

## THE IMPACT OF UNMOWN REFUGE-STRIPS ON THE BREEDING SITE FIDELITY OF COMMON QUAIL (*Coturnix coturnix*) – A CASE STUDY

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### ABSTRACT

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Common Quail (*Coturnix coturnix*) mainly breed in cereal crops and fallow grasslands, where they are threatened by harvesting/mowing. The aim of this research was to assess the impact of unmown refuge areas to the density and movements of Common Quails in the Moson Project, Northwest Hungary. The selected 80 ha study area was first visited shortly before mowing, while the second survey was carried out two weeks after mowing operation ended. Calling males of Common Quail were documented and their vocalizations were recorded. During the first survey (before mowing), a total of 18 Common Quails were recorded, while the second survey (after mowing) resulted in a lower number (14) of birds detected. In terms of density, the initially determined 2.25 calling males/10 ha decreased to 1.75 males/10 ha. For individual recognition bioacoustic methods were used. A total of six time and frequency-based variables were measured and were subjected to discriminant function analysis (DFA). A total of 9 males were re-identified with high probability, proving that the 15-20 m wide unmown refuge-srips can still provide optimal habitats for quails.

**KEY WORDS:** farmland birds, mowing, refuge areas, agricultural practices, habitat selection

### KIVONAT

NÉMETH T.M. & WINKLER D.: BÚVÓSÁVOK HATÁSA A FÜRJ (*Coturnix coturnix*) TERÜLETHŰSÉGÉRE FÉSZKELÉSI IDŐSZAKBAN – ESETTANULMÁNY. *Magyar Ápróvad Közlemények* **13**: 289–296. <http://dx.doi.org/10.17243/mavk.2017.289>

A fürj (*Coturnix coturnix*) a mezei élőhelyek jellegzetes fészkelő madara. Költ a különböző mezőgazdasági kultúrákban, ugar jellegű élőhelyeken, ahol a betakarítás illetve kaszálás nagy veszélyt jelent a fészkelési időszakban. Jelen kutatás a kaszálatlanul hagyott búvósávok szerepét vizsgálja a fürjek denzitása valamint területhűsége vonatkozásában egy északnyugat-magyarországi élőhelyen (MOSON Project). A fürjek felmérésére a vizsgálatokhoz kiválasztott mintegy 80 hektár nagyságú területen először a kaszálás időpontja előtt, majd ezt követően a kaszálás befejezése után két héttel került sor. Az éneklő kakasok számának feljegyzése mellett hangfelvételeket is készítettünk. Az első felmérés során összesen 18, majd a kaszálás után 14 hím egyedet detektáltunk, így a denzitás 2.25 éneklő kakas/10 ha-ról 1.75 éneklő kakas/10 ha értékre csökkent a kaszálást követően. Az éneklő hím egyedek elkülönítése bioakusztikai módszerekkel történt, amihez a fürj hangját jól reprezentáló hat hangfizikai változó mérését végeztük el a rögzített hangmintákon. A hangfizikai változók adatmátrixát diszkriminancia-analízis (DFA) segítségével elemeztük. Összesen 9 fürj kakas újraazonosítása sikerült nagy biztonsággal, amely alapján azt a következtetést vonhatjuk le, hogy a kaszálatlanul hagyott 15-20 m széles búvósávok megfelelő élőhelyet tudnak biztosítani a fürj számára.

**KEY WORDS:** mezei madárfajok, kaszálás, búvósávok, mezőgazdasági üzemmód, élőhelyválasztás

## 1. INTRODUCTION

The Common Quail (*Coturnix coturnix*) is a widely distributed breeding species mostly associated with farmland areas in Europe (CRAMP, 1980; MCGOWAN *et al.* 2013). The population trend declining in many European countries (BIRDLIFE INTERNATIONAL, 2016) owing to the intensive agricultural practices, the use of pesticides and heightened mortality during migration (SANDERSON *et al.*, 2009, KOSICKI *et al.*, 2014). Additionally, Common Quails are subjected to significant hunting pressure during the autumn migration period, especially in the Mediterranean countries (TUCKER & HEATH, 1994; GALLEGO *et al.*, 1997; PUIGSERVER *et al.*, 1998). In Hungary, the Common Quail is a protected species. Nevertheless, its population shows moderate decline both locally and countrywide (NÉMETH *et al.*, 2014; MAGYAR MADÁRTANI ÉS TERMÉSZETVÉDELMI EGYESÜLET, 2017).

In Hungary, major threats are mainly linked to agricultural practices (MÁRKUS, 1998; BÁLDI & BATÁRY, 2011; NÉMETH *et al.*, 2014; FARAGÓ, 2015). Since harvesting and mowing often takes place during the breeding season, both chicks and nesting birds are often killed by these processes (BROYER 1996, RODRÍGUEZ–TEIJEIRO *et al.*, 2009). Some studies revealed, the harvesting and mowing processes may be pushing Common Quails to search new breeding habitats (PUIGSERVER *et al.*, 1999; RODRÍGUEZ–TEIJEIRO *et al.*, 2009). Unmown refuge-strips can, however, have a positive effect on the survival of farmland birds, and can henceforth provide breeding habitat for the birds in the same site (BROYER 2003).

The main goal of this study was to assess the impact of the unmown refuge-strips on the density and breeding site fidelity of the Common Quail in an extensively managed area (MOSON Project) in Western Hungary.

## 2. MATERIAL AND METHODS

### 2.1. STUDY AREA

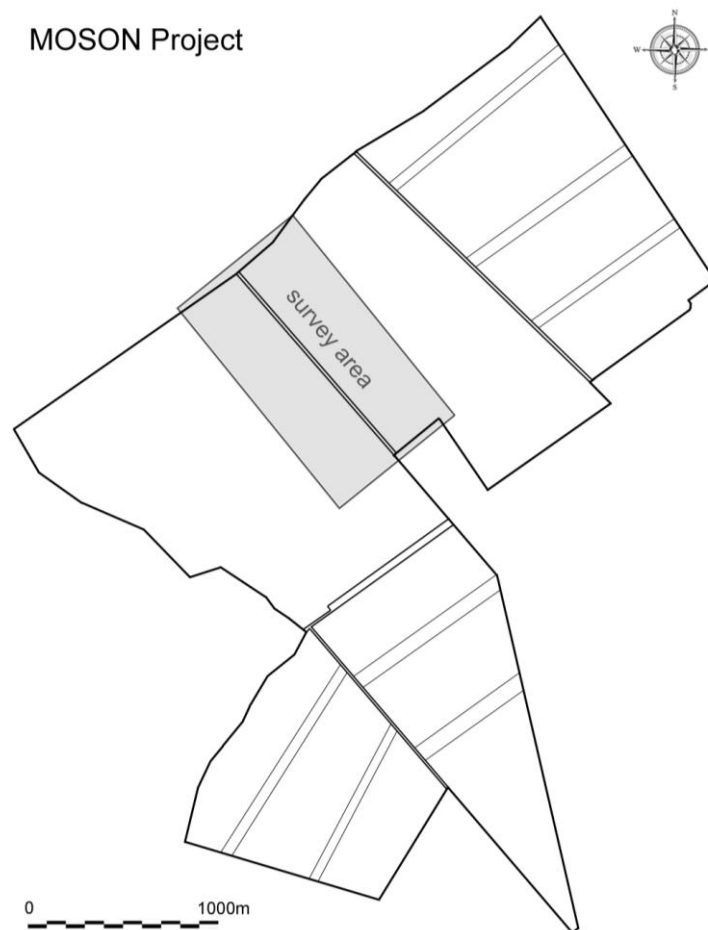
The research was carried out in the area of the MOSON Project (**Fig. 1**), situated in the Little Hungarian Plain (Northwest Hungary) underlain by alluvial deposits (mainly gravel from the River Danube) and silty loess, which result in thin, poor soils (DÖVÉNYI, 2010). The MOSON Project was launched on a former agricultural production site and covers 880 hectares. The main goal of the project was to increase the population of Great Bustard (*Otis tarda*) and Grey Partridge (*Perdix perdix*) and simultaneously of other farmland birds like the Common Quail, the Pheasant (*Phasianus colchicus*) or the passerine Corn Bunting (*Emberiza calandra*) by cultivating the field with ecologically sustainable methods (FARAGÓ & GICZI, 1997; FARAGÓ & KALMÁR, 2006). In the area traditional plant production systems with regular fallowing are dominant. About 80% of the project area is left fallow each year. The use of pesticides is restricted and there is no cultivation after April until harvesting (OECD 2008).

For the survey an approximately 80 ha area was selected, maintained by partial mowing leaving 15–20 m wide unmown strips in the field.

### 2.2. SURVEY OF COMMON QUAILS

Vocal individuality has been proved to be a useful tool for accurate bird censi (MCGREGOR & PEAKE, 1998; GILBERT *et al.*, 1994; WINKLER *et al.*, 2014; XIA *et al.*, 2017). Individual differences in Common Quail calls were demonstrated by COLLINS & GOLDSMITH

(1998), their method of analysis was roughly followed in our study. Common Quail surveys were conducted twice during the breeding season of 2015. The first survey was carried out at the end of June, one week prior to the beginning of mowing, while the second survey was undertaken mid June, nearly two weeks after mowing operation ended. By walking on the NW–SE cart-road bisecting the survey area (**Fig. 1**), Common Quail vocalizations were recorded using a linear PCM recorder (Olympus LS-5) and shotgun microphone (type RØDE NTG4+). The accuracy of the calling males' position was estimated to be within 50 m and was also mapped.

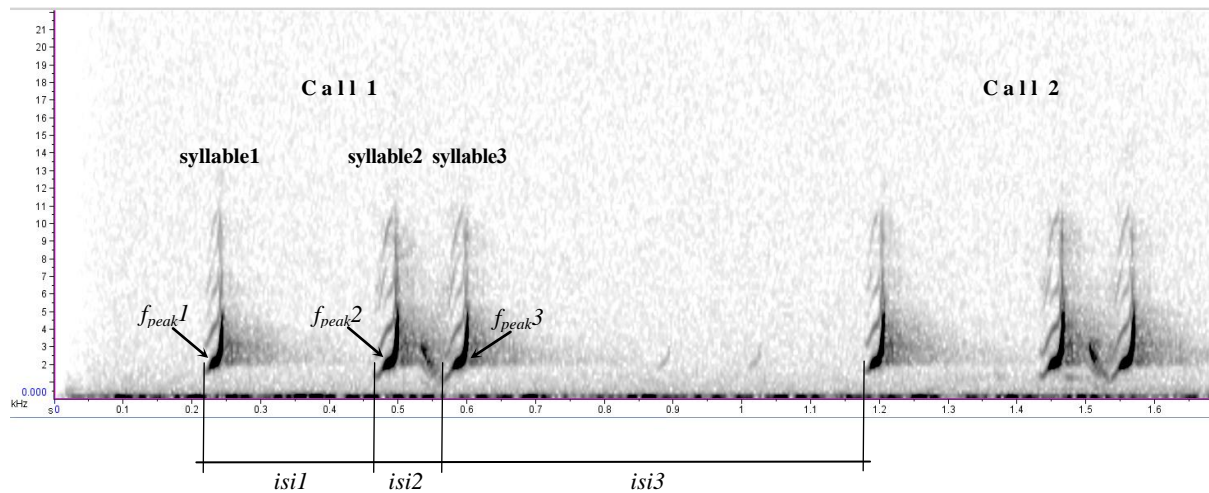


**Figure 1: Study area within the MOSON Project**

### 2.3. ANALYSIS OF RECORDINGS

Common Quail vocalizations were recorded in lossless .wav format with a sampling rate of 16bit – 44.1 kHz. Sonograms of the recorded calls were analysed using the software Adobe Audition 3.0. COLLINS & GOLDSMITH (1998) used a total of seven frequency and time variables to characterize Common Quail vocalization: apart from the fundamental frequency and duration of each syllable, the time gaps between syllable 1 and syllable 2 were determined. In our analyses, the following six variables were used: intersyllable intervals (*isi1*, *isi2* and *isi3*, respectively) and peak frequency (frequency with the most energy) of each syllables ( $f_{peak1}$ ,  $f_{peak2}$  and  $f_{peak3}$ , respectively) (**Fig. 2**). These parameters are easily measurable with high accuracy, while the duration of syllable 1 and 3, as well as the gap

between the first two syllables used by COLLINS & GOLDSMITH (1998) can hardly be determined in case when background noises (e.g. wind noise, song of passerines, orthopterans) are present. Durations were measured using the Time Selection Tool, while peak frequencies were determined with the help of the Frequency Analysis Tool within the Adobe Audition software.



**Figure 2: Sonogram of a Common Quail call recorded in the MOSON Project**

( $isi1$  – interval between syllable 1 and 2;  $isi2$  – interval between syllable 2 and 3;  $isi3$  – interval between syllable 3 and syllable 1 of the consecutive call;  $f_{peak1-3}$  – peak frequency of syllable 1-3)

### 2.3. QUANTITATIVE ANALYSIS

The measured parameters were used to perform a stepwise cross-validated discriminant function analysis (DFA). A total of 20 consecutive Common Quail calls were measured of each character for every individuals, so as to meet the requirement for the recommended adequate number (at least three times as large as the number of the measured parameters) of calls (WILLIAMS & TITUS, 1988). For re-identification of individual birds, Euclidean distance measure was used. Individual variables of re-identified calls were tested with paired samples  $t$  test. Statistical analyses were performed using SPSS Version 22 (IBM Corp., Armonk, New York).

### 3. RESULTS

During the first survey (before mowing – BM), a total of 18 Common Quails were recorded, while the second survey (after mowing – AM) resulted in a lower number (14) of birds detected. In terms of density, the initially determined 2.25 calling males/10 ha decreased to 1.75 males/10 ha.

Parameter datasets of recorded Common Quail calls are presented in **Tab. 2**. To exclude the possibility of double counts within the same survey, and to re-identify individual birds recorded in the second survey, discriminant function analysis (DFA) was applied. The analyses classified more than 94% of the Common Quail calls to the correct individuals in both surveys. The stepwise discrimination selected all six variables originally entered in the

analyses. The first three discriminant functions explained more than 95% of the total variance. Variables that contributed the most to the discrimination were the peak frequency of the second syllable ( $F_{peak2}$ ), the intersyllable interval between the first and second syllables (*isi1*) and the intersyllable interval between the second and the third syllables (*isi2*).

**Table 1: Values (Mean±SD) of the recorded Common Quail call parameters**

(For abbreviations see legend of Fig. 2)

Code of quails	<i>isi1</i> (sec)	<i>isi2</i> (sec)	<i>isi3</i> (sec)	$f_{peak1}$ (Hz)	$f_{peak2}$ (Hz)	$f_{peak3}$ (Hz)
	quails recorded before mowing (BM)					
BM1	0.215±0.002	0.127±0.002	0.630±0.013	2090.1±2.997	2131.2±1.581	2040.8±1.885
BM2	0.256±0.004	0.103±0.002	0.606±0.016	2088.7±4.400	2040.6±5.423	2045.1±2.232
BM3	0.213±0.003	0.111±0.001	0.667±0.018	1962.9±2.825	1967.8±12.620	1967.2±11.805
BM4	0.322±0.006	0.113±0.002	0.598±0.019	2132.8±1.553	2130.8±2.167	2170.0±2.619
BM5	0.244±0.004	0.126±0.001	0.649±0.020	2006.7±1.488	2005.7±2.493	2002.8±3.603
BM6	0.191±0.004	0.104±0.001	0.679±0.021	2130.6±4.596	2129.6±1.768	2088.0±4.036
BM7	0.160±0.003	0.085±0.001	0.572±0.013	2110.3±5.731	2130.3±3.777	2132.0±4.209
BM8	0.201±0.002	0.118±0.001	0.577±0.030	2006.0±2.268	2000.1±9.125	2004.3±4.534
BM9	0.266±0.003	0.118±0.002	0.700±0.061	2002.8±2.949	2003.7±3.770	2002.7±4.268
BM10	0.183±0.002	0.137±0.002	0.615±0.025	2049.6±4.438	2049.2±3.284	2051.0±6.000
BM11	0.187±0.004	0.102±0.001	0.607±0.016	1996.5±2.878	1994.6±2.138	1994.0±4.309
BM12	0.235±0.005	0.150±0.003	0.593±0.025	1992.7±4.166	1994.7±3.370	1991.6±4.406
BM13	0.234±0.002	0.129±0.001	0.621±0.020	2043.7±3.105	2042.5±3.423	2042.0±1.690
BM14	0.267±0.004	0.127±0.002	0.566±0.015	1997.0±2.138	1997.4±1.669	1996.3±1.923
BM15	0.213±0.003	0.099±0.002	0.723±0.043	2048.3±3.503	2047.7±3.327	1961.7±3.926
BM16	0.183±0.003	0.093±0.002	0.776±0.055	2049.4±4.811	2054.1±5.027	2086.0±14.590
BM17	0.258±0.004	0.130±0.002	0.683±0.019	1896.2±3.655	1897.1±4.581	1901.6±3.420
BM18	0.238±0.004	0.118±0.002	0.707±0.031	2261.6±3.249	2259.6±1.847	2261.7±1.753
	quails recorded after mowing (AM)					
AM1	0.257±0.003	0.102±0.002	0.603±0.015	2087.1±2.800	2041.3±3.105	2044.6±2.722
AM2	0.212±0.002	0.110±0.001	0.679±0.019	1961.5±2.777	1963.7±4.991	1965.1±3.727
AM3	0.237±0.003	0.121±0.003	0.629±0.018	1839.8±1.885	1838.8±1.885	1848.2±2.816
AM4	0.220±0.003	0.108±0.002	0.805±0.052	2088.5±1.927	2087.0±1.927	2000.1±7.553
AM5	0.192±0.003	0.104±0.001	0.683±0.023	2131.8±2.748	2129.8±1.727	2086.7±4.367
AM6	0.320±0.003	0.112±0.002	0.601±0.017	2133.2±1.165	2132.2±1.553	2171.0±2.828
AM7	0.271±0.002	0.116±0.004	0.583±0.034	2041.5±2.204	2042.2±1.685	2041.7±1.282
AM8	0.257±0.002	0.130±0.002	0.689±0.021	1897.4±2.507	1898.6±3.335	1902.8±2.031
AM9	0.186±0.003	0.102±0.001	0.616±0.022	1995.2±2.188	1995.5±2.138	1994.2±2.915
AM10	0.235±0.004	0.151±0.001	0.600±0.021	1993.9±3.137	1995.1±2.532	1988.5±3.857
AM11	0.265±0.003	0.127±0.001	0.573±0.023	1996.1±1.959	1997.3±1.302	1995.2±1.832
AM12	0.159±0.003	0.085±0.001	0.566±0.016	2112.2±3.834	2132.1±2.696	2131.6±2.504
AM13	0.253±0.004	0.102±0.002	0.656±0.020	2196.5±12.843	2068.8±21.570	2105.7±31.486
AM14	0.203±0.005	0.114±0.002	0.840±0.069	2032.7±1.581	2034.1±3.991	2036.8±2.835

Discrimination of the Common Quail calls recorded in the different surveys (before and after mowing) showed high degree of similarity in 9 cases based on the Euclidean distances between group centroids in the multidimensional space. According to these results, recognition of individuals was possible in the following cases (for codes see **Tab. 1**): BM2–AM1; BM3–AM2; BM4–AM6; BM6–AM5; BM7–AM12; BM11–AM9; BM12–AM10; BM14–AM11; BM17–AM8. Subsequently, we compared all measured parameters (paired

samples  $t$  test) to confirm the individual recognition (**Tab. 2**). Comparisons resulted in no significant differences with the exception of two cases. Peak frequency of syllable 2 differed significantly between calls/birds coded BM4 and AM6, while interval between syllable 1 and 2 varied significantly while comparing calls BM11 and AM9. In both cases only a single variable was affected. As randomly selected set of calls showed similar within-individual variation for these two variables, we can conclude that the 9 quails were re-identified with high probability. The individuals recorded after moving (AM) mostly aggregated in the refuge-strips, only two birds were calling in the mowed open area.

**Table 2: Comparison of call variables (paired sample  $t$  test) of the individuals re-identified with discrimination** (For abbreviations see legend of Fig. 2)

Code of quails	$isi1$	$isi2$	$isi3$	$f_{peak1}$	$f_{peak2}$	$f_{peak3}$
BM2 – AM1	$t = -0.382$ $p = 0.714$	$t = -0.491$ $p = 0.638$	$t = 0.910$ $p = 0.393$	$t = 1.328$ $p = 0.226$	$t = -0.741$ $p = 0.638$	$t = 0.404$ $p = 0.699$
BM3 – AM2	$t = 0.798$ $p = 0.451$	$t = 1.323$ $p = 0.228$	$t = -1.737$ $p = 0.126$	$t = 1.760$ $p = 0.123$	$t = 0.932$ $p = 0.383$	$t = 0.569$ $p = 0.587$
BM4 – AM6	$t = 1.416$ $p = 0.199$	$t = 1.798$ $p = 0.451$	$t = -0.957$ $p = 0.370$	$t = -2.049$ $p = 0.079$	$t = -2.246$ <b><math>p = 0.049</math></b>	$t = -1.871$ $p = 0.104$
BM6 – AM5	$t = -1.825$ $p = 0.111$	$t = 2.049$ $p = 0.080$	$t = -0.571$ $p = 0.586$	$t = -1.193$ $p = 0.272$	$t = -1.528$ $p = 0.171$	$t = 0.967$ $p = 0.366$
BM7 – AM12	$t = 1.174$ $p = 0.279$	$t = 0.798$ $p = 0.451$	$t = 1.134$ $p = 0.294$	$t = -1.618$ $p = 0.149$	$t = -1.571$ $p = 0.160$	$t = 0.414$ $p = 0.691$
BM11 – AM9	$t = 2.826$ <b><math>p = 0.026</math></b>	$t = 1.821$ $p = 0.112$	$t = -1.331$ $p = 0.225$	$t = 1.452$ $p = 0.189$	$t = 1.871$ $p = 0.104$	$t = -1.091$ $p = 0.311$
BM12 – AM10	$t = 0.832$ $p = 0.439$	$t = -0.856$ $p = 0.421$	$t = -1.527$ $p = 0.171$	$t = -1.515$ $p = 0.174$	$t = -0.532$ $p = 0.612$	$t = 1.917$ $p = 0.097$
BM14 – AM11	$t = 1.644$ $p = 0.144$	$t = -1.488$ $p = 0.180$	$t = -1.117$ $p = 0.300$	$t = 1.507$ $p = 0.176$	$t = 1.426$ $p = 0.197$	$t = 1.688$ $p = 0.135$
BM17 – AM8	$t = 1.476$ $p = 0.183$	$t = 1.871$ $p = 0.104$	$t = -1.441$ $p = 0.193$	$t = -1.488$ $p = 0.180$	$t = -1.426$ $p = 0.197$	$t = -1.452$ $p = 0.189$

#### 4. DISCUSSION

As result of our surveys revealed, leaving uncut refuge-strips in grassland area can have positive impact both on the survival and the site fidelity of Common Quails. As the second survey results indicated, there was only a slight decrease in calling males density. Based on the recorded calls, several birds have been re-identified proving that the uncut strips can still provide suitable foraging and nesting habitat for Common Quails, despite the fact that this species is reported to be highly nomadic during the breeding season in some regions (HERRMANN & DASSOW, 2006). As foreign researches showed, in the course of harvesting and mowing Common Quails are consequently moving to new optimal habitats for breeding (PUIGSERVER *et al.*, 1999; RODRÍGUEZ-TEJEIRO *et al.*, 2009). The five birds detected only during the second survey after mowing might have arrived from completely harvested or mown neighbouring areas.

Several studies emphasized the positive effects of unmown refuge areas on the survival of farmland birds (VICKERY *et al.* 2001; BROYER 2003; ARBEITER *et al.* 2017). In the MOSON Project, apart from the Common Quails, bird species that benefit from the uncut grass strips include the vulnerable Great Bustard (*Otis tarda*), the Grey Partridge (*Perdix*

*perdix*), the Pheasant (*Phasianus colchicus*) as well as passerines like the Skylark (*Alauda arvensis*), the Yellow Wagtail (*Motacilla flava*), the Whinchat (*Saxicola rubetra*) and the Corn Bunting (*Emberiza calandra*). On the other hand, unmown refuge-strips are attractive enough also for predator species (LÓRÁNT *et al.*, 2008). Apart from birds of prey, species of corvid like the Magpie (*Pica pica*) and the Hooded Crow (*Corvus cornix*), as well as mammal predators such as the Red Fox (*Vulpes vulpes*) are efficient predators in the area and can therefore affect the beneficial effects of the unmown refuge-strips.

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