

Supplementary table S1

Primers used for PCR and qRT-PCR analysis. References and accession numbers.

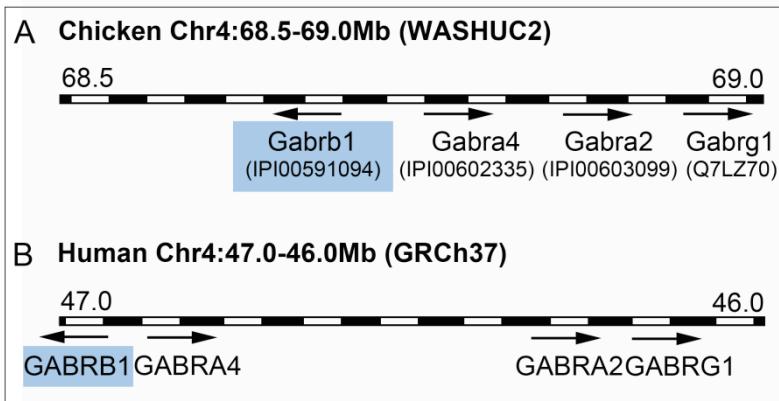
Henrik Ring et al.

Title: Increased A-to-I RNA editing of the transcript for $\alpha 3$ GABA_A receptor subunit during chick retinal development[”]

Subunit (chick)	Forward Primer	Reverse Primer	Accession number (NCBI)	Ensembl Gene	Chicken	Human orthologue
$\alpha 1$	5'-tgcgaggaactcactccctaag-3'	5'-gcacaccgtataaaccatcc-3'	NM_204318.1	ENSGALG00000001698	GBRA1_CHICK ¹⁾	GABRA1 ²³⁾
$\alpha 2$	5'-tcgcaggtagcgcttg-3'	5'-cagacagcgataaaccatcca-3'	XM_001233849.1	ENSGALG00000014206	IPI00603099.3	GABRA2 ²⁾
$\alpha 3$ (qRT PCR)	5'-cgccgtgaaccaaatatgacctg-3'	5'-tgcaccatctgtgtccaaac-3'	XM_420268.2	ENSGALG00000007269	IPI00583603.2	GABRA3 ²⁾
$\alpha 3$ PCR (sanger)	5'-ggtgtcaccactgtgctcacc-3'	5'-gcagtagcagtggcagcagcac-3'				
$\alpha 3$ PCR (pyroseq)	5'-gcaagaaactcgttaccaaagtggcgt-3'	5'-(Bio)-gcttggtaagtagttgacagtggcaaa-3'				
$\alpha 3$ (pyroseq)	5'-cgacggccatggactgggt-3'					
$\alpha 4$	5'-cacacacgagtacagatgcca-3'	5'-tgtcaattcgactcccacca-3'	XM_420724.2	ENSGALG00000014202	IPI00602335.3	GABRA4 ³⁾
$\alpha 5$	5'-acaacccctgtgggtgtggct-3'	5'-agtccccacagttgtcccatc-3'	XM_416880.2	ENSGALG00000016744	IPI00812061.1	GABRA5 ⁴⁾
$\alpha 6$	5'-caccgttggaggcacaagtaa-3'	5'-caaattccactggaaagagga-3'	NM_205058.1	ENSGALG00000001695	GBRA6_CHICK ⁵⁾	GABRA6 ²⁴⁾
$\beta 1$ (foot- note fig 1, 3)	5'-cacatatgcctccacgttgat-3'	5'-attccaagagccactctggct-3'		ENSGALG00000014200	IPI00591094.3	GABRB1 ⁶⁾
$\beta 4$ (foot- note fig 2, 3)	5'-ctggatataacaacgagctgc-3'	5'-gttcgagagtcgccaacact-3'		ENSGALG00000007255	GBRB4_CHICK ⁷⁾	-
$\beta 2$	5'-cttcgtttcatggctctgcta-3'	5'-gcttcttgacgctgaggc-3'	XM_001232377.1	ENSGALG00000001690	Q90590_CHICK ⁸⁾	GABRB2 ²⁵⁾
			XM_414492.2			
$\beta 3$	5'-aaccgaatgattcgccctca-3'	5'-tccatcatacaagctgctgtgg-3'	NM_205346.1	ENSGALG00000016745	GBRB3_CHICK ⁹⁾	GABRB3 ²⁶⁾
$\gamma 1$	5'-ggccctgcaccaaagattcat-3'	5'-tctggccgaagcttattgtcat-3'	XM_420725.2	ENSGALG00000020143	Q7LZ70_CHICK ¹⁰⁾	GABRG1 ¹¹⁾
$\gamma 2$	5'-tgtttctctgactggtaa-3'	5'-cttggcttctgtgtgctgacaa-3'	NM_205345.2	ENSGALG00000001706	GBRG2_CHICK ¹²⁾	GABRG2 ²⁷⁾
$\gamma 3$	5'-cagcaacatgggtgttgatc-3'	5'-cagtgtgcttctgtgtcttg-3'	XM_001233420.1	ENSGALG00000019144	IPI00811977.2	GABRG3 ¹³⁾
ϵ ($\gamma 4$)	5'-atgaagatgacgaccctggct-3'	5'-tgcagcagaagaaccctctaca-3'	NM_205245.1	ENSGALG00000020292	GBRG4_CHICK ¹⁴⁾	GABRE ²⁸⁾
δ	5'-ggattggaggccctctgttaa-3'	5'-gagatgtggtaatgctggta-3'	XM_001234040.1	ENSGALG00000001282	IPI00599008.2	GABRD ¹⁵⁾
π	5'-aagcttcacactggatgctcg-3'	5'-gtcaggaaggacccttgac-3'	XM_414507.2	ENSGALG00000002152	IPI00582090.2	GABRP ¹⁶⁾
NKCC1	5'-ctggcgacaaaagtgacatgaga-3'	5'-cctccctttagggcgatgacaa-3'	XR_027218.1	ENSGALG00000014690	IPI00600831.3	SLC12A2 ¹⁷⁾
KCC2	5'-acggctgtgaagctcaatgag-3'	5'-ttatcccccttccgggtg-3'		ENSGALG00000006930	IPI00571618.3	SLC12A5 ¹⁸⁾
ADAR1	5'-cacctgaacagccgttgaa-3'	5'-agctcccaacacccttgat-3'	XM_001232161.1	ENSGALG00000021475	Q2P9U9_CHICK ¹⁹⁾	ADAR ²⁹⁾
ADAR2	5'-atccctgtgcgtctactacca-3'	5'-acccaatacgttccacctgc-3'	NM_001111074.1	ENSGALG00000007527	ADARB1 ²⁰⁾	ADARB1 ³⁰⁾
β actin	5'-aggtcatcaccattggcaatg-3'	5'-cccaagaaagatggctggaa-3'	NM_205518.1		ACTB ²¹⁾	

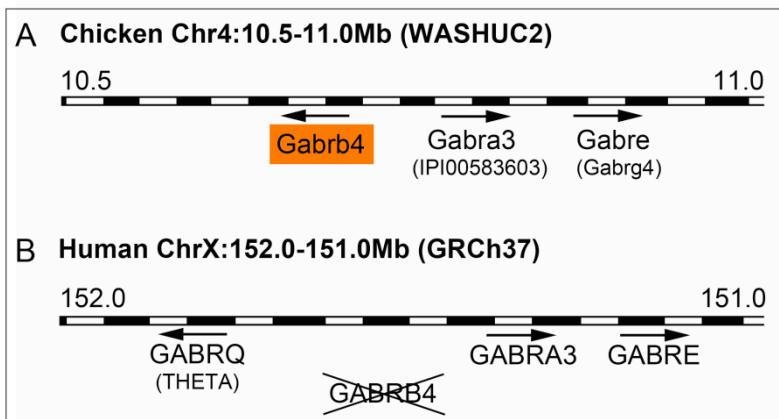
TBP	5'-tagcccgatgtatgcgtat-3'	5'-gttccctgtgtcgcttc-3'	NM_205103.1	ENSGALG00000011171	TBP_CHICK ²²⁾	TBP ³¹⁾
β2 microglobulin	5'-aaggagccgcaggctac-3'	5'-cttgctcttgccgtcatac-3'	NM_001001750.1	ENSGALG00000002160	B2MG_CHICK ³²	B2M ³³
GAPDH	5'-ggaaagcttactggaatggct-3'	5'-ggcaggcaggtaacaaca-3'	NM_204305.1	ENSGALG00000014442	G3P_CHICK ³⁴	GAPDH ³⁵

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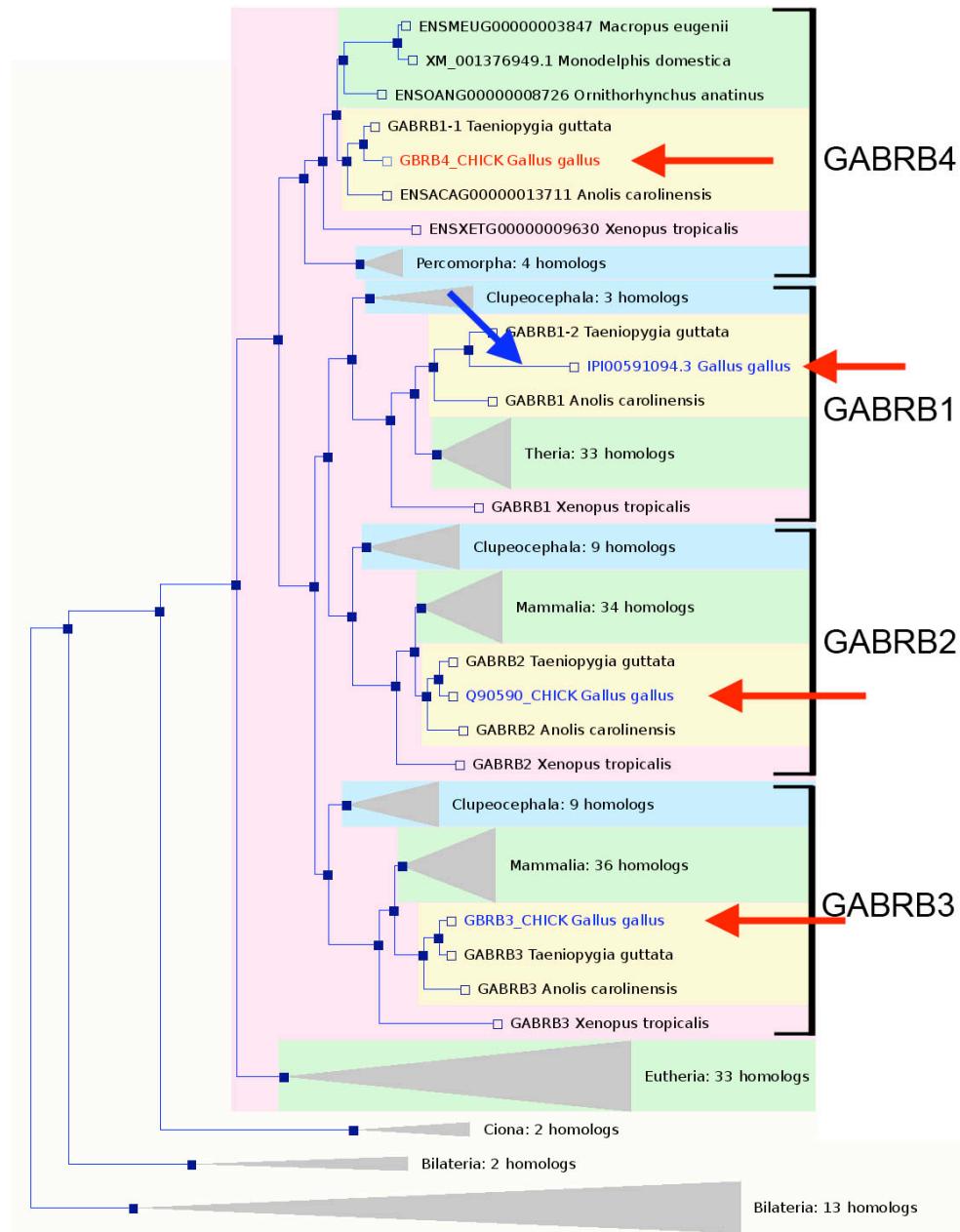


Foot-note figure 1. Conserved synteny in the gene cluster holding GABRB1 in chick and human genomes. Schematic diagram of the localization of GABRB1 in relation to GABRA4, GABRA2 and GABRG1 on A) chick chromosome 4 and B) human chromosome 4. Compiled from Ensembl release 57. Mb; Mega bases

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Foot-note figure 2. Schematic diagram of the conserved synteny in the gene cluster holding several GABR genes on A) chick chromosome 4 with GABRB4, GABRA3 and GABRE. B) The orthologous segment in the human genome lacks the human orthologue of GABRB4 but have GABRA3 and GABRE. Compiled from Ensembl release 57. Mb; Mega bases



Foot-note figure 3. Phylogenetic analysis of GABRB1, B2, B3 and B4 subunit GABA_A receptor sub-family. Maximum likelihood phylogenetic tree based on all available vertebrate GABA_A receptor β subunit sequences showing the existence and relation between chick Gabrb1 and B4. Note the chick sequences indicated by the red arrows. Blue arrow points at the relatively long branch length for chick Gabrb1, indicating a relaxed evolutionary constraint compared to chick B2, B3 and B4. Note also that only 3 mammalian (1 proto-, 2 metatherian) B4 orthologs exist.

Phylogenetic analysis using Gene Tree according to:
Vilella AJ et al. 2008 EnsemblCompara GeneTrees:
Complete, duplication-aware phylogenetic trees in
vertebrates. Genome Res 19:327-35.

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