Electrophoresis 2019, *40*, 1897–1903 1897

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- Received November 15, 2018 Revised February 11, 2019 Accepted February 11, 2019

1 Introduction

Rasagiline (RAS, chemically known as (*R*)-*N*-(prop-2-ynyl)- 2,3-dihydro-1*H*-inden-1-amine, Fig. 1, used as a mesylate salt) is an irreversible inhibitor of monoamine oxidase B, used for the treatment of the signs and symptoms of Parkinson's disease [1]. RAS has a single asymmetric carbon, resulting in two optical isomers (*R*- and *S-*enantiomers). Only the *R*enantiomer is active, therefore, the drug is marketed as a single enantiomeric agent, while the *S*-enantiomer could be present as a chiral impurity [2]. Determination of chiral purity of single enantiomeric agents is not only a necessity, but also

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Abbreviations: 2D ROESY, two-dimensional phase-sensitive rotating frame nuclear Overhauser effect spectroscopy; **CM**, carboxy-methylated; **DS**, degree of substitution; **EMO**, enantiomer migration order; **HP**, hydroxypropyl; **RAS**, rasagiline; SBE, sulfobutylether; TRIMEB, heptakis(2,3,6-tri-*O*-methyl)-β-CD

Research Article

Chiral separation of rasagiline using sulfobutylether-β-cyclodextrin: capillary **electrophoresis, NMR and molecular modeling study**

Pressure-assisted stereospecific capillary electrophoresis method was developed for the determination of enantiomeric purity of the antiparkinsonian agent *(R*)-rasagiline. The optimized method, 50 mM glycine-HCl buffer pH 2, supplied with 30 mM sulfobutylether- --cyclodextrin, at 35°C, applying 12 kV in reversed polarity, and –8 mbar pressure (vacuum), short-end injection with -25 mbar \times 2 s, was successful for baseline separation of rasagiline enantiomers ($R_s = 3.5 \pm 0.1$) in a short analysis time. The method was validated according to current guidelines and proved to be reliable, linear, precise and accurate for determination of 0.15% *S*-enantiomer as chiral impurity in *R*-rasagiline sample, as well as quantification of the eutomer. Method application was tested on a commercial tablet formulation. Determination of spatial structure of diastereomeric associates was based on 1H and 2D ROESY NMR, indicating that the aromatic moiety of the molecule can enter the cyclodextrin cavity. NMR titration and molecular modeling revealed that S-rasagiline formed a more stable inclusion complex with sulfobutylether-ß-cyclodextrin, than its antipode, which is in agreement with electrophoretic results.

Keywords:

Azilect $^{\circledR}$ / Chiral separation / Cyclodextrin / Enantioseparation / Migration order DOI 10.1002/elps.201800482

> a regulatory requirement. A literature survey revealed several direct liquid chromatographic methods for the chiral separation of RAS enantiomers (Table 1) using polysaccharide [3,4], crown ether [5] or protein-based chiral stationary phases [6]. To the best of our knowledge, CE was yet to be applied for the chiral separation and subsequent enantiomeric quality control of RAS.

> Although liquid chromatography is the predominant technique in chiral analysis, CE has also several advantages in this field, due to its low solvent and analyte consumption, high efficiency, and rapid analysis time. Moreover, the combination of CE with other techniques, especially NMR spectroscopy and theoretical calculations can help gain insight into the spatial structure and stoichiometry of diastereomeric selector-selectand complexes and offer valuable information upon the chiral discrimination process [7–11]. The aim of the present study was to develop, optimize and validate a fast and cost-effective CE method for the enantiomeric quality control of *R*-RAS using CDs as chiral selector. Moreover ¹H and two-dimensional phase-sensitive rotating frame nuclear overhauser effect spectroscopy (2D ROESY) NMR as well as molecular modeling study were also applied for the characterization of non-covalent intermolecular interactions

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Figure 1. Constitutional formula and numbering of rasagilin.

of RAS-CD diastereomeric complexes and elucidation of the possible enantiorecognation mechanism.

2. Materials and methods

2.1 Materials

R- and *S*-RAS, DMSO, D₂O, DCl, glycine, sulfated-β-CD with degree of substitution $(DS) \sim 11$ and gradient grade methanol (MeOH) were purchased from Sigma-Aldrich (Budapest, Hungary). α-, β-, γ-CD, hydroxypropyl- α -CD (HP- α -CD) DS~4.5, HP-β-CD DS~4.5, HP-γ-CD DS~4.5, methyl-α-CD DS~11, methyl-β-CD DS~12, methylγ-CD DS~13, heptakis(2,6-di-*O*-methyl)-β-CD content 95%, DS~14, heptakis(2,3,6-tri-O-methyl)-β-CD, carboxymethylated-α-CD (CM-α-CD) DS~3.5, CM-β-CD DS~4, CMγ-CD DS~4 and sulfobutyl-ether-β-CD DS~6.3 were products of Cyclolab (Budapest, Hungary). Tris, HCl and NaOH were of analytical grade and purchased from commercial suppliers. The deionized water was prepared by a Milli-Q Direct 8 Millipore system. Azilect \mathbb{R} 1 mg tablets (TEVA Pharma GmbH) were obtained from a local pharmacy in Budapest, Hungary.

2.2 CE measurements

CE experiments were performed on an Agilent 7100 CE instrument (Agilent Technologies, Waldbronn, Germany), equipped with a photodiode array detector and ChemStation software for data handling. An untreated fused-silica capillary (50 µm id, 48.5 cm total and 40 cm effective length) from Agilent was used for the separation. Conditioning of new capillaries was conducted by flushing with 1 M NaOH for 30 min followed by 0.1 M NaOH and water for 60 min each. Prior to all runs the capillary was preconditioned by flushing with 0.1 M NaOH (2 min), water (1 min), and BGE (3 min). UV detection was performed at 200 nm. If not stated otherwise, injections were carried out by applying a pressure of 25 mbar for 3 s.

2.3 Preparation of BGE, stock, and sample solutions

For preliminary experiments 50 mM phosphate buffer adjusted to pH 6 was used; the BGE contained CDs in various concentrations (3–30 mM), the applied voltage was 20 kV and the capillary was maintained at 25°C.

Application of the method was tested on a commercial tablet formulation (Azilect[®]). Twenty tablets were weighed and afterwards ground and mixed in a mortar. Then 10 mL MeOH was added to an accurately weighed portion of this powder equivalent to 10 mg of RAS. The sample was sonicated for 30 min and then centrifuged for 10 min applying 4000 rpm (Sartorius 2–16P benchtop centrifuge, Goettingen, Germany). The clear supernatant was filtered through a 0.22 µm pore size syringe filter. An appropriate dilution was made with water to obtain the final concentrations (100 μ g/mL). For the determination of chiral impurity, water was used as an extraction solvent, while the powder blend was weighed accurately as to obtain a theoretical concentration of 2666 µg/mL *R-*RAS.

R- and *S*-RAS stock solutions were prepared in methanol at 1 mg/mL and were further diluted with water. The sample solution used during the preliminary CD screening experiments contained approximately 150 µg/mL *R*-RAS and 30 µg/mL *S*-RAS. Other conditions including validation process are given in Results and Discussion section.

2.4 NMR experiments

All NMR measurements were carried out on an Agilent Varian Unity Inova DDR spectrometer (599.9 MHz for ¹H) with a 5 mm inverse-detection gradient probehead at 25°C. Standard pulse sequences and processing routines available

Table 1. Direct liquid chromatographic methods for the chiral separation of RAS enantiomers

Stationary phase/Chiral selector	Mobile phase	Literature
Chiralcel OJ-H ((based on cellulose tris(4-methylbenzoate))	hexane:isopropyl alcohol:ethanol:diethyl amine (96:2:2:0.01)	$[3]$
Chiralpak AD-RH ((based on amylose tris(3,5-dimethylphenylcarbamate))	20 mM potassium dihydrogen phosphate in water-acetonitrile (65:35, v/v)	$[4]$
$(+)$ -(18-crown-6)-2,3,11,12-tetracarboxylic acid	ethanol/acetonitrile/acetic acid/triethylamine (80:20:0.2:0.3, v/v/v/v)	$[5]$
Chiralpak AGP (based on α 1-acid glycoprotein)	ammonium acetate and isopropyl alcohol $(90:10, v/v)$	[6]

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in VnmrJ 2.2C/Chempack 4 were used. The molecular geometry of the complex was investigated by two-dimensional phase-sensitive rotating frame nuclear Overhauser effect spectroscopy (2D ROESY). The structures of the inclusion complexes were determined applying a spinlock of 3 kHz for a mixing time of 300 ms.

The stability constants of RAS enantiomers were determined by ¹H NMR titration in D_2O . The pH was set at pH 2 using DCl. A stock solution containing 5 mM of both enantiomers was prepared separately, while the $\texttt{SBE-B-CD}$ stock solution was 30 mM. 30 µL of RAS stock solution was mixed with different volumes of SBE-β-CD stock solution and filled to a total volume of 600 µL. The observed chemical shift ($\delta_{\rm obs}$) of a given nucleus can be expressed using the following formula:

$$
\begin{aligned} \delta_{\text{obs.}}\;&=\delta_{RAS} \\ &\qquad \qquad +\Delta\delta \frac{\left[\text{RAS}\right]_{T}+\left[\text{CD}\right]_{T}+\frac{1}{K}-\sqrt{\left(\left[\text{RAS}\right]_{T}+\left[\text{CD}\right]_{T}+\frac{1}{K}\right)^{2}-4\left[\text{RAS}\right]_{T}\left[\text{CD}\right]_{T}}}\left(1\right) \\ &\qquad \qquad +\Delta\delta \frac{\left[\text{RAS}\right]_{T}}&\qquad \qquad \end{aligned}
$$

where $\Delta \delta = \delta_{\text{RAS}-\text{CD}} - \delta_{\text{RAS}}$

The total concentration of RAS enantiomers ($[RAS]_T$) was constant during the titration.

Stability constants (*K*) of the inclusion complexes were calculated by non-linear parameter fitting of Eq. (1) to the $\delta_{\rm obs}$ versus $[CD]_T$ datasets using OriginPro 8 program based on previous works [12–14].

2.5 Molecular modeling

In order to elucidate the atomic details of the enantioseparation, the interaction of RAS enantiomers and SBE-β-CD was studied by using computational chemical tools. The model structures were built by using GaussView 5.0. As the actual configuration of SBE-β-CD is unknown (random substitution of the OH groups), three different isomeric forms were created to mimic the applied chiral selector based on the work of Jain et al. [15]. In each case, seven SBE groups were added to the β -CD skeleton, as the degree of substitution was \sim 7 in the experimental system (Fig. 2). Furthermore, the SBE groups

were properly distributed on the β -CD skeleton to represent the induced steric hindrance [15]. Energy minimization of the SBE - B -CD isomers were performed by using the optimized potentials for liquid simulations-AA forcefield in combination with the Generalized Born implicit solvation model. The resulted three optimized structures have been used as hosts in docking simulations. The RAS enantiomers were docked and their affinity towards the optimized SBE-ß-CD isomers were studied by applying a "blind docking" protocol [16] which was carried out by using AutoDock Vina [17]. The size of the grid box was set to be $40 \times 40 \times 40$ Å which is much bigger than the actual size of the isomers (host molecules). The exhaustiveness and the maximum number of binding modes was set to 8 and 9, respectively. The best binding mode from each simulation was selected and the corresponding structures and binding affinities were compared.

3 Results and discussion

3.1 CE method development

Sixteen different CDs were applied as potential chiral selectors in the scouting phase. Through the application of a high-number of different CD derivatives, the effect of differences in the cavity size, nature of the substituents and charge upon enantiorecognition was also tracked. During the initial scouting phase, the chiral recognition ability of each selector was checked at five concentration levels (3-30 mM) at pH 6 in a 50 mM phosphate buffer. The capillary temperature was kept at 25°C and 20 kV voltage was applied. From the sixteen studied CDs eight CD derivatives (CM-α-CD, CM-β-CD, CM-γ-CD, HP-β-CD, methyl-β-CD, DIMEB, sulfated-β-CD and SBE-ß-CD) showed chiral interactions with the analyte, with SBE- β -CD at 30 mM delivering the most promising results. Higher concentration of chiral selector resulted in better enantioseparation, however concentration above 30 mM could not be applied because of the high current generated.

In all cases the migration order was not the ideal one, because the eutomer *R-*RAS was migrating first, followed by the *S*-enantiomer. For further method development our aim

Figure 2. 2D structures of sulfobutylether-ß-cyclodextrin isomers, where R = sulfobutylether group.

was to reverse the enantiomer migration order (EMO), while still maintaining baseline separation.

One of the main advantages of CE enantioseparation is the ease of reversing EMO, when compared to chromatographic methods. In our recent study, we have successfully reversed the enantiomer migration order for praziquantel, by polarity switching and EOF reduction at strongly acidic BGE [18]. The same simple approach was employed in this study also, together with reduction of the effective length of the separation capillary to 8.5 cm, by employing short-end injection technique. Using 50 mM phosphate buffer pH 2.0, supplemented with 30 mM SBE-B-CD at 25 °C, hydrodynamic injection at the near-detector-end of the capillary (-25 mbar x 3 sec), and applying 15 kV (reversed polarity), the EMO was reversed and the ideal situation, when the distomer, *S-*RAS was the first migrating enantiomer, followed by the *R*-RAS was obtained.

Further, method optimization was undertaken to improve peak shape and further reduce analysis time. Higher concentrations than 30 mM of the ionized SBE-ß-CD and/or 50 mM phosphate buffer could not be utilized, because of the high currents generated. In the following, several variations of voltage and temperature were employed to further optimize the method, but still maintain currents below \sim 100-110 µA. Best results were obtained at a higher capillary temperature (35 °C), which implied lowering the voltage to 12 kV. However, using these conditions, the total analysis time was still longer than 17 minutes and peak shapes and efficiencies were still not ideal. In order to overcome these shortcomings, vacuum was applied to the outlet vial. This pressure-assisted approach lead to better peak shapes, overall higher efficiencies and reduced analysis time, but also decreased *Rs* values

Figure 3. Effect of applied pressure on the enantioseparation of.RAS enantiomers. (A) 0 mbar; (B) -5 mbar*; (C) -7 mbar*; (D) -8 mbar*; (E) -10 mbar*; (F) -15 mbar*. Other experimental conditions: BGE: 50 mM phosphate buffer pH 2.0, 30 mM SBE-β-CD; temperature: 35°C; 25 mbar x 2 sec hydrodynamic injection at the short-end of the capillary; applied voltage: 12 kV (reversed polarity, cathode at inlet). *Negative pressure indicates that vacuum was applied at the end of the capillary, in order to pull the sample plug through the detector.

(Fig. 3). A good compromise between reduced analysis time, but still sufficient enantioselectivity was obtained by applying -8 mbar pressure. As a last step, the influence of changing the BGE buffer system was investigated. Phosphate buffer, Tris-HCl and glycine-HCl all in a 50 mM concentration at pH 2.0 were employed as BGE constituents. No difference in migration times were recorded, however peak shapes were better using glycine-HCl buffer. Using these optimized conditions (50 mM glycine-HCl buffer pH 2.0, 30 mM SBE-ß-CD (DS 6.3), capillary temperature: 35°C, applied voltage 12 kV (reversed polarity) and – 8 mbar pressure, short-end injection with -25 mbar \times 2 sec) high resolution (R_s = 3.5 \pm 0.1) was achieved within 8 min. Therefore, further method optimization was not applied.

3.2 Method validation

Our developed method was validated for the determination of *S*-RAS as a potential optical impurity in *R*-RAS sample and for quantification of *R*-RAS following the International Council on Harmonization (ICH) guidelines in terms of sensitivity (LOD, LOQ), linearity, precision and accuracy.

3.2.1 Method validation for the determination of *S***-RAS as a potential optical impurity**

Method sensitivity was evaluated for the determination of *S*-RAS, by sequentially diluting sample solutions. The LOD of *S*-RAS was determined as the concentration yielding a signal three times the baseline noise while LOQ was determined at 10:1 signal to noise ratio in the presence of *R*-RAS solution. The LOQ was determined at 4 µg/mL (corresponding 0.15% impurity in 2666 µg/mL *R*-RAS sample), while LOD of *S*-RAS was 2 μ g/mL (corresponding 0.075%) impurity in 2666 µg/mL *R*-RAS sample). Calibration curve was constructed at six different concentration levels between 4 and 48 µg/mL corresponding to 0.15%-1.8% enantiomeric impurity in 2666 µg/mL *R*-RAS solution. Regression analysis shows a linear relationship in the above mentioned concentration ranges with the following equations: y $= 1.1809x + 0.2919$ ($r^2 = 0.9994$). Accuracy (calculated as recovery%) and precision (repeatability based on RSD% of the migration time and peak area) were analyzed by performing intra-and interday evaluation of three concentration levels 4, 16 and 40 µg/mL for *S*-RAS in the presence of *R*-RAS. In the intraday repeatability study, the RSD% of migration time was lower than 0.5%, In the interday repeatability (measured in two consecutive days) the RSD% of migration time was lower than 1.5%. The accuracy data and RSD% of peak areas are summarized in Table 2.

3.2.2 Method validation for quantification of *R***-RAS**

Our method was also validated and applied for quantification of *R*-RAS. In this case calibration curve was constructed **Table 2.** The accuracy data and RSD% of peak areas of *S*-RAS as a potential optical impurity in presence of *R*-RAS

S-RAS concentration in 2666 μ g/mL R-RAS	Precision (RSD%)		Accuracy (%)	
	Intradav	Interday	Intradav	Interday
4μ g/mL (0.15%)	3.1	4.5	96.1	96.3
16 μ g/mL (0.6%)	1.2	1.6	101.5	99.7
$40 \mu g/mL (1.5%)$	4.1	2.3	99.8	102.3

Table 3. The accuracy data and RSD% of peak areas of *R*-RAS

between 80 and 120 µg/mL of *R*-RAS. The results indicate a linear relationship with the following regression equation: $y = 1.472x + 0.0964$ ($r^2 = 0.9996$). Like the validation of determination of potential chiral impurity, the accuracy and precision of the method was tested. Quality control samples were prepared at concentration of 80, 100 and 120 µg/mL for *R*-RAS. The RSD% of migration time was lower than 0.2% and 0.9% in the case of intra- and interday repeatability, respectively. Further results regarding accuracy and precision are summarized in Table 3.

3.3 Application of the method

Our optimized and validated method was applied for the assay and enantiomeric quality control of *R*-RAS in Azilect® tablets (containing 1 mg *R*-RAS in mesylate salt form). In the analyzed samples, no chiral impurity could be detected, however after spiking the sample with the distomer to achieve 0.15% chiral impurity, *S*-RAS could be determined (Fig. 4). Moreover, during the assay of *R*-RAS, a good agreement between the value claimed by the manufacturer and that determined by our method (1.03 \pm 0.03 mg/tablet) was obtained. The real sample analysis further underlines the applicability of our method.

3.4 NMR and molecular modeling for characterization of enantioselective complexation

At first in the 1 H NMR measurements, spectrums were recorded using a solution containing 3 mM sample with 7:1 *R*:*S* enantiomeric ratio, with increasing concentrations of SBE-β-CD (0, 5, 15, 20,30 mM) to check a possible enantioselective differences. In the 1H NMR spectra the complexation-induced chemical shift is clearly indicative of selector-selectand complexation, moreover some minor

Figure 4. Electropherogram of a solution of Azilect[®] tablets containing 2666 µg/mL *R*-RAS spiked with 0.15% *S*-RAS chiral impurity. Conditions: BGE: 50 mM glycine-HCl buffer pH 2.0, 30 mM SBE-β-CD; temperature: 35°C; 25 mbar x 2 sec hydrodynamic injection at the short-end of the capillary; applied voltage: 12 kV (reversed polarity, cathode at inlet).

signal splitting was also observed due to nonequivalence of complexation induced chemical shifts of enantiomers in the aromatic region applying higher selector concentration (>15 mM) (Fig. 5A). The chemical shift differences for *S*-RAS were higher than *R*-RAS, indicating that the *S*-enantiomer formed more stable complex with SBE-ß-CD. To determine the exact stability constant ${}^{1}H$ NMR titration was applied. The stability constant for *R*-RAS and *S*-RAS SBE-ß-CD were 2.18 \pm 0.02 and 2.35 \pm 0.01, respectively. These data are in good accordance with the observed EMO in CE measurements.

2D ROESY NMR measurement was applied for determination of spatial structure of the RAS enantiomer- SBE-ß-CD complexes. Crosspeaks between RAS aromatic protons and inner $H³$ and $H⁵$ protons of the CD clearly show that the phenyl ring can enter the CD cavity. Based on ROESY experiments enantiomer specific complexation was not observed (Fig. 5B).

Docking simulation was used to aid in the understanding of the mechanism of the separation. The tridimensional structures of the diastereomeric complexes between RAS enantiomers and three different SBE-ß-CD isomers are presented in Fig. 5, together with the calculated relative binding affinities of complex formation. Results of the

Figure 5. (A) The aromatic part of ¹H NMR spectrum of *R*-enantiomer-enriched RAS (3 mM sample with 7:1 *R*:*S* enantiomeric ratio) without $SBE-\beta$ -CD (down) and with 30 mM SBE- β -CD (top). (B) 2D ROESY spectrum of a sample containing 3 mM SBE- β -CD, 2.5 mM *R*-RAS and 1.5 mM *S*-RAS, showing ROE crosspeaks between the aromatic protons (H-6, H-7 and H-8) of RAS with inner $H³$ and $H⁵$ protons of $SBE- β -CD.$

 R - and S-Rasagiline / SBE- β -CD complexes

Figure 6. 3D structures of the diastereomeric complexes between RAS enantiomers and $SBE- β -CD isomers to$ gether with the calculated relative binding affinities $(\Delta_{S-R}E_A,$ $kcal/mol)$. $\Delta_{S-R}E_A = E_A(S)$ – $E_A(R)$, where $E_A(S)$ and $E_A(R)$ are the binding affinity of the complex formed with *S*-RAS and *R*-RAS enantiomers, resepectively. *S*-RAS and *R*-RAS indicated with light green and orange color.

molecular modeling study are in line with experimental observations: a more stable complex was formed between the chiral selector, SBE-ß-CD and *S-*RAS, causing the later to reach the detector faster in the reversed polarity setup, than *R*-RAS. The study also confirmed the preferential inclusion of the aromatic part of the analyte in the cavity of the CD, the *N*-propargyl side-chain of RAS points outside of the cavity. In the case of *R*-RAS this positively charged side-chain points towards the apolar side-chain of the sulfobutyl group, whereas for *S*-RAS, the *N*-propargyl group points away. Energetically, the latter case is more favorable. We hypothesize that the electrostatic interactions between the negatively charged CD and the positively charged analyte are primary responsible for anchoring the enantiomers (leading interactions) and bringing the asymmetric moieties of the selector and selectand in close proximity [19, 20]. Enantiomeric discrimination most probably occurs afterwards, when the differences in secondary interactions, give rise to a more stable complex for *S*-RAS. Based on molecular modeling study it is evident, that *S*-RAS established more secondary

interactions $(H\text{-bonding}, -NH—O-;$ and $\Pi\text{-interactions}$ with the SBE-β-CD isomers in each case, than its *R*-RAS counterpart (Fig. 6).

4 Concluding remarks

A novel, rapid enantioselective pressure-assisted CE method using SBE-β-CD as chiral selector has been established for the determination of enantiomeric purity of RAS for the first time. Applying the optimized conditions 50 mM glycine-HCl buffer pH 2.0, supplied with 30 mM SBE-β-cyclodextrin, at 35°C, applying 12 kV in reversed polarity, and –8 mbar pressure (vacuum), short-end injection with -25 mbar \times 2 s, baseline separation of RAS enantiomers was achieved within 8 min. Our optimized and validated method can be successfully applied for the quality control of *R*-RAS. Using NMR and molecular modeling the SBE-ß-CD RAS inclusion complexation was characterized. Different binding affinities of the individual enantiomers towards SBE-ß-CD were identified with both NMR and molecular modeling, in favor of the *S*enantiomer, which are in accordance with the observed EMO. We assume that the secondary interactions with smaller energies could play a bigger role in the enantiomer recognition mechanism. In our work we proved that by combined use of CE, nuclear magnetic resonance spectroscopy and molecular modeling calculations can help to gain deeper understanding of the chiral recognition mechanism of CD chiral selectors.

The financial support from New National Excellence Program of the Ministry of Human Capacity (UNKP-17-4) and ´ Semmelweis Innovation Found (STIA-M-17, STIA-18-KF) is highly appreciated. The GITDA (Governmental Information-Technology Development Agency, Hungary) is gratefully acknowledged for allocating computing resources used in this work. BF thanks the support from the EU, Hungary and the European Regional Development Fund within the framework of the GINOP-2.3.4-15-2016-00004 project. Z.-I.Sz. and G.T. would like to thank the Transylvanian Museum Society for their support.

The authors have declared no conflict of interest.

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