

Ioannis Tsiripidis, Stavroula Stamellou, Aristotelis C. Papageorgiou, Andreas D. Drouzas

Supplementary materials

Table S1. Synoptic table of taxa relative frequencies in the five vegetation assemblages of *Fagus sylvatica* on Mt. Menikio (A to E). Relative frequencies with bold typescript indicate that taxa differentiate positively the corresponding vegetation assemblages, with italic typescript that they differentiate them negatively and with regular typescript that they do not differentiate the corresponding vegetation groups. Taxa with relative frequency smaller than 20% in any column have been omitted.

Species assemblage	Species	A	B	C	D	E
No of plots	abbrev.	10	22	9	7	12
<u>Diagnostic taxa of assemblage A (calcicolous and thermophilous species)</u>						
<i>Epipactis helleborine</i>	EpipHel	90	36	11	29	17
<i>Epipactis microphylla</i>	EpipMic	40	9	11	0	0
<i>Hypopitys monotropa</i>	HypoMon	60	9	11	29	50
<i>Brachypodium sylvaticum</i> ssp. <i>sylvaticum</i>	BracS-S	30	14	11	0	0
<u>Diagnostic taxa of assemblages A and C (calcicolous species)</u>						
<i>Cephalanthera rubra</i>	CephRub	90	23	67	14	0
<i>Lilium martagon</i>	LiliMar	30	0	67	0	0
<i>Cephalanthera damasonium</i>	CephDam	80	18	56	14	0
<i>Helleborus odorus</i> ssp. <i>cyclophyllus</i>	HellO-C	50	27	56	14	0
<i>Lonicera xylosteum</i>	LoniXyl	20	5	22	0	0
<u>Diagnostic taxa of assemblage B (thermophilous, nitrophilous species)</u>						
<i>Pteridium aquilinum</i>	PterAqu	30	95	44	86	17
<i>Rubus hirtus</i>	RubuHir	20	95	11	57	25
<i>Carex sylvatica</i>	CareSyl	0	32	0	0	0
<i>Sambucus nigra</i>	SambNig	0	36	11	0	0

<i>Urtica dioica</i>	UrtiDio	10	27	0	0	0
<i>Epilobium lanceolatum</i>	EpilLan	0	27	11	0	8
<i>Carpinus orientalis</i>	CarpOri	0	23	0	14	0
<u>Diagnostic taxa of assemblages B and C (mostly species of beech forests)</u>						
<i>Actaea spicata</i>	ActaSpi	10	32	56	0	0
<i>Rosa canina</i>	RosaCan	0	45	33	0	8
<i>Fragaria vesca</i>	FragVes	10	36	22	0	0
<i>Geranium robertianum</i>	GeraRob	10	32	22	0	0
<i>Epilobium montanum</i>	EpilMon	0	32	22	0	17
<i>Galium odoratum</i>	GaliOdo	10	41	44	29	8
<u>Diagnostic taxa of assemblage C (mostly species of ravine forests, calcicolous, oceanic and moist indicator species)</u>						
<i>Mercurialis perennis</i>	MercPer	40	5	100	0	0
<i>Lathyrus venetus</i>	LathVen	10	9	78	14	0
<i>Tilia platyphyllos</i>	TiliPla	0	0	44	0	0
<i>Ilex aquifolium</i>	IlexAqu	0	5	44	0	0
<i>Acer platanoides</i>	AcerPla	0	9	44	0	0
<i>Aquilegia vulgaris</i>	AquiVul	0	0	33	0	0
<i>Fritillaria pontica</i>	FritPon	0	0	33	0	0
<i>Euonymus latifolius</i>	EuonLat	20	68	100	29	25
<i>Acer hyrcanum</i>	AcerHyr	30	9	78	29	8
<i>Viola sieheana</i>	ViolSie	0	14	44	0	0
<i>Asplenium adiantum-nigrum</i>	AsplAdi	0	5	67	14	33
<i>Epipactis leptochila</i> ssp. <i>naousaensis</i>	EpipL-N	0	5	33	0	0
<i>Fraxinus ornus</i>	FraxOrn	20	55	100	43	75
<i>Platanthera chlorantha</i>	PlatChl	0	5	44	14	0
<i>Acer campestre</i>	AcerCam	0	9	44	14	0
<i>Ajuga reptans</i>	AjugRep	0	14	44	0	8
<i>Heracleum sphondylium</i>	HeraSph	20	5	44	0	0
<i>Milium effusum</i>	MiliEff	0	18	33	0	0
<i>Euonymus europaeus</i>	EuonEur	0	9	22	0	0
<i>Malus sylvestris</i>	MaluSyl	0	9	22	0	0

Diagnostic taxa of assemblages A, B and C (oceanic, thermophilous and moist indicator species)

<i>Hedera helix</i>	HedeHel	40	73	100	0	17
<i>Daphne laureola</i>	DaphLau	60	36	78	0	0
<i>Clematis vitalba</i>	ClemVit	20	45	67	0	0

Diagnostic taxa of assemblages A, B, C and D (species of deciduous broadleaved forests; mainly beech forests)

<i>Aremonia agrimonoides</i> ssp. <i>agrimonoides</i>	AremA-A	80	64	89	71	0
<i>Viola riviniana</i>	ViolRiv	80	86	100	86	25
<i>Physospermum cornubiense</i>	PhysCor	40	23	89	86	0
<i>Rosa arvensis</i>	RosaArv	60	59	67	100	0
<i>Euphorbia amygdaloides</i> ssp. <i>amygdaloides</i>	EuphA-A	30	59	56	100	0
<i>Polygonatum odoratum</i>	PolyOdo	40	41	78	71	0
<i>Cardamine bulbifera</i>	CardBul	40	82	67	43	8
<i>Sanicula europaea</i>	SaniEur	20	50	44	57	0

Diagnostic taxa of assemblages B, C and D (species of deciduous broadleaved forests; mainly oak forests)

<i>Melica uniflora</i>	MeliUni	10	32	67	100	0
<i>Prunus avium</i>	PrunAvi	0	50	56	57	17
<i>Lathyrus laxiflorus</i>	LathLax	10	41	67	57	8
<i>Veronica chamaedrys</i> ssp. <i>chamaedrys</i>	VeroC-C	0	32	33	29	8
<i>Viola alba</i> et <i>odorata</i>	Viola#O	20	36	56	43	8

Diagnostic taxon of assemblages B, C, D and E

(indicator species of oak and thermophilous beech forests)

<i>Luzula forsteri</i>	LuzuFor	0	32	33	71	42
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Diagnostic taxa of assemblages C and D (species of oak and thermophilous ravine forests, mostly moist indicator species)

<i>Sorbus torminalis</i>	SorbTor	10	5	100	86	0
<i>Primula vulgaris</i>	PrimVul	0	18	67	86	0
<i>Cornus mas</i>	CornMas	30	41	100	71	8
<i>Lathyrus niger</i>	LathNig	0	5	33	71	8

<i>Carpinus betulus</i>	CarpBet	10	5	33	71	0
<i>Dioscorea communis</i>	DiosCom	0	5	56	29	0
<i>Corylus avellana</i>	CoryAve	50	82	100	100	33
<i>Campanula trachelium</i> ssp. <i>athoia</i>	CampT-A	0	9	33	43	0
<i>Scutellaria columnae</i> ssp. <i>columnae</i>	ScutC-C	0	5	33	29	0
<i>Tilia tomentosa</i>	TiliTom	10	5	33	43	0
<i>Aegopodium podagraria</i>	AegoPod	0	9	33	29	0
<i>Campanula rapunculoides</i>	CampRap	20	9	44	43	0
<i>Cyclamen hederifolium</i>	CyclHed	0	14	33	29	0
<i>Bromus benekenii</i>	BromBen	0	14	33	29	0

Diagnostic taxa of the assemblage D (indicator species of mesic and thermophilous beech forests mainly of NE Greece)

<i>Pulmonaria rubra</i>	PulmRub	0	5	0	71	0
<i>Melittis melissophyllum</i>	MeliMel	0	0	22	71	0
<i>Quercus petraea</i> ssp. <i>polycarpa</i>	QuerP-P	10	23	11	100	42
<i>Potentilla micrantha</i>	PoteMic	10	59	44	100	33
<i>Symphytum tuberosum</i>	SympTub	0	18	0	43	0
<i>Festuca heterophylla</i>	FestHet	30	5	11	43	0
<i>Sedum cepaea</i>	SeduCep	0	5	0	29	8
<i>Lathyrus vernus</i>	LathVer	0	5	11	29	0

Diagnostic taxon of assemblages D and E (diagnostic taxon of beech forests of NE Greece)

<i>Carex digitata</i>	CareDig	50	50	22	86	92
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Diagnostic taxa of assemblage E (acidic or thermophilous species)

<i>Avenella flexuosa</i>	AvenFle	0	0	0	0	50
<i>Hieracium transiens</i> ssp. <i>erythrocarpum</i> et <i>H. rechingerorum</i>	HierE#R	0	0	0	0	50
<i>Luzula luzuloides</i>	LuzuLuz	20	55	11	14	83
<i>Veronica officinalis</i>	VeroOff	0	36	0	0	42
<i>Polypodium vulgare</i>	PolyVul	30	14	56	14	83
<i>Campanula persicifolia</i>	CampPer	0	0	0	14	42
<i>Hieracium olympicum</i> et <i>racemosum</i>	HierR#O	10	5	11	0	50
<i>Viscaria atropurpurea</i>	ViscAtr	0	5	0	0	33

<i>Hypericum cerastioides</i>	HypeCer	0	0	0	0	25
<i>Hieracium murorum</i>	HierMur	10	9	22	14	58
<i>Campanula patula</i> ssp. <i>patula</i>	CampP-P	0	18	11	14	50
<i>Quercus coccifera</i>	QuerCoc	10	9	0	0	33
<i>Juniperus communis</i>	JuniCom	0	9	0	14	33
<i>Orthilia secunda</i>	OrthSec	20	9	0	0	25
<i>Prunus cerasifera</i>	PrunCer	0	23	11	14	42
<u>Companion and common species</u>						
<i>Fagus sylvatica</i>	FaguSyl	100	100	100	100	100
<i>Neottia nidus-avis</i>	NeotNid	80	77	67	57	58
<i>Poa nemoralis</i>	Poa Nem	30	73	78	57	83
<i>Lactuca muralis</i>	LactMur	10	55	67	29	42
<i>Castanea sativa</i>	CastSat	10	32	33	29	33
<i>Dryopteris filix-mas</i>	DryoFil	20	36	33	14	8
<i>Moehringia trinervia</i>	MoehTri	0	18	33	43	42
<i>Euonymus verrucosus</i>	EuonVer	40	23	22	0	17
<i>Populus tremula</i>	PopuTre	30	18	0	43	8
<i>Asplenium trichomanes</i>	AsplTri	10	14	22	29	17
<i>Ligustrum vulgare</i>	LiguVul	20	18	22	0	0
<i>Acer pseudoplatanus</i>	AcerPse	10	9	33	14	0
<i>Dactylis glomerata</i>	DactGlo	0	18	22	0	8
<i>Crataegus monogyna</i>	CratMon	0	14	11	29	0
<i>Lamium maculatum</i>	LamiMac	10	9	22	0	0
<i>Ostrya carpinifolia</i>	OstrCar	20	9	11	0	0
<i>Cystopteris fragilis</i>	CystFra	0	0	22	14	0
<i>Luzula sylvatica</i>	LuzuSyl	20	0	11	0	0

Table S2: Fragments' length (in base pairs), assignment and frequencies of the *Fagus sylvatica* haplotypes identified by three polymorphic microsatellites (ccmp10, ccmp7, ccmp4) on Mt. Menikio.

Haplotype	ccmp 10	ccmp 7	ccmp 4	Observations	%
h1	110	148	114	3	1.7
h2	115	146	115	9	5.1
h3	110	147	115	10	5.6
h4	110	148	115	129	72.6
h5	110	149	115	14	7.8
h6	110	150	115	8	4.5
h7	110	151	115	1	0.5
h8	110	148	116	2	1.1
h9	115	147	115	2	1.1
Total				178	100

Table S3: Analysis of Molecular Variance (AMOVA) for geographic groups (GGs) of *Fagus sylvatica* on Mt. Menikio using haplotype distance based on fragment length differences.

Partition of diversity (%)	Distance between haplotypes	Differences in fragment length
Among GGs		35.22
Among plots within GGs		20.92
Within plots		43.86
Φ_{st} (p-value)		0.352 (0.000)

Table S4: Analysis of Molecular Variance (AMOVA) for plant assemblages of *Fagus sylvatica* on Mt. Menikio using haplotype distance based on fragment length differences.

Partition of diversity (%)	Distance between haplotypes	Differences in fragment length
Among assemblages		42.72
Among plots within assemblages		20.07
Within plots		37.21
Φ_{st} (p-value)		0.427 (0.000)