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Avian interspecific differences in VKOR activity and inhibition: insights from amino acid
sequence and mRNA expression ratio of VKORC1 and VKORC1L1

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ABSTRACT

Worldwide use of anticoagulant rodenticides (ARs) for rodents control has frequently led to secondary poisoning of non-target animals, especially raptors. In order to suggest some factors that may help considering the mechanism of the incidents, this study focused on the avian vitamin K 2, 3-epoxide reductase (VKOR) that is the target protein of ARs. We addressed the interspecific differences in VKOR activity and inhibition related to amino acid sequence and mRNA expression of VKORC1 and VKORC1-like1 (VKORC1L1). Poultry have been considered to be more tolerant to ARs than mammals. However, VKOR activity of owls, hawks, falcon and surprisingly, canaries, was lower and inhibited by warfarin more easily than that of chickens and turkeys. The amino acid sequence of VKORC1 and VKORC1L1 implied that the value of K_i for VKOR activity to ARs could depend on the amino acid at position 140 in the TYX warfarin-binding motif in VKORC1, and other amino acid mutations in VKORC1L1. The mRNA expression ratio of VKORC1:VKORC1L1 differed between turkey (8:1) and chicken (2:3) liver. VKORC1L1 has been reported to be resistant to warfarin compared to VKORC1. Hence, both the K_i of specific VKORC1 and VKORC1L1, and the mRNA expression ratio would cause avian interspecific difference of the VKOR inhibition. Our study also suggested the high inhibition of VKOR activities in raptors and surprisingly that in canaries as well. These factors are the most likely to contribute to the high sensitivity to ARs found in raptors.

Keywords: anticoagulant rodenticides, VKORC1, VKORC1L1, raptors,

INTRODUCTION

Worldwide use of anticoagulant rodenticides (ARs) for vertebrate pest control has led to the unintentional exposure of non-target animals, especially raptors, to these poisons (López-Perea et al., 2018). Exposure pathways of ARs to raptors have been presumed to be prey on target rodents or non-target animals. Rattner et al. (2018a) summarize the median lethal dose (LD₅₀) for anticoagulant rodenticides (ARs) among animals. Avian species seem more tolerant to ARs than mammals. However, the LD₅₀ for the first-generation AR diphacinone in the American kestrel (*Falco sparverius*) is lower than in poultry (Rattner et al., 2011), and a comparative risk model found that second-generation ARs pose a far greater risk to predatory and scavenging birds (Erickson et al., 2004; Anderson et al., 2011). Considering both the low values of LD₅₀ in raptors and the frequent poisoning of non-target wild birds, especially raptors, implies raptors have different susceptibility to ARs than poultry.

Interspecific difference in sensitivity to ARs could be caused by differences in the both AR metabolism by cytochrome P450 (CYP) and the Vitamin K 2, 3-epoxide reductase (VKOR) inhibition, which is the pharmacological target of AR. However, those data are limited to very few avian species. Watanabe et al. (2010) used warfarin as a model compound of ARs and reported that owls in particular showed very low CYP-dependent warfarin metabolic activity compared with other avian species and rats. Rattner et al. (2014) showed that the eliminated half-life of diphacinone in the liver was longer in eastern screech-owls (*Megascops asio*) than in mammals. These studies imply that owls may possess a low ability to detoxify ARs. On the other hand, there are few studies on VKOR characterization in raptors. The VKOR IC₅₀ of several ARs in hepatic microsomes of the American kestrel has been described (Rattner et al., 2018b), i.e., brodifacoum (0.22 µM), and preliminary warfarin (177 µM) and chlorophacinone (5.1 µM). Brodifacoum was reported to have similar inhibition efficiency for VKOR containing hepatic microsomes to mammals (IC₅₀ of 0.15-0.26 µM in mammals).

VKOR catalyzes the reduction of vitamin K 2, 3-epoxide (VKO) to vitamin K

79 quinone (vitamin K) (Fig. S1). This reaction is inhibited by ARs such as warfarin. Vitamin
80 K is reduced to vitamin K hydroquinone (VKH₂) by vitamin K quinone reductase (VKR).
81 VKH₂ is a cofactor for the γ -glutamyl carboxylation of glutamate (Glu) in vitamin K
82 dependent proteins (VKDPs). Some of the most studied VKDPs are clotting factors II,
83 VII, IX, and X in mammals (Ferland, 1998). Although it was thought that avian clotting
84 factors were different from that of mammals (Walz et al., 1975; Frost et al., 1999), recent
85 studies show that avian plasma possesses functional coagulation factors (Thomson et al.,
86 2002), and that avian prothrombin times (chicken, 9.7 s – 27 s) are similar to those of
87 mammals (human, 14 s – 17 s) (Frost et al., 1999; Webster 2009). The inhibiting VKOR
88 by ARs impedes carboxylation of blood clotting factors which can result in hemorrhage
89 in both birds and mammals (Cain et al., 1998; Rattner et al., 2012; Rattner et al., 2014;
90 Rattner et al., 2015).

91 There are two paralogous multisubunit membrane protein complexes that
92 perform VKOR activity, VKOR complex 1 (VKORC1) and VKORC1-like1
93 (VKORC1L1) (Tie and Stafford, 2016). In mammals, the focus has principally been on
94 VKORC1. Spohn et al. (2009) demonstrated that VKORC1 knockout caused early
95 postnatal lethality due to severe bleeding in mice. Hammed et al. (2013) and Caspers et
96 al. (2015) reported that the mRNA expression of VKORC1 was over 10-fold higher than
97 that of VKORC1L1 in the liver of rats and mice. Therefore, VKORC1 has been
98 considered as the main protein supporting VKOR activity in liver of mammals. However,
99 VKORC1L1 was described to be involved in VKOR activity in some extrahepatic tissues
100 (Hammed et al., 2013). Moreover, warfarin inhibition constant K_i for rat VKORC1L1
101 was reported to be higher than that for rat VKORC1. The authors demonstrated that
102 mRNA expression of VKORC1L1 was higher than that of VKORC1 in rat testis, and the
103 VKOR activity of the testis was not inhibited as easily by warfarin compared to VKOR
104 activity of the liver. These facts imply that a high mRNA expression of VKORC1L1,
105 which has a high K_i , caused warfarin resistance. Hence, warfarin susceptibility could be
106 caused by both the mRNA expression ratio, and the individual K_i of VKORC1 and

VKORC1L1.

Although the warfarin-binding site of VKOR has not been clearly identified, amino acids at the 138 to 140 positions in the rat VKORC1 “TYX motif” may be the warfarin-binding site. In rats and mice, tyrosine 139 mutations in VKORC1 exhibit high resistance to warfarin compared to other mutations (Lasseur et al., 2006a; Lasseur et al., 2006b; Rost et al., 2009). The Thr-Tyr-Ala motif also exists in NAD(P)H quinone oxidoreductase (NQOR) that is sensitive to warfarin (Rost et al., 2005). The TYX warfarin-binding site could be important to the K_i for VKOR activity in response to ARs.

There are few studies on avian VKOR. It was reported that the V_{max} (maximum velocity) of VKOR activity in chickens and ostriches was 3- to 7-fold lower than that of rats, and K_i for VKOR activity after warfarin treatment was 17- to 40-fold higher in chickens than in ostriches or rats (Watanabe et al., 2010). These facts were the first to indicate low VKOR activity in poultry compared to mammals and the high warfarin resistance of chickens. Nevertheless, it was not enough to elucidate broad avian interspecific differences in sensitivity to ARs, especially the high susceptibility to ARs found in raptors.

This study addressed avian interspecific differences in VKOR activity and sensitivity to warfarin, one of the most common ARs in the world. The cause of avian interspecific differences in VKOR activity and sensitivity were discussed by comparing amino acid sequence and mRNA expression of VKORC1 and VKORC1L1 among avian species, including raptors. The aim of this study was to reveal the factors contributing to the high sensitivity to ARs found in raptors, which are frequently reported as non-target poisoning instances.

MATERIALS AND METHODS

Chemicals

HEPES was purchased from Dojindo Laboratories (Kumamoto, Japan). Vitamin K₁ was from Kanto Chemicals (Tokyo, Japan). Diethyl ether and RNA*later*® were from Sigma-Aldrich Co. (St. Louis, MO). Racemic warfarin, dithiothreitol (DTT), isopropyl alcohol, hexane, ethanol, methanol, H₂O₂, K₂HPO₄, KH₂PO₄, Na₂CO₃, and NaOH were purchased from Wako Pure Chemical Industries (Osaka, Japan). Isoflurane was purchased from DS Pharma Animal Health Co. (Osaka, Japan).

Animals

All avian species including raptors used in this study are shown in Table 1. Chickens (13 months old) were provided from the farm at Hokkaido University (Sapporo, Japan). Turkeys (20 months old) were purchased from Sankyo Labo Service Corporation, Inc. (Tokyo, Japan). Canaries (six months old) were purchased from a local pet shop (Sapporo, Japan). These three species were included in this study because the genome information of both VKORC1 and VKORC1L1 are available. Turkeys and canaries were acclimatized to the environment at a constant temperature (22°C ± 1°C) with a 12:12 h light:dark cycle, and given food and water *ad libitum* for 1 week before commencement of the experiment. Chickens, turkeys and canaries were anesthetized using isoflurane and euthanized by CO₂. After euthanasia, nine different tissues were dissected, including liver, pancreas, spleen, testis or ovary, kidney, lung, heart, brain, and muscle. The excised tissues were cut into small pieces and stored in RNA*later*® (Sigma-Aldrich) at –20°C after an overnight incubation at 4°C. The rests of the tissues were immediately frozen in liquid nitrogen and stored at –80°C until use.

Livers of raptors were provided by the Institute for Raptor Biomedicine Japan (Kushiro, Japan), and Sapporo Maruyama Zoo (Sapporo, Japan). After the raptors were dead due to illness or traffic accidents, some raptor individuals were dissected immediately, while dead body of some individuals were kept at 4°C in a few days, then

dissected. After dissection, liver samples were immediately frozen in liquid nitrogen and stored at -80°C until use.

All experiments using animals were performed under the supervision and with the approval of the Institutional Animal Care and Use Committee of Hokkaido University, Japan (approval number 14-0119).

Preparation of liver microsomes

Livers were taken from animals and liver microsomes were prepared as described previously (Watanabe et al., 2010). Because the livers of canaries were very small, two pools were made from six livers for the preparation of microsomes. For other species, a microsome was made from an individual. Briefly, the livers were homogenized with three times their volume of potassium phosphate buffer (KPB, 0.1M, pH7.4). The homogenates were centrifuged at $9000 \times g$ at 4°C for 20 min. The supernatant was decanted to an ultracentrifugation tube and centrifuged at $105000 \times g$ at 4°C for 60 min. The pellet was homogenized in KPB in ice and then centrifuged again at $105000 \times g$ at 4°C for 60 min for washing. The resultant microsomal pellets were homogenized in KPB again. The suspensions were transferred to 1.5 mL tubes and stored at -80°C until use. The protein concentration of hepatic microsomes was measured with the BCA protein assay kit (Thermo Fisher Scientific, Waltham, MA) according to the manufacture's instruction.

Preparation of VKO

VKO was produced as previously described (Tishler et al., 1940). Briefly, 100 mg of vitamin K_1 was dissolved in 5 mL of ethanol and pre-incubated at 75°C for 5 min. 1 mL of 30% hydrogen peroxide and 250 μL of water containing 100 mg of sodium carbonate were added to the vitamin K solution and the mixture was kept at 75°C for 15 min. The mixture was cooled, diluted with 10 mL of water and 40 mL of diethyl ether, and centrifuged at $1000 \times g$ for 10 min. The supernatant was evaporated under nitrogen

and dissolved in methanol. The solution was refined by high-pressure liquid chromatography (HPLC) using the following components: a PU-980 pump (Jasco, Tokyo, Japan); an Inertsil PREP-ODS column, 30.0 × 250 mm (GL Science Inc., Tokyo, Japan); a Mightysil, RP-18GP Aqua guard column (Kanto Chemical Co. Inc., Tokyo, Japan); and a 4.6 × 5 mm, 5 µm SPD-6AV detector (SHIMADZU, Kyoto, Japan). The detection wavelength was 270 nm, the flow rate 10.0 ml/min, and the mobile phase 3% double distilled water (DDW) in methanol. The refined solution was evaporated under nitrogen and dissolved in ethanol. The VKO was kept at 4°C and shielded from light until use.

The concentrations of VKO and vitamin K₁ were determined spectrophotometrically using a molar absorption coefficient of 30,800 M⁻¹ cm⁻¹ at 266 nm and of 18,900 M⁻¹ cm⁻¹ at 249 nm each (Wallin and Martin, 1987).

VKOR activity assay

VKOR activity was assayed in duplicate for each sample as described previously with slight modifications (Watanabe et al., 2010; Hammed et al., 2013). The reaction mixture (500 µL, total volume) contained microsomes (1.0 mg/ml, final concentration), HEPES buffer (pH 7.4, 0.1 M), and VKO (12.5, 100 or 400 µM). After 5 min pre-incubation, the reaction was started by the addition of 2 mM DTT solution. The reaction was performed for 5 min and stopped by adding 1 mL of an iced 1:1 isopropyl alcohol/hexane solution.

The temperatures for pre-incubation and reaction were 41.5°C for turkeys, 42°C for chickens, 38.5°C for canaries, 40.5°C for a peregrine falcon, sparrowhawks, goshawk, Steller's sea eagle and white-tailed eagle, and 39.5°C for snowy owls and great horned owls (McNab 1966; Richards 1971; Siegfried et al., 1975; Chaplin et al., 1984; Herrero and Barja, 1998).

To perform the liquid-liquid extraction of vitamin K, 2 mL of 1:1 isopropyl alcohol/hexane solution and 1 mL HEPES buffer were added. After centrifugation at 750 ×g for 10 min, the organic layer was collected and dried under nitrogen. The dry residue

was dissolved in 80 μ L of methanol, and the reaction product was analyzed by HPLC coupled with UV-VIS detector quantified at 270 nm (Pump: LC-20AB, Detector: SPD-20A; SHIMADZU, Kyoto, Japan). Separation was achieved on an Inertsil ODS-3, 2.1 $\times$ 150 mm, 5 μ m analytical column (GL Sciences, Kyoto, Japan) run at 40°C at 0.5 mL/min in 99.8% methanol. Vitamin K concentration in all samples except for *Haliaeetus* (Steller's sea eagle and white-tailed eagle) was higher than the limit of quantitation (0.0378 μ M).

Inhibition of VKOR activity

The assays were done using the same procedure as for the VKOR activity assay as described above. The reaction was performed at concentrations of 100 μ M of substrate (VKO) and 1 or 10 μ M of warfarin-sodium.

Total RNA extraction and cDNA synthesis

Total RNA was extracted from ten different tissues (liver, pancreas, spleen, testis or ovary, kidney, lung, heart, brain and muscle) of chickens and turkeys, and the livers of night heron, snowy owl, and great horned owl, using NucleoSpin® RNA II (TAKARA BIO INC., Tokyo, Japan). The purity and quantity of RNA were determined by electrophoresis as well as spectrophotometry using NanoDrop ND-1000 (Thermo Scientific, DE). $A_{260/280}$ and $A_{260/230}$ were generally ≥ 2 . Total RNA (10 μ g) was reverse transcribed using ReverTra Ace (TOYOBO, Osaka, Japan) in a final volume of 100 μ L, according to manufacturer's instructions.

Partial cloning of VKORC1L1

VKORC1L1 of night heron, snowy owl and great horned owl were cloned in this study. The cDNA was amplified by polymerase chain reaction (PCR) using specific primers shown in Table S1. The PCR was performed using *Ex Taq*® (Takara, Tokyo, Japan). The PCR cycle program was on one cycle of 30 s at 94°C, 40 cycles of 30 s at

94°C, 30 s at 56.8°C, and 30 s at 72°C, with one cycle of final extension for 1 min at 72°C.

Plasmids were constructed with the PCR products and pCR2.1-TOPO vector using a TOPO TA Cloning Kit (Invitrogen, CA) and transformed into DH5 α -competent cells (TOYOBO, Osaka, Japan). Plasmids were purified using a plasmid miniprep spin kit (Qiagen, Tokyo, Japan). Inserts were sequenced using a BigDye Terminator version 1.1 (Applied Biosystems, Foster City, CA). Ethanol precipitation was performed after the amplification reaction, and the plasmid sequence was analyzed by an automated DNA sequencer, ABI Prism 310 Genetic Analyzer (Thermo Fisher Scientific Inc., Kanagawa, Japan), following the manufacturer's instructions.

Amino acid sequence alignment of VKORC1 and VKORC1L1

The VKORC1 and VKORC1L1 genes of animals including various avian species were retrieved using an NCBI BLAST search (https://blast.ncbi.nlm.nih.gov/Blast.cgi?CMD=Web&PAGE_TYPE=BlastHome; Table S2). Amino acid sequences were aligned by MUSCLE (Edgar 2004).

Phylogenetic analysis

The phylogenetic relationship of VKORC1 and VKORC1L1 was inferred by using the Maximum Likelihood method based on the Le_Gascuel_2008 model (Le and Gascuel, 2008). The model was selected based on Akaike information criterion with a correction for finite sample sizes (AICc). The bootstrap consensus tree inferred from 500 replicates (Felsenstein 1985) [32]. Branches with poor bootstrap values (less than 60%) are collapsed. Initial tree(s) for the heuristic search were constructed by the Neighbor-Joining method with a JTT model. A discrete Gamma distribution was used to model evolutionary rate differences among sites (5 categories (+G, parameter = 2.8254)). The rate variation model allowed for some sites to be evolutionarily invariable ([+I], 8.9717% sites). The analysis involved 42 amino acid sequences and 93 positions. Only positions

with more than 85% site coverage were analyzed. Phylogenetic analyses were performed by Molecular Evolutionary Genetics Analysis (MEGA) ver. 6.06 (Tamura et al., 2013).

Plasmid constructions for quantitative real-time PCR

The cDNA of chickens and turkeys was amplified by PCR using specific primers for VKORC1, VKORC1L1, glyceraldehyde-3-phosphate dehydrogenase (GAPDH), and actin beta (β -actin) shown in Table S3. Because VKORC1 of chickens and turkeys could not be cloned using the same primers as for quantitative real-time PCR, VKORC1 was cloned using the alternative primers. The PCR was performed using SapphireAmp® (Takara, Tokyo, Japan). The PCR cycle program was on one cycle of 30 s at 94°C, 40 cycles of 5 s at 98°C, 5 s at 63°C, and 5 s at 72°C, with one cycle of final extension for 1 min at 72°C.

Plasmids were constructed with the PCR products and pCR2.1-TOPO vector using a TOPO TA Cloning Kit (Invitrogen). All sequences inserted into the plasmids include an amplicon of the quantitative real-time PCR products.

Quantitative real-time PCR

Gene-specific quantitative real-time PCR primers (Table S3) were synthesized by Sigma- Aldrich (Tokyo, Japan). The efficiency of all primers was 89%–107%. Quantitative real-time PCR (qPCR) was performed with the StepOnePlus Real-Time PCR system (Applied Biosystems, Foster City, CA). The 10 μ l PCR reaction mixtures consisted of the Fast SYBR Green Master Mix (Applied Biosystems), forward and reverse primers (200 nM each), and cDNA derived from 80 ng of total RNA. Plasmids containing each amplicon were used for the calibration curves. All samples, including cDNA derived from the tissues of chickens and turkeys and the plasmid standards, were analyzed in duplicate using the following protocol: 95°C for 20 s followed by 40 cycles of 95°C for 3 s and 60°C for 30 s. At the end of each PCR run, melt curve analysis was performed in the range of 60°C–95°C. PCR products were confirmed as single fragments

by electrophoresis and direct or plasmid sequencing methods. For the negative control, DDW was added in the reaction mixture instead of cDNA, and confirmed no contamination

Both the GAPDH and the β -actin tested in this study were not appropriate as housekeeping genes among various tissues, since the differences of Ct values among various tissues varied 5-10, resulting in the wide variation of copy numbers (2^5 – 2^{10} copies). In addition, because appropriate housekeeping were not found in any other avian genes (Olias et al., 2014), mRNA expression of VKORC1 and VKORC1L1 were not compensated and shown as “copy number/ng total RNA” calculated from the standard curve with plasmids.

Statistical analysis

For the comparison of VKOR activity and inhibition assays among avian species, significant difference among avian groups were analyzed using a Steel-Dwass test because the data were not parametric. All the statistical analysis was performed with a significance level of $p < 0.05$, using JMP software (version 12.0; SAS Institute, Cary, NC). Because of the small sample size, goshawk and sparrowhawks, and snowy owls and great horned owls were grouped for statistical testing (*Accipiter* and *Bubo*, respectively).

RESULTS

Comparison of VKOR activity among avian species

Results of VKOR activity among eight avian species are shown in Fig. S2 and Table 2. At every concentration of VKO (12.5, 100, and 400 μ M), there was no significant difference between male (78 ± 28 , 124 ± 50 , and 153 ± 57 pmol/min/mg protein respectively, n=3) and female (49 ± 31 , 97 ± 66 , and 103 ± 65 pmol/min/mg protein respectively, n=3) chickens. Therefore, both sexes from the same species were analyzed together in other avian samples.

VKOR activities of raptors and canaries were lower than that of turkeys and chickens (Table 2, Fig. S2). At 400 μ M VKO, VKOR activities were following orders; turkeys (270 ± 17 pmol/min/mg protein, n=3), male chickens (153 ± 57 pmol/min/mg protein, n=3), female chickens (103 ± 65 pmol/min/mg protein, n=3), *Bubo* (snowy owls and great horned owls; 83 ± 21 pmol/min/mg protein, n=4), and *Accipiter* (goshawk and sparrowhawks; 45 ± 5 pmol/min/mg protein, n=3). The VKOR activity of peregrine falcon was only 14 pmol/min/mg protein (n=1). That of canaries was 85 pmol/min/mg protein (2 pools). VKOR activity in raptors ranged to a small fraction (5.2–81%) of that observed in turkey and chicken hepatic microsomes. VKOR activity in peregrine falcon accounted for 5.2% of that in turkeys, and VKOR activity in *Bubo* accounted for 81% of that in female chickens. VKOR activities of *Haliaeetus* (Steller's sea eagle and white-tailed eagle) were undetectable.

Comparison of VKOR activity inhibited by warfarin among avian species

Fig. 1A shows VKOR activity without warfarin, and VKOR activity with 1 or 10 μ M warfarin incubation at 100 μ M of VKO. With 1 μ M warfarin, VKOR inhibition did not occur in chickens and turkeys. However, VKOR inhibition was found in other avian species. With 10 μ M warfarin, inhibition of VKOR activity occurred in every avian species. For male chickens (n=3), VKOR activity without warfarin, with 1 μ M warfarin, and with 10 μ M warfarin were 124 ± 50 , 132 ± 53 , and 83 ± 27 pmol/min/mg protein,

respectively. For female chickens (n=3), VKOR activity without warfarin, with 1 μ M warfarin, and with 10 μ M warfarin were 97 ± 66 , 104 ± 71 , and 72 ± 44 pmol/min/mg protein, respectively. For turkeys (n=3), those were 222 ± 36 , 207 ± 30 , and 117 ± 24 pmol/min/mg protein, respectively. For a goshawk, those were 38, 29, and 26 pmol/min/mg protein, respectively. For sparrowhawks (n=2), those were 36, 29, and 20 pmol/min/mg protein, respectively. For snowy owls (n=2), those were 79, 56, and 35 pmol/min/mg protein, respectively. For great horned owls (n=2), those were 61, 52, and 19 pmol/min/mg protein, respectively. For a peregrine falcon, those were 14, 12, and 9 pmol/min/mg protein, respectively. For canaries (2 pools), those were 75, 33, and 9 pmol/min/mg protein, respectively.

Fig. 1B shows the percentage of VKOR activity with 1 or 10 μ M warfarin incubation compared to VKOR activity of untreated microsomes. For male and female chickens, and turkeys, 1 μ M warfarin inhibited control activity by $106 \pm 4.6\%$ and $106 \pm 16\%$, and $93 \pm 1.7\%$, respectively. For *Accipiter* (a goshawk and sparrowhawks) and *Bubo* (snowy owls and great horned owls), 1 μ M warfarin inhibited control activity by $80 \pm 13\%$ and $79 \pm 11\%$, respectively. 1 μ M warfarin inhibited control activity for *Bubo* (snowy owls and great horned owls) compared to male and female chickens. For a peregrine falcon and canaries, 1 μ M warfarin inhibited control activity by 85% and 45%, respectively.

For male and female chickens, and turkeys, 10 μ M warfarin inhibited control activity by $69 \pm 8.8\%$, $75 \pm 16\%$, and $53 \pm 2.2\%$, respectively. For *Accipiter* (a goshawk and sparrowhawks) and *Bubo* (snowy owls and great horned owls), 10 μ M warfarin inhibited control activity by $60 \pm 13\%$ and $38 \pm 15\%$, respectively. 10 μ M warfarin inhibited control activity for *Bubo* (snowy owls and great horned owls) compared to male and female chickens. For a peregrine falcon and canaries, 10 μ M warfarin inhibited control activity by 62% and 12%, respectively.

Amino acid sequence alignment of VKORC1 and VKORC1L1 in avian species

The nucleotide sequence of VKORC1 has been defined for only nine avian species in the NCBI database, despite the fact that it has been defined for more than 500 mammals. The nucleotide sequence of raptor VKORC1 has not been clarified. Fig. S3 describes the phylogenetic tree of the VKOR protein. Fig. S4-A shows the amino acid sequence alignment of VKORC1 for nine avian species (turkey, brown roatelo, chicken, ostrich, canary, hummingbird, sandgrouse, crested ibis and emperor penguin) and six other animals (human, rat, mouse, turtle, frog and fugu). The CXXC motif, which is supposed to be the active site of the VKORC1 (Rost et al., 2005; Wajih et al., 2005; Quan et al., 2007), had the following variations: CIVC (human, rat and mouse); CLVC (turkey, brown roatelo, chicken, ostrich, canary, crested ibis, emperor penguin and turtle); CPVC (hummingbird); CVIC (sandgrouse and frog); and CMVC (fugu).

Among avian species, there were three types variations at 68th, 76th, and 143rd amino acid, respectively. The 68th amino acid mutation causes warfarin resistance in human (Hodroge et al., 2012). The 76th amino acid mutation causes warfarin resistance in rats (Tanaka et al., 2012). The 143rd amino acid mutation causes high VKOR activity in rats (Rost et al., 2009). There were two variations among avian species at the 141st amino acid, whose mutation causes low VKOR activity in rats (Rost et al., 2009). There were five variations among avian species at the 140th amino acid in the warfarin-binding site: alanine (turkey and brown roatelo); valine (chicken); glycine (ostrich, canary and hummingbird); isoleucine (sandgrouse); and leucine (crested ibis and emperor penguin).

The nucleotide sequence of VKORC1L1 has been defined for more than 100 avian species and more than 200 mammals in the NCBI database. Fig. S4-B shows the amino acid sequence alignment of VKORC1L1 of seven raptors, ten other avian species, and six other animals. The CXXC motif had the only three variations: CIIC (human, rat, mouse and turtle); CIVC (avian species); and CVIC (frog and fugu).

Few studies have reported the amino acid mutations of VKORC1L1 that contribute to VKOR activity. Hünérberg (2009) used HEK cells expressing recombinant human VKORC1L1, and reported that 36th or 65th amino acid mutations caused warfarin

resistance of VKOR activity, and 135th or 146th amino acid mutations caused low VKOR activity. There was no diversity among avian species in the present study at 36th, 65th, 135th, or 146th amino acid. However, the 36th amino acid of avian species was different to that of mammals (leucine in avian and valine in mammalian species; Fig S4-B). In the warfarin-binding site, the 147th amino acid of avian species was also different to that of mammals (leucine in avian and valine in mammalian species).

Distribution of VKORC1 and VKORC1L1 mRNA in tissues of turkey and chicken

Fig. 2A shows the patterns of the mRNA expression of VKORC1 and VKORC1L1 among nine different tissues from male turkeys. VKORC1 expression in liver was the highest of all tissues (i.e. 4-fold higher than in kidney and testis, 7-fold higher than in lung, 9-fold higher than in spleen, 11-fold higher than in brain, 15-fold higher than in muscle and pancreas, and 16-fold higher than in heart). VKORC1 was also expressed at higher levels than VKORC1L1 in six additional tissues. The expression ratio of VKORC1:VKORC1L1 was 8:1 in liver, 4:1 in pancreas, 3:1 in kidney, 3:2 in spleen, 5:4 in muscle and lung, 1:1 in heart and testis, and 2:3 in brain (Fig. 2B).

Fig. 3A shows the patterns of the mRNA expression of VKORC1 and VKORC1L1 among ten different tissues from male and female chickens. There was no significant difference in the mRNA expression levels between male and female chickens. VKORC1 expression in ovaries was 2-fold higher than in liver. VKORC1 expression in the other tissues was lower than in liver (i.e., 2-fold lower in lung, spleen and brain, 3-fold lower in kidney, 5-fold lower in testis and heart, 9-fold lower in muscle, and 23-fold lower in pancreas). VKORC1L1 was expressed at higher levels than VKORC1 in all tissues except for spleen. The expression ratio of VKORC1:VKORC1L1 was 5:4 in spleen, 4:5 in pancreas, 2:3 in liver, 1:2 in heart, ovary and lung, 1:4 in brain and muscle, 1:9 in kidney, and 1:19 in testis (Fig. 3B).

DISCUSSION

1. Avian interspecific differences in VKOR activity

1-1. Comparison of VKOR activity among avian species

This study suggested hepatic VKOR activities of raptors were lower than that of turkeys and chickens. VKOR plays an important role in the vitamin K cycle for activations of vitamin K-dependent coagulation factors. Inhibiting VKOR by ARs induces lethal hemorrhages in raptors (Murray, M, 2018; Rattner et al., 2012; Rattner et al., 2014; Rattner et al., 2015). Because the present study focused on comparing the maximum VKOR activity between avian species, 12.5, 100, and 400 μ M of VKO were used as substrates for 1.0 mg/mL liver microsomes. The estimated concentration of VKO is 12.5, 100, and 400 μ mol/g liver microsomes, respectively. However, the physiological concentration of VKO is approximately 13 pmol/g in chicken liver (Will et al., 1992) and 0.1 pmol/g in mice liver (Okano et al., 2008). The concentration of VKO used for the VKOR activity assay in the present study is much higher than physiological concentrations. It is also important to investigate the activity under physiological concentrations of the substrate.

It is reported that concentration of vitamin K in liver of chickens is less than that of rats, while the concentration of VKO in liver is higher in chickens than in rats (Will et al., 1992). This indicates that chickens have a poor ability to reduce VKO in the vitamin K cycle, so they require high doses of vitamin K in their diets. Similarly, it is possible that raptors require high doses of dietary vitamin K. K vitamins are divided into two major types. Vitamin K₁ (phylloquinone) is synthesized in plants and is the primary dietary source of vitamin K. Vitamin K₂ (menaquinones, MK) represents a family of many different subtypes. One subtype, MK₄, is converted from phylloquinone and is the major physiological vitamin K form (Van Horn 2013). Although many other menaquinones are converted by gut bacteria, the menaquinones of gut origin make a relatively minor contribution to the hepatic stores of vitamin K. Phylloquinone and MK₄ are also available from meat, including rats and mice, which are diets of raptors (Will et

al., 1992; Elder et al., 2006; Okano et al., 2008). Considering these previous studies, raptors are presumed to gain vitamin K from rodents and other prey in the wild or menaquinones synthesized in gut to respond to the dose of required vitamin K.

1-2. VKOR activity related to amino acid sequence and mRNA expression of VKORC1 and VKORC1L1

This study suggested that not only VKORC1, but also VKORC1L1 is important for VKOR activity in avian species. It was reported that the $V_{\max}$ of VKORC1 was 16-fold higher than that of VKORC1L1 in rats (Hammed et al., 2013). Moreover, mRNA expression of VKORC1 was over 10-fold higher than that of VKORC1L1 in rat and mouse livers (Hammed et al., 2013; Caspers et al., 2015). Therefore, VKORC1 may support 96% of VKOR activity in mammals. Conversely, the individual $V_{\max}$ of VKORC1 and VKORC1L1 have been unknown in avian species. Because the avian CXXC motif was different between VKORC1 (CLVC, Fig. S4-A) and VKORC1L1 (CIVC, Fig. S4-B), it is possible that the $V_{\max}$ values could be different between avian VKORC1 and VKORC1L1. While mRNA of VKORC1 was expressed higher than that of VKORC1L1 in turkey liver (Fig. 2), both mRNA of VKORC1 and VKORC1L1 were equivalently expressed in chicken liver (Fig. 3). Therefore, in contrast to mammals, it is possible that avian VKOR activity is supported by both VKORC1 and VKORC1L1, and the mRNA expression ratio could cause avian interspecific differences in VKOR activity.

Although it is difficult to compare the absolute amount of VKORC1 and VKORC1L1 between turkeys and chickens, the total expression amount of VKORC1 and VKORC1L1 was 2.5-fold higher in turkey liver (total amount was 3050, VKORC1 was 2700, and VKORC1L1 was 350 copy number/ng total RNA) than in chicken liver (total amount was 1200, VKORC1 was 500, and VKORC1L1 was 700 copy number/ng total RNA). The total expression amount of VKORC1 and VKORC1L1 could also cause avian interspecific differences in VKOR activity. Information on the mRNA expression ratio and the total quantities of VKORC1 and VKORC1L1 from various avian species might

provide some basis to account for interspecific differences in VKOR activity.

2. Avian interspecific differences in sensitivity of VKOR activity to warfarin

2-1. Comparison of VKOR activity inhibited by warfarin among avian species

This study suggested that the percentage of hepatic microsomal VKOR activity remaining after warfarin incubation had a rank order of chickens > turkeys, *Accipiter* (goshawk and sparrowhawks) and peregrine falcon > *Bubo* (snowy owls and great horned owls) > canaries. This result indicates that the VKOR activity of raptors, especially *Bubo*, is inhibited by warfarin more easily than that of chickens, which is supported in part by the observation of 20-30-fold greater toxicity of the AR diphacinone in American kestrel compared to that described for Northern bobwhite and mallards (Rattner et al., 2018a). Therefore, it is not recommended to extrapolate poultry data to raptors for risks assessment of ARs in raptors. Moreover, because there were interspecific differences between raptors and canaries, it is also important to examine individual species data about the inhibited rate of VKOR activity for accurate risk assessment of ARs in non-target avian species.

2-2. Amino acid sequence of VKORC1

For rats, the mutation of tyrosine at amino acid 139 in the sequence of VKORC1 shows resistance to warfarin, and it has been inferred that the Thr-Tyr-Ala (TYA) motif (at amino acid 138 to 140) is the warfarin-binding site (Rost et al., 2005). Hence, the warfarin-binding site of VKORC1 sequence was compared among avian species. The motifs in chickens, turkeys, and canaries were TYV, TYA, TYG, respectively (Fig. S4-A). The VKOR activity of chickens was more resistant to warfarin compared to that of turkeys (Fig. 1B). It was reported that chickens had a 40-fold higher K_i than rats for VKOR activity in response to warfarin (Watanabe et al., 2010). Combined with the fact that the warfarin-binding site of VKORC1 is TYV in chickens and TYA in turkeys and rats (Fig. S4-A), warfarin resistance might be higher with TYV than TYA.

Surprisingly, the VKOR activity of canaries was strongly inhibited compared to other avian species, including raptors. In canaries, the warfarin-binding site of VKORC1 is TYG as well as ostriches (Fig. S4-A). A previous study reported that the VKOR activity of ostriches was also strongly inhibited: inhibited VKOR activity was 29% at 100 μ M of VKO with 1 μ M of warfarin (Watanabe et al., 2010). These data suggest that TYG is a significant determinant of sensitivity to warfarin and possibly other ARs. The amino acid at position 140 of VKORC1 could be important for avian interspecific difference in VKOR inhibition. Information about VKORC1 sequences and structures of other avian species, especially raptors, are needed for more accurate avian VKOR characterization and risk assessments of ARs.

2-3. Amino acid sequence of VKORC1L1

Because the sequence of VKORC1L1 in all the avian species shown in Fig. S4-B had the same warfarin-binding site (TYL), the warfarin-binding site of VKORC1L1 does not cause avian interspecific differences in VKOR susceptibility. There is little information about amino acid mutations in VKORC1L1. It was reported that warfarin resistance was caused by a V36L mutation in VKORC1L1 obtained from HEK-293-EBNA cell microsomes expressing recombinant human VKORC1L1 (Hünerberg 2009). As shown in Fig. S4-C, the 36th amino acid was valine in the VKORC1L1 of human and rats, and in VKORC1 of human, rats, and avian species. However, interestingly the 36th amino acid was leucine in avian VKORC1L1. Therefore, it is possible that the leucine at the 36th amino acid position in avian VKORC1L1 causes warfarin resistance in VKOR activity. More information about avian VKORC1L1 is needed, including mutations.

2-4. mRNA expression of VKORC1 and VKORC1L1

In avian species, the K_i for VKOR activity in response to warfarin is unknown for both VKORC1 and VKORC1L1. It was reported that the K_i for VKOR activity in response to warfarin was 50-fold higher in VKORC1L1 than in VKORC1 obtained from

yeast microsomes expressing recombinant proteins from rats (Hammed et al., 2013). Moreover, VKOR activity in the testis (predominantly VKORC1L1 mRNA) was more resistant to warfarin compared to that of the liver (predominantly VKORC1 mRNA). Therefore, high expression of VKORC1L1 may induce warfarin resistance in mammals. The VKOR activity of chickens was more resistant to warfarin than that of turkeys (Fig. 1B). The mRNA expression ratio of VKORC1:VKORC1L1 differed between turkey (8:1, Fig. 2B) and chicken liver (2:3, Fig. 3B). This result suggests that expression ratio of VKORC1:VKORC1L1 could account for interspecific differences in VKOR inhibition by ARs among various species of birds. Both the K_i of avian individual specific VKORC1 and VKORC1L1 and expression ratio should be considered for the risk assessment of ARs in wild avian species.

Although our study focused on avian interspecific differences in VKOR, which is the target molecule of ARs, the warfarin-binding capacity of albumin in serum is also an important factor for avian interspecific differences in susceptibility to ARs. Chicken albumin may have a greater warfarin-binding capacity (9-fold greater than human), resulting in a longer half-life and less toxicity despite high metabolic ability (Rajaian et al., 1997; Watanabe et al., 2015). Although there is no information on raptor albumin, albumin in raptors and chickens may be less identical than in other avian species such as turkey, ostrich, and zebra finch (91%-70% identical amino acid sequences). Lower warfarin binding capacity of albumin in some species of birds (perhaps raptors) might contribute to differences in AR sensitivity.

3. mRNA distribution of VKORC1 and VKORC1L1 in turkey and chicken tissues

Interestingly, the mRNA distribution in turkey tissues was similar to that in rat and mouse tissues: VKORC1 was expressed higher than VKORC1L1 in liver, kidney and lung; and VKORC1L1 was expressed higher than VKORC1 in testis and brain (Fig. 2) (Caspers et al., 2015; Hammed et al., 2013). In contrast, VKORC1L1 was expressed higher than VKORC1 in most chicken tissues, except for spleen (Fig. 3). These results

suggest that interspecific difference in mRNA distribution of VKORC1 and VKORC1L1 may be greater in birds than mammals, and VKORC1L1 can support VKOR activity in liver as well as in extra hepatic tissues in some avian species. In mammals, the focus has principally been on VKORC1. In birds, however, both VKORC1 and VKORC1L1 should be considered.

CONCLUSIONS

In conclusion, our study demonstrated that VKOR activity of raptors is both lower and more readily inhibited by the prototypic AR warfarin compared to poultry. Moreover, both the K_i of VKORC1 and VKORC1L1, and the mRNA expression ratio, may contribute to interspecific difference in AR inhibition of VKOR in birds. The value of K_i for VKOR to ARs could depend on the amino acid at position 140 in the TYX warfarin-binding motif of VKORC1. Further, the value of K_i for VKOR to ARs could depend on other amino acid mutations in VKORC1L1. Therefore, further information about raptor VKORC1 and VKORC1L1, including K_i , mRNA expression ratio and amino acid sequences, are needed to better assess the risk of ARs to raptors. Such data may help elucidate the molecular and genetic factors that contribute to the seemingly greater sensitivity of raptorial and scavenging birds to AR intoxication.

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Table 1. Avian species used in this study.

Common name	Scientific name	Sex	Sample size	Source
Chicken (Rhode island red)	<i>Gallus gallus domesticus</i>	Male	3	a
		Female	3	a
Turkey (Bronze)	<i>Meleagris gallopavo</i>	Male	3	b
Canary (Lemon)	<i>Serinus canaria</i>	Male	6	c
Peregrine falcon	<i>Falco peregrinus pealei</i>	Female	1	d
Sparrowhawk	<i>Accipiter nisus</i>	Female	2	d, e
	<i>nisosimilis</i>			
Goshawk	<i>Accipiter gentilis</i>	Female	1	d
	<i>fuiyamae</i>			
Steller's sea eagle	<i>Haliaeetus pelagicus</i>	Male	1	d
White-tailed eagle	<i>Haliaeetus albicilla</i>	Female	1	d
Snowy owl	<i>Bubo scandiacus</i>	Male	2	e
Great horned owl	<i>Bubo virginianus</i>	Male	1	e
		Female	1	e
Night heron	<i>Nycticorax nycticorax</i>	Female	1	e

a: Farm at Hokkaido University (Sapporo, Japan)

b: Sankyo Labo Service Corporation, Inc. (Tokyo, Japan)

c: Local pet shop (Sapporo, Japan)

d: Institute for Raptor Biomedicine Japan (Kushiro Shitsugen Wildlife Center, Kushiro, Japan)

e: Sapporo Maruyama Zoo (Hokkaido, Japan)

611 Table 2. VKOR activity (produced vitamin K, pmol/min/mg protein) at 12.5, 100 and 400 μ M of VKO in turkeys, chickens, canaries and
612 5 species of raptors. The upper row in each VKO concentration shows individual data, and the bottom row shows the mean $\pm$ S.D.

613

VKO [μ M]	Turkeys	Male chickens	Female chickens	Canaries	Snowy owls ^a	Great horned owls ^a	Goshawk ^b	Sparrow- hawks ^b	Peregrine falcon
12.5				32, 45	51, 51	20, 31	26	23, 28	
	115 $\pm$ 22	78 $\pm$ 28	49 $\pm$ 31	38	38 $\pm$ 15		26 $\pm$ 3		9
100				79, 72	89, 70	44, 79	38	33, 38	
	222 $\pm$ 36	124 $\pm$ 50	97 $\pm$ 66	75	70 $\pm$ 19		36 $\pm$ 3		14
400				77, 94	100, 94	52, 84	42	42, 51	
	270 $\pm$ 17	153 $\pm$ 57	103 $\pm$ 65	85	83 $\pm$ 21		45 $\pm$ 5		14

614 No significant differences in VKOR activity at each substrate (VKO) concentration among turkeys, male chickens, female chickens, *Bubo*^a
615 (snowy owls and great horned owls) and *Accipiter*^b (a goshawk and sparrowhawks) were observed (Steel-Dwass test, $P < 0.05$).
616 VKOR activities in *Haliaeetus* (Steller's sea eagle and white-tailed eagle) were undetectable.

617 Table S1. Primers for avian VKORC1L1 cloning.

618 Primers were designed based on the VKORC1L1 nucleotide sequences.

Gene		Sequence (5'→3') *	Amplicon size (bp)
Avian VKORC1L1	Forward	CTTGGTATGACAGCAAGTGC	225
Avian VKORC1L1	Reverse	CTGTTTGGGTTGGRAGTTGC	

619 * The letters R encodes A and G.

620 Table S2. Accession numbers of VKORC1 and VKORC1L1 genes.

621

Common name	Scientific name	Accession number	
		VKORC1	VKORC1L1
Chicken	<i>Gallus gallus</i>	NM_206807	NM_001001328
Ostrich	<i>Struthio camelus australis</i>	-	XM_009675917
Turkey	<i>Meleagris gallopavo</i>	XM_010726800	XM_003211735
Anna's hummingbird	<i>Calypte anna</i>	XM_008503547	XM_008490706
Yellow-throated sandgrouse	<i>Pterocles gutturalis</i>	XM_010087012	XM_010074393
Brown roatelo	<i>Mesitornis unicolor</i>	XM_010190461	XM_010179204
Crested ibis	<i>Nipponia nippon</i>	XM_009474946	XM_009469732
Night heron	<i>Nycticorax nycticorax</i>	-	LC097088
Emperor penguin	<i>Aptenodytes forsteri</i>	XM_009285145	XM_009287681
Golden eagle	<i>Aquila chrysaetos</i> <i>canadensis</i>	-	XM_011581635
Barn owl	<i>Tyto alba</i>	-	XM_009973792
Snowy owl	<i>Bubo scandiacus</i>	-	LC097089
Great horned owl	<i>Bubo virginianus</i>	-	LC097090

Bald eagle	<i>Haliaeetus leucocephalus</i>	-	XM_010573578
White-tailed eagle	<i>Haliaeetus albicilla</i>	-	XM_009928642
Peregrine falcon	<i>Falco peregrinus</i>	-	XM_005240548
Common canary	<i>Serinus canaria</i>	XM_009100016	XM_009094710
Human	<i>Homo sapiens</i>	NM_024006	NM_173517
Norway rat	<i>Rattus norvegicus</i>	NM_203335	NM_203338
House mouse	<i>Mus musculus</i>	NM_178600	NM_027121
Western painted turtle	<i>Chrysemys picta bellii</i>	XM_005279372	XM_005283424
Western clawed frog	<i>Xenopus tropicalis</i>	NM_001006927	NM_001016769
Fugu rubripes	<i>Takifugu rubripes</i>	NM_001032666	NM_001032768
Japanese medaka	<i>Oryzias latipes</i>	XM_004080314	XM_004076602
Cephalochordata	<i>Branchiostoma floridae</i>	XM_002611843, XM_002587531*	
Purple sea urchin	<i>Strongylocentrotus</i> <i>purpuratus</i>	XM_001181369 *	

622 * Cephalochordata and Purple sea urchin possesses ancestral type of VKOR gene.

623 Table S3. Primers for cloning and qPCR in turkeys and chickens.

624

Gene		Sequence (5'→3')	Amplicon size (bp)	Efficiency (%)	Accession number	Reference
Chicken_VKORC1	Forward	TTTGTGGGTCGGGACAGCGCCA	218	95.6	NM_206807	This study
	Reverse	ACGACGTAGGTGCTGAGGCAGA				
Chicken_VKORC1 _ for cloning	Forward	GTCGGGACAGCGCCATCAACGT	215	Not tested	NM_206807	This study
	Reverse	GTTGACGTAGGTGCTGAG				
Chicken_ VKORC1L1	Forward	AGAAGGGCCGCGATCTCCACTACCA	106	89.2	NM_001001328	This study
	Reverse	CCCAACAGACCGAATCCTCGACCCCAT				
Chicken_GAPDH	Forward	CTCTGTTGTTGACCTGACCT	125	98.8	NM_204305	Watanabe et al., 2013
	Reverse	CAACCTGGTCCTCTGTGTAT				
Chicken_β-actin	Forward	GAGAAATTGTGCGTGACATCA	152	95.3	NM_205518	This study
	Reverse	CCTGAACCTCTCATTGCCA				
Turkey_VKORC1	Forward	TTTGTGGGTCGGGACAGCGCCA	218	107.2	XM_010726800	This study
	Reverse	ACGGCGTAGGTGCTGAGGCAGA				
Turkey_VKORC1_ for cloning	Forward	GTTTGTGGGTCGGGACAGCGCCA	219	Not tested	XM_010726800	This study
	Reverse	ACGGCGTAGGTGCTGAGGCAGA				

Turkey_	Forward	GAAGGGGCGCGATCTCCACTACCA	106	104.0	XM_003211735	This study
VKORC1L1	Reverse	CCCAACAGCCCGAATCCTCGACCCCAT				
Turkey_GAPDH	Forward	CTCTGTTGTTGACCTGACCT	125	98.0	NM_001303179	This study
	Reverse	CAACCTGGTCCTCTGTGTAT				
Turkey_β-actin	Forward	GAGAAATTGTGCGTGACATCA	152	104.0	NM_001303173	This study
	Reverse	CCTGAACCTCTCATTGCCA				

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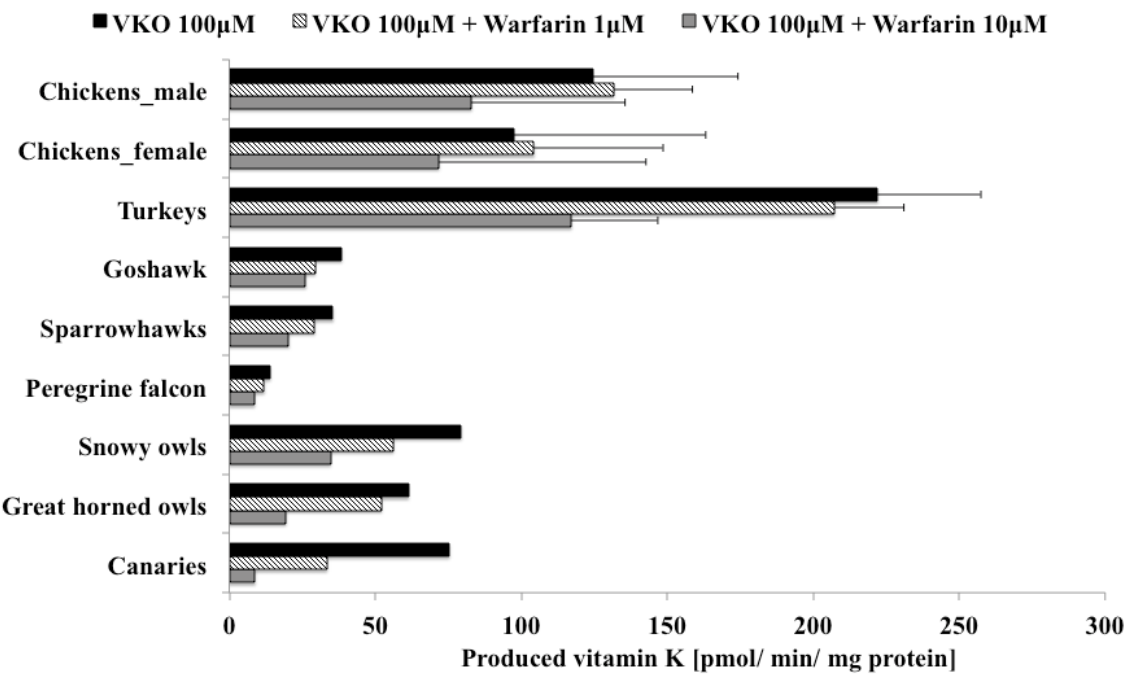
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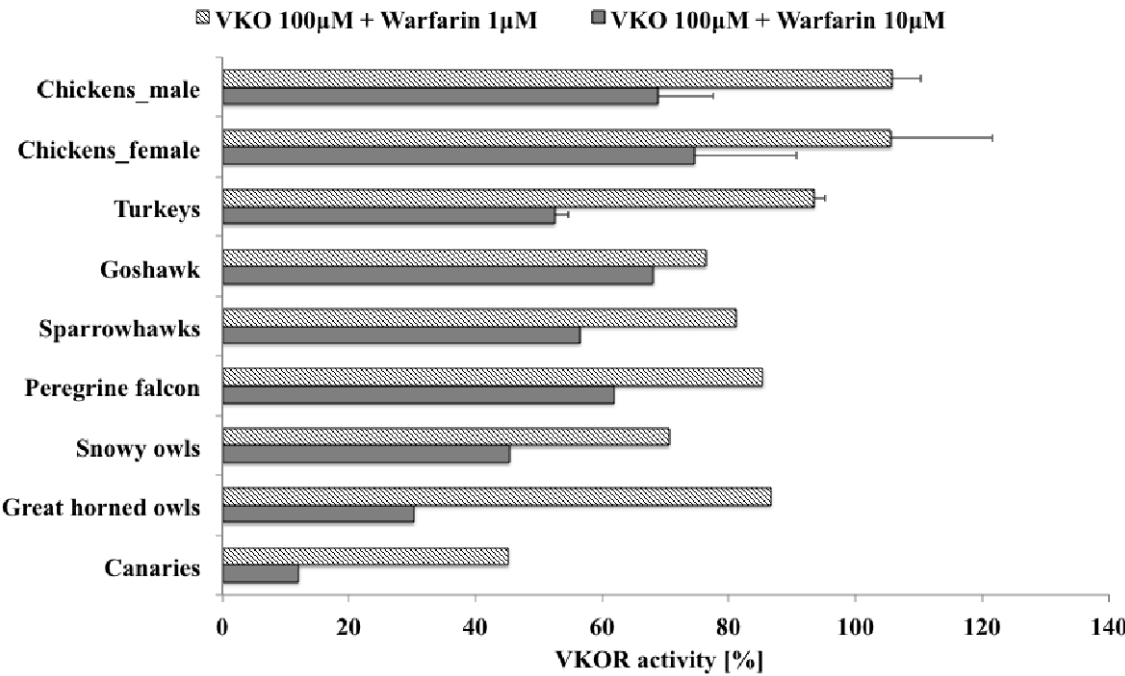
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640 B



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642

643 Fig. 1. VKOR activity inhibited by 1 and 10 µM warfarin at 100 µM of VKO (A) and
644 percentage of inhibited activity (B) in turkeys, chickens, canaries and five species of

645 raptors.

646 The percentage (%) was calculated using the following formula;

647 $\text{VKOR activity (\%)} = \text{VKOR activity with warfarin} / \text{that without warfarin} * 100$

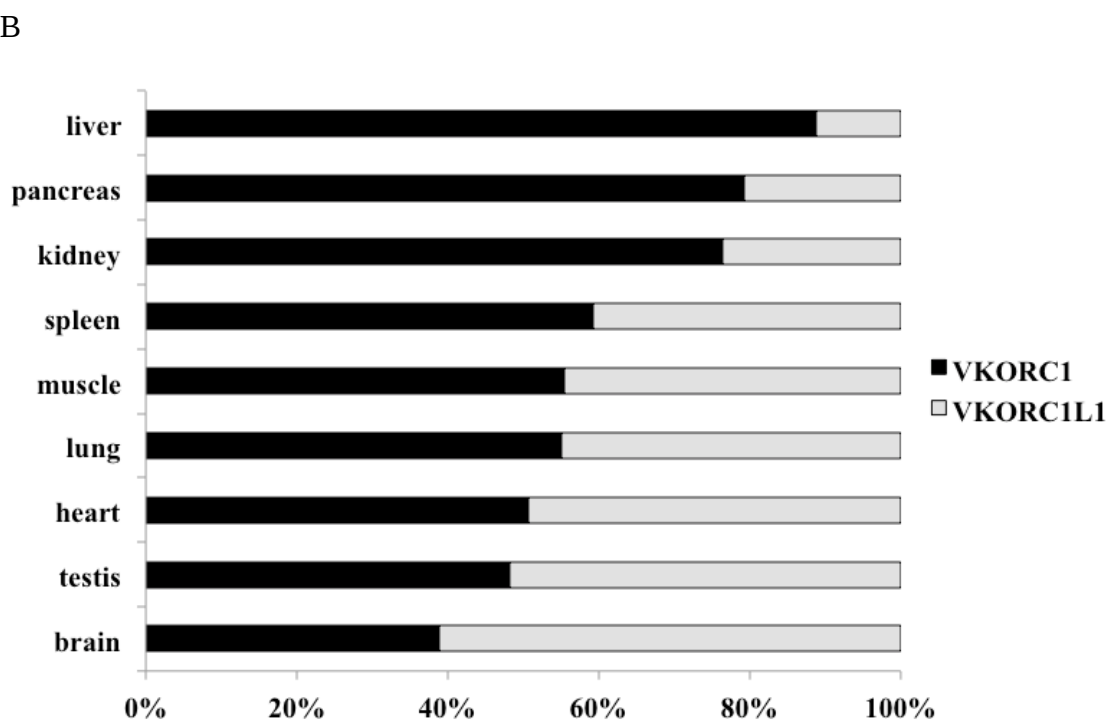
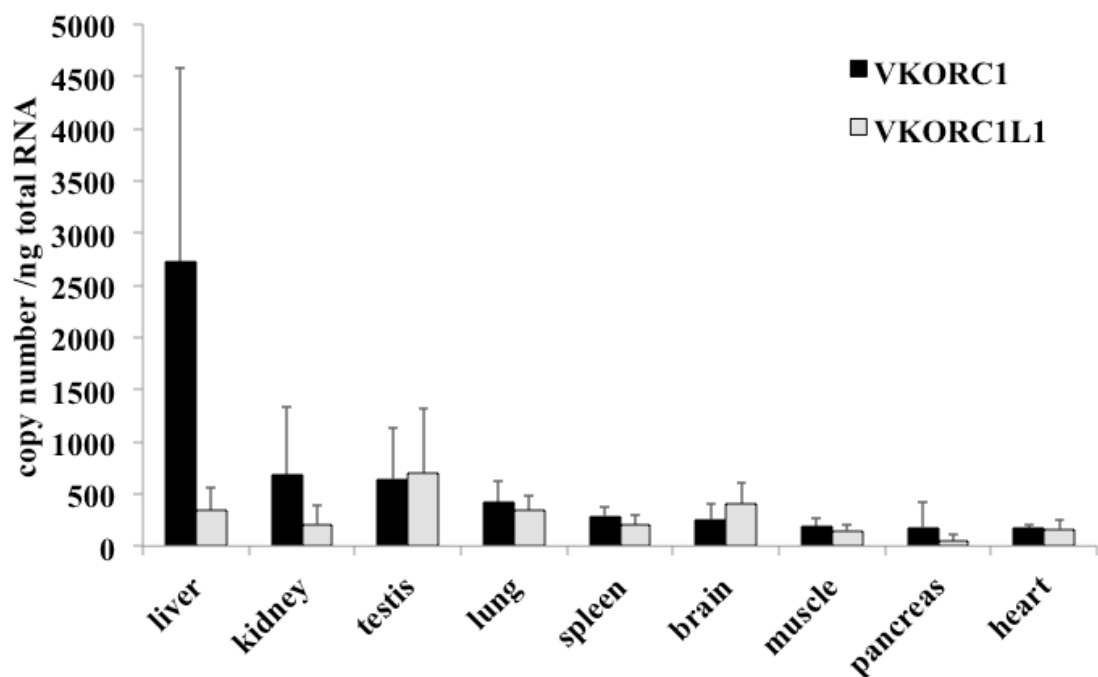
648 Each data point from turkeys and chickens represents the mean of three animals $\pm$ S.D.

649 (error bars). The numbers of other species were less than three. Therefore there are no

650 S.D. values. No significant differences in percentage of VKOR activity at each warfarin

651 concentration among chickens, *Accipiter*^a (a goshawk and sparrowhawks), turkeys and

652 *Bubo*^b (snowy owls and great horned owls) were observed (Steel-Dwass test, $P < 0.05$).



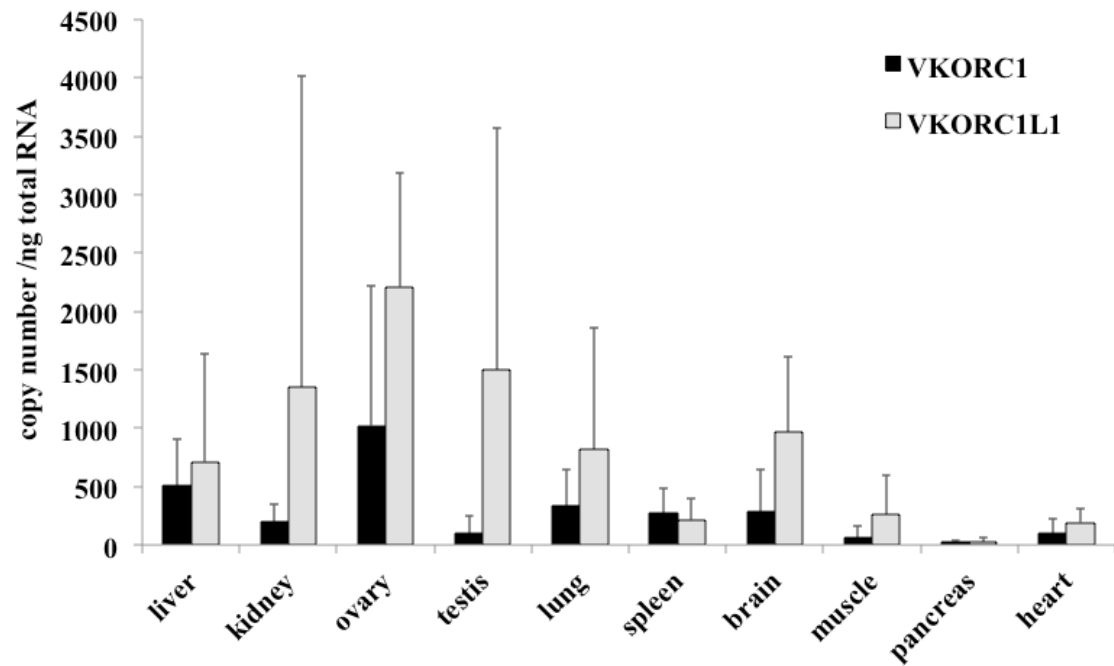
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657 Fig. 2. mRNA expression of VKORC1 and VKORC1L1 (A) and their ratios (B) in nine

658 different tissues of male turkeys. Each data represents the mean of three animals $\pm$ S.D.

659 (error bars).

A



B

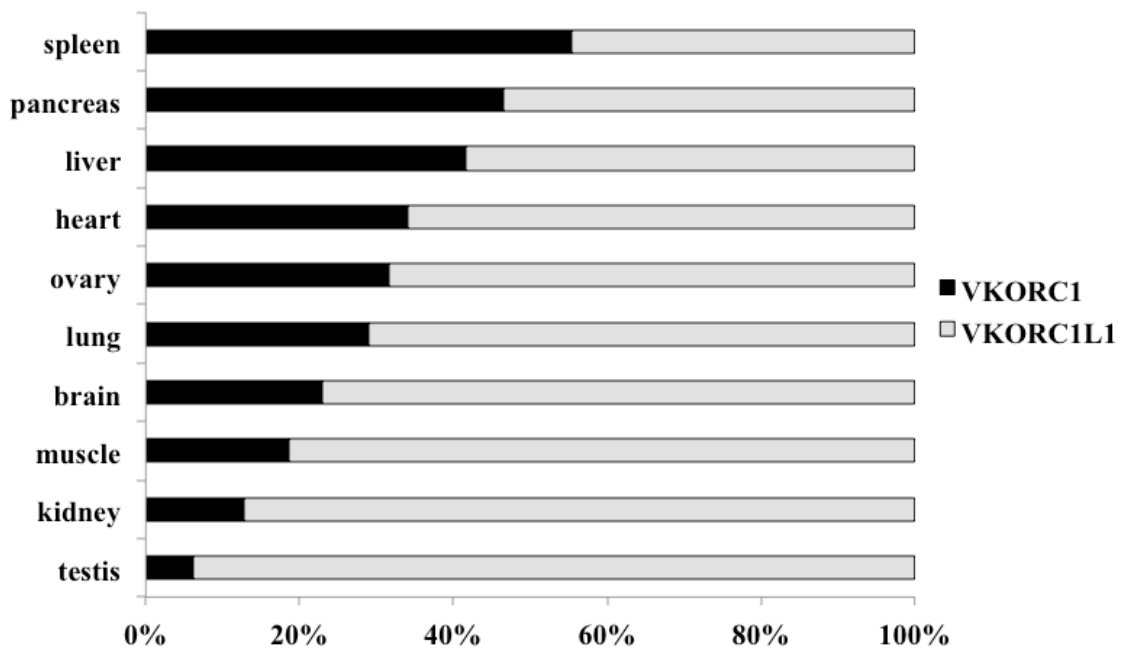


Fig. 3. mRNA expression of VKORC1 and VKORC1L1 (A) and their ratios (B) in ten different tissues of three male and female chickens. Each data point represents the mean of six animals $\pm$ S.D. (error bars) except for ovary and testis. Data of ovary and testis represents the mean of three animals $\pm$ S.D. (error bars).

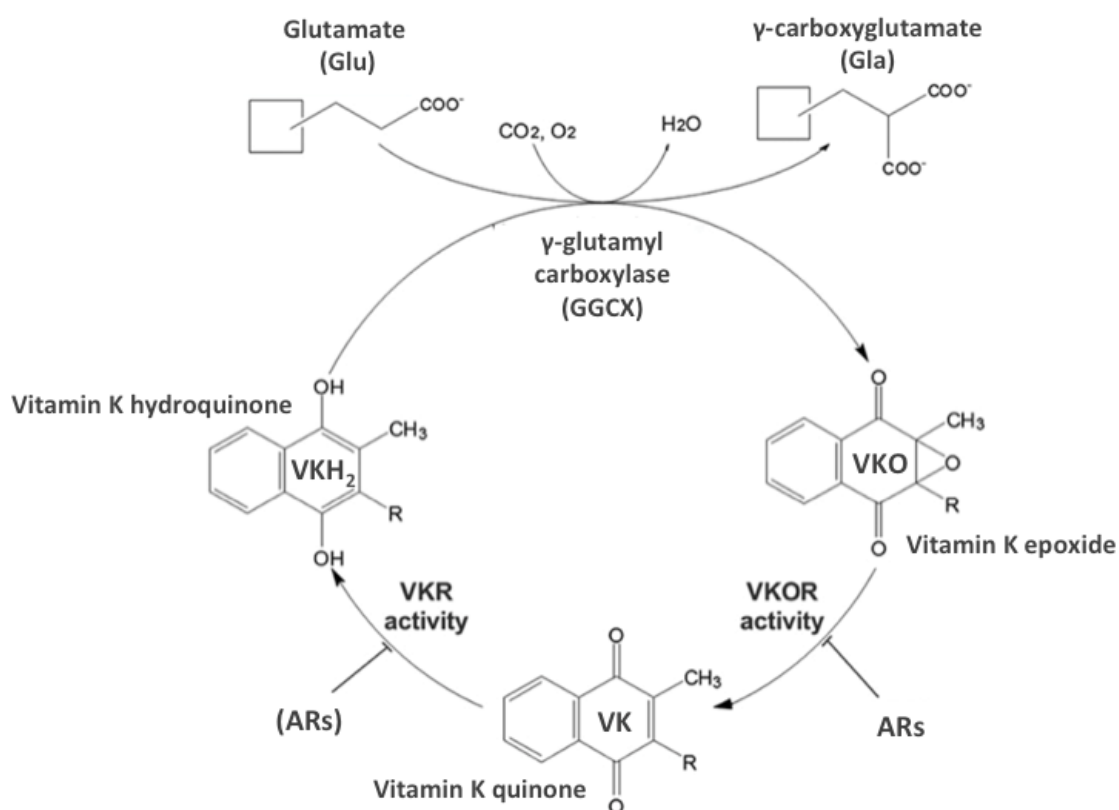


Fig. S1. Modified vitamin K cycle (Hammed et al., 2013; Tie et al., 2013). During vitamin K dependent carboxylation, vitamin K hydroquinone is oxidized to vitamin K 2, 3-epoxide (VKO) by γ -glutamyl carboxylase. VKO is reduced to vitamin K by VKOR. This reaction is inhibited by ARs (e.g. warfarin).

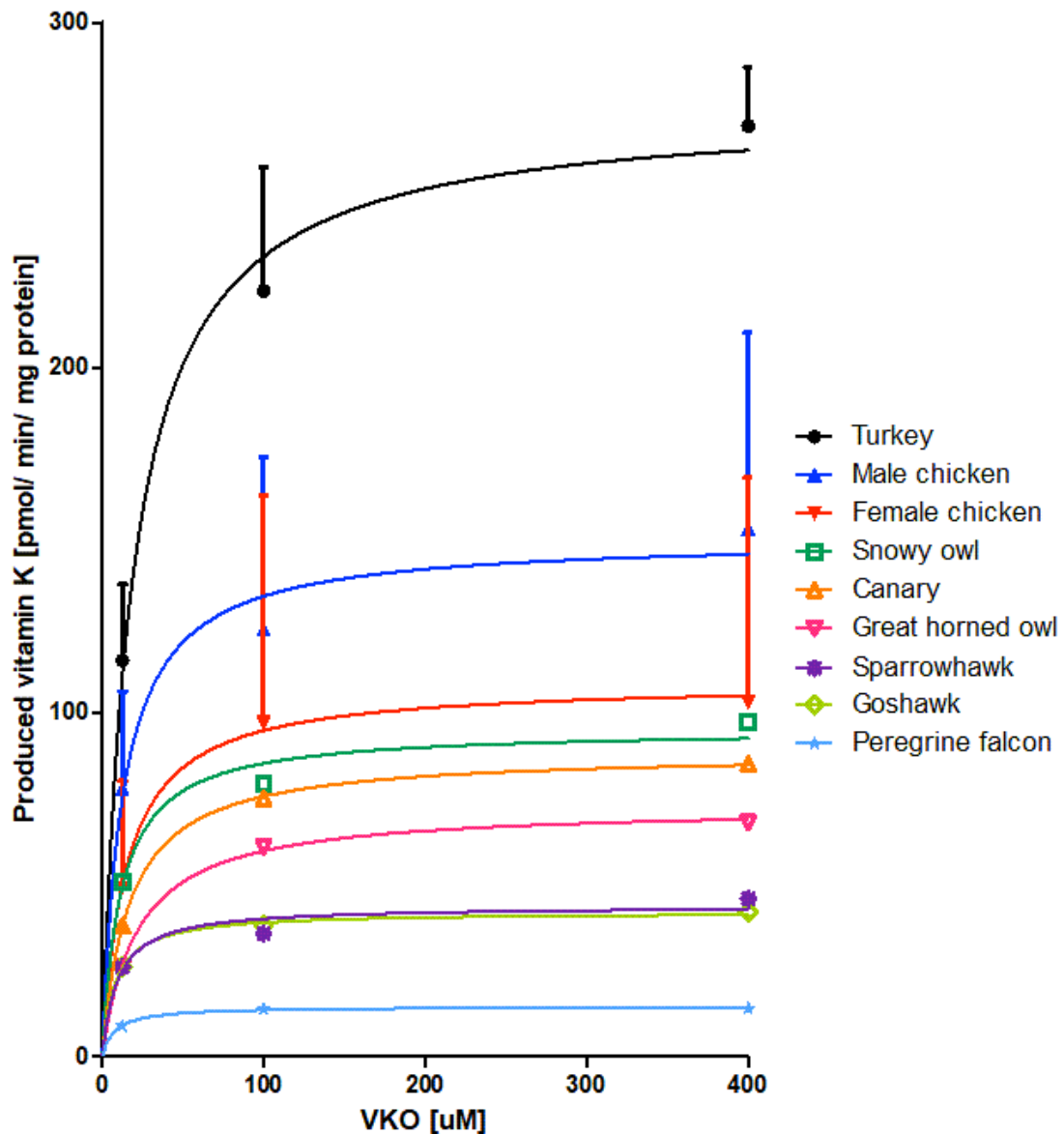


Fig. S2. VKOR activity: vitamin K produced versus 12.5, 100 and 400 μ M of VKO in turkeys, chickens, canaries and five species of raptors.

VKOR activities of *Haliaeetus* (Steller's sea eagle and white-tailed eagle) were undetectable. Each data point of turkeys and chickens represents the mean of three animals $\pm$ S.D. (error bars). The numbers of other species were less than three, therefore there are no S.D. values.

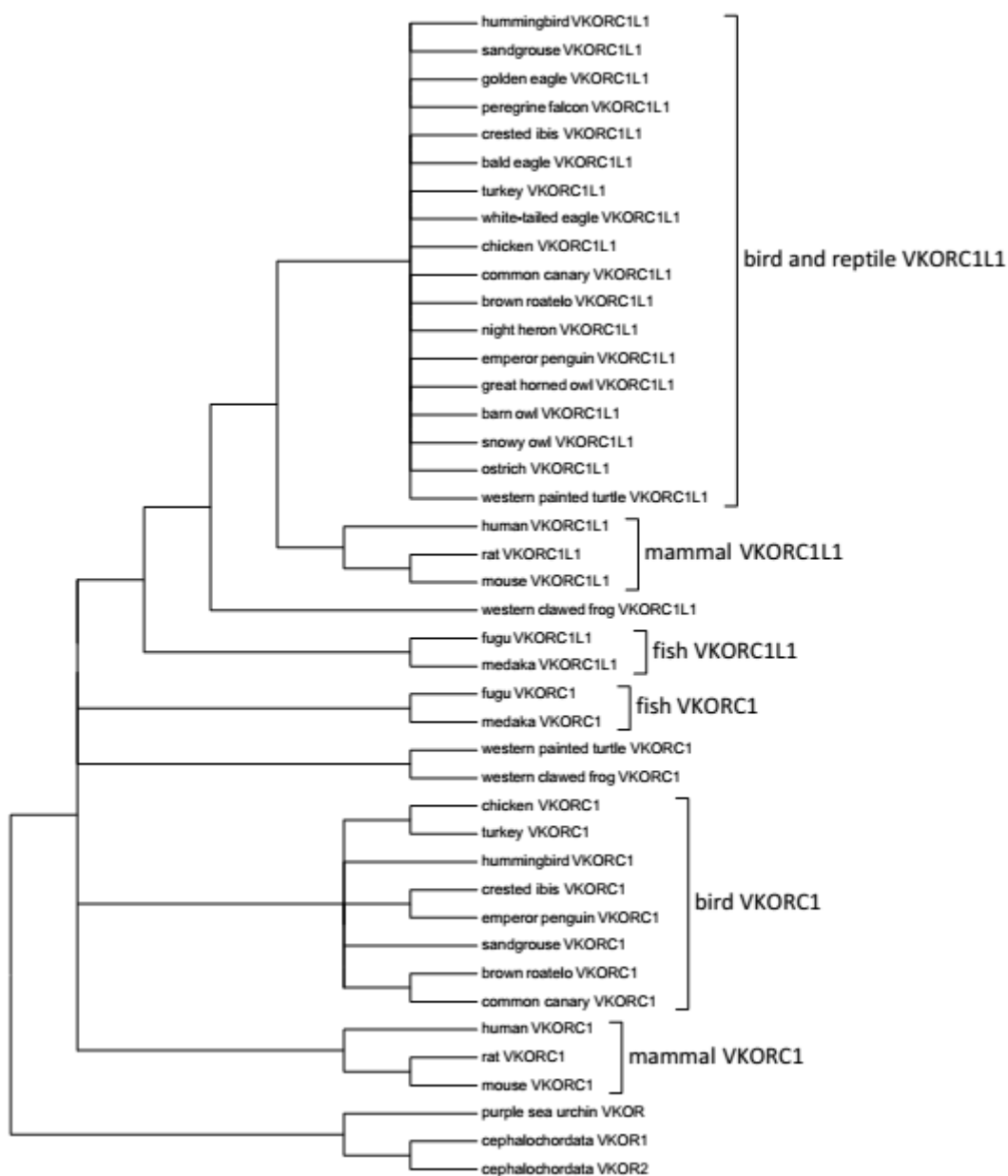


Fig. S3. Phylogenetic tree of the VKOR protein family. In all vertebrates, VKORC1L1 made one cluster. VKORC1 were collapsed because of low bootstrap value (less than 60%). In VKORC1, 4 clusters; fish, bird, mammal, and other animal VKOC1, were made.

692 et al., 2012; Tanaka et al., 2012; Müller et al., 2014).

	1			36					
human_C1L1	MAAPVLLRVS	VPRWERVARY	AVCAAGILLS	IYAYHVEREK	ERDPEHRALC	DLGPWVKCSA	ALASRWGRGF	GLLGSIFGKD	GVLNQPNVSF
rat_C1L1	MAAPVLLRVS	VPRWERVARY	AVCAAGILLS	IYAYHVEREK	ERDPEHRALC	DLGPWVKCSA	ALASRWGRGF	GLLGSIFGKD	GVLNQPNVSF
mouse_C1L	MAAPVLLRVS	VPRWERVARY	AVCAAGILLS	IYAYHVEREK	ERDPEHRALC	DLGPWVKCSA	ALASRWGRGF	GLLGSIFGKD	GVLNQPNVSF
ostrich_C1L1							----RWGRGF	GLLGSIFGKD	SAINQPNVSF
chicken_C1L1	MAAPVLLRVS	VPRWERVARS	AVCAAGILLS	LYACHLEREK	GRDLHYQALC	DLSEVRCSA	AITSRWGRGF	GLLGSIFGKD	SAINQSNVSF
turkey_C1L1	MAAPVLLRVS	VPRWERVARS	AVCAAGILLS	LYACHLERER	GRDLHYQALC	DLSEVRCSA	AITSRWGRGF	GLLGSIFGKD	SAINQSNVSF
hummingbird_C1L1							----RWGRGF	GLLGSIFGKD	SAINQSNVSF
broen_roate1o_C1L1							----WGRGF	GLLGSIFGKD	SAINQSNVSF
sandgrouse_C1L1							----RWGRGF	GLLGSIFGKD	SAINQSNVSF
crested_ibis_C1L1							----RWGRGF	GLLGSIFGKD	SAINQSNVSF
night_heron_C1L1									
emper1r_penguin_C1L1							----RWGRGF	GLLGSIFGKD	SAMNQSNVSF
golden_eagle_C1L1	MAAPVLLRVS	VPRWERVARS	AVCAAGILLS	LYACHLEREK	GRDLHYQALC	DLSEVRCSA	AITSRWGRGF	GLLGSIFGKD	SAINQSNVSF
bald_eagle_C1L1	MAAPVLLRVS	VPRWERVARS	AVCAAGILLS	LYACHLEREK	GRDLHYQALC	DLSEVRCSA	AITSRWGRGF	GLLGSIFGKD	SAINQSNVSF
white_tailed_eagle_C1L1							----WGRGF	GLLGSIFGKD	SAINQSNVSF
barn_owl_C1L1							----RWSRGF	GLLGSIFGKD	SAVNQSNVSF
snowy_owl_C1L1									
great_horned_owl_C1L1									
peregrine_falcon_C1L1							----RWGRGF	GLLGSIFGKD	SAINQSNVSF
canary_C1L1	MAAPVLLRVS	VPRWERVARS	AVCAAGILLS	LYACHLEREK	GRDSHYQALC	DLSEVRCSA	AITSRWGRGF	GLLGSIFGKD	SAINQSNVSF
turtle_C1L1	MAAPVLLRVS	VPRWERVARY	VCAAGILLS	LYACHLEREK	GLDLHYRALC	DISERVCSA	AIASRWGRGF	GLLGSIFGKD	SAINQPNVSF
frog_C1L1	MAAPVM-RVS	VPRWESGARY	AVCVLGIVLS	IYAFHVEREK	ERDPGYKAIC	DFNEWVHCST	VLSSRWGRGF	GMLGSIFGKD	SLLNQPNVSF
fugu_C1L1	MAAPVL-RVS	TPRWERIARV	LVCLLGILLS	LYAFHVEREH	ARDPSYKALC	DVSSSISCSK	VFGSRWGRGF	GLLGSIFGND	SALNQPNSVY

	91				147				
human_C1L1	GLIFYILQLL	LGMTASAVAA	LILMTSSIMS	VVGSLYLAYI	LYFVLKEFCI	ICVITYVLNF	LLLIINYKRL	VYLNEAWKRQ	LQPKQD*
rat_C1L1	GLIFYILQLL	LGMTASAVAA	LVLMTSSIVS	VVGSLYLAYI	LYFVLKEFCI	ICVITYVLNF	LLLIINYKRL	VYLNEAWKRQ	LQPKED*
mouse_C1L	GLIFYILQLL	LGMTASAVAA	LVLMTSSIVS	VVGSLYLAYI	LYFVLKEFCI	ICVITYVLNF	LLLIINYKRL	VYLNEAWKRQ	LQPKED*
ostrich_C1L1	GLVFIYLQML	LGMTASAVAA	LILMTSSIVS	VVGSLYLAYI	LYFVLKEFCI	VCVITYLLNF	ILFIINYKRL	VYLNEAWKRQ	LQPKQE*
chicken_C1L1	GLVFIYLQML	LGMTASAVAA	LILMTSSIVS	VVGSLYLAYI	LYFVLKEFCI	VCVITYLLNF	ILFIINYKRL	VYLNEAWKRQ	LQPKQE*
turkey_C1L1	GLVFIYLQML	LGMTASAVAA	LILMTSSIVS	VVGSLYLAYI	LYFVLKEFCI	VCVITYLLNF	ILFIINYKRL	VYLNEAWKRQ	LQPKQE*
hummingbird_C1L1	GLVFIYLQML	LGMTASAVAA	LILMTSSIVS	VVGSLYLAYI	LYFVLKEFCI	VCVITYLLNF	ILFIINYKRL	VYLNEAWKRQ	LQPKQE*
broen_roate1o_C1L1	GLVFIYLQML	LGMTASAVAA	LILMTSSIVS	VVGSLYLAYI	LYFVLKEFCI	VCVITYLLNF	ILFIVNYKRL	VYLNEAWKRQ	LQPKQE*
sandgrouse_C1L1	GLVFIYLQML	LGMTASAVAA	LILMTSSIVS	VVGSLYLAYI	LYFVLKEFCI	VCVITYLLNF	ILFIINYKRL	VYLNEAWKRQ	LQPKQE*
crested_ibis_C1L1	GLVFIYLQML	LGMTASAVAA	LILMTSSIVS	VVGSLYLAYI	LYFVLKEFCI	VCVITYLLNF	ILFIINYKRL	VYLNEAWKRQ	LQPKQE*
night_heron_C1L1		LGMTASAVAA	LILMTSSIVS	VLGSLYLAYI	LYFVLKEFCI	VCVITYLLNF	ILFIVNYKRL	VYLNEAWKRQ	L-----
emper1r_penguin_C1L1	GLVFIYLQML	LGMTASAVAA	LILMTSSIVS	VVGSLYLAYI	LYFVLKEFCI	VCVITYLLNF	ILFIINYKRL	VYLNEAWKRQ	LQPKQE*
golden_eagle_C1L1	GLVFIYLQML	LGMTASAVAA	LILMTSSIVS	VVGSLYLAYI	LYFVLKEFCI	VCVITYLLNF	ILFIINYKRL	VYLNEAWKRQ	LQPKQE*
bald_eagle_C1L1	GLVFIYLQML	LGMTASAVAA	LILMTSSIVS	VVGSLYLAYI	LYFVLKEFCI	VCVITYLLNF	ILFIINYKRL	VYLNEAWKRQ	LQPKQE*
white_tailed_eagle_C1L1	GLVFIYLQML	LGMTASAVAA	LILMTSSIVS	VVGSLYLAYI	LYFVLKEFCI	VCVITYLLNF	ILFIINYKRL	VYLNEAWKRQ	LQPKQE*
barn_owl_C1L1	GLVFIYLQML	LGMTASAVAA	LILMTSSIVS	VVGSLYLAYI	LYFVLKEFCI	VCVITYLLNF	ILFIINYKRL	VYLNEAWKRQ	LQPKQE*
snowy_owl_C1L1		LGMTASAVAA	LILMTSSIVS	VVGSLYLAYI	LYFVLKEFCI	VCVITYLLNF	ILFIINYKRL	VYLNEAWKQQ	L-----
great_horned_owl_C1L1		LGMTASAVAA	LILMTSSIVS	VVGSLYLAYI	LYFVLKEFCI	VCVITYLLNF	ILFIINYKRL	VYLNEAWKQQ	L-----
peregrine_falcon_C1L1	GLVFIYLQML	LGMTASAVAA	LILMTSSIVS	VVGSLYLAYI	LYFVLKEFCI	VCVITYLLNF	ILFIINYKRL	VYLNEAWKRQ	LQPKQE*
canary_C1L1	GLVFIYLQML	LGMTASAVAA	LILMTSSIVS	VVGSLYLAYI	LYFVLKEFCI	VCVITYLLNF	ILFIINYKRL	VYLNEAWKRQ	LQPKQE*
turtle_C1L1	GLVFIYLQML	LGMTASAVAA	LILMTSSIVS	VVGSLYLAYI	LYFVLKEFCI	ICVITYLLNF	ILFIINYKRL	VYLNEAWKRQ	LQPKQE*
frog_C1L1	GLVFIYLQML	LGMTASAVAA	LVLMTSSIVS	VVGSVLYLAYI	LYFVLKDFCV	ICVITYLLNF	ILLIINYKRL	VYLNEAWKRQ	LQDKQE*
fugu_C1L1	GIVFYAFQLL	LGMTVSAMAA	LILMTSSIMS	VVGSVLYGYI	LYFVLKDL CV	ICVITYLLNF	ILFVLNYKRL	VYLNEAWKQK	LQAKQD*

694 Fig. S4-B. Amino acid sequence alignment of VKORC1L1 of different species.
695 Grey box indicates the catalytic CXXC motif, which is supposed to be the active site of the VKORC1L1 protein. White box indicates the
696 warfarin-binding site. Brown or sky blue shading highlights amino acids whose mutations give warfarin resistance or low VKOR activity,
697 respectively (Hünerberg, 2009).

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