

Supplementary Material

Visual pigments in a palaeognath bird, the emu *Dromaius novaehollandiae*: implications for spectral sensitivity and the origin of ultraviolet vision

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Running title: Emu photoreceptors and visual pigments

Key words: Spectral tuning, opsin evolution, microspectrophotometry, oil droplets, Casuariiformes

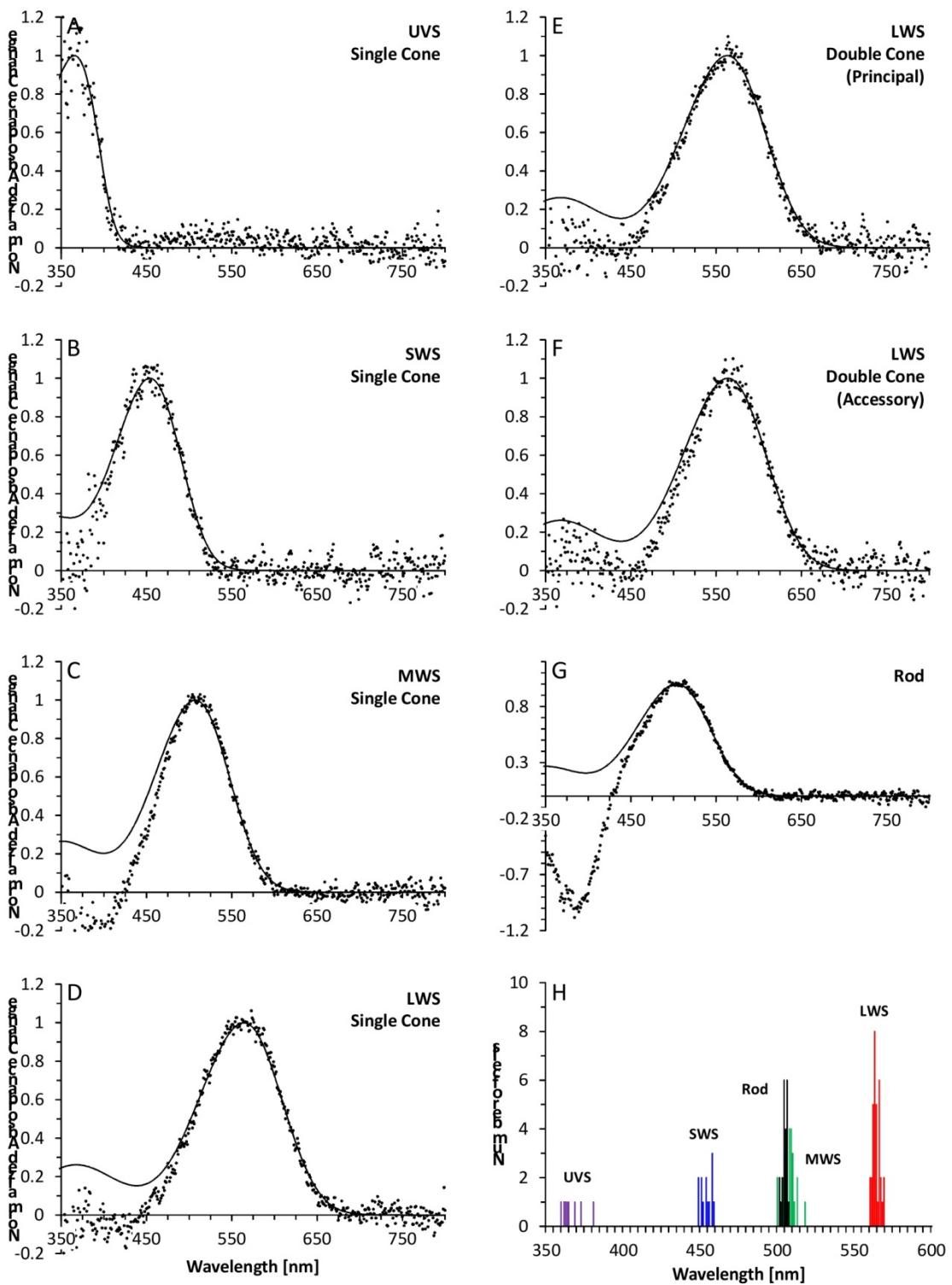


Figure S1. (a-g) Normalised mean bleaching difference spectra (black circles) of visual pigments measured using microspectrophotometry from emu photoreceptor outer segments. Difference spectra are overlayed with best-fit rhodopsin (vitamin A1) templates (black line). (h) Histogram shows the spectral distribution of the wavelength of maximum absorbance change (λ_{\max}) values for individual photoreceptor cell outer segments that were used to generate the mean spectra. The λ_{\max} value distribution of long-wavelength-sensitive (LWS) pigments includes measurements from LWS single cones as well as both the principal and accessory members of double cones. UVS, ultraviolet-sensitive; SWS, short-wavelength-sensitive; MWS, medium-wavelength-sensitive.

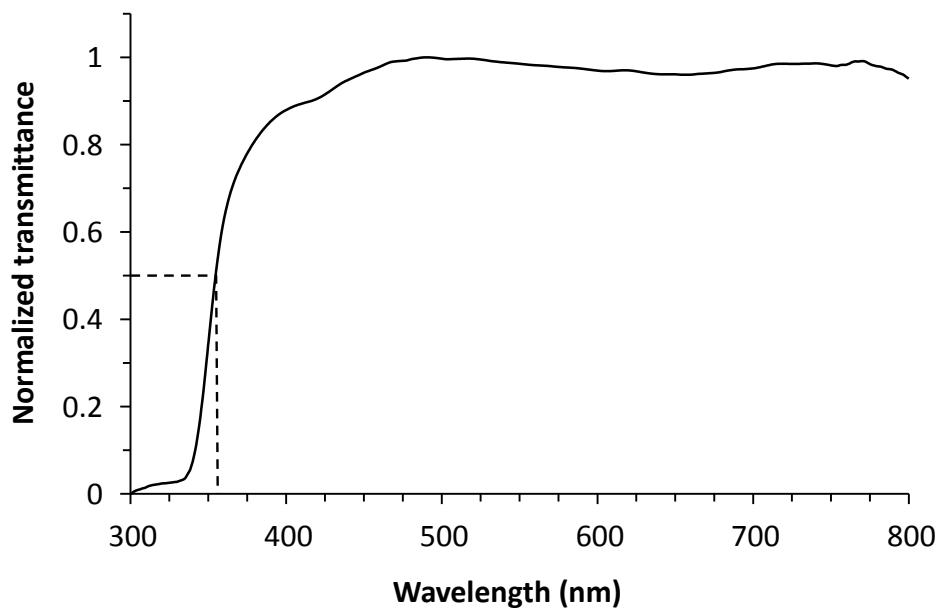


Figure S2. Normalised mean spectral transmittance of the combined ocular media (cornea, aqueous humour and lens) of the emu ($n = 2$ eyes from different birds). The wavelength at 0.5 normalised transmittance ($\lambda_{T0.5}$) is 355 nm and is indicated by the dashed line.

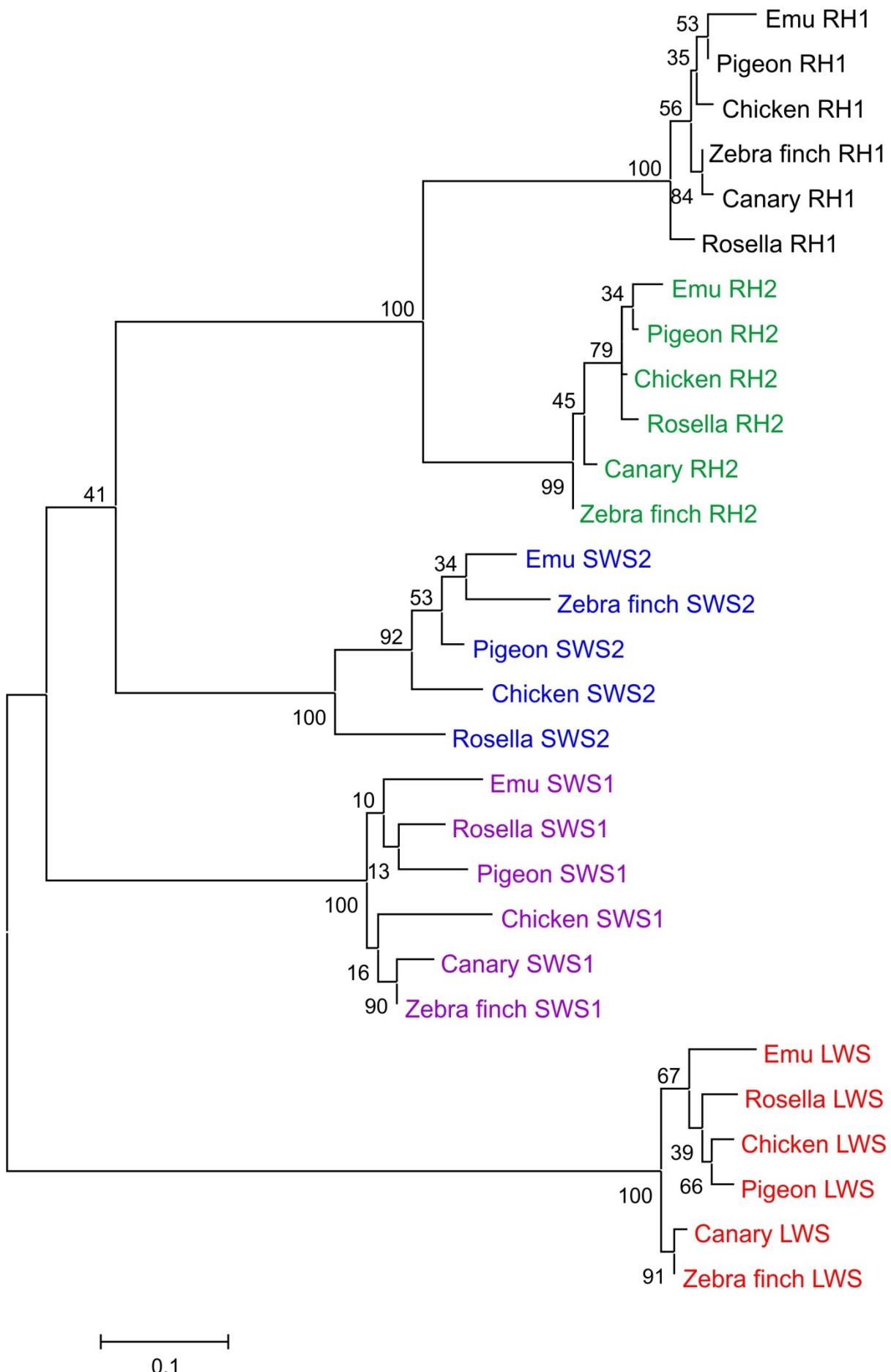


Figure S3. Maximum Likelihood phylogenetic analysis of emu (*Dromaius novaehollandiae*) LWS, SWS1, SWS2, RH2 and RH1 visual opsins compared to orthologues present in other representative birds with *Drosophila* RH1 opsin (GenBank accession no. X65877) as an outgroup (not shown). The degree of support for internal branching is expressed as a percentage with the scale bar indicating the number of amino acid substitutions per site. The

sequences used for generating the tree are as follows: (1) RH1 opsin class: emu (*Dromaius novaehollandiae*), KU568456; pigeon (*Columba livia*), AH007730; chicken (*Gallus gallus*), NM001030606; zebra finch (*Taeniopygia guttata*), NM001076695; canary (*Serinus canaria*), AJ277926; crimson rosella (*Platycercus elegans*), KF134487; (2) RH2 opsin class: emu (*Dromaius novaehollandiae*), KU568455; pigeon (*Columba livia*), AH007731; chicken (*Gallus gallus*), M92038; crimson rosella (*Platycercus elegans*), KF134489; canary (*Serinus canaria*), AJ277924; zebra finch (*Taeniopygia guttata*), NM001076696; (3) SWS2 opsin class: emu (*Dromaius novaehollandiae*), KU568454; zebra finch (*Taeniopygia guttata*), NM001076697; pigeon (*Columba livia*), AH007799; chicken (*Gallus gallus*), NM205517; crimson rosella (*Platycercus elegans*), KF134491; (4) SWS1 opsin class: emu (*Dromaius novaehollandiae*), KU568453; crimson rosella (*Platycercus elegans*), KF134492; pigeon (*Columba livia*), AH007798; chicken (*Gallus gallus*), NM205438; canary (*Serinus canaria*), AJ277922; zebra finch (*Taeniopygia guttata*), NM001076704; and (5) LWS opsin class: emu (*Dromaius novaehollandiae*), KU568452; crimson rosella (*Platycercus elegans*), KF134493; chicken (*Gallus gallus*), NM205440; pigeon (*Columba livia*), AH007800; canary (*Serinus canaria*), AJ277925; zebra finch (*Taeniopygia guttata*), NM001076702.

SWS1 sequences

	SWS1 sequences			SWS2 sequences		
Zebra finch	M-DEEEFYLFKNQSSVGPWDPGPQYHIAPMWAFYLQTIFMGLVFVAGTPLNAAIVLIVTIKY	Zebra finch	MPPKPRMRDDELPEDFYIPMSLETPNTALSPFLVPOHLGSPGI FKAMAAFMFLLIVLIGV			
Canary	M-DEEEFYLFKNQSSVGPWDPGPQYHIAPMWAFYXQJTFMGLVFVAGTPLNAAIVLIVTIKY	Chicken	MHPPRPTTD-LPEDFYIPMADAPNLTLASPLFLVPOTHLGSPLFRAMAAMFELLIALIGV			
Chicken	MGGDEEFYLFKNQSSVGPWDPGPQYHIAPMWAFYLQTAFAMFVFEVNGTFLNAIVLIVTIKY	Pigeon	MQRAREARDELPEDFYIPMADAPNLTLASPLFLVPOTHLGSPLFRAMAAMFELLIALIGV			
Pigeon	MGGDEEFYLFKNQSSVGPWDPGPQYHIAPMWAFYLQTAFAMFVFEVNGTFLNAIVLIVTIKY	Rosella	-----PQTHLGSAGLFAAMAAFMVIALVAIGV			
Emu	-----SSVCPWDPGEQYHIAPMWAFYLQTAFAMFFFVVGTPLNAAIVLIVTIKY	Emu	-----			
Zebra finch	86 KKLRLQPINLYILVNISVSGLMCCVCI 90 TVFIASSCGYVFVKGHIMCAFFEGPAGATGGLVIG	Zebra finch	PINALTVIOTAKYKKLRSHLNYILVNLAANLIVCVGTTIAFYFSQSQYMFALGPACKI			
Canary	KKLRLQPINLYILVNISVSGLMCCVCI 93 TVFVASSCGYVFVKGHMCFFEGPAGATGGVIG	Chicken	PINALTVIOTAKYKKLRSHLNYILVNLAANLIVCVGTTIAFYFSQSQYMFALGPACKI			
Chicken	KKLRLQPINLYILVNISASGPFV SCVCI FVVASCGYVFVKGKRCVELEAVGTHGGLVIG	Pigeon	PINALTVIOTAKYKKLRSHLNYILVNLAANLIVCVGTTIAFYFSQSQYMFALGPACKI			
Pigeon	KKLRLQPINLYILVNISASGPFV SCVCI FVVASCGYVFVKGKRCVELEAVGTHGGLVIG	Rosella	PINALTVIOTAKYKKLRSHLNYILVNLAANLIVCVGTTIAFYFSQSQYMFALGPACKI			
Emu	KKLRLQPINLYILVNISASGPFV SCVCI FVVASCGYVFVKGKRCVELEAVGTHGGLVIG	Emu	PINALTVIOTAKYKKLRSHLNYILVNLAANLIVCVGTTIAFYFSQSQYMFALGPACKI			
Zebra finch	WSLAFLAFERYIVICKPFGNRFRNSRHALLVAAATTWIVGVVAIPPPFGNSRYIPEGLC	Zebra finch	-----			
Canary	WSLAFLAFERYIVICKPFGNRFRNSRHALLVAAATTWIVGVVAIPPPFGNSRYIPEGLC	Chicken	EGFTATLGGMWSIWSLAVVAPERFLVICKPLGNFTFRGSHAVLGCAITWIFGLIASAPPL			
Chicken	WSLAFLAFERYIVICKPFGNRFRNSRHALLVAAATTWIVGVVAIPPPFGNSRYIPEGLC	Pigeon	EGFAATLGGMWSIWSLAVVAPERFLVICKPLGNFTFRGSHAVLGCAITWIFGLIASAPPL			
Pigeon	WSLAFLAFERYIVICKPFGNRFRNSRHALLVAAATTWIVGVVAIPPPFGNSRYIPEGLC	Rosella	EGFMATLGGMWSIWSLAVVAPERFLVICKPLGNFTFRGSHALLGCAITWACGLAAAPPL			
Emu	WSLAFLAFERYIVICKPFGNRFRNSRHALLVAAATTWIVGVVAIPPPFGNSRYIPEGLC	Emu	-----GGMWSIWSLAVVAPERFLVICKPLGNFTFRGSHALLGCAITWACGLAAAPPL			
Zebra finch	SCGPDWYTGTKYKS EYYTWLFIFCPIVPLSLIIFSYSQLISALRAVAQQESATTQK	Zebra finch	FGWSRYIPEGLCSCGPDWYTNNWHEENEYVLFETFCFCGVPLAIIVFSYGRLLLTIR			
Canary	SCGPDWYTGTKYKS EYYTWLFIFCPIVPLSLIIFSYSQLISALRAVAQQESATTQK	Chicken	FGWSRYIPEGLCSCGPDWYTNNWHEENEYVLFETFCFCGVPLAIIVFSYGRLLLTIR			
Chicken	SCGPDWYTGTKYKS EYYTWLFIFCPIVPLSLIIFSYSQLISALRAVAQQESATTQK	Pigeon	FGWSRYIPEGLCSCGPDWYTNNWHEENEYVLFETFCFCGVPLAIIVFSYGRLLLTIR			
Pigeon	SCGPDWYTGTKYKS EYYTWLFIFCPIVPLSLIIFSYSQLISALRAVAQQESATTQK	Rosella	LGWSRYIPEGLCSCGPDWYTNNWHEENEYVLFETFCFCGVPLAIIVFSYGRLLLTIR			
Emu	SCGPDWYTGTKYKS EYYTWLFIFCPIVPLSLIIFSYSQLISALRAVAQQESATTQK	Emu	FGWSRYIPEGLCSCGPDWYTNNWHEENEYVLFETFCFCGVPLAIIVFSYGRLLLTIR			
Zebra finch	AEREVSRMVVVMGSFCMCYVPAALAMYMVNNREHHIDLRLVTI PAFSKSSCVYNPII	Zebra finch	VAKQHQQSATQKAEREVTVKMMVVNVLGFLVCLWPCSFALWWVTHRGRFDLGLASIPS	269		
Canary	AEREVSRMVVVMGSFCMCYVPAALAMYMVNNREHHIDLRLVTI PAFSKSSCVYNPII	Chicken	VARCQHQQSATQKAEREVTVKMMVVNVLGFLVCPWYTAFAWMVVTHRGRFDLGLASIPS			
Chicken	AEREVSRMVVVMGSFCMCYVPAALAMYMVNNREHHIDLRLVTI PAFSKSSCVYNPII	Pigeon	VARCQHQQSATQKAEREVTVKMMVVNVLGFLVCPWYSAFALMWVTHRGRFDLGLASIPS			
Pigeon	AEREVSRMVVVMGSFCMCYVPAALAMYMVNNREHHIDLRLVTI PAFSKSSCVYNPII	Rosella	VARCQHQQSATQKAEREVTVKMMVVNVLGFLVCPWYAAFAWMVVTHRGRFDLGLASIPS			
Emu	A- -----	Emu_SWS2	VAKQHQQSATQKAEREVTVKMMVVNVLGFLVCPWYSAFALMWVTHRGRFDLGLASIPS			
Zebra finch	YCFMNMKQF RAC IMET VGRPM TIDSE VS SSSA QRTE VSSV S S QVGPS-----	Zebra finch	VFSKASTVNP <small>I</small> IYVFMNKQF RSCMLKLVFCGRSPFGDEDVDSGSQSATQVSSV-SSS--			
Canary	YCFMNMKQF RAC IMET VGRPM TIDSE VS SSSA QRTE VSSV S S QVGPS-----	Chicken	VFSKASTVNP <small>I</small> IYVFMNKQF RSCMLKLVFCGRSPFGDEDVDSGSQSATQVSSV-SSS--			
Chicken	YCFMNMKQF RAC IMET VGRPM TIDSE VS SSSA QRTE VSSV S S QVGPS-----	Pigeon	VFSKASTVNP <small>I</small> IYVFMNKQF RSCMLKLVFCGRSPFGDEDVDSGSQSATQVSSV-SSS--			
Pigeon	YCFMNMKQF RAC IMET VGRPM TIDSE VS SSSA QRTE VSSV S S QVGPS-----	Rosella	VFSKASTVNP <small>I</small> IYVFMNKQF RSCMLKLVFCGRSPFGDEDVDSGSQSATQVSSV-SSS--			
Emu	-----	Emu	VFSKASTVNP <small>I</small> IYVFMNKQF RSCMLKLVFCGRSPFGDEDVDSGSQSATQVSSV-SSS--			
RH2 sequences						
Zebra finch	MNGTEGNIYVYVPMNSNKTGVVRSPFEPQYLAEPWKYRVLVCCYIFFLISTGFNFNLTLL	Zebra finch	QVSPA			
Canary	-----PMNSKTGVVRSPFEPQYLAEPWKYRVLVCCYIFFLISTGFNFNLTLL	Chicken	HVAPA			
Chicken	MNGTEGNIYVYVPMNSNKTGVVRSPFEPQYLAEPWKYRVLVCCYIFFLISTGFNFNLTLL	Pigeon	QVAPA			
Pigeon	-----PMNSKTGVVRSPFEPQYLAEPWKYRVLVCCYIFFLISTGFNFNLTLL	Rosella	-----			
Emu	-----	Emu	-----			
Zebra finch	83 VTFKHKKLRLQPINLYILVNLAVID 86 ADLMACF CGT VTV FTYAANGYFVFGPPIGCAVEGFFATLG	Zebra finch	LWS sequences			
Canary	VTFKHKKLRLQPINLYILVNLAVIDADLMACF CGT VTV FTYAANGYFVFGPPIGCAVEGFFATLG	Chicken	Zebra finch			
Chicken	VTFKHKKLRLQPINLYILVNLAVIDADLMACF CGT VTV FTYAANGYFVFGPPIGCAVEGFFATLG	Pigeon	Zebra finch			
Pigeon	VTFKHKKLRLQPINLYILVNLAVIDADLMACF CGT VTV FTYAANGYFVFGPPIGCAVEGFFATLG	Rosella	Zebra finch			
Emu	-----	Emu	Zebra finch			
Zebra finch	GQVALWSIVLVLAIERYIVICKPMGNSRFSAHLMGIAFTWVMAISCAAPPLEGWSRYIP	Zebra finch	IYVVAASVFTNLGVLVATAFKKLRLRHPNWLIVLNLAVALDGETVIASI SVVNQIIFYI			
Canary	GQVALWSIVLVLAIERYIVICKPMGNSRFSAHLMGIAFTWVMAISCAAPPLEGWSRYIP	Canary	IYVVAASVFTNLGVLVATAFKKLRLRHPNWLIVLNLAVALDGETVIASI SVVNQIIFYI			
Chicken	GQVALWSIVLVLAIERYIVICKPMGNSRFSAHLMGIAFTWVMAISCAAPPLEGWSRYIP	Chicken	IYVVAASVFTNLGVLVATAFKKLRLRHPNWLIVLNLAVALDGETVIASI SVVNQIIFYI			
Pigeon	GQVALWSIVLVLAIERYIVICKPMGNSRFSAHLMGIAFTWVMAISCAAPPLEGWSRYIP	Pigeon	IYVVAASVFTNLGVLVATAFKKLRLRHPNWLIVLNLAVALDGETVIASI SVVNQIIFYI			
Emu	GQVALWSIVLVLAIERYIVICKPMGNSRFSAHLMGIAFTWVMAISCAAPPLEGWSRYIP	Rosella	IYVVAASVFTNLGVLVATAFKKLRLRHPNWLIVLNLAVALDGETVIASI SVVNQIIFYI			
Zebra finch	EGMQCS CGF DYT YTHNPD FHNE S YVLYMVF HII FVII IFFS YGRLK VKEVREAAAQQES	Zebra finch	164 LGHPMCVI BEYTVSACGITALWS LAI IS WERWFVWCKPFGNIKFDGKLVAGVLSWWS			
Canary	EGMQCS CGF DYT YTHNPD FHNE S YVLYMVF HII FVII IFFS YGRLK VKEVREAAAQQES	Canary	LGHPMCVI BEYTVSACGITALWS LAI IS WERWFVWCKPFGNIKFDGKLVAGVLSWWS			
Chicken	EGMQCS CGF DYT YTHNPD FHNE S YVLYMVF HII FVII IFFS YGRLK VKEVREAAAQQES	Chicken	LGHPMCVI BEYTVSACGITALWS LAI IS WERWFVWCKPFGNIKFDGKLVAGVLSWWS			
Pigeon	EGMQCS CGF DYT YTHNPD FHNE S YVLYMVF HII FVII IFFS YGRLK VKEVREAAAQQES	Pigeon	LGHPMCVI BEYTVSACGITALWS LAI IS WERWFVWCKPFGNIKFDGKLVAGVLSWWS			
Emu	EGMQCS CGF DYT YTHNPD FHNE S YVLYMVF HII FVII IFFS YGRLK VKEVREAAAQQES	Rosella	LGHPMCVI BEYTVSACGITALWS LAI IS WERWFVWCKPFGNIKFDGKLVAGVLSWWS			
Zebra finch	AT TQKAKEVTRNVLVILMVLGMLAWTPYAVAWIFTNGADFTATLMAVPAFFSKSSSL	Zebra finch	CWATAP PI FGWSRYWPH 261 LKTS CSGPDVFSGSTDGVQSYMMVLMTCFFPLAVI IF CYL			
Canary	AT TQKAKEVTRNVLVILMVLGMLAWTPYAVAWIFTNGADFTATLMAVPAFFSKSSSL	Canary	CWATAP PI FGWSRYWPH 261 LKTS CSGPDVFSGSTDGVQSYMMVLMTCFFPLAVI IF CYL			
Chicken	AT TQKAKEVTRNVLVILMVLGMLAWTPYAVAWIFTNGADFTATLMAVPAFFSKSSSL	Chicken	CWATAP PI FGWSRYWPH 261 LKTS CSGPDVFSGSTDGVQSYMMVLMTCFFPLAVI IF CYL			
Pigeon	AT TQKAKEVTRNVLVILMVLGMLAWTPYAVAWIFTNGADFTATLMAVPAFFSKSSSL	Pigeon	CWATAP PI FGWSRYWPH 261 LKTS CSGPDVFSGSTDGVQSYMMVLMTCFFPLAVI IF CYL			
Emu	AT TQKAKEVTRNVLVILMVLGMLAWTPYAVAWIFTNGADFTATLMAVPAFFSKSSSL	Rosella	CWATAP PI FGWSRYWPH 261 LKTS CSGPDVFSGSTDGVQSYMMVLMTCFFPLAVI IF CYL			
Zebra finch	YNPIIYVIMMNKQPRNCM IT II CGKPNPFGDEETSSTVSQSKEVSSVSSQVSPA	Zebra finch	QWLAIRAAVAQQES ESTOKAKEEVSRMVAIYLCFCWGP <small>262</small> TIFACFAAANF GYAFH			
Canary	YNPIIYVIMMNKQPRNCM IT II CGKPNPFGDEETSSTVSQSKEVSSVSSQVSPA	Canary	QWLAIRAAVAQQES ESTOKAKEEVSRMVAIYLCFCWGP <small>262</small> TIFACFAAANF GYAFH			
Chicken	YNPIIYVIMMNKQPRNCM IT II CGKPNPFGDEETSSTVSQSKEVSSVSSQVSPA	Chicken	QWLAIRAAVAQQES ESTOKAKEEVSRMVAIYLCFCWGP <small>262</small> TIFACFAAANF GYAFH			
Pigeon	YNPIIYVIMMNKQPRNCM IT II CGKPNPFGDEETSSTVSQSKEVSSVSSQVSPA	Pigeon	QWLAIRAAVAQQES ESTOKAKEEVSRMVAIYLCFCWGP <small>262</small> TIFACFAAANF GYAFH			
Emu	YNPIIYVIMMNKQPRNCM IT II CGKPNPFGDEETSSTVSQSKEVSSVSSQVSPA	Rosella	QWLAIRAAVAQQES ESTOKAKEEVSRMVAIYLCFCWGP <small>262</small> TIFACFAAANF GYAFH			
Zebra finch	YNPIIYVIMMNKQPRNCM IT II CGKPNPFGDEETSSTVSQSKEVSSVSSQVSPA	Zebra finch	QWLAIRAAVAQQES ESTOKAKEEVSRMVAIYLCFCWGP <small>262</small> TIFACFAAANF GYAFH			
Canary	YNPIIYVIMMNKQPRNCM IT II CGKPNPFGDEETSSTVSQSKEVSSVSSQVSPA	Canary	QWLAIRAAVAQQES ESTOKAKEEVSRMVAIYLCFCWGP <small>262</small> TIFACFAAANF GYAFH			
Chicken	YNPIIYVIMMNKQPRNCM IT II CGKPNPFGDEETSSTVSQSKEVSSVSSQVSPA	Chicken	QWLAIRAAVAQQES ESTOKAKEEVSRMVAIYLCFCWGP <small>262</small> TIFACFAAANF GYAFH			
Pigeon	YNPIIYVIMMNKQPRNCM IT II CGKPNPFGDEETSSTVSQSKEVSSVSSQVSPA	Pigeon	QWLAIRAAVAQQES ESTOKAKEEVSRMVAIYLCFCWGP <small>262</small> TIFACFAAANF GYAFH			
Emu	YNPIIYVIMMNKQPRNCM IT II CGKPNPFGDEETSSTVSQSKEVSSVSSQVSPA	Rosella	QWLAIRAAVAQQES ESTOKAKEEVSRMVAIYLCFCWGP <small>262</small> TIFACFAAANF GYAFH			
Zebra finch	YNPIIYVIMMNKQPRNCM IT II CGKPNPFGDEETSSTVSQSKEVSSVSSQVSPA	Zebra finch	QWLAIRAAVAQQES ESTOKAKEEVSRMVAIYLCFCWGP <small>262</small> TIFACFAAANF GYAFH			
Canary	YNPIIYVIMMNKQPRNCM IT II CGKPNPFGDEETSSTVSQSKEVSSVSSQVSPA	Canary	QWLAIRAAVAQQES ESTOKAKEEVSRMVAIYLCFCWGP <small>262</small> TIFACFAAANF GYAFH			
Chicken	YNPIIYVIMMNKQPRNCM IT II CGKPNPFGDEETSSTVSQSKEVSSVSSQVSPA	Chicken	QWLAIRAAVAQQES ESTOKAKEEVSRMVAIYLCFCWGP <small>262</small> TIFACFAAANF GYAFH			
Pigeon	YNPIIYVIMMNKQPRNCM IT II CGKPNPFGDEETSSTVSQSKEVSSVSSQVSPA	Pigeon	QWLAIRAAVAQQES ESTOKAKEEVSRMVAIYLCFCWGP <small>262</small> TIFACFAAANF GYAFH			
Emu	YNPIIYVIMMNKQPRNCM IT II CGKPNPFGDEETSSTVSQSKEVSSVSSQVSPA	Rosella	QWLAIRAAVAQQES ESTOKAKEEVSRMVAIYLCFCWGP <small>262</small> TIFACFAAANF GYAFH			
Zebra finch	-----	Zebra finch	SVSPA			
Canary	-----	Canary	SVSPA			
Chicken	-----	Chicken	SVSPA			
Pigeon	-----	Pigeon	SVSPA			
Rosella	-----	Rosella	SVSPA			
Emu	-----	Emu	SVSPA			

Figure S4. Alignment of the amino acid sequences of four cone opsins (SWS1, SWS2, RH2 and LWS) expressed in the emu (*Dromaius novaehollandiae*) compared to orthologues found in other bird species, including zebra finch (*Taeniopygia guttata*), canary (*Serinus canaria*), chicken (*Gallus gallus*), pigeon (*Columba livia*), and the crimson rosella (*Platycercus elegans*). Gaps inserted to maintain a high degree of sequence identity and unsequenced regions are indicated by dashes (-). Spectral tuning sites mentioned in the text are highlighted in red.

Zebra finch	MNGTEGQDFYVPMNSNKTGVVRSPFEYPQYLAEPWKFSALAAYMFMLILLGFPINFLTLY
Canary	-----PFYEYPQYLAEPWKFSALAAYMFMLILLGFPINFLTLY
Chicken	MNGTEGQDFYVPMNSNKTGVVRSPFEYPQYLAEPWKFSALAAYMFMLILLGFPVNFLTLY
Pigeon	MNGTEGQDFYVPMNSNKTGVVRSPFEYPQYLAEPWKFSALAAYMFMLILLGFPVNFLTLY
Rosella	-----TLY
Emu	-----PISNKTTGVVRSPFEYPQYLAEPWKFSALAAYMFMLILLGFPINFLTLY
Bovine	MNGTEGPNFYVPFSNKTGVVRSPFEAPQYLAEPWQFSMLAAYMFLLIMLGFPINFLTLY
 Zebra finch	
Canary	VTIQHKKLRTPLN YILLNLAVADLFMVFGGFTTMYTSMNGYFVFGVTGCYIEGFFATLG
Chicken	VTIQHKKLRTPLN YILLNLAVADLFMVFGGFTTMYTSMNGYFVFGVTGCYIEGFFATLG
Pigeon	VTIQHKKLRTPLN YILLNLVVA DLFMVFGGFTTMYTSMNGYFVFGVTGCYIEGFFATLG
Rosella	VTIQHKKLRTPLN YILLNLAIADLFMVFGGFTTMYTSMNGYFVFGVTGCYIEGFFATLG
Emu	VTIQHKKLRTPLN YILLNLAVADLFMVFGGFTTMYTSMNGYFVFGVTGCYIEGFFATLG
Bovine	VTIQHKKLRTPLN YILLNLAVANLFMVFGGFTTLYTSMHGYFVFGVTGCYIEGFFATLG
 Zebra finch	GEIALWSLVLAIERYVVVCKPMNSNFRGENHAIMGVA FSWIMALACAAPP LFGWSRYIP
Canary	GEIALWSLVLAIERYVVVCKPMNSNFRGENHAIMGVA FSWIMALACAAPP LFGWSRYIP
Chicken	GEIALWSLVLAVERYVVVCKPMNSNFRGENHAIMGVA FSWIMAMACAAPP LFGWSRYIP
Pigeon	GEIALWSLVLAIERYVVVCKPMNSNFRGENHAIMGVA FSWIMALACAAPP LFGWSRYIP
Rosella	GEIALWSLVLAIERYVVVCKPMNSNFRGENHAIMGVA FSWIMALACAAPP LFGWSRYIP
Emu	GEIALWSLVLAIERYVVVCKPMVSNFRGENHAIMGVA FSWIMALACAAPP LFGWSRYIP
Bovine	GEIALWSLVLAIERYVVVCKPMNSNFRGENHAIMGVA FSWIMALACAAPP LFGWSRYIP
 Zebra finch	EGMQCSCGIDYYTLKPEVNNE SFVIYMFVVFHFMIPLSIIFFCYGNLVC TVKEAAAQQQES
Canary	EGMQCSCGIDYYTLKPEVNNE SFVIYMFVVFHFMIPLLIIFFCYGNLVC TVKEAAAQQQES
Chicken	EGMQCSCGIDYYTLKPEINNESFVIYMFVVFHFMIPLA VIIFFCYGNLVC TVKEAAAQQQES
Pigeon	EGMQCSCGIDYYTLKPEINNESFVIYMFVVFHFMIPLMVIIFFCYGNLVC TVKEAAAQQQES
Rosella	EGMQCSCGIDYYTLKPEINNESFVIYMFVVFHFMIPLMIIIFFCYGNLVC TVKEAAAQQQES
Emu	EGMQCSCGIDYYTLKPEVNNE SFVIYMFVVFHTIPLMVIFFCYGNLVC TVKEAAAQQQES
Bovine	EGMQCSCGIDYYTLKPEETNNESFVIYMFVVFHFIPLIVIIFFCYQQLVFTVKEAAAQQQES
 Zebra finch	ATTQKAKEVTRMVIIMVIAFLICWV PYASVAFYIFTNQGSDFGP IFMTIPAFFAKSSAI
Canary	ATTQKAKEVTRMVIIMVISFLICWV PYASVAFYIFTNQGSDFGP IFMTIPAFFAKSSAI
Chicken	ATTQKAKEVTRMVIIMVIAFLICWV PYASVAFYIFTNQGSDFGP IFMTIPAFFAKSSAI
Pigeon	ATTQKAKEVTRMVIIMVIAFLICWL PYASVAFYIFTNQGSDFGP IFMTIPAFFAKSSAI
Rosella	ATTQKAKEVTRMVIIMVIAFLICWL PYASVAFYIFTNQGSDFGP IFMTIPAFFAKSSAI
Emu	ATTQKAKEVTRMVIIMVIAFLICWL PYASVAFYIFTNQGSDFGP IFMTIPAFFAKSSAI
Bovine	ATTQKAKEVTRMVIIMVIAFLICWL PYAGVAFYIFTHQGSDFGP IFMTIPAFFAKTS AV
 Zebra finch	YNPVIYIVMNKQFRNCMITTLCGKNPLGDEDTSAGKTETSSVSTS QVSPA
Canary	YNPVIYIVMNKQFRNCMITTLCGKNPLGDEDTSAGKTETSS-----
Chicken	YNPVIYIVMNKQFRNCMITTLCGKNPLGDEDTSAGKTETSSVSTS QVSPA
Pigeon	YNPVIYIVMNKQFRNCMITTLCGKNPLGDEDTSAGKTETSSVSTS QVSPA
Rosella	FNPLIYVFRDKQFRNCMITTLCGKNPLGDEDTSAGKTETSSVSTS QVSPA
Emu	YNPVIYIVMNKQFRNCMI-----
Bovine	YNPVIYIMMNKOFRNCMVTTLCCGKNPLGDEAST---TVSKTETSOVAPA

Figure S5. Alignment of the amino acid sequences of the emu (*Dromaius novaehollandiae*) rod opsin (RH1) compared to orthologues found in other bird species, including zebra finch (*Taeniopygia guttata*), canary (*Serinus canaria*), chicken (*Gallus gallus*), pigeon (*Columba livia*), and the crimson rosella (*Platycercus elegans*), as well as the common cow (*Bos taurus*). Gaps inserted to maintain a high degree of sequence identity and unsequenced regions are indicated by dashes (–). Spectral tuning sites mentioned in the text are highlighted in red.

Supplementary Tables:**Table S1.** Primers used to isolate and amplify emu opsin gene sequences from retinal cDNA.

Primer	Sequence
LWS_F1	AAGCGTATTYAYTTAYACCRACASCAACAA
LWS_R1	CATCCTBGACACYTCCYTCTVGCCTCTG
LWS_F2	AGTGTCACTCAACCAGWTCTYBGGSTAYTT
LWS_R2	CATCATCCACYYTTSCTCRAASAGCTGCA
SWS1_F1	TCCCATGTCCGGAGAVGAVGABTTYTACCT
SWS1_R1	CACCACSACCATSCGVGASACCTCCCGCTC
SWS1_F2	GGCCTTCGARCGHTACATYGTYATCTGCAA
SWS1_R2	TTAGCTGGGCGYACYTGRCTGGAGGACAC
SWS2_F1	CAACATCACRRCSTSAGCCCBTTCTGGT
SWS2_R1	CAGGAAGCCCADSACCATSACYACYACCAT
SWS2_F2	CTGCAAGATAGAGGGNTTYDCBGMACGCT
SWS2_R2	AAGAATTTABGCBGGGMSACBTGGCTGG
RH2_F1	ATCAACATCCTCACCYTVYTKGTSACCTTC
RH2_R1	CAAGGAGGAATCCMADCACCATSARRATCA
RH2_F2	CTTCTCTGCCACTCAYGCCWTRWTRGGCAT
RH2_R2	CACTGGCTGGAAGARAYRGAVGAKACCTC
RH1_F1	GTCAAAATTCTAYRTBCCCWTKTCCAACA
RH1_R1	ACAGTGCAGACAAGRYKYCCRTAGCAGAAG
RH1_F2	AATAGGATGCWRCWTYGARGGCTTCTTGC
RH1_R2	ATTCTTCCACARCARAGRGTBRTGATCAT

Table S2. Spectral absorption characteristics of retinal photoreceptors in the emu measured using MSP. λ_{\max} , wavelength of maximum absorbance/absorbance change; λ_{cut} , cut-off wavelength; λ_{mid} , wavelength of half maximum absorptance. Values are ± 1 standard deviation. Avian rods do not contain oil droplets and no oil droplets were observed in the accessory members of double cones in the emu (although a diffuse pigmentation was present). T-, C-, Y-, R- and P-type oil droplets are located in the UVS, SWS, MWS and LWS single cones, and the principal members of the LWS double cones, respectively. D, dorsal retina; V, ventral retina; N, number of cells used in the analysis. Subscripts 1 and 2 identify two different spectral types of P-type oil droplet located in the dorsal retina.

Visual Pigments	Single Cones				Double Cones				Rods					
	UVS	SWS	MWS	LWS	Principal		Accessory							
Mean λ_{\max} of prebleach spectra (nm)	367.0 \pm 2.5	453.2 \pm 2.3	502.3 \pm 2.8	562.0 \pm 2.0	562.8 \pm 2.6		563.4 \pm 2.0		501.0 \pm 1.0					
λ_{\max} of mean prebleach spectrum (nm)	366.8	453.1	501.2	562.0	562.9		562.1		500.7					
Mean transverse absorbance at λ_{\max}	0.018 \pm 0.008	0.012 \pm 0.003	0.016 \pm 0.005	0.018 \pm 0.004	0.013 \pm 0.004		0.012 \pm 0.003		0.023 \pm 0.004					
Mean λ_{\max} of difference spectra (nm)	366.6 \pm 6.8	453.7 \pm 3.5	507.2 \pm 4.0	564.2 \pm 1.9	562.8 \pm 2.8		563.5 \pm 2.3		504.0 \pm 1.7					
λ_{\max} of mean difference spectrum (nm)	366.5	454.0	507.0	564.0	563.0		563.7		503.5					
N prebleach (difference)	9 (8)	9 (13)	21 (25)	10 (14)	12 (12)		7 (8)		21 (22)					
Oil Droplets	T-type		C-type		Y-type		R-type		P-type		A-type			
	D	V	D	V	D	V	D	V	D ₁	D ₂	V	D	V	
Mean λ_{cut} of absorptance spectra (nm)	-	-	408.4 \pm 4.3	408.5 \pm 3.5	508.1 \pm 1.9	507.0 \pm 4.0	558.8 \pm 3.8	559.9 \pm 4.2	404.0 \pm 2.0	475.9 \pm 11.8	491.1 \pm 3.8	480.2 \pm 3.9	479.5 \pm 2.0	-
λ_{cut} of mean absorptance spectrum (nm)	-	-	409.3	407.5	507.6	506.8	559.2	560.6	403.5	479.2	492.3	479.2	480.1	-
Mean λ_{mid} of absorptance spectra (nm)	-	-	426.0 \pm 1.9	432.0 \pm 4.9	524.6 \pm 3.4	526.8 \pm 4.2	580.9 \pm 3.8	582.2 \pm 4.2	433.5 \pm 5.9	499.1 \pm 6.2	507.4 \pm 2.3	494.4 \pm 3.5	491.9 \pm 0.6	-
λ_{mid} of mean absorptance spectrum (nm)	-	-	426.2	430.9	524.3	527.3	581.4	582.8	432.9	500.6	508.1	494.6	492.1	-
Mean maximum absorptance	0.04 \pm 0.02	0.05 \pm 0.03	0.54 \pm 0.09	0.47 \pm 0.11	0.84 \pm 0.08	0.84 \pm 0.06	0.85 \pm 0.08	0.83 \pm 0.08	0.46 \pm 0.10	0.66 \pm 0.14	0.58 \pm 0.10	0.21 \pm 0.13	0.20 \pm 0.08	-
Mean diameter (μm)	2.4 \pm 0.2	3.0 \pm 0.4	3.3 \pm 0.8	3.7 \pm 0.6	3.3 \pm 0.6	3.6 \pm 0.4	3.2 \pm 0.3	3.5 \pm 0.5	2.8 \pm 0.3	3.1 \pm 0.3	3.1 \pm 0.4	-	-	-
N	5	4	12	11	18	22	13	19	13	11	20	9	7	-

Table S3. Amino acid differences between ultraviolet-sensitive (UVS) and violet-sensitive (VS) short-wavelength-sensitive-1 (SWS1) visual pigments at known or potential tuning sites, including residues 46, 49, 52, 114 and 118 [1]; 261, 269 and 292 [2]; 86 and 90 [3, 4]; and 93 [5]. Sequence data for avian SWS1 pigments are derived from this study and published articles [6-13], and are compared to residues found in the green anole (*Anolis carolinensis*) SWS1 pigment (GenBank Accession Number AH007736).

Pigment	Tuning sites											
	46	49	52	86	90	93	114	118	164	261	269	292
Avian UVS	Phe Leu Val	Leu Met Val	Thr	Ala Cys	Cys	Thr	Gly	Ala	Gly	Phe	Ala	Ala
Avian VS	Phe Ile	Ala Leu	Thr	Cys Ser	Ser	Thr Val	Ala Gly	Ala Thr	Gly	Phe	Ala	Ala
Anole UVS	Phe	Phe	Thr	Phe	Ser	Thr	Ala	Ser	Gly	Phe	Ala	Ala
Emu UVS	Phe	Phe	Thr	Phe	Cys	Met	Gly	Ser	Gly	Phe	Ala	Ala

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