) Inferred from mtDNA Sequences of *COI* and *COII* and Nuclear Sequences of *EF1-α*. *Karyotype Diversification and Species Radiation. Systematic Biology* 53(2): 278–298. <https://doi.org/10.1080/10635150490423692>
- Kandul NP, Lukhtanov VA, Pierce NE (2007) Karyotypic diversity and speciation in *Agrodiaetus* butterflies. *Evolution* 61(3): 546–559. <https://doi.org/10.1111/j.1558-5646.2007.00046.x>
- Lukhtanov VA (1989) Karyotypes of some blue butterflies of the *Agrodiaetus* species groups. *Annales Entomologici Fennici* 55: 137–144.
- Lukhtanov VA (2015) The blue butterfly *Polyommatus (Plebicula) atlanticus* (Lepidoptera, Lycaenidae) holds the record of the highest number of chromosomes in the non-polyploidy eukaryotic organisms. *Comparative Cytogenetics* 9(4): 683–690. <https://doi.org/10.3897/CompCytogen.v9i4.5760>
- Lukhtanov VA, Dantchenko AV (2002a) Principles of highly ordered metaphase I bivalent arrangement in spermatocytes of *Agrodiaetus* (Lepidoptera). *Chromosome Research* 10(1): 5–20. <https://doi.org/10.1023/A:1014249607796>
- Lukhtanov VA, Dantchenko AV (2002b) Descriptions of new taxa of the genus *Agrodiaetus* Hübner, [1822] based on karyotype investigation (Lepidoptera, Lycaenidae). *Atalanta* 33(1/2): 81–107, 224–225.

- Lukhtanov VA, Dantchenko AV, Kandul NP (1997) Die Karyotypen von *Polyommatus (Agrodiaetus) damone damone* und *P. (A.) damocles rossicus* nebst einigen Problemen bei *Agrodiaetus* (Lepidoptera: Lycaenidae). Nachrichten des Entomologischen Vereins Apollo Supplement 16: 43–48.
- Lukhtanov VA, Kandul NP, Plotkin JB, Dantchenko AV, Haig D, Pierce NE (2005) Reinforcement of pre-zygotic isolation and karyotype evolution in *Agrodiaetus* butterflies. Nature 436(7049): 385–389. <https://doi.org/10.1038/nature03704>
- Lukhtanov VA, Vila R, Kandul NP (2006) Rearrangement of the *Agrodiaetus dolus* species group (Lepidoptera, Lycaenidae) using a new cytological approach and molecular data. Insect and Systematic and Evolution 37(3): 325–334. <https://doi.org/10.1163/187631206788838563>
- Lukhtanov VA, Shapoval NA, Dantchenko AV (2008) *Agrodiaetus shahkuhensis* sp. n. (Lepidoptera, Lycaenidae), a cryptic species from Iran discovered by using molecular and chromosomal markers. Comparative Cytogenetics 2(2): 99–114.
- Lukhtanov VA, Sourakov A, Zakharov EV, Hebert PDN (2009) DNA barcoding Central Asian butterflies: increasing geographical dimension does not significantly reduce the success of species identification. Molecular Ecology Resources 9: 1302–1310. <https://doi.org/10.1111/j.1755-0998.2009.02577.x>
- Lukhtanov VA, Shapoval NA, Dantchenko AV (2014) Taxonomic position of several enigmatic *Polyommatus (Agrodiaetus)* species (Lepidoptera, Lycaenidae) from Central and Eastern Iran: insights from molecular and chromosomal data. Comparative Cytogenetics 8(4): 313–322. <https://doi.org/10.3897/CompCytogen.v8i4.8939>
- Lukhtanov VA, Shapoval NA, Anokhin BA, Saifitdinova AF, Kuznetsova VG (2015) Homoploid hybrid speciation and genome evolution via chromosome sorting. Proceedings of the Royal Society B – Biological Sciences 282(1807): 20150157. <https://doi.org/10.1098/rspb.2015.0157>
- Lukhtanov VA, Sourakov A, Zakharov EV (2016) DNA barcodes as a tool in biodiversity research: testing pre-existing taxonomic hypotheses in Delphic Apollo butterflies (Lepidoptera, Papilionidae). Systematics and Biodiversity 14(6): 599–613. <https://doi.org/10.1080/14772000.2016.1203371>
- Maddison WP, Maddison DR (2015) Mesquite: a modular system for evolutionary analysis. Version 3.04. <http://mesquiteproject.org>
- Przybyłowicz Ł, Lukhtanov V, Lachowska-Cierlik D (2014) Towards the understanding of the origin of the Polish remote population of *Polyommatus (Agrodiaetus) ripartii* (Lepidoptera: Lycaenidae) based on karyology and molecular phylogeny. Journal of Zoological Systematics and Evolutionary Research 52(1): 44–51. <https://doi.org/10.1111/jzs.12040>
- Rambaut A, Drummond AJ, Xie D, Baele G, Suchard MA (2018) Posterior summarisation in Bayesian phylogenetics using Tracer 1.7. Systematic Biology 67: 901–904. <https://doi.org/10.1093/sysbio/syy032>
- Ronquist F, Teslenko M, van der Mark P, Ayres DL, Darling A, Höhna S, Larget B, Liu L, Suchard MA, Huelsenbeck JP (2012) MrBayes 3.2: efficient Bayesian phylogenetic inference and model choice across a large model space. Systematic Biology 61: 539–542. <https://doi.org/10.1093/sysbio/sys029>
- Staudinger O (1899) Ueber die Arten und Formen der *Lycaena damon*-Gruppe. Deutsche Entomologische Zeitschrift Iris 12: 137–155.

- Vershinina AO, Lukhtanov VA (2010) Geographical distribution of the cryptic species *Agrodiaetus alcestitis alcestitis*, *A. alcestitis karacetinae* and *A. demavendi* (Lepidoptera, Lycaenidae) revealed by cytogenetic analysis. *Comparative Cytogenetics* 4(1): 1–11. <https://doi.org/10.3897/compcytogen.v4i1.21>
- Vershinina AO, Anokhin BA, Lukhtanov VA (2015) Ribosomal DNA clusters and telomeric (TTAGG)_n repeats in blue butterflies (Lepidoptera, Lycaenidae) with low and high chromosome numbers. *Comparative Cytogenetics* 9(2): 161–171. <https://doi.org/10.3897/CompCytogen.v9i2.4715>
- Vershinina AO, Lukhtanov VA (2017) Evolutionary mechanisms of runaway chromosome number change in *Agrodiaetus* butterflies. *Scientific Reports* 7: e8199. <https://doi.org/10.1038/s41598-017-08525-6>
- Vishnevskaya MS, Saifitdinova AF, Lukhtanov VA (2016) Karyosystematics and molecular taxonomy of the anomalous blue butterflies (Lepidoptera, Lycaenidae) from the Balkan Peninsula. *Comparative Cytogenetics* 10(5): 1–85. <https://doi.org/10.3897/CompCytogen.v10i5.10944>
- Vodolazhsky D, Stradomsky B (2012) Comparative genital and molecular genetic analysis of blue butterflies (Lepidoptera: Lycaenidae) of Rostov-on-Don area. In: Troitsky AV, Rusin LY, Aleoshin VV (Eds) *Molecular Phylogenetics*. Moscow, Torus Press, 70–71.
- Vodolazhsky DI, Yakovlev R, Stradomsky B (2011) Study of taxonomic status of some specimens of subgenus *Agrodiaetus* (Lepidoptera: Lycaenidae: *Polyommatus*) from Western Mongolia based on mtDNA markers. *Caucasian Entomological Bulletin* 7(1): 81–82. <https://doi.org/10.23885/1814-3326-2011-7-1-81-82>
- Wiemers M (2003) Chromosome Differentiation and the Radiation of the Butterfly Subgenus *Agrodiaetus* (Lepidoptera: Lycaenidae: *Polyommatus*) a Molecular Phylogenetic Approach. PhD Thesis. University of Bonn, Bonn, 203 pp. <http://hss.ulb.uni-bonn.de/2003/0278/0278.htm>