Supplementary Information

GABA$_A$ receptor-mediated seizure liabilities: A mixed-methods screening approach

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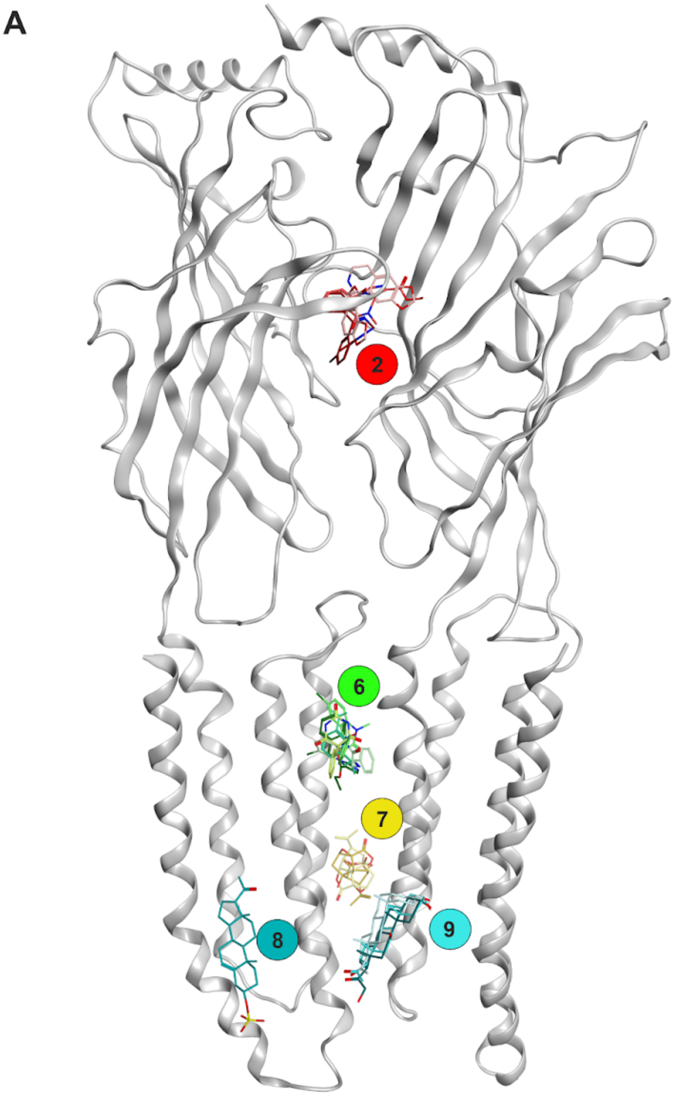
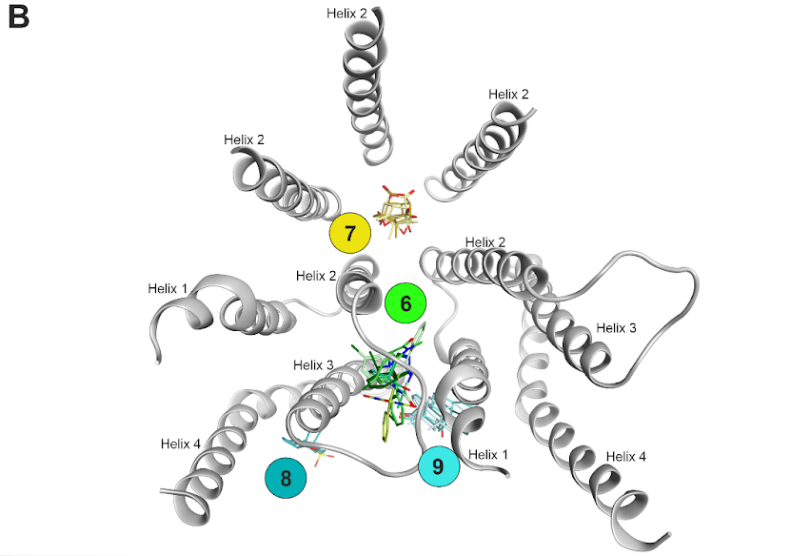
Nr	PDB ID	Protein	Ligand	Subunits	Puthenkalam et al., 2016	References
1	5AFJ	AChBP	fragment 1	α7	1	Spurny et al., 2015
2	5VDH *	Glycine	AM-3607	α3+α3-	-	Huang et al., 2017
	6HJX *	ELIC	2-Ethanesulfonic acid		-	Henault et al., 2019
3	6HUP	GABA-A	GABA	β3+α1-	Site 1	Masiulis et al., 2019
			Diazepam	α1+γ2-		
	6HUO		GABA	β3+α1-		
			Alprazolam	α1+γ2-		
	6HUK *		Bicuculline methochloride	β3+α1-		
	6A96		GABA	β3+α5-		Liu et al., 2018
	6DW0, 6DW1		GABA	β1+α1-		Phulera et al., 2018
	6DW0, 6DW1		GABA	α1+β1-		
	6D6T, 6D6U		GABA	β2+α1-		Zhu et al., 2018
			Flumazenil	α1+γ2-		
	4COF		Benzamidine	β3+β3-		Miller and Aricescu., 2014
	6QFA		Histamine	β3+β3-		Uchanski et al., 2021
	7PC0		alpha-Cobratoxin	β3+α1-		Kasaragod et al., 2022
	7QN9, 7QN8, 7QNC		histamine	β3+β3-		Sente et al., 2022
	7QND, 7QNC		gaboxadol	δ+β3-		Sente et al., 2022
	7QNB		GABA	β3+γ2-		Sente et al., 2022
	7QNE		Ro15-1413	α1+γ2-		Sente et al., 2022
	4TWF, 4TWD	ELIC	memantine, bromomemantine			Ulens et al., 2014
	4F8H *	GLIC	Ketamine		Site 2	Pan et al., 2012a
4	2YN6 *	ELIC	Ba-Atom		Site 3	Zimmermann et al., 2012
5	5LG3 *	ELIC	Chlorpromazine	β1-	Site 5	Nys et al., 2016
	5LID		Bromopromazine			
	3ZKR		Br- atom			Spurny et al., 2013
6	5MUR *	GLIC	Propofol		Site 8	Sauguet et al., 2018
	3P4W *	GLIC	Desflurane		Site 8	Nury et al., 2011
L	6D6T, 6D6U *	GABA-A	Cholesterol	β2+	-	Zhu et al., 2018
7	6HUP *	GABA-A	Diazepam	β3+α1-	Site 7	Masiulis et al., 2019
	5MVM	GLIC	Propofol			Fourati et al., 2018
	4HFE	GLIC	Ethanol			Sauget et al., 2013
	6X3V	GABA-A	Etomidate	β2+α1-		Kim et al., 2020
	6X3T	GABA-A	Propofol			
	6X3W	GABA-A	Phenobarbital			
	6X3X	GABA-A	Diazepam	β2+α1-, γ2+β2-		
	5VDH *	alpha-3 Glycine	Avermectin	α3+α3-		Huang et al., 2017
L	6D6T, 6D6U *	GABA-A	Cholesterol	β2+α1-		Zhu et al., 2018
L	6D6T *	GABA-A	Cholesterol	α1-	-	Zhu et al., 2018
L	6D6T	GABA-A	Cholesterol	α1+β2-	-	Zhu et al., 2018
	5OSA *	GLIC-GABAAR alpha-1 chimera	Cholesterol Hemisuccinate	α1+α1-	-	Laverty et al., 2017

L	6D6T, 6D6U *	GABA-A	Cholesterol	β2+α1-	Site 7	Zhu et al., 2018
L	6D6T *	GABA-A	Cholesterol	α1-		Zhu et al., 2018
L	6D6T	GABA-A	Cholesterol	α1+β2-		Zhu et al., 2018
	5OSA *	GLIC-GABAAR alpha-1 chimera	Cholesterol Hemisuccinate	α1+α1-		Laverty et al., 2017
8	6HUG, 6HUJ, 6X40 *	GABA-A	Picrotoxin	αβαγβ	-	Masiulis et al., 2019, Kim et al., 2020
	5MUO *	GLIC	Propofol		-	Sauget et al., 2018
L	6D6T,6D6U *	GABA-A	Cholesterol *	β2+	Site 10	Zhu et al., 2018
				α1+		
9	5OSC *	GLIC-GABAAR alpha-1 chimera	Pregnenolone Sulfate *	α1+		Laverty et al., 2017
L	3EAM *	GLIC	Lipid *			Bocquet et al., 2008
10	6CDU *	chimeric human alpha-1 GABAA	Alphaxalone *	α1+α1-	overlap with Site 9	Chen et al., 2018
	5O8F *	chimeric beta-3 alpha-5 GABAA	Pregnenolone *	α5β3		Miller et al., 2017
	5OSB *	GLIC-GABAAR alpha-1 chimera	THDOC *	α1+α1-		Laverty et al., 2017
L	6D6T *	GABA-A	Cholesterol *	β2-	Site 10	Zhu et al., 2018
L	6D6T, 6D6U *	GABA-A	Cholesterol *	α1+γ2-	-	Zhu et al., 2018
L	6I53 *	GABA-A	PIP2 *	α1+	overlap with site 10	Laverty et al., 2019
	6HUO, 6HUK, 6HUJ, 6HUG, 6HUP					Masiulis et al., 2019
11	4TWD, 4TWF	ELIC	memantine, bromomemantine		-	Ulens et al., 2014

Supplementary Table S1: Overview of binding sites of $GABA_A$ receptors, including homologous proteins as displayed in Figure 2. Binding sites were compared to Puthenkalam et al. and assigned to the new binding site nomenclature (Bocquet *et al.*, 2009; Nury *et al.*, 2011; Pan *et al.*, 2012; Zimmermann *et al.*, 2012; Sauguet *et al.*, 2013; Spurny *et al.*, 2013; Miller *et al.*, 2014; Ulens *et al.*, 2014; Spurny *et al.*, 2015; Nys *et al.*, 2016; Huang *et al.*, 2017; Laverty *et al.*, 2017; Miller *et al.*, 2017; Chen *et al.*, 2018; Fourati *et al.*, 2018; Liu *et al.*, 2018; Phulera *et al.*, 2018; Zhu *et al.*, 2018; Hénault *et al.*, 2019; Masiulis *et al.*, 2019; Kim *et al.*, 2020; Uchański *et al.*, 2021; Kasaragod *et al.*, 2022; Sente *et al.*, 2022; Zhu *et al.*, 2022). Numbers correspond to Figure 2, “L” stands for “lipid or cholesterol derived ligand”.

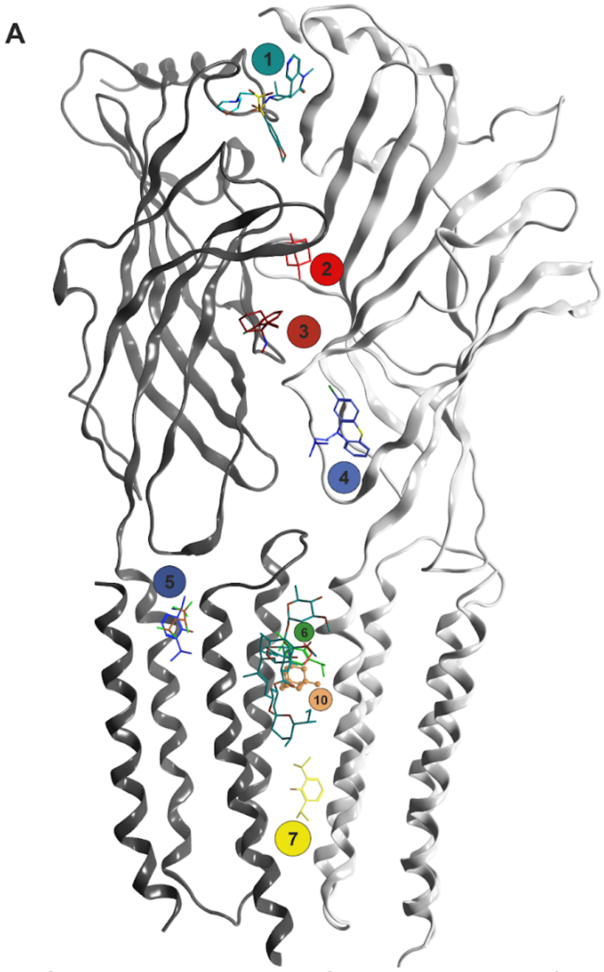


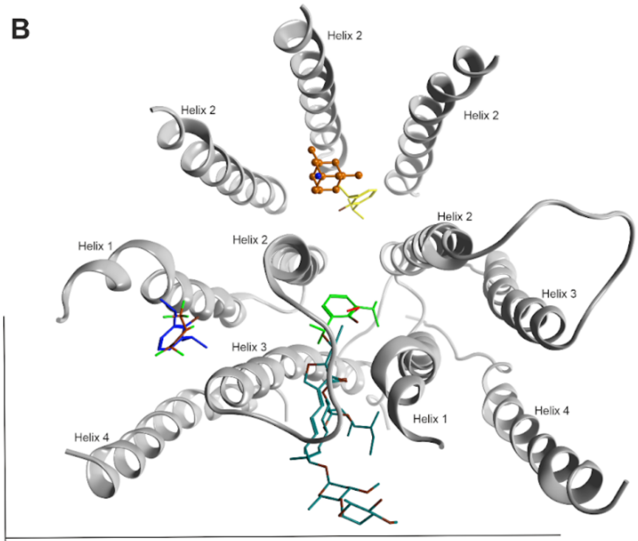


C

	Ligand
2	GABA, Diazepam, Alprazolam, Flumazenil, Bicuculline methochloride, histamine, α-Cobratoxin
6	Diazepam, Propofol, Etomidate, Phenobarbital
7	Picrotoxin
8	Pregnenolone Sulfate, PIP2, lipid
9	Alphaxalone, Pregnenolone, THDOC

Supplementary Figure S1: Binding sites found in GABA$_A$ receptor atomic resolution structures. (a) Side view and (**b**) top view of a superposition of the PDB files shown in (**c**). The ligands are represented in stick representation (GABA in red, diazepam in red for the ECD binding site (2), bicuculline methochloride in pink, flumazenil in red, diazepam in the TMD binding site (6) in light green, etomidate in green, phenobarbital in yellow-green, propofol in dark green, picrotoxin in yellow, alphaxalone in light cyan, THDOC in cyan, pregnanolone in dark cyan, pregnenolone sulfate in cyan). Binding sites are coloured and numbered according to their localization in the protein. Sites 2,6 and 9 are interface-located, whereas sites 8-9 are intrasubunit-located. Site 7 is located within the channel pore. (**c**) Table detailing all binding sites and their respective ligands. The corresponding binding site identification numbers reported in Puthenkalam *et. al.* are also depicted (Puthenkalam *et al.*, 2016).

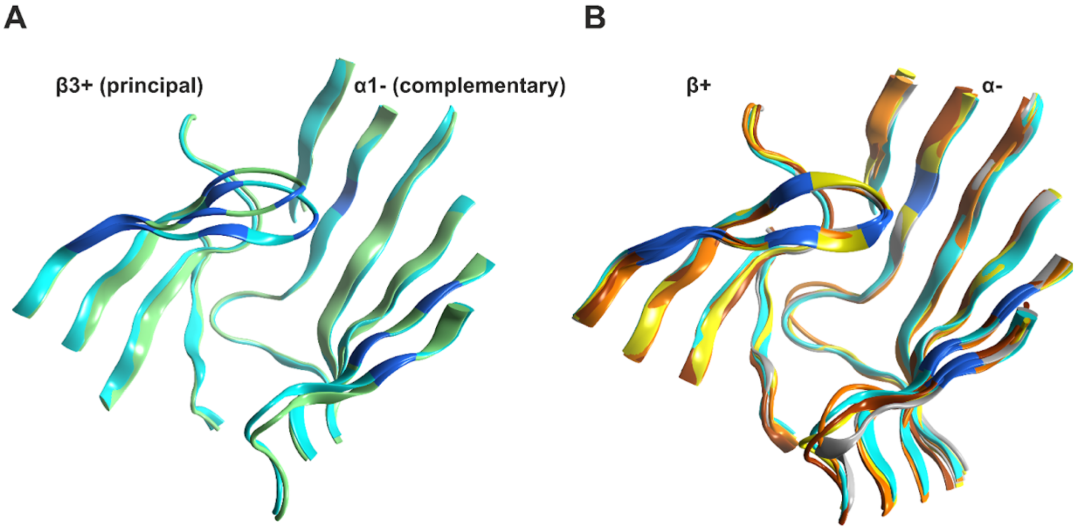




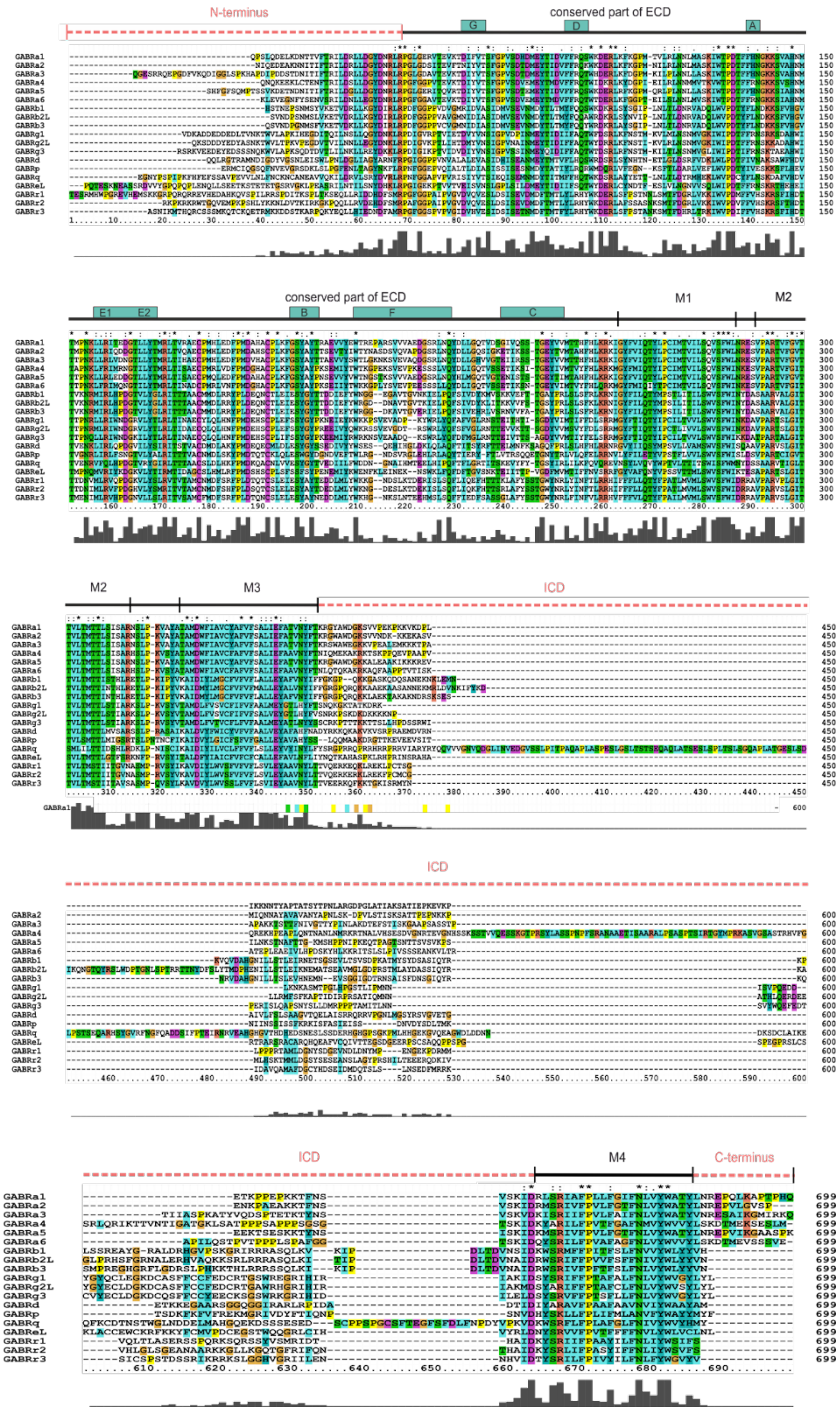
C

	Ligand
1	AM-3607, 2-Ethanesulfonic acid
2	Memantine, Bromomemantine
3	Ketamine
4	Chlorpromazine, Bromopromazine
5	Propofol, Desflurane
6	Avermectin, Ethanol, Propofol
7	Propofol
10	Memantine, Bromomemantine

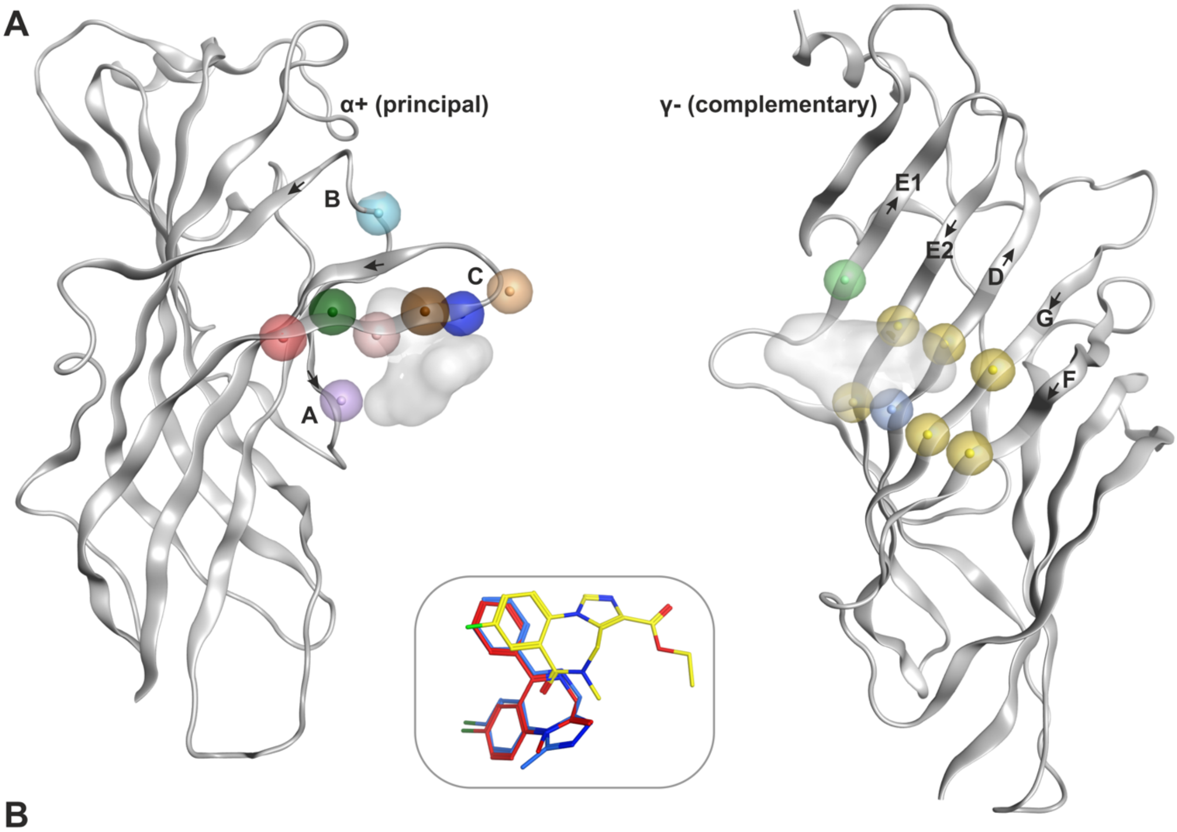
Supplementary Figure S2. Binding sites found in homologous GABA$_A$ receptor atomic resolution structures. (a) Side view and (**b**) top view of a superposition of the protein data bank identities (PDB IDs) shown in (**c**). Two subunits of 6HUP are shown in grey ribbon representation. The ligands are represented in stick representation (AM-3607 in cyan, memantine in the ECD site (2) in red, ketamine in dark red, chlorpromazine in blue, propofol in dark blue, avermectin in green, propofol in the channel blocker site (7), memantine in the channel blocker site (10) in orange). Binding sites are coloured and numbered according to their localization in the protein. Sites 1-3 and 6 are interface-located, whereas sites 4, 5 are intrasubunit-located. Site 7 and 10 is located within the channel pore. (**c**) Table detailing all binding sites and their respective ligands and PDB IDs. The corresponding binding site identification numbers reported in Puthenkalam *et. al.* are also depicted (Puthenkalam *et al.*, 2016). ECD: extracellular domain, TMD: transmembrane domain.



Supplementary Figure S3: GABA binding site (A) Comparison of the bicuculline bound state (6HUK in green) and GABA/picrotoxin bound state (6HUJ in cyan). Highlighted blue residues are showing the different residues in the b1, b2 and b3 subunits as in Figure 3A and 3D. (B) Superposition of b1 (6DW0 in orange), b2 (6D6T in yellow) and b3 (6HUJ in cyan, 7QNA in grey, 6A96 in brown) principal site and a1 (6HUJ, 6DW0, 6D6T), a4 (7QNA) and a5 (6A96) complementary site.



Supplementary Figure S4: Sequence alignment of the 19 human subunits with annotated segments A-G, M1-M4.



B

Loops	A	B	C
α-1	FHN	YAYT	GIVQSSTGEY
α-2	FHN	YAYT	ETIKSSTGEY
α-3	FHN	YAYT	EIIRSSTGEY
α-4	FRN	YAYP	ENIKSITGEY
α-5	FHN	YAYP	ETISTSTGEY
α-6	FRN	YAYP	ETIKSNTGEY

	E1	E2	D	G	F
γ-1	RLLR	YTLR	IFAQ	TDVY	VEV
γ-2	RMLR	YTLR	FFAQ	TDMY	VEV
γ-3	QLLR	YTLR	FFAQ	VDIY	VEA

Supplementary Figure S5: ECD interface high affinity binding site for benzodiazepines: (A) Side view of the benzodiazepine binding site (α+/γ− interface) from a PDBeFold superposition of selected atomic resolution structures (PDB IDs: 6HUP – diazepam, 6HUO – alprazolam, 6D6T/ 6D6U – flumazenil). The subunits are displayed individually to view also the deeper parts of the pocket. The direction of the beta strands is indicated by arrows on the ribbon. Ligands are displayed as shadows on the protein, for visualization purposes diazepam and alprazolam on the principal (+) side and flumazenil on the complementary (-) side. The insert box in the middle depicts the binding modes of diazepam (red), alprazolam (blue) and flumazenil (yellow). The coloured spheres on the ribbon rendering identify the variable positions, color matched with panel (B), on the so-called "loops" (segments) A-G. In loop A, highlighted in purple, are the amino acid differences between α4,6 and α1,2,3,5 subunits that render the former diazepam-insensitive (DI) and the latter diazepam-sensitive (DS). In loop B, highlighted in light blue, is the amino acid difference between α4,5,6 and α1,2,3 subunits that render the former zolpidem-insensitive (ZI) and the latter zolpidem-sensitive (ZS). Several differences are observed in loop C: Amino acid positions which are unique for any isoform are highlighted in different colors: in red and pink the unique amino acids of the α1 subunit; in brown the amino acids which are unique for the α1, α3 and α5 subunits, respectively; in dark blue the unique amino acid of the α5 subunit and in light orange the amino acids unique to α4 and 6 subunits, respectively. On the complementary side of the pocket (γ−) the binding site contributing amino acids are highlighted: in yellow all the positions that are conserved in all three γ isoforms; in cyan the unique amino acid in the γ1 subunit and in light green the unique amino acid in the γ2 subunit. More variable positions are found on segment F (Supplementary Figure S4). (B) provides the partial alignment for the binding site forming segments, more loop F information is in S4.

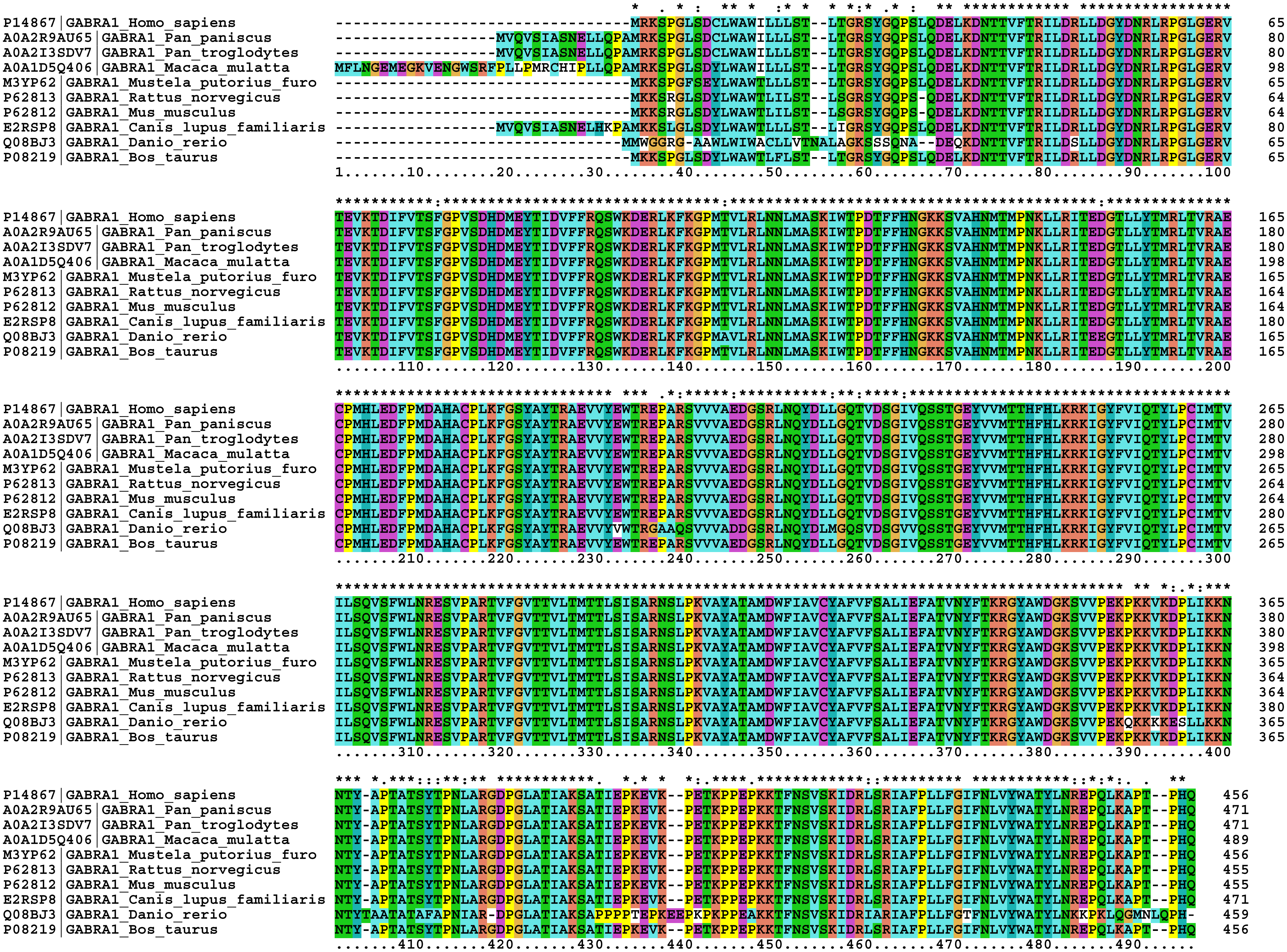
Scientific name of species	Species
Homo sapiens	Human
Rattus norvegicus	Rat
Mus musculus	Mouse
Pan paniscus	Bonobo, historically called : pygmy chimpanzee
Pan troglodytes	Common chimpanzee
Mustela putorius furo	Ferret
Macaca mulatta	Rhesus macaque
Canis lupus familiaris	Domestic dog
Danio rerio	Zebrafish
Bos taurus	Cattle

Supplementary Table S2: Names of species that were used in this study for protein sequence comparisons.

Supplementary Figure S6: Overview of pairwise differences between the human subunits and nine orthologs in the ECD. Substitution scores are plotted for pairwise alignments between human sequences and their orthologs as described in the methods. The colors reflect the different species as indicated in the legend. Full alignments and the correspondences for Danio rerio (which lacks an epsilon subunit) are in Supplementary Figure S8. The upwards spikes correlate with the (normalized) substitution score for amino acid exchanges from the blosum90 matrix, and downwards deflections indicate insertions/deletions (INDELs). The score graphs are aligned along the cys-loop, the second cysteine is represented by the vertical bar. The binding site forming segments G, D, A, E, B, F and C (from N- to C-terminal) are indicated by cyan bars below the graph for each subunit.

Supplementary Figure S7: Overview of pairwise differences between the human subunits and nine orthologs in the TMD and ICD. Substitution scores are plotted for pairwise alignments between human sequences and their orthologs as described in the methods. The colors reflect the different species as indicated in the legend. Full alignments and the correspondences for Danio rerio (which lacks an epsilon subunit) are in Supplementary Figure S8. The upwards spikes correlate with the (normalized) substitution score for amino acid exchanges from the blosum90 matrix, and downwards deflections indicate INDELs. The score graphs are aligned at the beginning of TM1, the four TM segments are indicated by red bars below each score graph.

The next pages provide per subunit alignments from up to 10 species as Supplementary Figure S8



P14867|GABRA1_Homo_sapiens ----------------------------------MRKSPGLSDCLWAWILLLST--LTGRSYGQPSLQDELKDNTTVFTRILDRLLDGYDNRLRPGLGERV 65
A0A2R9AU65|GABRA1_Pan_paniscus ------------------MVQVSIASNELLQPAMRKSPGLSDCLWAWILLLST--LTGRSYGQPSLQDELKDNTTVFTRILDRLLDGYDNRLRPGLGERV 80
A0A2I3SDV7|GABRA1_Pan_troglodytes ------------------MVQVSIASNELLQPAMRKSPGLSDCLWAWILLLST--LTGRSYGQPSLQDELKDNTTVFTRILDRLLDGYDNRLRPGLGERV 80
A0A1D5Q406|GABRA1_Macaca_mulatta MFLNGEMEGKVENGWSRFPLLPMRCHIPLLQPAMRKSPGLSDYLWAWILLLST--LTGRSYGQPSLQDELKDNTTVFTRILDRLLDGYDNRLRPGLGERV 98
M3YP62|GABRA1_Mustela_putorius_furo ----------------------------------MKKSPGFSEYLWAWTLLLST--LTGRSYGQPSLQDELKDNTTVFTRILDRLLDGYDNRLRPGLGERV 65
P62813|GABRA1_Rattus_norvegicus ----------------------------------MKKSRGLSDYLWAWTLILST--LSGRSYGQPS-QDELKDNTTVFTRILDRLLDGYDNRLRPGLGERV 64
P62812|GABRA1_Mus_musculus ----------------------------------MKKSRGLSDYLWAWTLILST--LSGRSYGQPS-QDELKDNTTVFTRILDRLLDGYDNRLRPGLGERV 64
E2RSP8|GABRA1_Canis_lupus_familiaris ------------------MVQVSIASNELHKPAMKKSLGLSDYLWAWTLLLST--LIGRSYGQPSLQDELKDNTTVFTRILDRLLDGYDNRLRPGLGERV 80
Q08BJ3|GABRA1_Danio_rerio ---------------------------------MMWGGRG-AAWLWIWACLLVTNALAGKSSSQNA--DEQKDNTTVFTRILDSLLDGYDNRLRPGLGERV 65
P08219|GABRA1_Bos_taurus ----------------------------------MKKSPGLSDYLWAWTLFLST--LTGRSYGQPSLQDELKDNTTVFTRILDRLLDGYDNRLRPGLGERV 65
1.......10........20........30........40........50........60........70........80........90.......100
P14867|GABRA1_Homo_sapiens TEVKTDIFVTSFGPVSDHDMEYTIDVFFRQSWKDERLKFKGPMTVLRLNNLMASKIWTPDTFFHNGKKSVAHNMTMPNKLLRITEDGTLLYTMRLTVRAE 165
A0A2R9AU65|GABRA1_Pan_paniscus TEVKTDIFVTSFGPVSDHDMEYTIDVFFRQSWKDERLKFKGPMTVLRLNNLMASKIWTPDTFFHNGKKSVAHNMTMPNKLLRITEDGTLLYTMRLTVRAE 180
A0A2I3SDV7|GABRA1_Pan_troglodytes TEVKTDIFVTSFGPVSDHDMEYTIDVFFRQSWKDERLKFKGPMTVLRLNNLMASKIWTPDTFFHNGKKSVAHNMTMPNKLLRITEDGTLLYTMRLTVRAE 180
A0A1D5Q406|GABRA1_Macaca_mulatta TEVKTDIFVTSFGPVSDHDMEYTIDVFFRQSWKDERLKFKGPMTVLRLNNLMASKIWTPDTFFHNGKKSVAHNMTMPNKLLRITEDGTLLYTMRLTVRAE 198
M3YP62|GABRA1_Mustela_putorius_furo TEVKTDIFVTSFGPVSDHDMEYTIDVFFRQSWKDERLKFKGPMTVLRLNNLMASKIWTPDTFFHNGKKSVAHNMTMPNKLLRITEDGTLLYTMRLTVRAE 165
P62813|GABRA1_Rattus_norvegicus TEVKTDIFVTSFGPVSDHDMEYTIDVFFRQSWKDERLKFKGPMTVLRLNNLMASKIWTPDTFFHNGKKSVAHNMTMPNKLLRITEDGTLLYTMRLTVRAE 164
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E2RSP8|GABRA1_Canis_lupus_familiaris TEVKTDIFVTSFGPVSDHDMEYTIDVFFRQSWKDERLKFKGPMTVLRLNNLMASKIWTPDTFFHNGKKSVAHNMTMPNKLLRITEDGTLLYTMRLTVRAE 180
Q08BJ3|GABRA1_Danio_rerio TEVKTDIFVTSIGPVSDHDMEYTIDVFFRQSWKDERLKFKGPMAVLRLNNLMASKIWTPDTFFHNGKKSVAHNMTMPNKLLRITEEGTLLYTMRLTVRAE 165
P08219|GABRA1_Bos_taurus TEVKTDIFVTSFGPVSDHDMEYTIDVFFRQSWKDERLKFKGPMTVLRLNNLMASKIWTPDTFFHNGKKSVAHNMTMPNKLLRITEDGTLLYTMRLTVRAE 165
.......110.......120.......130.......140.......150.......160.......170.......180.......190.......200
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A0A2I3SDV7|GABRA1_Pan_troglodytes CPMHLEDFPMDAHACPLKFGSYAYTRAEVVYEWTREPARSVVVAEDGSRLNQYDLLGQTVDSGIVQSSTGEYVVMTTHFHLKRKIGYFVIQTYLPCIMTV 280
A0A1D5Q406|GABRA1_Macaca_mulatta CPMHLEDFPMDAHACPLKFGSYAYTRAEVVYEWTREPARSVVVAEDGSRLNQYDLLGQTVDSGIVQSSTGEYVVMTTHFHLKRKIGYFVIQTYLPCIMTV 298
M3YP62|GABRA1_Mustela_putorius_furo CPMHLEDFPMDAHACPLKFGSYAYTRAEVVYEWTREPARSVVVAEDGSRLNQYDLLGQTVDSGIVQSSTGEYVVMTTHFHLKRKIGYFVIQTYLPCIMTV 265
P62813|GABRA1_Rattus_norvegicus CPMHLEDFPMDAHACPLKFGSYAYTRAEVVYEWTREPARSVVVAEDGSRLNQYDLLGQTVDSGIVQSSTGEYVVMTTHFHLKRKIGYFVIQTYLPCIMTV 264
P62812|GABRA1_Mus_musculus CPMHLEDFPMDAHACPLKFGSYAYTRAEVVYEWTREPARSVVVAEDGSRLNQYDLLGQTVDSGIVQSSTGEYVVMTTHFHLKRKIGYFVIQTYLPCIMTV 264
E2RSP8|GABRA1_Canis_lupus_familiaris CPMHLEDFPMDAHACPLKFGSYAYTRAEVVYEWTREPARSVVVAEDGSRLNQYDLLGQTVDSGIVQSSTGEYVVMTTHFHLKRKIGYFVIQTYLPCIMTV 280
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P08219|GABRA1_Bos_taurus CPMHLEDFPMDAHACPLKFGSYAYTRAEVVYEWTREPARSVVVAEDGSRLNQYDLLGQTVDSGIVQSSTGEYVVMTTHFHLKRKIGYFVIQTYLPCIMTV 265
.......210.......220.......230.......240.......250.......260.......270.......280.......290.......300
P14867|GABRA1_Homo_sapiens ILSQVSFWLNRESVPARTVFGVTTVLTMTTLSISARNSLPKVAYATAMDWFIAVCYAFVFSALIEFATVNYFTKRGYAWDGKSVVPEKPKKVKDPLIKKN 365
A0A2R9AU65|GABRA1_Pan_paniscus ILSQVSFWLNRESVPARTVFGVTTVLTMTTLSISARNSLPKVAYATAMDWFIAVCYAFVFSALIEFATVNYFTKRGYAWDGKSVVPEKPKKVKDPLIKKN 380
A0A2I3SDV7|GABRA1_Pan_troglodytes ILSQVSFWLNRESVPARTVFGVTTVLTMTTLSISARNSLPKVAYATAMDWFIAVCYAFVFSALIEFATVNYFTKRGYAWDGKSVVPEKPKKVKDPLIKKN 380
A0A1D5Q406|GABRA1_Macaca_mulatta ILSQVSFWLNRESVPARTVFGVTTVLTMTTLSISARNSLPKVAYATAMDWFIAVCYAFVFSALIEFATVNYFTKRGYAWDGKSVVPEKPKKVKDPLIKKN 398
M3YP62|GABRA1_Mustela_putorius_furo ILSQVSFWLNRESVPARTVFGVTTVLTMTTLSISARNSLPKVAYATAMDWFIAVCYAFVFSALIEFATVNYFTKRGYAWDGKSVVPEKPKKVKDPLIKKN 365
P62813|GABRA1_Rattus_norvegicus ILSQVSFWLNRESVPARTVFGVTTVLTMTTLSISARNSLPKVAYATAMDWFIAVCYAFVFSALIEFATVNYFTKRGYAWDGKSVVPEKPKKVKDPLIKKN 364
P62812|GABRA1_Mus_musculus ILSQVSFWLNRESVPARTVFGVTTVLTMTTLSISARNSLPKVAYATAMDWFIAVCYAFVFSALIEFATVNYFTKRGYAWDGKSVVPEKPKKVKDPLIKKN 364
E2RSP8|GABRA1_Canis_lupus_familiaris ILSQVSFWLNRESVPARTVFGVTTVLTMTTLSISARNSLPKVAYATAMDWFIAVCYAFVFSALIEFATVNYFTKRGYAWDGKSVVPEKPKKVKDPLIKKN 380
Q08BJ3|GABRA1_Danio_rerio ILSQVSFWLNRESVPARTVFGVTTVLTMTTLSISARNSLPKVAYATAMDWFIAVCYAFVFSALIEFATVNYFTKRGYAWDGKSVVPEKQKKVKKESLLKKN 365
P08219|GABRA1_Bos_taurus ILSQVSFWLNRESVPARTVFGVTTVLTMTTLSISARNSLPKVAYATAMDWFIAVCYAFVFSALIEFATVNYFTKRGYAWDGKSVVPEKPKKVKDPLIKKN 365
.......310.......320.......330.......340.......350.......360.......370.......380.......390.......400
P14867|GABRA1_Homo_sapiens NTY-APTATSYTPNLARGDPGLATIAKSATIEPKEVK--PETKPPEPKKTFNSVSKIDRLSRIAFPLLFGIFNLVYWATYLNREPQLKAPT--PHQ 456
A0A2R9AU65|GABRA1_Pan_paniscus NTY-APTATSYTPNLARGDPGLATIAKSATIEPKEVK--PETKPPEPKKTFNSVSKIDRLSRIAFPLLFGIFNLVYWATYLNREPQLKAPT--PHQ 471
A0A2I3SDV7|GABRA1_Pan_troglodytes NTY-APTATSYTPNLARGDPGLATIAKSATIEPKEVK--PETKPPEPKKTFNSVSKIDRLSRIAFPLLFGIFNLVYWATYLNREPQLKAPT--PHQ 471
A0A1D5Q406|GABRA1_Macaca_mulatta NTY-APTATSYTPNLARGDPGLATIAKSATIEPKEVK--PETKPPEPKKTFNSVSKIDRLSRIAFPLLFGIFNLVYWATYLNREPQLKAPT--PHQ 489
M3YP62|GABRA1_Mustela_putorius_furo NTY-APTATSYTPNLARGDPGLATIAKSATIEPKEVK--PETKPPEPKKTFNSVSKIDRLSRIAFPLLFGIFNLVYWATYLNREPQLKAPT--PHQ 456
P62813|GABRA1_Rattus_norvegicus NTY-APTATSYTPNLARGDPGLATIAKSATIEPKEVK--PETKPPEPKKTFNSVSKIDRLSRIAFPLLFGIFNLVYWATYLNREPQLKAPT--PHQ 455
P62812|GABRA1_Mus_musculus NTY-APTATSYTPNLARGDPGLATIAKSATIEPKEVK--PETKPPEPKKTFNSVSKIDRLSRIAFPLLFGIFNLVYWATYLNREPQLKAPT--PHQ 455
E2RSP8|GABRA1_Canis_lupus_familiaris NTY-APTATSYTPNLARGDPGLATIAKSATIEPKEVK--PETKPPEPKKTFNSVSKIDRLSRIAFPLLFGIFNLVYWATYLNREPQLKAPT--PHQ 471
Q08BJ3|GABRA1_Danio_rerio NTYTAATATAFAPNIAR-DPGLATIAKSAPPPTEPKEEPKPKPPEAKKTFNSVSKIDRIARIAFPLLFGTFNLVYWATYLNKKPKLQGMNLQPH- 459
P08219|GABRA1_Bos_taurus NTY-APTATSYTPNLARGDPGLATIAKSATIEPKEVK--PETKPPEPKKTFNSVSKIDRLSRIAFPLLFGIFNLVYWATYLNREPQLKAPT--PHQ 456
.......410.......420.......430.......440.......450.......460.......470.......480.......490......

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                                                               .   *::.: *   .        .*  *****:********************* :*** *:*******
P47869|GABRA2_Homo_sapiens                     ----------------------MKTKLNIYNMQFLLFVFLVWDPARLVLANIQEDEAKNNITIFTRILDRLLDGYDNRLRPGLGDSITEVFTNIYVTSFG   78
XP_003816076.1|GABRA2_Pan_paniscus             ----------------------MKTKLNIYNMQFLLFVFLVWDPARLVLANIQEDEAKNNITIFTRILDRLLDGYDNRLRPGLGDSITEVFTNIYVTSFG   78
M3Y0V9|GABRA2_Mustela_putorius_furo            MVQLNVILSVFYFSCMALQRAAMKTKLNTYNVQLLLVFLVWDPARLVLANIQEDEAKNNITIFTRILDRLLDGYDNRLRPGLGDSITEVFTNIYVTSFG  100
P23576|GABRA2_Rattus_norvegicus                ----------------------MRTKLSTCNVWFPLLVLLVWNPARLVLANIQEDEAKNNITIFTRILDRLLDGYDNRLRPGLGDSITEVFTNIYVTSFG   78
P26048|GABRA2_Mus_musculus                     ----------------------MKTKLSTCNVWSLLLVLLVWDPVRLVLANIQEDEAKNNITIFTRILDRLLDGYDNRLRPGLGDSITEVFTNIYVTSFG   78
A0A5F4CYJ7|GABRA2_Canis_lupus_familiaris       -------------------------------MRLLLLVVLLWDPARLVLANIQEDEAKNNITIFTRILDRLLDGYDNRLRPGLGDSITEVFTNIYVTSFG   69
H2QPE5|GABRA2_Pan_troglodytes                  ----------------------MKTKLNIYNMQFLLFVFLVWDPARLVLANIQEDEAKNNITIFTRILDRLLDGYDNRLRPGLGDSITEVFTNIYVTSFG   78
E7F635|GABRA2a_Danio_rerio                     ---------------------MILRRGHSHGFLIHLFLLVLW-RACSEAEVSGSDAFKNNITLFTRILDRLLDGYDNRLRPGLGDRVTEVKTDIYVTSFG   78
XP_028703913.1|GABRA2_Macaca_mulatta           ----------------------MKTKLNIYNMQLLLFVFLVWDPARLVLANIQEDEAKNNITIFTRILDRLLDGYDNRLRPGLGDSITEVFTNIYVTSFG   78
P10063|GABRA2_Bos_taurus                       ----------------------MKTKLNSSNMQLLLFVFLAWDPARLVLANIQEDEAKNNITIFTRILDRLLDGYDNRLRPGLGDSITEVFTNIYVTSFG   78
                                               1.......10........20........30........40........50........60........70........80........90......100

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P47869|GABRA2_Homo_sapiens                     PVSDTDMEYTIDVFFRQKWKDERLKFKGPMNILRLNNLMASKIWTPDTFFHNGKKSVAHNMTMPNKLLRIQDDGTLLYTMRLTVQAECPMHLEDFPMDAH  178
XP_003816076.1|GABRA2_Pan_paniscus             PVSDTDMEYTIDVFFRQKWKDERLKFKGPMNILRLNNLMASKIWTPDTFFHNGKKSVAHNMTMPNKLLRIQDDGTLLYTMRLTVQAECPMHLEDFPMDAH  178
M3Y0V9|GABRA2_Mustela_putorius_furo            PVSDTDMEYTIDVFFRQKWKDERLKFKGPMNILRLNNLMASKIWTPDTFFHNGKKSVAHNMTMPNKLLRIQDDGTLLYTMRLTVQAECPMHLEDFPMDAH  200
P23576|GABRA2_Rattus_norvegicus                PVSDTDMEYTIDVFFRQKWKDERLKFKGPMNILRLNNSMASKIWTPDTFFHNGKKSVAHNMTMPNKLLRIQDDGTLLYTMRLTVQAECPMHLEDFPMDAH  178
P26048|GABRA2_Mus_musculus                     PVSDTDMEYTIDVFFRQKWKDERLKFKGPMNILRLNNLMASKIWTPDTFFHNGKKSVAHNMTMPNKLLRIQDDGTLLYTMRLTVQAECPMHLEDFPMDAH  178
A0A5F4CYJ7|GABRA2_Canis_lupus_familiaris       PVSDTDMEYTIDVFFRQKWKDERLKFKGPMNILRLNNLMASKIWTPDTFFHNGKKSVAHNMTMPNKLLRIQDDGTLLYTMRLTVQAECPMHLEDFPMDAH  169
H2QPE5|GABRA2_Pan_troglodytes                  PVSDTDMEYTIDVFFRQKWKDERLKFKGPMNILRLNNLMASKIWTPDTFFHNGKKSVAHNMTMPNKLLRIQDDGTLLYTMRLTVQAECPMHLEDFPMDAH  178
E7F635|GABRA2a_Danio_rerio                     PVSDTDMEYTVDVFFRQRWTDERLKFHGPMNILRLNNLMASKIWTPDTFFHNGKKSVAHNMTMPNKLLRIMENGTLLYTMRLTVQAECPMHLEDFPMDFH  178
XP_028703913.1|GABRA2_Macaca_mulatta           PVSDTDMEYTIDVFFRQKWKDERLKFKGPMNILRLNNLMASKIWTPDTFFHNGKKSVAHNMTMPNKLLRIQDDGTLLYTMRLTVQAECPMHLEDFPMDAH  178
P10063|GABRA2_Bos_taurus                       PVSDTDMEYTIDVFFRQKWKDERLKFKGPMNILRLNNLMASKIWTPDTFFHNGKKSVAHNMTMPNKLLRIQDDGTLLYTMRLTVQAECPMHLEDFPMDAH  178
                                               .......110.......120.......130.......140.......150.......160.......170.......180.......190.......200

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P47869|GABRA2_Homo_sapiens                     SCPLKFGSYAYTTSEVTYIWTYNASDSVQVAPDGSRLNQYDLLGQSIGKETIKSSTGEYTVMTAHFHLKRKIGYFVIQTYLPCIMTVILSQVSFWLNRES  278
XP_003816076.1|GABRA2_Pan_paniscus             SCPLKFGSYAYTTSEVTYIWTYNASDSVQVAPDGSRLNQYDLLGQSIGKETIKSSTGEYTVMTAHFHLKRKIGYFVIQTYLPCIMTVILSQVSFWLNRES  278
M3Y0V9|GABRA2_Mustela_putorius_furo            SCPLKFGSYAYTTSEVTYIWTYNASDSVQVAPDGSRLNQYDLLGQSIGKETIKSSTGEYTVMTAHFHLKRKIGYFVIQTYLPCIMTVILSQVSFWLNRES  300
P23576|GABRA2_Rattus_norvegicus                SCPLKFGSYAYTTSEVTYIWTYNPSDSVQVAPDGSRLNQYDLLGQSIGKETIKSSTGEYTVMTAHFHLKRKIGYFVIQTYLPCIMTVILSQVSFWLNRES  278
P26048|GABRA2_Mus_musculus                     SCPLKFGSYAYTTSEVTYIWTYNASDSVQVAPDGSRLNQYDLLGQSIGKETIKSSTGEYTVMTAHFHLKRKIGYFVIQTYLPCIMTVILSQVSFWLNRES  278
A0A5F4CYJ7|GABRA2_Canis_lupus_familiaris       SCPLKFGSYAYTTSEVTYIWTYNASDSVQVAPDGSRLNQYDLLGQSIGKETIKSSTGEYTVMTAHFHLKRKIGYFVIQTYLPCIMTVILSQVSFWLNRES  269
H2QPE5|GABRA2_Pan_troglodytes                  SCPLKFGSYAYTTSEVTYIWTYNASDSVQVAPDGSRLNQYDLLGQSIGKETIKSSTGEYTVMTAHFHLKRKIGYFVIQTYLPCIMTVILSQVSFWLNRES  278
E7F635|GABRA2a_Danio_rerio                     SCPLKFGSYAYTLTEVTYVWTRNATLSVEVAPDGSRLNQYDLMGQTVGKETIKSSTGEYTVMTAHFHLKRKIGYFVIQTYLPCIMTVILSQVSFWLNRES  278
XP_028703913.1|GABRA2_Macaca_mulatta           SCPLKFGSYAYTTSEVTYIWTYNASDSVQVAPDGSRLNQYDLLGQSIGKETIKSSTGEYTVMTAHFHLKRKIGYFVIQTYLPCIMTVILSQVSFWLNRES  278
P10063|GABRA2_Bos_taurus                       SCPLKFGSYAYTTSEVTYIWTYNASDSVQVAPDGSRLNQYDLPGQSIGKETIKSSTGEYTVMTAHFHLKRKIGYFVIQTYLPCIMTVILSQVSFWLNRES  278
                                               .......210.......220.......230.......240.......250.......260.......270.......280.......290.......300

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P47869|GABRA2_Homo_sapiens                     VPARTVFGVTTVLTMTTLSISARNSLPKVAYATAMDWFIAVCYAFVFSALIEFATVNYFTKRGWAWDGKSVVNDKKKEKASVMIQNNAYAVAVANYAPNL  378
XP_003816076.1|GABRA2_Pan_paniscus             VPARTVFGVTTVLTMTTLSISARNSLPKVAYATAMDWFIAVCYAFVFSALIEFATVNYFTKRGWAWDGKSVVNDKKKEKASVMIQNNAYAVAVANYAPNL  378
M3Y0V9|GABRA2_Mustela_putorius_furo            VPARTVFGVTTVLTMTTLSISARNSLPKVAYATAMDWFIAVCYAFVFSALIEFATVNYFTKRGWAWDGKSVVNDKKKEKASVMIQNNAYAVAVANYAPNL  400
P23576|GABRA2_Rattus_norvegicus                VPARTVFGVTTVLTMTTLSISARNSLPKVAYATAMDWFIAVCYAFVFSALIEFATVNYFTKRGWAWDGKSVVNDKKKEKGSVMIQNNAYAVAVANYAPNL  378
P26048|GABRA2_Mus_musculus                     VPARTVFGVTTVLTMTTLSISARNSLPKVAYATAMDWFIAVCYAFVFSALIEFATVNYFTKRGWAWDGKSVVNDKKKEKGSVMIQNNAYAVAVANYAPNL  378
A0A5F4CYJ7|GABRA2_Canis_lupus_familiaris       VPARTVFGVTTVLTMTTLSISARNSLPKVAYATAMDWFIAVCYAFVFSALIEFATVNYFTKRGWAWDGKSVVNDKKKEKASVMIQNNAYAVAVANYAPNL  369
H2QPE5|GABRA2_Pan_troglodytes                  VPARTVFGVTTVLTMTTLSISARNSLPKVAYATAMDWFIAVCYAFVFSALIEFATVNYFTKRGWAWDGKSVVNDKKKEKASVMIQNNAYAVAVANYAPNL  378
E7F635|GABRA2a_Danio_rerio                     VPARTVFGVTTVLTMTTLSISARNSLPKVAYATAMDWFIAVCYAFVFSALIEFATVNYFTKRGWAWDGKSIVNDKKKE-TSVMKKNNAYAVAVANYAPHI  377
XP_028703913.1|GABRA2_Macaca_mulatta           VPARTVFGVTTVLTMTTLSISARNSLPKVAYATAMDWFIAVCYAFVFSALIEFATVNYFTKRGWAWDGKSVVNDKKKEKASVMIQNNAYAVAVANYAPNL  378
P10063|GABRA2_Bos_taurus                       VPARTVFGVTTVLTMTTLSISARNSLPKVAYATAMDWFIAVCYAFVFSALIEFATVNYFTKRGWAWDGKSVVNDKKKEKASVMIQNNAYAVAVANYAPNL  378
                                               .......310.......320.......330.......340.......350.......360.......370.......380.......390.......400

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P47869|GABRA2_Homo_sapiens                     SKDPVLSTISKSATTPEPNKKPENKPAEAKKTFNSVSKIDRMSRIVFPVLFGTFNLVYWATYLNREPVL-GVSP--   451
XP_003816076.1|GABRA2_Pan_paniscus             SKDPVLSTISKSATTPEPNKKPENKPAEAKKTFNSVSKIDRMSRIVFPVLFGTFNLVYWATYLNREPVL-GVSP--   451
M3Y0V9|GABRA2_Mustela_putorius_furo            SKDPVLSTISKSATTPEPNKKPENKPAEAKKTFNSVSKIDRMSRIVFPVLFGTFNLVYWATYLNREPVL-GVSP--   473
P23576|GABRA2_Rattus_norvegicus                SKDPVLSTISKSATTPEPNKKPENKPAEAKKTFNSVSKIDRMSRIVFPVLFGTFNLVYWATYLNREPVL-GVSP--   451
P26048|GABRA2_Mus_musculus                     SKDPVLSTISKSATTPEPNKKPENKPAEAKKTFNSVSKIDRMSRIVFPVLFGTFNLVYWATYLNREPVL-GVSP--   451
A0A5F4CYJ7|GABRA2_Canis_lupus_familiaris       SKDPVLSTISKSATTPEPNKKPENKPAEAKKTFNSVSKIDRMSRIVFPVLFGTFNLVYWATYLNREPVL-GVSP--   442
H2QPE5|GABRA2_Pan_troglodytes                  SKDPVLSTISKSATTPEPNKKPENKPAEAKKTFNSVSKIDRMSRIVFPVLFGTFNLVYWATYLNREPVL-GVSP--   451
E7F635|GABRA2a_Danio_rerio                     AKDSALPTVSKSA-TPEPS-KAQPVAKETKKTFNSVSKIDRMSRIIFPVLFGSFNLVYWATYLNREPVIKNMVPSP   451
XP_028703913.1|GABRA2_Macaca_mulatta           SKDPVLSTISKSATTPEPNKKPENKPAEAKKTFNSVSKIDRMSRIVFPVLFGTFNLVYWATYLNREPVL-GVSP--   451
P10063|GABRA2_Bos_taurus                       SKDPVLSTISKSATTPEPNKKPENKPAEAKKTFNSVSKIDRMSRIVFPVLFGTFNLVYWATYLNREPVL-GVSP--   451
                                               .......410.......420.......430.......440.......450.......460.......470.....
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P34903|GABRA3_Homo_sapiens           MIITQTSHCYMTSLGILFLINILPGTTGQGESRRQEPGDFVKQDIGGLSPKHAPDIPDDSTDNITIFTRILDRLLDGYDNRLRPGLGDAVTEVKTDIYVT  100
A0A2R9AHX2|GABRA3_Pan_paniscus       MIITQTSHCYMTSLGILFLINILPGTTGQGESRRQEPGDFVKQDIGGLSPKHAPDIPDDSTDNITIFTRILDRLLDGYDNRLRPGLGDAVTEVKTDIYVT  100
H2QZ88|GABRA3_Pan_troglodytes        MIITQTSHCYMTSLGILFLINILPGTTGQGESRRQEPGDFVKQDIGGLSPKHAPDIPDDSTDNITIFTRILDRLLDGYDNRLRPGLGDAVTEVKTDIYVT  100
F7G8R5|GABRA3_Macaca_mulatta         MIITQMSHYYMTSLGILFLINILPGTTGQGESRRQEPGDFVKQDIGGLSPKHAPDIPDDSTDNITIFTRILDRLLDGYDNRLRPGLGDAVTEVKTDIYVT  100
M3YJX4|GABRA3_Mustela_putorius_furo  MIVTQMSHFYTTSLGFLFLINIFPGITGQGESRRQEPGDFVKQDIGGLSPKHAPDIPDDSTDNITIFTRILDRLLDGYDNRLRPGLGDAVTEVKTDIYVT  100
P20236|GABRA3_Rattus_norvegicus      MITTQMWHFYVTRVGLLLLISILPGTTGQGESRRQEPGDFVKQDIGGLSPKHAPDIPDDSTDNITIFTRILDRLLDGYDNRLRPGLGDAVTEVKTDIYVT  100
P26049|GABRA3_Mus_musculus           MIITQMWHFYVTRVVLLLLISILPGTTSQGESRRQEPGDFVKQDIGGLSPKHAPDIPDDSTDNITIFTRILDRLLDGYDNRLRPGLGDAVTEVKTDIYVT  100
E2QVJ7|GABRA3_Canis_lupus_familiaris MIVTQMSHFYTIILGLLFLINIFPGTTGQGESRRQEPGDFVKQDIGGLSPKHAPDIPDDSTDNITIFTRILDRLLDGYDNRLRPGLGDAVTEVKTDIYVT  100
A0A2R8QC15|GABRA3_Danio_rerio        -----------------------------------------HRAVTG----HREPIPDDSKDNITIFTRILDRLLDGYDNRLRPGLGESVTEVRTNIYVT   55
P10064|GABRA3_Bos_taurus             MIITQMSQFYMAGLGLLFLINILPGTTGQVESRRQEPGDFVKQDIGGLSPKHAPDIPDDSTDNITIFTRILDRLLDGYDNRLRPGLGDAVTEVKTDIYVT  100
                                     1.......10........20........30........40........50........60........70........80........90.......100

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P34903|GABRA3_Homo_sapiens           SFGPVSDTDMEYTIDVFFRQTWHDERLKFDGPMKILPLNNLLASKIWTPDTFFHNGKKSVAHNMTTPNKLLRLVDNGTLLYTMRLTIHAECPMHLEDFPM  200
A0A2R9AHX2|GABRA3_Pan_paniscus       SFGPVSDTDMEYTIDVFFRQTWHDERLKFDGPMKILPLNNLLASKIWTPDTFFHNGKKSVAHNMTTPNKLLRLVDNGTLLYTMRLTIHAECPMHLEDFPM  200
H2QZ88|GABRA3_Pan_troglodytes        SFGPVSDTDMEYTIDVFFRQTWHDERLKFDGPMKILPLNNLLASKIWTPDTFFHNGKKSVAHNMTTPNKLLRLVDNGTLLYTMRLTIHAECPMHLEDFPM  200
F7G8R5|GABRA3_Macaca_mulatta         SFGPVSDTDMEYTIDVFFRQTWHDERLKFDGPMKILPLNNLLASKIWTPDTFFHNGKKSVAHNMTTPNKLLRLVDNGTLLYTMRLTIHAECPMHLEDFPM  200
M3YJX4|GABRA3_Mustela_putorius_furo  SFGPVSDTDMEYTIDVFFRQTWHDERLKFDGPMKILPLNNLLASKIWTPDTFFHNGKKSVAHNMTTPNKLLRLVDNGTLLYTMRLTIHAECPMHLEDFPM  200
P20236|GABRA3_Rattus_norvegicus      SFGPVSDTDMEYTIDVFFRQTWHDERLKFDGPMKILPLNNLLASKIWTPDTFFHNGKKSVAHNMTTPNKLLRLVDNGTLLYTMRLTIHAECPMHLEDFPM  200
P26049|GABRA3_Mus_musculus           SFGPVSDTDMEYTIDVFFRQTWHDERLKFDGPMKILPLNNLLASKIWTPDTFFHNGKKSVAHNMTTPNKLLRLVDNGTLLYTMRLTIHAECPMHLEDFPM  200
E2QVJ7|GABRA3_Canis_lupus_familiaris SFGPVSDTDMEYTIDVFFRQTWHDERLKFDGPMKILPLNNLLASKIWTPDTFFHNGKKSVAHNMTTPNKLLRLVDNGTLLYTMRLTIHAECPMHLEDFPM  200
A0A2R8QC15|GABRA3_Danio_rerio        SFGPVSDTDMEYTIDVFFRQSWRDERLKFDGPMQVLPLNNLLASKIWTPDTFFHNGKKSVAHNMTTPNKLLRLVDNGTLLYTMRLTIHAECPMHLEDFPM  155
P10064|GABRA3_Bos_taurus             SFGPVSDTDMEYTIDVFFRQTWHDERLKFDGPMKILPLNNLLASKIWTPDTFFHNGKKSVAHNMTTPNKLLRLVDNGTLLYTMRLTIHAECPMHLEDFPM  200
                                     .......110.......120.......130.......140.......150.......160.......170.......180.......190.......200

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P34903|GABRA3_Homo_sapiens           DVHACPLKFGSYAYTTAEVVYSWTLGKNKSVEVAQDGSRLNQYDLLGHVVGTEIIRSSTGEYVVMTTHFHLKRKIGYFVIQTYLPCIMTVILSQVSFWLN  300
A0A2R9AHX2|GABRA3_Pan_paniscus       DVHACPLKFGSYAYTTAEVVYSWTLGKNKSVEVAQDGSRLNQYDLLGHVVGTEIIRSSTGEYVVMTTHFHLKRKIGYFVIQTYLPCIMTVILSQVSFWLN  300
H2QZ88|GABRA3_Pan_troglodytes        DVHACPLKFGSYAYTTAEVVYSWTLGKNKSVEVAQDGSRLNQYDLLGHVVGTEIIRSSTGEYVVMTTHFHLKRKIGYFVIQTYLPCIMTVILSQVSFWLN  300
F7G8R5|GABRA3_Macaca_mulatta         DVHACPLKFGSYAYTTAEVVYSWTLGKNKSVEVAQDGSRLNQYDLLGHVVGTEIIRSSTGEYVVMTTHFHLKRKIGYFVIQTYLPCIMTVILSQVSFWLN  300
M3YJX4|GABRA3_Mustela_putorius_furo  DVHACPLKFGSYAYTTAEVVYSWTLGKNKSVEVAQDGSRLNQYDLLGHVVGTEIIRSSTGEYVVMTTHFHLKRKIGYFVIQTYLPCIMTVILSQVSFWLN  300
P20236|GABRA3_Rattus_norvegicus      DVHACPLKFGSYAYTKAEVIYSWTLGKNKSVEVAQDGSRLNQYDLLGHVVGTEIIRSSTGEYVVMTTHFHLKRKIGYFVIQTYLPCIMTVILSQVSFWLN  300
P26049|GABRA3_Mus_musculus           DVHACPLKFGSYAYTKAEVIYSWTLGKNKSVEVAQDGSRLNQYDLLGHVVGTEIIRSSTGEYVVMTTHFHLKRKIGYFVIQTYLPCIMTVILSQVSFWLN  300
E2QVJ7|GABRA3_Canis_lupus_familiaris DVHACPLKFGSYAYTTAEVVYSWTLGKNKSVEVAQDGSRLNQYDLLGHVVGTEIIRSSTGEYVVMTTHFHLKRKIGYFVIQTYLPCIMTVILSQVSFWLN  300
A0A2R8QC15|GABRA3_Danio_rerio        DAHACPLKFGSYAYTNNEVVYIWTADDEKSVSVAPDGSRLNQYDLLGHVIGKETISSSTGEYVVMTTYFHLKRKIGYFVIQTYLPCIMTVILSQVSFWLN  255
P10064|GABRA3_Bos_taurus             DVHACPLKFGSYAYTTAEVVYSWTLGKNKSVEVAQDGSRLNQYDLLGHVVGTEIIRSSTGEYVVMTTHFHLKRKIGYFVIQTYLPCIMTVILSQVSFWLN  300
                                     .......210.......220.......230.......240.......250.......260.......270.......280.......290.......300

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P34903|GABRA3_Homo_sapiens           RESVPARTVFGVTTVLTMTTLSISARNSLPKVAYATAMDWFIAVCYAFVFSALIEFATVNYFTKRSWAWEGKKVPEALEMKKKTPAAPAKKTSTTFNIVG  400
A0A2R9AHX2|GABRA3_Pan_paniscus       RESVPARTVFGVTTVLTMTTLSISARNSLPKVAYATAMDWFIAVCYAFVFSALIEFATVNYFTKRSWAWEGKKVPEALEMKKKTPAAPAKKTSTTFNIVG  400
H2QZ88|GABRA3_Pan_troglodytes        RESVPARTVFGVTTVLTMTTLSISARNSLPKVAYATAMDWFIAVCYAFVFSALIEFATVNYFTKRSWAWEGKKVPEALEMKKKTPAAPAKKTSTTFNIVG  400
F7G8R5|GABRA3_Macaca_mulatta         RESVPARTVFGVTTVLTMTTLSISARNSLPKVAYATAMDWFIAVCYAFVFSALIEFATVNYFTKRSWAWEGKKVPEALEMKKKTPAAPAKKTSTTFNIVG  400
M3YJX4|GABRA3_Mustela_putorius_furo  RESVPARTVFGVTTVLTMTTLSISARNSLPKVAYATAMDWFIAVCYAFVFSALIEFATVNYFTKRSWAWEGKKVPEALEMKKKTPAAPTKKTSTTFNIVG  400
P20236|GABRA3_Rattus_norvegicus      RESVPARTVFGVTTVLTMTTLSISARNSLPKVAYATAMDWFIAVCYAFVFSALIEFATVNYFTKRSWAWEGKKVPEALEMKKKTPAAPTKKTSTTFNIVG  400
P26049|GABRA3_Mus_musculus           RESVPARTVFGVTTVLTMTTLSISARNSLPKVAYATAMDWFIAVCYAFVFSALIEFATVNYFTKRSWAWEGKKVPEALEMKKKTPAAPTKK-NTTFNIVG  399
E2QVJ7|GABRA3_Canis_lupus_familiaris RESVPARTVFGVTTVLTMTTLSISARNSLPKVAYATAMDWFIAVCYAFVFSALIEFATVNYFTKRSWAWEGKKVPEALEMKKKTPAAPTKKTSTTFNIVG  400
A0A2R8QC15|GABRA3_Danio_rerio        RESVPARTVFGKS-----------------------------------------------FF--------------------------------------  270
P10064|GABRA3_Bos_taurus             RESVPARTVFGVTTVLTMTTLSISARNSLPKVAYATAMDWFMAVCYAFVFSALIEFATVNYFTKRSWAWEGKKVPEALEMKKKTPAVPTKKTSTTFNIVG  400
                                     .......310.......320.......330.......340.......350.......360.......370.......380.......390.......400

                                       **                                                                  * :**
P34903|GABRA3_Homo_sapiens           TTYPINLAKDTEFSTISK-GAAPSASSTPTIIASPKATYVQDSPTETKTYNSVSKVDKISRIIFPVLFAIFNLVYWATYVNRESAIKGMIRKQ  492
A0A2R9AHX2|GABRA3_Pan_paniscus       TTYPINLAKDTEFSTISK-GAAPSASSTPTIIASPKATYVQDSPTETKTYNSVSKVDKISRIIFPVLFAIFNLVYWATYVNRESAIKGMIRKQ  492
H2QZ88|GABRA3_Pan_troglodytes        TTYPINLAKDTEFSTISK-GAAPSASSTPTIIASPKATYVQDSPTETKTYNSVSKVDKISRIIFPVLFAIFNLVYWATYVNRESAIKGMIRKQ  492
F7G8R5|GABRA3_Macaca_mulatta         TTYPINLAKDTEFSTISK-GAAPSASSTPTIIASPKATYVQDSPTETKTYNSVSKVDKISRIIFPVLFAIFNLVYWATYVNRESAIKGMIRKQ  492
M3YJX4|GABRA3_Mustela_putorius_furo  TTYPINLAKDTEFSTISK-GAAPSASSTPTIIASPKTTYVQDIPTETKTYNSVSKVDKISRIIFPVLFAIFNLVYWATYVNRESAIKGMIRKH  492
P20236|GABRA3_Rattus_norvegicus      TTYPINLAKDTEFSTISKAAAAPSASSTPTVIASPKTTYVQDSPAETKTYNSVSKVDKISRIIFPVLFAIFNLVYWATYVNRESAIKGMIRKQ  493
P26049|GABRA3_Mus_musculus           TTYPINLAKDTEFSTISKSAAAPSASSTPTAIASPKATYVQDSPAETKTYNSVSKVDKISRIIFPVLFAIFNLVYWATYVNRESAIKGMIRKQ  492
E2QVJ7|GABRA3_Canis_lupus_familiaris TTYPINLAKDTEFSTISK-GAAPSASSTPTIIASPKTTYVQDIPTETKTYNSVSKVDKISRIIFPVLFAIFNLVYWATYVNRESAIKGMIRKH  492
A0A2R8QC15|GABRA3_Danio_rerio        --YP------------------------------------------------------------------FAVVY------------------  277
P10064|GABRA3_Bos_taurus             TTYPINLAKDTEFSAISK-GAAPSTSSTPTIIASPKTTCVQDIPTETKTYNSVSKVDKISRIIFPVLFAIFNLVYWATYVNRESAIKGMIRKQ  492
                                     .......410.......420.......430.......440.......450.......460.......470.......480.......490...
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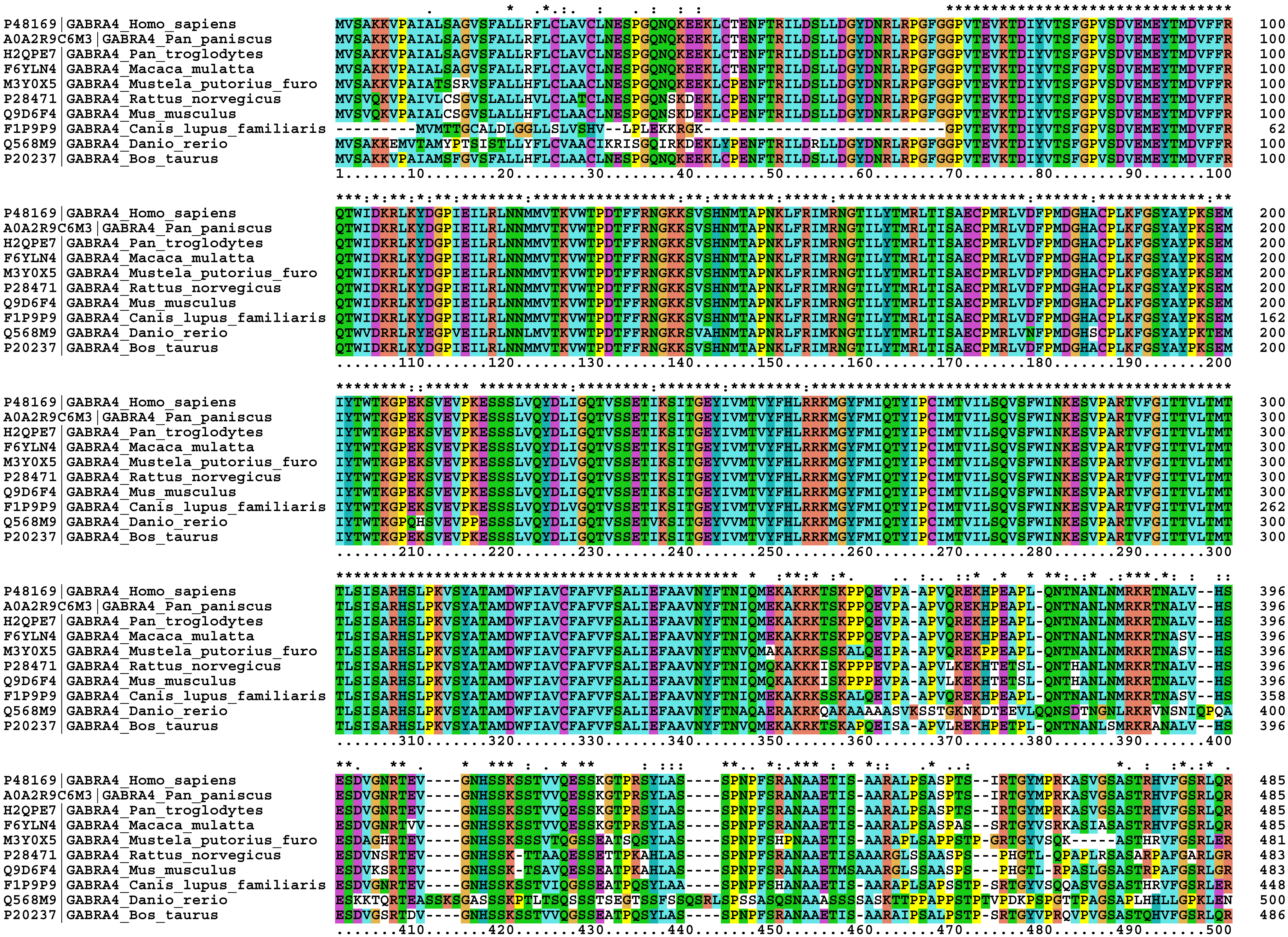
P48169|GABRA4_Homo_sapiens
A0A2R9C6M3|GABRA4_Pan_paniscus
H2QPE7|GABRA4_Pan_troglodytes
F6YLN4|GABRA4_Macaca_mulatta
M3Y0X5|GABRA4_Mustela_putorius_furo
P28471|GABRA4_Rattus_norvegicus
Q9D6F4|GABRA4_Mus_musculus
F1P9P9|GABRA4_Canis_lupus_familiaris
Q568M9|GABRA4_Danio_rerio
P20237|GABRA4_Bos_taurus

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                                     **          : .* : . .:.*..* *:****:********:******************* . :
P48169|GABRA4_Homo_sapiens           IKTTVNTIGATGKLSA-TPPPSAPPPSGSGTSKIDKYARILFPVTFGAFNMVYWVVYLSKDTMEKSESLM   554
A0A2R9C6M3|GABRA4_Pan_paniscus       IKTTVNTIGATGKLSA-TPPPSTPPPSGSGTSKIDKYARILFPVTFGAFNMVYWVVYLSKDTMEKSESLM   554
H2QPE7|GABRA4_Pan_troglodytes        IKTTVNTIGATGKLSA-TPPPSAPPPSGSGTSKIDKYARILFPVTFGAFNMVYWVVYLSKDTMEKSESLM   554
F6YLN4|GABRA4_Macaca_mulatta         IKTTVNTIGATGKLSA-TPPPSAPPPSGSGTSKIDKYARILFPVTFGAFNMVYWVVYLSKDTMEKSESLM   554
M3Y0X5|GABRA4_Mustela_putorius_furo  IKTTVNTTGAPGKLSATTPPPPAPPPPGSGTSKIDKYARILFPVTFGAFNMVYWVVYLSKDTMEKSESLM   551
P28471|GABRA4_Rattus_norvegicus      IKTTVNTTGVPGNVSA-TPPPSAPPPSGSGTSKIDKYARILFPVTFGAFNMVYWVVYLSKDTMEKSESLM   552
Q9D6F4|GABRA4_Mus_musculus           IKTTVNTTGAAGNVSA-TPPPPAPPPSGSGTSKIDKYARILFPVTFGAFNMVYWVVYLSKDTMEKSESLM   552
F1P9P9|GABRA4_Canis_lupus_familiaris IKTTVNTIGASGKLSA-TTPPSAPPPSGSGTSKIDKYARILFPVTFGAFNMVYWVVYLSKDTMEKSESLM   517
Q568M9|GABRA4_Danio_rerio            IK----------KKEAKAQPQPAAPATG-GASKIDEYARILFPVSFGAFNMVYWVVYLSKDTMEAKGA--   557
P20237|GABRA4_Bos_taurus             IKTTVNSIGTSGKLSA-TTTPSAPPPSGSGTSKIDKYARILFPVTFGAFNMVYWVVYLSKDTMEKSESLM   555
                                     .......510.......520.......530.......540.......550.......560.......570
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                                      *. *  .   * :.* :  : *.*:.*:.::* .*.: :  * :****:*********************::*:::*:*:******************
P31644|GABRA5_Homo_sapiens            MDNGMFSGFIMIKNLLLFCISMNLSSHFGFSQMPTSSVKD-ETNDNITIFTRILDGLLDGYDNRLRPGLGERITQVRTDIYVTSFGPVSDTEMEYTIDVF   99
A0A2R9BFW8|GABRA5_Pan_paniscus        MDNGMFSGFIMIKNLLLFCISMNLSSHFGFSQMPTSSVKD-ETNDNITIFTRILDGLLDGYDNRLRPGLGERITQVRTDIYVTSFGPVSDTEMEYTIDVF   99
H2Q919|GABRA5_Pan_troglodytes         MDNGMFSGFIMIKNLLLFCISMNLSSHFGFSQMPTSSVKD-ETNDNITIFTRILDGLLDGYDNRLRPGLGERITQVRTDIYVTSFGPVSDTEMEYTIDVF   99
G7MW76|GABRA5_Macaca_mulatta          MDNGMFSGFIMIKNLLLFCISMNLSSHFGFSQMPTSSVKD-ETNDNITIFTRILDGLLDGYDNRLRPGLGERITQVRTDIYVTSFGPVSDTEMEYTIDVF   99
M3XNJ0|GABRA5_Mustela_putorius_furo   MDNGMFSSFIMIKNLLLFCISMNLASHFGFSQMPTSSVKD-ETNDNITIFTRILDGLLDGYDNRLRPGLGERITQVRTDIYVTSFGPVSDTEMEYTIDVF   99
P19969|GABRA5_Rattus_norvegicus       MDNGMLSRFIMTKTLLVFCISMTLSSHFGFSQMPTSSVQD-ETNDNITIFTRILDGLLDGYDNRLRPGLGERITQVRTDIYVTSFGPVSDTEMEYTIDVF   99
Q8BHJ7|GABRA5_Mus_musculus            MDNGMLSRFIMTQTLLVFCISMTLSSHFGFSQMPTSSVQD-ETNDNITIFTRILDGLLDGYDNRLRPGLGERITQVRTDIYVTSFGPVSDTEMEYTIDVF   99
E2RG38|GABRA5_Canis_lupus_familiaris  MDNGMFSSFIMIKNLLLFCISMNLASHFGFAQMPTSSVKD-ETNDNITIFTRILDGLLDGYDNRLRPGLGERITQVRTDIYVTSFGPVSDTEMEYTIDVF   99
E9QE70|GABRA5_Danio_rerio             MGYGPHCSGKM-RSLWVGVLFMTLTCHLSLSQSTTGTPKESELNDNITVFTRILDGLLDGYDNRLRPGLGEKVTEIKTNIFVTSFGPVSDTEMEYTIDVF   99
Q08E50|GABRA5_Bos_taurus              MDNGMFSSFIMIKNLLLFCISMNLASHFGFSQMPTSSVKA-ETDDNITIFTRILDGLLDGYDNRLRPGLGERITQVRTDIYVTSFGPVSDTEMEYTIDVF   99
                                      1.......10........20........30........40........50........60........70........80........90.......100

                                      ********** *****: *********:******** *******************:************************************** *
P31644|GABRA5_Homo_sapiens            FRQSWKDERLRFKGPMQRLPLNNLLASKIWTPDTFFHNGKKSIAHNMTTPNKLLRLEDDGTLLYTMRLTISAECPMQLEDFPMDAHACPLKFGSYAYPNS  199
A0A2R9BFW8|GABRA5_Pan_paniscus        FRQSWKDERLRFKGPMQRLPLNNLLASKIWTPDTFFHNGKKSIAHNMTTPNKLLRLEDDGTLLYTMRLTISAECPMQLEDFPMDAHACPLKFGSYAYPNS  199
H2Q919|GABRA5_Pan_troglodytes         FRQSWKDERLRFKGPMQRLPLNNLLASKIWTPDTFFHNGKKSIAHNMTTPNKLLRLEDDGTLLYTMRLTISAECPMQLEDFPMDAHACPLKFGSYAYPNS  199
G7MW76|GABRA5_Macaca_mulatta          FRQSWKDERLRFKGPMQRLPLNNLLASKIWTPDTFFHNGKKSIAHNMTTPNKLLRLEDDGTLLYTMRLTISAECPMQLEDFPMDAHACPLKFGSYAYPNS  199
M3XNJ0|GABRA5_Mustela_putorius_furo   FRQSWKDERLRFKGPMQRLPLNNLLASKIWTPDTFFHNGKKSIAHNMTTPNKLLRLEDDGTLLYTMRLTISAECPMQLEDFPMDAHACPLKFGSYAYPNS  199
P19969|GABRA5_Rattus_norvegicus       FRQSWKDERLRFKGPMQRLPLNNLLASKIWTPDTFFHNGKKSIAHNMTTPNKLLRLEDDGTLLYTMRLTISAECPMQLEDFPMDAHACPLKFGSYAYPNS  199
Q8BHJ7|GABRA5_Mus_musculus            FRQSWKDERLRFKGPMQRLPLNNLLASKIWTPDTFFHNGKKSIAHNMTTPNKLLRLEDDGTLLYTMRLTISAECPMQLEDFPMDAHACPLKFGSYAYPNS  199
E2RG38|GABRA5_Canis_lupus_familiaris  FRQSWKDERLRFKGPMQRLPLNNLLASKIWTPDTFFHNGKKSIAHNMTTPNKLLRLEDDGTLLYTMRLTISAECPMQLEDFPMDAHACPLKFGSYAYPNS  199
E9QE70|GABRA5_Danio_rerio             FRQSWKDERLCFKGPMEMLPLNNLLASNIWTPDTFFLNGKKSIAHNMTTPNKLLRLKDDGTLLYTMRLTISAECPMQLEDFPMDAHACPLKFGSYAYPIS  199
Q08E50|GABRA5_Bos_taurus              FRQSWKDERLRFKGPMQRLPLNNLLASKIWTPDTFFHNGKKSIAHNMTTPNKLLRLEDDGTLLYTMRLTISAECPMQLEDFPMDAHACPLKFGSYAYPNS  199
                                      .......110.......120.......130.......140.......150.......160.......170.......180.......190.......200

                                      **:* **:*. *******:*********:*:*.***:**** *:**:* ***:*************:**:*****************************
P31644|GABRA5_Homo_sapiens            EVVYVWTNGSTKSVVVAEDGSRLNQYHLMGQTVGTENISTSTGEYTIMTAHFHLKRKIGYFVIQTYLPCIMTVILSQVSFWLNRESVPARTVFGVTTVLT  299
A0A2R9BFW8|GABRA5_Pan_paniscus        EVVYVWTNGSTKSVVVAEDGSRLNQYHLMGQTVGTENISTSTGEYTIMTAHFHLKRKIGYFVIQTYLPCIMTVILSQVSFWLNRESVPARTVFGVTTVLT  299
H2Q919|GABRA5_Pan_troglodytes         EVVYVWTNGSTKSVVVAEDGSRLNQYHLMGQTVGTENISTSTGEYTIMTAHFHLKRKIGYFVIQTYLPCIMTVILSQVSFWLNRESVPARTVFGVTTVLT  299
G7MW76|GABRA5_Macaca_mulatta          EVVYVWTNGSTKSVVVAEDGSRLNQYHLMGQTVGTENISTSTGEYTIMTAHFHLKRKIGYFVIQTYLPCIMTVILSQVSFWLNRESVPARTVFGVTTVLT  299
M3XNJ0|GABRA5_Mustela_putorius_furo   EVVYVWTNGTTKSVVVAEDGSRLNQYHLMGQTVGTENISTSTGEYTVMTAHFHLKRKIGYFVIQTYLPCIMTVILSQVSFWLNRESVPARTVFGVTTVLT  299
P19969|GABRA5_Rattus_norvegicus       EVVYVWTNGSTKSVVVAEDGSRLNQYHLMGQTVGTENISTSTGEYTIMTAHFHLKRKIGYFVIQTYLPCIMTVILSQVSFWLNRESVPARTVFGVTTVLT  299
Q8BHJ7|GABRA5_Mus_musculus            EVVYVWTNGSTKSVVVAEDGSRLNQYHLMGQTVGTENISTSTGEYTIMTAHFHLKRKIGYFVIQTYLPCIMTVILSQVSFWLNRESVPARTVFGVTTVLT  299
E2RG38|GABRA5_Canis_lupus_familiaris  EVVYVWTNGTTKSVVVAEDGSRLNQYHLMGQTVGTENISTSTGEYTIMTAHFHLKRKIGYFVIQTYLPCIMTVILSQVSFWLNRESVPARTVFGVTTVLT  299
E9QE70|GABRA5_Danio_rerio             EVVYKWTKGPGKSVVVAEEGSRLNQYHLIGHTAGTEDISTSRGQYTVMMAHFYLKRKIGYFVIQTYMPCFMTVILSQVSFWLNRESVPARTVFGVTTVLT  299
Q08E50|GABRA5_Bos_taurus              EVIYVWTNGTAKSVVVAEDGSRLNQYHLMGQTVGTENISTSTGEYTIMTAHFHLKRKIGYFVIQTYLPCIMTVILSQVSFWLNRESVPARTVFGVTTVLT  299
                                      .......210.......220.......230.......240.......250.......260.......270.......280.......290.......300

                                      ********************************************.******** ** : ***   : :.*..* :*:       ** *:   .
P31644|GABRA5_Homo_sapiens            MTTLSISARNSLPKVAYATAMDWFIAVCYAFVFSALIEFATVNYFTKRGWAWDGKKALEAAKIKKK-REVILNKSTNAFTTGKMSHPPNIPKEQTPAGTS  398
A0A2R9BFW8|GABRA5_Pan_paniscus        MTTLSISARNSLPKVAYATAMDWFIAVCYAFVFSALIEFATVNYFTKRGWAWDGKKALEAAKIKKK-REVILNKSTNAFTTGKMSHPPNIPKEQTPAGTS  398
H2Q919|GABRA5_Pan_troglodytes         MTTLSISARNSLPKVAYATAMDWFIAVCYAFVFSALIEFATVNYFTKRGWAWDGKKALEAAKIKKK-REVILNKSTNAFTTGKMSHPPNIPKEQTPAGTS  398
G7MW76|GABRA5_Macaca_mulatta          MTTLSISARNSLPKVAYATAMDWFIAVCYAFVFSALIEFATVNYFTKRGWAWDGKKALEAAKIKKK-REVMLNKSTNAFTAGKMSHPPNIPKEQTPAGTS  398
M3XNJ0|GABRA5_Mustela_putorius_furo   MTTLSISARNSLPKVAYATAMDWFIAVCYAFVFSALIEFATVNYFTKRGWAWDGKKASEAAKVKKKERELILNKSTNAYTTGKMTHPPNIPKEQTPAGTS  399
P19969|GABRA5_Rattus_norvegicus       MTTLSISARNSLPKVAYATAMDWFIAVCYAFVFSALIEFATVNYFTKRGWAWDGKKALEAAKIKKKERELILNKSTNAFTTGKLTHPPNIPKEQLPGGTG  399
Q8BHJ7|GABRA5_Mus_musculus            MTTLSISARNSLPKVAYATAMDWFIAVCYAFVFSALIEFATVNYFTKRGWAWDGKKALEAAKIKKKERELILNKSTNAFTTGKLTHPPNIPKEQPPAGTA  399
E2RG38|GABRA5_Canis_lupus_familiaris  MTTLSISARNSLPKVAYATAMDWFIAVCYAFVFSALIEFATVNYFTKRGWAWDGKKASEAAKVKKKERELILNKSTNAYTTGKMTHPPNIPKEQTPAVTS  399
E9QE70|GABRA5_Danio_rerio             MTTLSISARNSLPKVAYATAMDWFIAVCYAFVFSALIEFATVNYFTKRSWAWDGKKALEAQQPKKK-DPLALSKKPNNFTA-------NINKDAAISTIS  391
Q08E50|GABRA5_Bos_taurus              MTTLSISARNSLPKVAYATAMDWFIAVCYAFVFSALIEFATVNYFTKRGWAWDGKKALEAAKIKKKERELTINKSTNAYTTGKMTHPPNIPKEQTPAGTT  399
                                      .......310.......320.......330.......340.......350.......360.......370.......380.......390.......400

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P31644|GABRA5_Homo_sapiens            NTT-SVSVKPSEEK-TSESKKTYNSISKIDKMSRIVFPVLFGTFNLVYWATYLNREPVIKGAASPK  462
A0A2R9BFW8|GABRA5_Pan_paniscus        NTT-SVSVKPSEEK-TSESKKTYNSISKIDKMSRIVFPVLFGTFNLVYWATYLNREPVIKGAASPK  462
H2Q919|GABRA5_Pan_troglodytes         NTT-SVSVKPSEEK-TSESKKTYNSISKIDKMSRIVFPVLFGTFNLVYWATYLNREPVIKGAASPK  462
G7MW76|GABRA5_Macaca_mulatta          NTT-SVSVKPSEEK-TPESKKTYNSISKIDKMSRIVFPVLFGTFNLVYWATYLNREPVIKGAASPK  462
M3XNJ0|GABRA5_Mustela_putorius_furo   NAS-SASVKPSEEK-TSENKKTYNSISKIDKMSRIVFPVLFGTFNLVYWATYLNREPVIKGAASPK  463
P19969|GABRA5_Rattus_norvegicus       NAVGTASIRASEEK-TSESKKTYNSISKIDKMSRIVFPILFGTFNLVYWATYLNREPVIKGATSPK  464
Q8BHJ7|GABRA5_Mus_musculus            NAP-TVSIKASEEK-TAESKKTYNSISKIDKMSRIVFPILFGTFNLVYWATYLNREPVIKGATSPK  463
E2RG38|GABRA5_Canis_lupus_familiaris  NAS-SASVKPSEEK-TSENKKTYNSISKIDKMSRIVFPVLFGTFNLVYWATYLNREPVIKGATSPK  463
E9QE70|GABRA5_Danio_rerio             NST-TVQLKSAEAKPAPDPKKTYNSVSKIDKMSRIVFPVLFGTFNLVYWATYLNREPVIKGAV---  453
Q08E50|GABRA5_Bos_taurus              NAS-SASVKP-EDK-ASENKKTYNSISKIDKMSRIIFPLLFGTFNLVYWATYLNREPVIKGATSPK  462
                                      .......410.......420.......430.......440.......450.......460......
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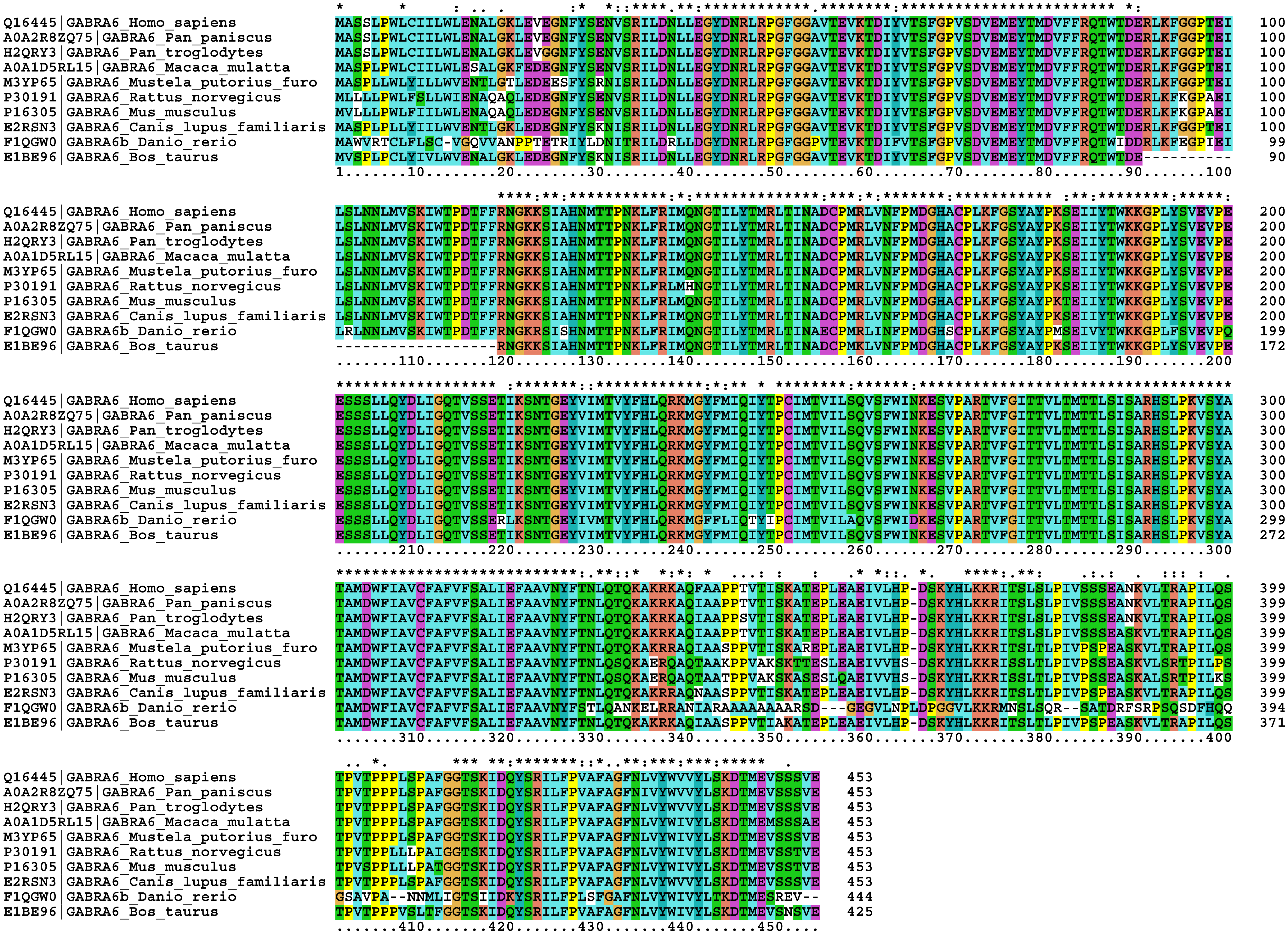
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H2QRY3|GABRA6_Pan_troglodytes
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M3YP65|GABRA6_Mustela_putorius_furo
P30191|GABRA6_Rattus_norvegicus
P16305|GABRA6_Mus_musculus
E2RSN3|GABRA6_Canis_lupus_familiaris
F1QGW0|GABRA6b_Danio_rerio
E1BE96|GABRA6_Bos_taurus

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                                        *  . .:   *::.: *::::.       **  .: **:.* *  .* :*..  : :           : : :  * :.* :. ***:********
P18505|GABRB1_Homo_sapiens              ----MWTVQNRESLGLLSFPVMITMVCC----AHSTNEPSNMSYVKETVDRLLKGYDIRLRPDFGGPPVDVGMRIDVASIDMVSEVNMDYTLTMYFQQSW   92
A0A2R9C4H8|GABRB1_Pan_paniscus          ----MWTVQNRESLGLLSFPVMITMVCC----AHSTNEPSNMSYVKETVDRLLKGYDIRLRPDFGGPPVDVGMRIDVASIDMVSEVNMDYTLTMYFQQSW   92
H2QPE8|GABRB1_Pan_troglodytes           ----MWTVQNRESLGLLSFPVMITMVCC----AHSTNEPSNMSYVKETVDRLLKGYDIRLRPDFGGPPVDVGMRIDVASIDMVSEVNMDYTLTMYFQQSW   92
G7MSU9|GABRB1_Macaca_mulatta            ----MWTAQNRESLGLLSFPVMITMVCC----AHSTNEPSNMSYVKETVDRLLKGYDIRLRPDFGGPPVDVGMRIDVASIDMVSEVNMDYTLTMYFQQSW   92
M3Y0Y5|GABRB1_Mustela_putorius_furo     ----MWTVQNRESLGLLSFPVMIAMVCC----AHSANEPSNMSYVKETVDRLLKGYDIRLRPDFGGPPVDVGMRIDVASIDMVSEVNMDYTLTMYFQQSW   92
P15431|GABRB1_Rattus_norvegicus         ----MWTVQNRESLGLLSFPVMVAMVCC----AHSSNEPSNMSYVKETVDRLLKGYDIRLRPDFGGPPVDVGMRIDVASIDMVSEVNMDYTLTMYFQQSW   92
P50571|GABRB1_Mus_musculus              ----MWTVQNRESLGLLSFPVMVAMVCC----AHSSNEPSNMSYVKETVDRLLKGYDIRLRPDFGGPPVDVGMRIDVASIDMVSEVNMDYTLTMYFQQSW   92
F1PEG2|GABRB1_Canis_lupus_familiaris    ----MWTAHSRESLGLLSFPVMIAMVCC----AHRVGK-SNLEYRKHITDDFLSSMMMMMM-----------IMMMMMIIIFFSFISQDYTLTMYFQQSW   80
F1QPW7|GABRB1_Danio_rerio               MMRGMRIAGDQWRAGIFCLLVLVALTLGRSPPAHSVNEPSNMSYVKVTVDKLLKGYDIRLRPDFGGPPVDVGMSIDISSIDMVSEVNMDYTITMYFQQSW  100
P08220|GABRB1_Bos_taurus                ----MWTVQNRESLGLLSFPVMIAMVCC----AHSANEPSNMSYVKETVDRLLKGYDIRLRPDFGGPPVDVGMRIDVASIDMVSEVNMDYTLTMYFQQSW   92
                                        1.......10........20........30........40........50........60........70........80........90.......100

                                        :******:*********************:**********************************************************************
P18505|GABRB1_Homo_sapiens              KDKRLSYSGIPLNLTLDNRVADQLWVPDTYFLNDKKSFVHGVTVKNRMIRLHPDGTVLYGLRITTTAACMMDLRRYPLDEQNCTLEIESYGYTTDDIEFY  192
A0A2R9C4H8|GABRB1_Pan_paniscus          KDKRLSYSGIPLNLTLDNRVADQLWVPDTYFLNDKKSFVHGVTVKNRMIRLHPDGTVLYGLRITTTAACMMDLRRYPLDEQNCTLEIESYGYTTDDIEFY  192
H2QPE8|GABRB1_Pan_troglodytes           KDKRLSYSGIPLNLTLDNRVADQLWVPDTYFLNDKKSFVHGVTVKNRMIRLHPDGTVLYGLRITTTAACMMDLRRYPLDEQNCTLEIESYGYTTDDIEFY  192
G7MSU9|GABRB1_Macaca_mulatta            KDKRLSYSGIPLNLTLDNRVADQLWVPDTYFLNDKKSFVHGVTVKNRMIRLHPDGTVLYGLRITTTAACMMDLRRYPLDEQNCTLEIESYGYTTDDIEFY  192
M3Y0Y5|GABRB1_Mustela_putorius_furo     KDKRLSYSGIPLNLTLDNRVADQLWVPDTYFLNDKKSFVHGVTVKNRMIRLHPDGTVLYGLRITTTAACMMDLRRYPLDEQNCTLEIESYGYTTDDIEFY  192
P15431|GABRB1_Rattus_norvegicus         KDKRLSYSGIPLNLTLDNRVADQLWVPDTYFLNDKKSFVHGVTVKNRMIRLHPDGTVLYGLRITTTAACMMDLRRYPLDEQNCTLEIESYGYTTDDIEFY  192
P50571|GABRB1_Mus_musculus              KDKRLSYSGIPLNLTLDNRVADQLWVPDTYFLNDKKSFVHGVTVKNRMIRLHPDGTVLYGLRITTTAACMMDLRRYPLDEQNCTLEIESYGYTTDDIEFY  192
F1PEG2|GABRB1_Canis_lupus_familiaris    KDKRLSYSGIPLNLTLDNRVADQLWVPDTYFLNDKKSFVHGVTVKNRMIRLHPDGTVLYGLRITTTAACMMDLRRYPLDEQNCTLEIESYGYTTDDIEFY  180
F1QPW7|GABRB1_Danio_rerio               RDKRLSYTGIPLNLTLDNRVADQLWVPDTYFINDKKSFVHGVTVKNRMIRLHPDGTVLYGLRITTTAACMMDLRRYPLDEQNCTLEIESYGYTTDDIEFY  200
P08220|GABRB1_Bos_taurus                KDKRLSYSGIPLNLTLDNRVADQLWVPDTYFLNDKKSFVHGVTVKNRMIRLHPDGTVLYGLRITTTAACMMDLRRYPLDEQNCTLEIESYGYTTDDIEFY  192
                                        .......110.......120.......130.......140.......150.......160.......170.......180.......190.......200

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P18505|GABRB1_Homo_sapiens              WNGGEGAVTGVNKIELPQFSIVDYKMVSKKVEFTTGAYPRLSLSFRLKRNIGYFILQTYMPSTLITILSWVSFWINYDASAARVALGITTVLTMTTISTH  292
A0A2R9C4H8|GABRB1_Pan_paniscus          WNGGEGAVTGVNKIELPQFSIVDYKMVSKKVEFTTGAYPRLSLSFRLKRNIGYFILQTYMPSTLITILSWVSFWINYDASAARVALGITTVLTMTTISTH  292
H2QPE8|GABRB1_Pan_troglodytes           WNGGEGAVTGVNKIELPQFSIVDYKMVSKKVEFTTGAYPRLSLSFRLKRNIGYFILQTYMPSTLITILSWVSFWINYDASAARVALGITTVLTMTTISTH  292
G7MSU9|GABRB1_Macaca_mulatta            WNGGEGAVTGVNKIELPQFSIVDYKMVSKKVEFTTGAYPRLSLSFRLKRNIGYFILQTYMPSTLITILSWVSFWINYDASAARVALGITTVLTMTTISTH  292
M3Y0Y5|GABRB1_Mustela_putorius_furo     WNGGEGAVTGVNKIELPQFSIVDYKMVSKKVEFTTGAYPRLSLSFRLKRNIGYFILQTYMPSTLITILSWVSFWINYDASAARVALGITTVLTMTTISTH  292
P15431|GABRB1_Rattus_norvegicus         WNGGEGAVTGVNKIELPQFSIVDYKMVSKKVEFTTGAYPRLSLSFRLKRNIGYFILQTYMPSTLITILSWVSFWINYDASAARVALGITTVLTMTTISTH  292
P50571|GABRB1_Mus_musculus              WNGGEGAVTGVNKIELPQFSIVDYKMVSKKVEFTTGAYPRLSLSFRLKRNIGYFILQTYMPSTLITILSWVSFWINYDASAARVALGITTVLTMTTISTH  292
F1PEG2|GABRB1_Canis_lupus_familiaris    WNGGEGAVTGVNKIELPQFSIVDYKMVSKKVEFTTGAYPRLSLSFRLKRNIGYFILQTYMPSTLITILSWVSFWINYDASAARVALGITTVLTMTTISTH  280
F1QPW7|GABRB1_Danio_rerio               WQGG-SSVTGVDNIELPQFSIIDYKTLSKKVVFATGSYPRLSLSFKLKRNIGYFILQTYMPSTLITILSWVSFWINYDASAARVALGITTVLTMTTINTH  299
P08220|GABRB1_Bos_taurus                WNGGEGAVTGVNKIELPQFSIVDYKMVSKKVEFTTGAYPRLSLSFRLKRNIGYFILQTYMPSTLITILSWVSFWINYDASAARVALGITTVLTMTTISTH  292
                                        .......210.......220.......230.......240.......250.......260.......270.......280.......290.......300

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P18505|GABRB1_Homo_sapiens              LRETLPKIPYVKAIDIYLMGCFVFVFLALLEYAFVNYIFFGKGP--QKKGASKQDQSANEKNKLEMNKVQVDAHGNILLSTLEIRNETSGSEVLTSVSDP  390
A0A2R9C4H8|GABRB1_Pan_paniscus          LRETLPKIPYVKAIDIYLMGCFVFVFLALLEYAFVNYIFFGKGP--QKKGASKQDQSANEKNKLEMNKVQVDAHGNILLSTLEIRNETSGSEVLTSVSDP  390
H2QPE8|GABRB1_Pan_troglodytes           LRETLPKIPYVKAIDIYLMGCFVFVFLALLEYAFVNYIFFGKGP--QKKGASKQDQSANEKNKLEMNKVQVDAHGNILLSTLEIRNETSGSEVLTSVSDP  390
G7MSU9|GABRB1_Macaca_mulatta            LRETLPKIPYVKAIDIYLMGCFVFVFLALLEYAFVNYIFFGKGP--QKKGASKQDQSANEKNKLEMNKVQVDAHGNILLSTLEIRNETSGSEVLTSVSDP  390
M3Y0Y5|GABRB1_Mustela_putorius_furo     LRETLPKIPYVKAIDIYLMGCFVFVFLALLEYAFVNYIFFGKGP--QKKGASKQDQSANEKNKLEMNKVQVDAHGNILLSTLEIRNETSGSEVLTGVSEP  390
P15431|GABRB1_Rattus_norvegicus         LRETLPKIPYVKAIDIYLMGCFVFVFLALLEYAFVNYIFFGKGP--QKKGASKQDQSANEKNKLEMNKVQVDAHGNILLSTLEIRNETSGSEVLTGVSDP  390
P50571|GABRB1_Mus_musculus              LRETLPKIPYVKAIDIYLMGCFVFVFLALLEYAFVNYIFFGKGP--QKKGASKQDQSANEKNRLEMNKVQVDAHGNILLSTLEIRNETSGSEVLTGVSDP  390
F1PEG2|GABRB1_Canis_lupus_familiaris    LRETLPKIPYVKAIDIYLMGCFVFVFLALLEYAFVNYIFFGKGP--QKKGASKQDQSANEKNKLEMNKVQVDAHGNILLSALELRTEASGSEVLTGASEP  378
F1QPW7|GABRB1_Danio_rerio               LRETLPKIPYVKAIDIYLMGCFVFVFLALLEYAFVNYIFFGRGPHLQKKVAEKAAKSNNEKSRLESNKVQVDVHGSIPLTNLDLR--LSGAEMLGALGDP  397
P08220|GABRB1_Bos_taurus                LRETLPKIPYVKAIDIYLMGCFVFVFLALLEYAFVNYIFFGKGP--QKKGAGKQDQSANEKNKLEMNKVQVDAHGNILLSTLEIRNETSGSEVLTGVGDP  390
                                        .......310.......320.......330.......340.......350.......360.......370.......380.......390.......400

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P18505|GABRB1_Homo_sapiens              K-ATMYSYDSASIQYRKPLSSREAYGR---ALDRHGVPSKGRIRRRASQLKVKIPDLTDVNSIDKWSRMFFPITFSLFNVVYWLYYVH  474
A0A2R9C4H8|GABRB1_Pan_paniscus          K-ATMYSYDSASIQYRKPLSSREAYGR---ALDRHGVPSKGRIRRRASQLKVKIPDLTDVNSIDKWSRMFFPITFSLFNVVYWLYYVH  474
H2QPE8|GABRB1_Pan_troglodytes           K-ATMYSYDSASIQYRKPLSSREAYGR---ALDRHGVPSKGRIRRRASQLKVKIPDLTDVNSIDKWSRMFFPITFSLFNVVYWLYYVH  474
G7MSU9|GABRB1_Macaca_mulatta            K-ATMYSYDSASIQYRKPLSSREAYGR---TLDRHGVPSKGRIRRRASQLKVKIPDLTDVNSIDKWSRMFFPITFSLFNVVYWLYYVH  474
M3Y0Y5|GABRB1_Mustela_putorius_furo     KATSMYSYDSASIQYRKPLSSREGYGR---ALDRHGAHSKGRIRRRASQLKVKIPDLTDVNSIDKWSRMFFPITFSLFNVVYWLYYVH  475
P15431|GABRB1_Rattus_norvegicus         K-ATMYSYDSASIQYRKPLSSREGFGR---GLDRHGVPGKGRIRRRASQLKVKIPDLTDVNSIDKWSRMFFPITFSLFNVVYWLYYVH  474
P50571|GABRB1_Mus_musculus              K-ATMYSYDSASIQYRKPLSSREGFGR---GLDRHGVPGKGRIRRRASQLKVKIPDLTDVNSIDKWSRMFFPITFSLFNVVYWLYYVH  474
F1PEG2|GABRB1_Canis_lupus_familiaris    QAAGVFAFDSASVQFRRPLSSRDAYGR---APDRLGAHGKGRIRRRASQLKVKIPDLTDVNSIDKWSRMFFPITFSLFNVVYWLYYVH  463
F1QPW7|GABRB1_Danio_rerio               R-NTMFSYDSASIQYRKPLTGRDLYGRPTASIERPLAPKKSRLRRRAAQLKVKIPDLTDVNAIDKWSRVIFPITYTFFNLVYWLYYVH  484
P08220|GABRB1_Bos_taurus                K-TTMYSYDSASIQYRKPMSSREGYGR---ALDRHGAHSKGRIRRRASQLKVKIPDLTDVNSIDKWSRMFFPITFSLFNVVYWLYYVH  474
                                        .......410.......420.......430.......440.......450.......460.......470.......480........
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                                            : *  :: : . ***:*..****:.*****.:**:*****:*************.**** ************************************.
P47870|GABRB2_Homo_sapiens                 MWRVRKRGYFGIWSFPLIIAAVCAQSVNDPSNMSLVKETVDRLLKGYDIRLRPDFGGPPVAVGMNIDIASIDMVSEVNMDYTLTMYFQQAWRDKRLSYNV   100
A0A2R9C8C8|GABRB2_Pan_paniscus             MWRVRKRGYFGIWSFPLIIAAVCAQSVNDPSNMSLVKETVDRLLKGYDIRLRPDFGGPPVAVGMNIDIASIDMVSEVNMDYTLTMYFQQAWRDKRLSYNV   100
A0A2J8NKG3|GABRB2_Pan_troglodytes          MWRVRKRGYFGIWSFPLIIAAVCAQSVNDPSNMSLVKETVDRLLKGYDIRLRPDFGGPPVAVGMNIDIASIDMVSEVNMDYTLTMYFQQAWRDKRLSYNV   100
M3YP67|GABRB2_Mustela_putorius_furo        MWRVRKKGYFGIWSFPLIIAAVCAQSVNDPSNMSLVKETVDRLLKGYDIRLRPDFGGPPVAVGMNIDIASIDMVSEVNMDYTLTMYFQQAWRDKRLSYNV   100
P63138|GABRB2_Rattus_norvegicus            MWRVRKRGYFGIWSFPLIIAAVCAQSVNDPSNMSLVKETVDRLLKGYDIRLRPDFGGPPVAVGMNIDIASIDMVSEVNMDYTLTMYFQQAWRDKRLSYNV   100
P63137|GABRB2_Mus_musculus                 MWRVRKRGYFGIWSFPLIIAAVCAQSVNDPSNMSLVKETVDRLLKGYDIRLRPDFGGPPVAVGMNIDIASIDMVSEVNMDYTLTMYFQQAWRDKRLSYNV   100
A0A5F4CKV8|GABRB2_Canis_lupus_familiaris   MWRVRKKGYFGIWSFPLIIAAVCAQSVNDPSNMSLVKETVDRLLKGYDIRLRPDFGGPPVAVGMNIDIASIDMVSEVNMDYTLTMYFQQAWRDKRLSYNV   100
D1LYT2|GABRB2_Macaca_mulatta               MWRVRKRGYFGIWSFPLIIAAVCAQSVNDPSNMSLVKETVDRLLKGYDIRLRPDFGGPPVAVGMNIDIASIDMVSEVNMDYTLTMYFQQAWRDKRLSYNV   100
Q9DDD9|GABRB2_Danio_rerio                  MESIGKTPHIFLLC-PLIVAVACAQSIRDPSNMPVVKDTVDRLMKGYDIRLRPDFGGAPVAVRMNIDIASIDMVSEVNMDYTLTMYFQQAWRDKRLSYSE    99
P0C2W5|GABRB2_Bos_taurus                   --RVRKKDYFGIWSFPLIIAAVCAQSVNDPSNMSLVKETVDRLLKGYDIRLRPDFGGPPVAVGMNIDIASIDMVSEVNMDYTLTMYFQQAWRDKRLSYNV    98
                                           1.......10........20........30........40........50........60........70........80........90.......100

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P47870|GABRB2_Homo_sapiens                 IPLNLTLDNRVADQLWVPDTYFLNDKKSFVHGVTVKNRMIRLHPDGTVLYGLRITTTAACMMDLRRYPLDEQNCTLEIESYGYTTDDIEFYWRGDDNAVT   200
A0A2R9C8C8|GABRB2_Pan_paniscus             IPLNLTLDNRVADQLWVPDTYFLNDKKSFVHGVTVKNRMIRLHPDGTVLYGLRITTTAACMMDLRRYPLDEQNCTLEIESYGYTTDDIEFYWRGDDNAVT   200
A0A2J8NKG3|GABRB2_Pan_troglodytes          IPLNLTLDNRVADQLWVPDTYFLNDKKSFVHGVTVKNRMIRLHPDGTVLYGLRITTTAACMMDLRRYPLDEQNCTLEIESYGYTTDDIEFYWRGDDNAVT   200
M3YP67|GABRB2_Mustela_putorius_furo        IPLNLTLDNRVADQLWVPDTYFLNDKKSFVHGVTVKNRMIRLHPDGTVLYGLRITTTAACMMDLRRYPLDEQNCTLEIESYGYTTDDIEFYWRGDDNAVT   200
P63138|GABRB2_Rattus_norvegicus            IPLNLTLDNRVADQLWVPDTYFLNDKKSFVHGVTVKNRMIRLHPDGTVLYGLRITTTAACMMDLRRYPLDEQNCTLEIESYGYTTDDIEFYWRGDDNAVT   200
P63137|GABRB2_Mus_musculus                 IPLNLTLDNRVADQLWVPDTYFLNDKKSFVHGVTVKNRMIRLHPDGTVLYGLRITTTAACMMDLRRYPLDEQNCTLEIESYGYTTDDIEFYWRGDDNAVT   200
A0A5F4CKV8|GABRB2_Canis_lupus_familiaris   IPLNLTLDNRVADQLWVPDTYFLNDKKSFVHGVTVKNRMIRLHPDGTVLYGLRITTTAACMMDLRRYPLDEQNCTLEIESYGYTTDDIEFYWRGDDNAVT   200
D1LYT2|GABRB2_Macaca_mulatta               IPLNLTLDNRVADQLWVPDTYFLNDKKSFVHGVTVKNRMIRLHPDGTVLYGLRITTTAACMMDLRRYPLDEQNCTLEIESYGYTTDDIEFYWRGDDNAVT   200
Q9DDD9|GABRB2_Danio_rerio                  IPLNLTLDNRVADQLWVPDTYFLNDKKSFVHGVTVKNRMIRLHPDGTVLYGLRITTTAACMMDLRRYPLDEQNCTLEIESYGYTTDDIEFYWRGGDGAVS   199
P0C2W5|GABRB2_Bos_taurus                   IPLNLTLDNRVADQLWVPDTYFLNDKKSFVHGVTVKNRMIRLHPDGTVLYGLRITTTTACMMDLRRYPLDEQNCTLEIESYGYTTDDIEFYWRGDDNAVT   198
                                           .......110.......120.......130.......140.......150.......160.......170.......180.......190.......200

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P47870|GABRB2_Homo_sapiens                 GVTKIELPQFSIVDYKLITKKVVFSTGSYPRLSLSFKLKRNIGYFILQTYMPSILITILSWVSFWINYDASAARVALGITTVLTMTTINTHLRETLPKIP   300
A0A2R9C8C8|GABRB2_Pan_paniscus             GVTKIELPQFSIVDYKLITKKVVFSTGSYPRLSLSFKLKRNIGYFILQTYMPSILITILSWVSFWINYDASAARVALGITTVLTMTTINTHLRETLPKIP   300
A0A2J8NKG3|GABRB2_Pan_troglodytes          GVTKIELPQFSIVDYKLITKKVVFSTGSYPRLSLSFKLKRNIGYFILQTYMPSILITILSWVSFWINYDASAARVALGITTVLTMTTINTHLRETLPKIP   300
M3YP67|GABRB2_Mustela_putorius_furo        GVTKIELPQFSIVDYKLITKKVVFSTGSYPRLSLSFKLKRNIGYFILQTYMPSILITILSWVSFWINYDASAARVALGITTVLTMTTINTHLRETLPKIP   300
P63138|GABRB2_Rattus_norvegicus            GVTKIELPQFSIVDYKLITKKVVFSTGSYPRLSLSFKLKRNIGYFILQTYMPSILITILSWVSFWINYDASAARVALGITTVLTMTTINTHLRETLPKIP   300
P63137|GABRB2_Mus_musculus                 GVTKIELPQFSIVDYKLITKKVVFSTGSYPRLSLSFKLKRNIGYFILQTYMPSILITILSWVSFWINYDASAARVALGITTVLTMTTINTHLRETLPKIP   300
A0A5F4CKV8|GABRB2_Canis_lupus_familiaris   GVTKIELPQFSIVDYKLITKKVVFSTGSYPRLSLSFKLKRNIGYFILQTYMPSILITILSWVSFWINYDASAARVALGITTVLTMTTINTHLRETLPKIP   300
D1LYT2|GABRB2_Macaca_mulatta               GVTKIELPQFSIVDYKLITKKVVFSTGSYPRLSLSFKLKRNIGYFILQTYMPSILITILSWVSFWINYDASAARVALGITTVLTMTTINTHLRETLPKIP   300
Q9DDD9|GABRB2_Danio_rerio                  GVERIELPQFSIVGYKLISKNVVFSTGSYPRLSLSFKLKRNIGYFILQTYMPSILITILSWVSFWINYDASAARVALGITTVLTMTTINTHLRETLPKIP   299
P0C2W5|GABRB2_Bos_taurus                   GVTKIELPQFSIVDYKLITKKVVFSTGSYPRLSLSFKLKRNIGYFILQTYMPSILITILSWVSFWINYDASAARVALGITTVLTMTTINTHLRETLPKIP   298
                                           .......210.......220.......230.......240.......250.......260.......270.......280.......290.......300

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P47870|GABRB2_Homo_sapiens                 YVKAIDMYLMGCFVFVFMALLEYALVNYIFFGRGPQRQKKAAEKAASANNEKMRLDVNK-----------IFYKDIKQNGTQ-YRSLWDPTGNLSPTRRT   388
A0A2R9C8C8|GABRB2_Pan_paniscus             YVKAIDMYLMGCFVFVFMALLEYALVNYIFFGRGPQRQKKAAEKAASANNEKMRLDVNK-----------IFYKDIKQNGTQ-YRSLWDPTGNLSPTRRT   388
A0A2J8NKG3|GABRB2_Pan_troglodytes          YVKAIDMYLMGCFVFVFMALLEYALVNYIFFGRGPQRQKKAAEKAASANNEKMRLDVNK-----------IFYKDIKQNGTQ-YRSLWDPTGNLSPTRRT   388
M3YP67|GABRB2_Mustela_putorius_furo        YVKAIDMYLMGCFVFVFMALLEYALVNYIFFGRGPQRQKKAAEKAASANNEKMRLDVNK-----------IFYKDFKQNGTQ-YRSLWDPTGNLSPTRRT   388
P63138|GABRB2_Rattus_norvegicus            YVKAIDMYLMGCFVFVFMALLEYALVNYIFFGRGPQRQKKAAEKAANANNEKMRLDVNK-----------------------------------------   359
P63137|GABRB2_Mus_musculus                 YVKAIDMYLMGCFVFVFMALLEYALVNYIFFGRGPQRQKKAAEKAANANNEKMRLDVNK-----------MFYKDIKQNGTQ-YRSLWDPTGDLSPTRRT   388
A0A5F4CKV8|GABRB2_Canis_lupus_familiaris   YVKAIDMYLMGCFVFVFMALLEYALVNYIFFGRGPQRQKKAAEKAASANNEKMRLDVNK-----------IFYKDIKQNGTQ-YRSLWDPTGNLSPTRRT   388
D1LYT2|GABRB2_Macaca_mulatta               YVKAIDMYLMGCFVFVFMALLEYALVNYIFFGRGPQRQKKAAEKAASANNEKMRLDVNK-----------IFYKDIKQNGTQ-YRSLWDPTGNLSPTRRT   388
Q9DDD9|GABRB2_Danio_rerio                  YVKAIDMYLMGCFVFVFLALLEYAFVNYIFFGRGPQRQKKAAEKAATANNEKLRPDPNKWLVGSVVQRDDALYARMKQREIDGYDSMWDP---IFAEDAA   396
P0C2W5|GABRB2_Bos_taurus                   YVKAIDMYLMGCFVFVFMALLEYALVNYIFFGRGPQRQKKAAEKAASANNEKMRLDVNK-----------------------------------------   357
                                           .......310.......320.......330.......340.......350.......360.......370.......380.......390.......400

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P47870|GABRB2_Homo_sapiens                 TNYDFSLYTMDPHENILLSTLEIKNEMATSEAVMGLGDPRSTMLAYDASSIQYRKAGLPRHSFGRNALERHVAQKKSRLRRRASQLKITIPDLTDVNAID   488
A0A2R9C8C8|GABRB2_Pan_paniscus             TNYDFSLYTMDPHENILLSTLEIKNEMATSEAVMGLGDPRSTMLAYDASSIQYRKAGLPRHSFGRNALERHVAQKKSRLRRRASQLKITIPDLTDVNAID   488
A0A2J8NKG3|GABRB2_Pan_troglodytes          TNYDFSLYTMDPHENILLSTLEIKNEMATSEAVMGLGDPRSTMLAYDASSIQYRKAGLPRHSFGRNALERHVAQKKSRLRRRASQLKITIPDLTDVNAID   488
M3YP67|GABRB2_Mustela_putorius_furo        TNYDFSLYTMDPHENILLSTLEIKNEMATSEAVMGLGDPRSTMLAYDATSIQYRKAGLPRHSFGRNALERHVAQKKSRLRRRASQLKITIPDLTDVNAID   488
P63138|GABRB2_Rattus_norvegicus            ---------MDPHENILLSTLEIKNEMATSEAVMGLGDPRSTMLAYDASSIQYRKAGLPRHSFGRNALERHVAQKKSRLRRRASQLKITIPDLTDVNAID   450
P63137|GABRB2_Mus_musculus                 TNYDFSLYTMDPHENILLSTLEIKNEMATSEAVMGLGDPRSTMLAYDASSIQYRKAGLPRHSFGRNALERHVAQKKSRLRRRASQLKITIPDLTDVNAID   488
A0A5F4CKV8|GABRB2_Canis_lupus_familiaris   TNYDFSLYTMDPHENILLSTLEIKNEMATSEAVMGLGDPRSTMLAYDATSIQYRKAGLPRHSFGRNALERHVAQKKSRLRRRASQLKITIPDLTDVNAID   488
D1LYT2|GABRB2_Macaca_mulatta               TNYDFSLYTMDPHENILLSTLEIKNEMATSEAVMGLGDPRSTMLAYDASSIQYRKAGLPRHSFGRNALERHVAQKKSRLRRRASQLKITIPDLTDVNAID   488
Q9DDD9|GABRB2_Danio_rerio                  LGLGDQRLKMTP-DDIRLTTVEMKNEMGPSDLSKGLGDPRSTMLAYDSSTLQYRRAAMARQNYGHSALERHATQKKSRLRRRASQLKVNIPDLSDVNSID   495
P0C2W5|GABRB2_Bos_taurus                   ---------MDPHENILLSTLEIKNEMATSEAVMGLGDPRSTMLAYDASSIQYRKAGLPRHSFWRNALERHVAQKKSRLRERASQLKITIPDLTDVNAID   448
                                           .......410.......420.......430.......440.......450.......460.......470.......480.......490.......500
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P47870|GABRB2_Homo_sapiens                       RWSRIFFPVVFSFFNIVYWLYYVN   512
A0A2R9C8C8|GABRB2_Pan_paniscus                   RWSRIFFPVVFSFFNIVYWLYYVN   512
A0A2J8NKG3|GABRB2_Pan_troglodytes                RWSRIFFPVVFSFFNIVYWLYYVN   512
M3YP67|GABRB2_Mustela_putorius_furo              RWSRIFFPVVFSFFNIVYWLYYVN   512
P63138|GABRB2_Rattus_norvegicus                  RWSRIFFPVVFSFFNIVYWLYYVN   474
P63137|GABRB2_Mus_musculus                       RWSRIFFPVVFSFFNIVYWLYYVN   512
A0A5F4CKV8|GABRB2_Canis_lupus_familiaris         RWSRIFFPVVFSFFNIVYWLYYVN   512
D1LYT2|GABRB2_Macaca_mulatta                     RWSRIFFPVVFSFFNIVYWLYYVN   512
Q9DDD9|GABRB2_Danio_rerio                        KWSRMIFPTLFSFFNIVYWLYYVH   519
P0C2W5|GABRB2_Bos_taurus                         RWSRIFFPVVFSFFNIVYWLYYVN   472
                                                 .......510.......520....
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P28472|GABRB3_Homo_sapiens           MWGLAGGRLFGIFSAPVLVAVVCCAQSVNDPGNMSFVKETVDKLLKGYDIRLRPDFGGPPVCVGMNIDIASIDMVSEVNMDYTLTMYFQQYWRDKRLAYS   100
A0A2R9BWE5|GABRB3_Pan_paniscus       ----------------------------------------------------------SP-------------------QDYTLTMYFQQYWRDKRLAYS   23
A0A2I3THQ1|GABRB3_Pan_troglodytes    MWGLAGGRLFGIFSAPVLVAVVCCAQSVNDPGNMSFVKETVDKLLKGYDIRLRPDFGGPPVCVGMNIDIASIDMVSEVNMDYTLTMYFQQYWRDKRLAYS   100
F6ZKJ4|GABRB3_Macaca_mulatta         MCSGLLALLLPIWLSWTLGTRGSEPRSVNDPGNMSFVKETVDKLLKGYDIRLRPDFGGPPVCVGMNIDIASIDMVSEVNMDYTLTMYFQQYWRDKRLAYS   100
M3XNI6|GABRB3_Mustela_putorius_furo  MWGFAGGRLFGIFSAPVLVAVVCCAQSVNDPGNMSFVKETVDKLLKGYDIRLRPDFGGPPVCVGMNIDIASIDMVSEVNMDYTLTMYFQQYWRDKRLAYS   100
P63079|GABRB3_Rattus_norvegicus      MWGFAGGRLFGIFSAPVLVAVVCCAQSVNDPGNMSFVKETVDKLLKGYDIRLRPDFGGPPVCVGMNIDIASIDMVSEVNMDYTLTMYFQQYWRDKRLAYS   100
P63080|GABRB3_Mus_musculus           MWGFAGGRLFGIFSAPVLVAVVCCAQSVNDPGNMSFVKETVDKLLKGYDIRLRPDFGGPPVCVGMNIDIASIDMVSEVNMDYTLTMYFQQYWRDKRLAYS   100
J9P3X1|GABRB3_Canis_lupus_familiaris ----------------------------------------------------------PPVCVGMNIDIASIDMVSEVNMDYTLTMYFQQYWRDKRLAYS   42
A0A0R4ILP2|GABRB3_Danio_rerio        MFGIQKRRLFGVLSLPVIVAVMCCAQSANEPGNMSFVKETVDKLLKGYDIRLRPDFGGAPVAVGMSIDVASIDMVSEVNMDYTLTMYFQQYWRDKRLAYT   100
A5D7U6|GABRB3_Bos_taurus             MWGFAGGRFFGIFSAPVLVAVVCCAQSVNDPGNMSFVKETVDKLLKGYDIRLRPDFGGPPVCVGMNIDIASIDMVSEVNMDYTLTMYFQQYWRDKRLAYS   100
                                     1.......10........20........30........40........50........60........70........80........90.......100

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P28472|GABRB3_Homo_sapiens           GIPLNLTLDNRVADQLWVPDTYFLNDKKSFVHGVTVKNRMIRLHPDGTVLYGLRITTTAACMMDLRRYPLDEQNCTLEIESYGYTTDDIEFYWRGGDKAV   200
A0A2R9BWE5|GABRB3_Pan_paniscus       GIPLNLTLDNRVADQLWVPDTYFLNDKKSFVHGVTVKNRMIRLHPDGTVLYGLRITTTAACMMDLRRYPLDEQNCTLEIESYGYTTDDIEFYWRGGDKAV   123
A0A2I3THQ1|GABRB3_Pan_troglodytes    GIPLNLTLDNRVADQLWVPDTYFLNDKKSFVHGVTVKNRMIRLHPDGTVLYGLRITTTAACMMDLRRYPLDEQNCTLEIESYGYTTDDIEFYWRGGDKAV   200
F6ZKJ4|GABRB3_Macaca_mulatta         GIPLNLTLDNRVADQLWVPDTYFLNDKKSFVHGVTVKNRMIRLHPDGTVLYGLRITTTAACMMDLRRYPLDEQNCTLEIESYGYTTDDIEFYWRGGDKAV   200
M3XNI6|GABRB3_Mustela_putorius_furo  GIPLNLTLDNRVADQLWVPDTYFLNDKKSFVHGVTVKNRMIRLHPDGTVLYGLRITTTAACMMDLRRYPLDEQNCTLEIESYGYTTDDIEFYWRGGDKAV   200
P63079|GABRB3_Rattus_norvegicus      GIPLNLTLDNRVADQLWVPDTYFLNDKKSFVHGVTVKNRMIRLHPDGTVLYGLRITTTAACMMDLRRYPLDEQNCTLEIESYGYTTDDIEFYWRGGDKAV   200
P63080|GABRB3_Mus_musculus           GIPLNLTLDNRVADQLWVPDTYFLNDKKSFVHGVTVKNRMIRLHPDGTVLYGLRITTTAACMMDLRRYPLDEQNCTLEIESYGYTTDDIEFYWRGGDKAV   200
J9P3X1|GABRB3_Canis_lupus_familiaris GIPLNLTLDNRVADQLWVPDTYFLNDKKSFVHGVTVKNRMIRLHPDGTVLYGLRITTTAACMMDLRRYPLDEQNCTLEIESYGYTTDDIEFYWRGGDKAV   142
A0A0R4ILP2|GABRB3_Danio_rerio        GIPLNLTLDNRVADQLWVPDTYFLNDKKSFVHGVTVKNRMIRLHPDGTVLYGLRITTTAACMMDLRRYPLDEQNCTLEIESYGYTTDDIEFYWKGGETAV   200
A5D7U6|GABRB3_Bos_taurus             GIPLNLTLDNRVADQLWVPDTYFLNDKKSFVHGVTVKNRMIRLHPDGTVLYGLRITTTAACMMDLRRYPLDEQNCTLEIESYGYTTDDIEFYWRGGDKAV   200
                                     .......110.......120.......130.......140.......150.......160.......170.......180.......190.......200

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P28472|GABRB3_Homo_sapiens           TGVERIELPQFSIVEHRLVSRNVVFATGAYPRLSLSFRLKRNIGYFILQTYMPSILITILSWVSFWINYDASAARVALGITTVLTMTTINTHLRETLPKI   300
A0A2R9BWE5|GABRB3_Pan_paniscus       TGVERIELPQFSIVEHRLVSRNVVFATGAYPRLSLSFRLKRNIGYFILQTYMPSILITILSWVSFWINYDASAARVALGITTVLTMTTINTHLRETLPKI   223
A0A2I3THQ1|GABRB3_Pan_troglodytes    TGVERIELPQFSIVEHRLVSRNVVFATGAYPRLSLSFRLKRNIGYFILQTYMPSILITILSWVSFWINYDASAARVALGITTVLTMTTINTHLRETLPKI   300
F6ZKJ4|GABRB3_Macaca_mulatta         TGVERIELPQFSIVEHRLVSRNVVFATGAYPRLSLSFRLKRNIGYFILQTYMPSILITILSWVSFWINYDASAARVALGITTVLTMTTINTHLRETLPKI   300
M3XNI6|GABRB3_Mustela_putorius_furo  TGVERIELPQFSIVEHRLVSRNVVFATGAYPRLSLSFRLKRNIGYFILQTYMPSILITILSWVSFWINYDASAARVALGITTVLTMTTINTHLRETLPKI   300
P63079|GABRB3_Rattus_norvegicus      TGVERIELPQFSIVEHRLVSRNVVFATGAYPRLSLSFRLKRNIGYFILQTYMPSILITILSWVSFWINYDASAARVALGITTVLTMTTINTHLRETLPKI   300
P63080|GABRB3_Mus_musculus           TGVERIELPQFSIVEHRLVSRNVVFATGAYPRLSLSFRLKRNIGYFILQTYMPSILITILSWVSFWINYDASAARVALGITTVLTMTTINTHLRETLPKI   300
J9P3X1|GABRB3_Canis_lupus_familiaris TGVERIELPQFSIVEHRLVSRNVVFATGAYPRLSLSFRLKRNIGYFILQTYMPSILITILSWVSFWINYDASAARVALGITTVLTMTTINTHLRETLPKI   242
A0A0R4ILP2|GABRB3_Danio_rerio        TGVSRIELPQFSIVDYKLVSRNVVFSTGAYPRLSLSFKLKRNIGYFILQTYMPSILITILSWVSFWINYDASAARVALGITTVLTMTTINTHLRETLPKI   300
A5D7U6|GABRB3_Bos_taurus             TGVERIELPQFSIVEHRLVSRNVVFATGAYPRLSLSFRLKRNIGYFILQTYMPSILITILSWVSFWINYDASAARVALGITTVLTMTTINTHLRETLPKI   300
                                     .......210.......220.......230.......240.......250.......260.......270.......280.......290.......300

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P28472|GABRB3_Homo_sapiens           PYVKAIDMYLMGCFVFVFLALLEYAFVNYIFFGRGPQRQKKLAEKTAKAKNDRSKSESNR---------------------------VDAHGNILLTSLE   373
A0A2R9BWE5|GABRB3_Pan_paniscus       PYVKAIDMYLMGCFVFVFLALLEYAFVNYIFFGRGPQRQKKLAEKTAKAKNDRSKSESNR---------------------------VDAHGNILLTSLE   296
A0A2I3THQ1|GABRB3_Pan_troglodytes    PYVKAIDMYLMGCFVFVFLALLEYAFVNYIFFGRGPQRQKKLAEKTAKAKNDRSKSESNR---------------------------VDAHGNILLTSLE   373
F6ZKJ4|GABRB3_Macaca_mulatta         PYVKAIDMYLMGCFVFVFLALLEYAFVNYIFFGRGPQRQKKLAEKTAKAKNDRSKSESNR---------------------------VDAHGNILLTSLE   373
M3XNI6|GABRB3_Mustela_putorius_furo  PYVKAIDMYLMGCFVFVFLALLEYAFVNYIFFGRGPQRQKKLAEKTAKAKNDRSKSESNR---------------------------VDAHGSILLTSLE   373
P63079|GABRB3_Rattus_norvegicus      PYVKAIDMYLMGCFVFVFLALLEYAFVNYIFFGRGPQRQKKLAEKTAKAKNDRSKSEINR---------------------------VDAHGNILLAPMD   373
P63080|GABRB3_Mus_musculus           PYVKAIDMYLMGCFVFVFLALLEYAFVNYIFFGRGPQRQKKLAEKTAKAKNDRSKSEINR---------------------------VDAHGNILLAPMD   373
J9P3X1|GABRB3_Canis_lupus_familiaris PYVKAIDMYLMGCFVFVFLALLEYAFVNYIFFGRGPQRQKKLAEKTAKAKNDRSKSESNR---------------------------VDAHGNILLTSLE   315
A0A0R4ILP2|GABRB3_Danio_rerio        PYVKAIDMYLMGCFVFVFLALLEYAFVNYIFFGRGPQMQKKLAEKAEKANNDRNKYDGNKSVQEGGNCKPSSVKQSNQVQGHRHSRGVDSQGNILLTTLE   400
A5D7U6|GABRB3_Bos_taurus             PYVKAIDMYLMGCFVFVFLALLEYAFVNYIFFGRGPQRQKKLAEKNAKAKNDRSKGDSNR---------------------------VDAHGNILLTSLE   373
                                     .......310.......320.......330.......340.......350.......360.......370.......380.......390.......400

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P28472|GABRB3_Homo_sapiens           VHNEM--NEVSGGIGDTRNS-AISFDNSGIQYRKQSMPREGHGRFLGDRSLPHKKTHLRRRSSQLKIKIPDLTDVNAIDRWSRIVFPFTFSLFNLVYWLY   470
A0A2R9BWE5|GABRB3_Pan_paniscus       VHNEM--NEVAGGIGDTRNS-AISFDNSGIQYRKQSMPREGHGRFLGDRSLPHKKTHLRRRSSQLKIKIPDLTDVNAIDRWSRIVFPFTFSLFNLVYWLY   393
A0A2I3THQ1|GABRB3_Pan_troglodytes    VHNEM--NEVAGGIGDTRNS-AISFDNSGIQYRKQSMPREGHGRFLGDRSLPHKKTHLRRRSSQLKIKIPDLTDVNAIDRWSRIVFPFTFSLFNLVYWLY   470
F6ZKJ4|GABRB3_Macaca_mulatta         VHNEM--NEVAGGVGDTRNS-AISFDNSGIQYRKQSMPREGHGRFLGDRSLPHKKTHLRRRSSQLKIKIPDLTDVNAIDRWSRIVFPFTFSLFNLVYWLY   470
M3XNI6|GABRB3_Mustela_putorius_furo  VHNEM--NEVTGGVGDTRNS-ARPFDNSGIQYRKQSMPREGHGRHMGDRSIPHKKTHLRRRSSQLKIKIPDLTDVNAIDRWSRIVFPFTFSLFNLVYWLY   470
P63079|GABRB3_Rattus_norvegicus      VHNEM--NEVAGSVGDTRNS-AISFDNSGIQYRKQSMPKEGHGRYMGDRSIPHKKTHLRRRSSQLKIKIPDLTDVNAIDRWSRIVFPFTFSLFNLVYWLY   470
P63080|GABRB3_Mus_musculus           VHNEM--NEVAGSVGDTRNS-AISFDNSGIQYRKQSMPKEGHGRYMGDRSIPHKKTHLRRRSSQLKIKIPDLTDVNAIDRWSRIVFPFTFSLFNLVYWLY   470
J9P3X1|GABRB3_Canis_lupus_familiaris VHNEM--NEVTGSVGDTRNS-ARPFDNSGIQYRKQSMPREGHGRHMGDRSIPHKKTHLRRRSSQLKIKIPDLTDVNAIDRWSRIVFPFTFSLFNLVYWLY   412
A0A0R4ILP2|GABRB3_Danio_rerio        IHNEVAGNEITTSVSEARNSTSMVFDNSGIQYRKQST-----ARHSMDRNAQSKKPRLRRRSSQLKIKIPDLTDVNAIDRWSRIIFPAVFSLFNLIYWLY   495
A5D7U6|GABRB3_Bos_taurus             VHNEM--NEVAGGVGDTRNS-ARSFDNSGIQYRKQSMPREGHGRHM-DRSVPHKKTHLRRRSSQLKIKIPDLTDVNAIDRWSRIVFPFTFSLFNLVYWLY   469
                                     .......410.......420.......430.......440.......450.......460.......470.......480.......490.......500
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                                             ***
P28472|GABRB3_Homo_sapiens                   YVN    473
A0A2R9BWE5|GABRB3_Pan_paniscus               YVN    396
A0A2I3THQ1|GABRB3_Pan_troglodytes            YVN    473
F6ZKJ4|GABRB3_Macaca_mulatta                 YVN    473
M3XNI6|GABRB3_Mustela_putorius_furo          YVN    473
P63079|GABRB3_Rattus_norvegicus              YVN    473
P63080|GABRB3_Mus_musculus                   YVN    473
J9P3X1|GABRB3_Canis_lupus_familiaris         YVN    415
A0A0R4ILP2|GABRB3_Danio_rerio                YVN    498
A5D7U6|GABRB3_Bos_taurus                     YVN    472
                                             ...
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                                                                       :. * :   :: *    *.:: *: ** : :** :*. .*.*.::::**** *** **:**
O14764|GABRD_Homo_sapiens                  -----------MDAPARLL--APLLLLCAQQLRGTRAMNDIGDYVGSNLEISWLPNLDGLIAGYARNFRPGIGGPPVNVALALEVASIDHISEANMEYTM   87
A0A2R9B726|GABRD_Pan_paniscus              MSEATPLDRNDSENTGGLISRPHPWDQSPSCVQEDRAMNDIGDYVGSNLEISWLPNLDGLIAGYARNFRPGIGGPPVNVALALEVASIDHISEANMEYTM  100
A0A2J8K8J4|GABRD_Pan_troglodytes           -----------MDAPARLL--APLLLLCAQQLRGTRAMNDIGNYVGSNLEISWLPNLDGLIAGYARNFRPGIGGPPVNVALALEVASIDHISEANMEYTM   87
F6QDC4|GABRD_Macaca_mulatta                -------------------------------------MNDIGDYVGSNLEISWLPNLDGLIAGYARNFRPGIGGPPVNVALALEVASIDHISEANMEYTM   63
XP_004768807.1|GABRD_Mustela_putorius_furo -----------MDTLIGLL--PPLLLLCAQQPRGARAMNDIGDYVGSNLEISWLPNLDGLMEGYARNFRPGIGGPPVNVALAIEVASIDHISEANMEYTM   87
P18506|GABRD_Rattus_norvegicus             -----------MDVLGWLL--LPLLLLCTQPHHGARAMNDIGDYVGSNLEISWLPNLDGLMEGYARNFRPGIGGPPVNVALALEVASIDHISEANMEYTM   87
P22933|GABRD_Mus_musculus                  -----------MDVLGWLL--LPLLLLCTQPHHGARAMNDIGDYVGSNLEISWLPNLDGLMEGYARNFRPGIGGAPVNVALALEVASIDHISEANMEYTM   87
E2R3M6|GABRD_Canis_lupus_familiaris        -----------MDALAWPL--PLLLLLCAQQPRGTRAMNDIGDYVGSNLEISWLPNLDGLMDGYARNFRPGIGGPPVNVALAIEVASIDHISEVNMEYTM   87
E9QHL0|GABRD_Danio_rerio                   -----------XRAM----------------------LSDIGDYVGTDIEISWLPNLDDLMKGYARNFRPGIGGPPVNVAMAIEVASIDHISEANMEYTM   67
A0A3Q1LQH4|GABRD_Bos_taurus                -----------MDELAWLL--PPLLLLCAQ-HRGARAMNDIGDYVGSNLEISWLPNLDGLIEGYARNFRPGIGGPAVNVALAIEVASIDHISEVNMEYTM   86
F1QZB5|GABRZ_Danio_rerio                   -----------MHVCSFVG--FLLLLLF----------ISRIMDGSVANTEEILPPTIQKLMKGYNKYLRPFFGNGPVTVGMSLDIASIDTISEINMDYTA   78
                                           1       10        20        30        40        50        60        70        80        90       100

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O14764|GABRD_Homo_sapiens                  TVFLHQSWRDSRLSYNHTNETLGLDSRFVDKLWLPDTFIVNAKSAWFHDVTVENKLIRLQPDGVILYSIRITSTVACDMDLAKYPMDEQECMLDLESYGY  187
A0A2R9B726|GABRD_Pan_paniscus              TVFLHQSWRDSRLSYNHTNETLGLDSRFVDKLWLPDTFIVNAKSAWFHDVTVENKLIRLQPDGVILYSIRITSTVACDMDLAKYPMDEQECMLDLESYGY  200
A0A2J8K8J4|GABRD_Pan_troglodytes           TVFLHQSWRDSRLSYNHTNETLGLDSRFVDKLWLPDTFIVNAKSAWFHDVTVENKLIRLQPDGVILYSIRITSTVACDMDLAKYPMDEQECMLDLESYGY  187
F6QDC4|GABRD_Macaca_mulatta                TVFLHQSWRDSRLSYNHTNETLGLDSRFVDKLWLPDTFIVNAKSAWFHDVTVENKLIRLQPDGVILYSIRITSTVACDMDLAKYPMDEQECMLELESYGY  163
XP_004768807.1|GABRD_Mustela_putorius_furo TVFLHQSWRDSRLAYNHTNETLGLDSRFVDKLWLPDTFIVNAKSAWFHDVTVENKLIRLQPDGVILYSIRITSTVACDMDLAKYPMDEQECMLLLESYGY  187
P18506|GABRD_Rattus_norvegicus             TVFLHQSWRDSRLSYNHTNETLGLDSRFVDKLWLPDTFIVNAKSAWFHDVTVENKLIRLQPDGVILYSIRITSTVACDMDLAKYPMDEQECMLDLESYGY  187
P22933|GABRD_Mus_musculus                  TVFLHQSWRDSRLSYNHTNETLGLDSRFVDKLWLPDTFIVNAKSAWFHDVTVENKLIRLQPDGVILYSIRITSTVACDMDLAKYPLDEQECMLDLESYGY  187
E2R3M6|GABRD_Canis_lupus_familiaris        TVFLHQSWRDSRLAYNHTNETLGLDSRFVDKLWLPDTFIVNAKSAWFHDVTVENKLIRLQPDGVILYSIRITSTVACDMDLAKYPMDEQECMLHLESYGY  187
E9QHL0|GABRD_Danio_rerio                   TVFLRQSWRDDRLSYNHTNKTLGLDSRFVDKLWLPDTFIVNAKSAWFHDVTVENKLIRLQPDGVILYSSRITSTVACDMDLTKYPMDEQECMLDLESYGY  167
A0A3Q1LQH4|GABRD_Bos_taurus                TVFLHQSWRDSRLSYNHTNETLGLDSRFVDKLWLPDTFIVNAKSAWFHDVTVENKLIRLQPDGVILYSIRITSTVACDMDLAKYPMDEQECMLHLESYGY  186
F1QZB5|GABRZ_Danio_rerio                   TIFLRQRWTDERLCFD-GNKSLSLDGRLVELLWVPDTFIVDSKKSFLHDITVENRLIRIFPNGTVLYALRITTTVACSMDLTKYPMDRQTCMLQLESWGY  177
                                           .......110.......120.......130.......140.......150.......160.......170.......180.......190.......200

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O14764|GABRD_Homo_sapiens                  SSEDIVYYWSESQEHIHGLDKLQLAQFTITSYRFTTELMNFKSAGQFPRLSLHFHLRRNRGVYIIQSYMPSVLLVAMSWVSFWISQAAVPARVSLGITTV  287
A0A2R9B726|GABRD_Pan_paniscus              SSEDIVYYWSESQEHIHGLDKLQLAQFTITSYRFTTELMNFKSAGQFPRLSLHFHLRRNRGVYIIQSYMPSVLLVAMSWVSFWISQAAVPARVSLGITTV  300
A0A2J8K8J4|GABRD_Pan_troglodytes           SSEDIVYYWSESQEHIHGLDKLQLAQFTITSYRFTTELMNFKSAGQFPRLSLHFHLRRNRGVYIIQSYMPSVLLVAMSWVSFWISQAAVPARVSLGITTV  287
F6QDC4|GABRD_Macaca_mulatta                SSEDIVYYWSESQEHIHGLDKLQLAQFTITSYRFTTELMNFKSAGQFPRLSLHFHLRRNRGVYIIQSYMPSVLLVAMSWVSFWISQAAVPARVSLGITTV  263
XP_004768807.1|GABRD_Mustela_putorius_furo SSEDVVYYWSENQERIHGLNELHLAQFTITSHRFTAELRNFKSAGQFPRLSLHFRLRRNRGVYIIQSYMPSVLLVAMSWVSFWISQAAVPARVSLGITTV  287
P18506|GABRD_Rattus_norvegicus             SSEDIVYYWSENQEQIHGLDRLQLAQFTITSYRFTTELMNFKSAGQFPRLSLHFQLRRNRGVYIIQSYMPSVLLVAMSWVSFWISQAAVPARVSLGITTV  287
P22933|GABRD_Mus_musculus                  SSEDIVYYWSENQEQIHGLDRLQLAQFTITSYRFTTELMNFKSAGQFPRLSLHFQLRRNRGVYIIQSYMPSVLLVAMSWVSFWISQAAVPARVSLGITTV  287
E2R3M6|GABRD_Canis_lupus_familiaris        SSEDIVYYWSESQEHIHGLNKLQLAQFTITSYHFATELMNFKSAGQFPRLSLHFHLRRNRGVYIIQSYMPSILLVAMSWVSFWISQAAVPARVSLGITTV  287
E9QHL0|GABRD_Danio_rerio                   SSEDIVYHWSESQKLIHGLDKLELSQFTITDYRFVTEMMNFKSAGRFPRLSLRFQLRRNRGVYIIQSYMPSILLVAMSWVSFWISQSAVPARVSLGITTV  267
A0A3Q1LQH4|GABRD_Bos_taurus                SSEDIVYYWSENQEQIHGLDKLQLAQFTITSYRFTTELMNFKSAGQFPRLSLHFHLRRNRGVYIIQSYMPSVLLVAMSWVSFWISQAAVPARVSLGITTV  286
F1QZB5|GABRZ_Danio_rerio                   NVKDVVFYWTRGNQSVSGLDHLQLAQYTLEDH-YTSESEAVYETGNYPKLIFHFKLKRSILYFILETYVPSSALVVLSWVSFWISQSSVPARICIGVTTV  276
                                           .......210.......220.......230.......240.......250.......260.......270.......280.......290.......300

                                           *******:.**:***.*.. ***:***: **: *:*.**:***.**: : :  :                 .   .    : .  :. :.  .     :   :  *
O14764|GABRD_Homo_sapiens                  LTMTTLMVSARSSLPRASA-IKALDVYFWICYVFVFAALVEYAFAHFNADYRKKQ----KAKVKVSRPRAEM--DVRNAIVLFSLSAAGVTQELAISRRQ  380
A0A2R9B726|GABRD_Pan_paniscus              LTMTTLMVSARSSLPRASA-IKALDVYFWICYVFVFAALVEYAFAHFNADYRKKQ----KAKVKVSRPRAEM--DVRNAIVLFSLSAAGVTQELAISRRQ  393
A0A2J8K8J4|GABRD_Pan_troglodytes           LTMTTLMVSARSSLPRASA-IKALDVYFWICYVFVFAALVEYAFAHFNADYRKKQ----KAKVKVSRPRAEM--DVRNAIVLFSLSAAGVTQELAISRRQ  380
F6QDC4|GABRD_Macaca_mulatta                LTMTTLMVSARSSLPRASA-IKALDVYFWICYVFVFAALVEYAFAHFNADYRKKQKA--KAKVKVSRPRAEM--DVRNAIVLFSLSAAGVTQELAISRRQ  358
XP_004768807.1|GABRD_Mustela_putorius_furo LTMTTLMVSARSSLPRASA-IKALDVYFWICYVFVFAALVEYAFAHFNADYRKKQ----KAKVKVKEQKAEL--DVKNAIVLLSLSAAGATQELAVSRRH  380
P18506|GABRD_Rattus_norvegicus             LTMTTLMVSARSSLPRASA-IKALDVYFWICYVFVFAALVEYAFAHFNADYRKKR----KAKVKVTKPRAEM--DVRNAIVLFSLSAAGVSQELAISRRQ  380
P22933|GABRD_Mus_musculus                  LTMTTLMVSARSSLPRASA-IKALDVYFWICYVFVFAALVEYAFAHFNADYRKKR----KAKVKVTKPRAEM--DVRNAIVLFSLSAAGVSQELAISRRQ  380
E2R3M6|GABRD_Canis_lupus_familiaris        LTMTTLMVSARSSLPRASA-IKALDVYFWICYVFVFAALVEYAFAHFNADYRKKQKAKAKAKVKVTEQRAEM--DVKNAIVLFSLSAAGVTQELAVSRRQ  384
E9QHL0|GABRD_Danio_rerio                   LTMTTLMVSARSSLPRASA-IKALDVYFWICYVFVFAALIEYAFAHYNADYSKKE----KAKVKTNKNAESMVKNGKQAMVLFSLSVAGMNQGLLISNRQ  362
A0A3Q1LQH4|GABRD_Bos_taurus                LTMTTLMVSARSSLPRASA-IKALDVYFWICYVFVFAALVEYAFAHFNADYRKKQKAKAKAKVKVKEQRAEM--DVKNAIVLFSLSAAGVTQELAVSRRP  383
F1QZB5|GABRZ_Danio_rerio                   LTMTTLMMGARTSLPNANCFIKAIDVYLGICFSFIFGALIEYAVAHFCTLHQPNA----ANAYMYGQEMQEREDEMNGIVTSFGSHALRARKREEMLSRM  372
                                           .......310.......320.......330.......340.......350.......360.......370.......380.......390.......400

                                                       .                                          . .     *                       .     .         * *   .
O14764|GABRD_Homo_sapiens                  RRVPGNLMGSYR---SVGVETGETK----------------------KEGAARSGGQGGIRAR-----------LRPIDADTIDIYARAVF-PAAFAAVN  443
A0A2R9B726|GABRD_Pan_paniscus              RRVPGNLMGSYR---SVGVETGETK----------------------KEGAARSGGQGGIRAR-----------LRPIDADTIDIYARAVF-PAAFAAVN  456
A0A2J8K8J4|GABRD_Pan_troglodytes           RRVPGNLMGSYR---SVGVETGETK----------------------KEGAARSGGQGGIRAR-----------LRPIDADTIDIYARAVF-PAAFAAVN  443
F6QDC4|GABRD_Macaca_mulatta                RRVPGNLMGSYR---SVGVETGEMK----------------------KEGAARSGGQGGIRAR-----------LRPIDADTIDIYARAVF-PAAFAAVN  421
XP_004768807.1|GABRD_Mustela_putorius_furo CRGAGNLMGSYR---CVEMETGQAE----------------------RPGAGRSGGRGGLRAL-----------FKPIDADTIDIYARVVF-PAAFVAVN  443
P18506|GABRD_Rattus_norvegicus             GRVPGNLMGSYR---SVEVE---AK----------------------KEGGSRPGGPGGIRSR-----------LKPIDADTIDIYARAVF-PAAFAAVN  440
P22933|GABRD_Mus_musculus                  GRVPGNLMGSYR---SVEVE---AK----------------------KEGGSRPGGPGGIRSR-----------LKPIDADTIDIYARAVF-PAAFAAVN  440
E2R3M6|GABRD_Canis_lupus_familiaris        CRRPGNLMGSYR---CVEVETGEAK----------------------KQRGARAGGQGGLRTL-----------FKPIDADTIDIYARVVF-PAAFVAVN  447
E9QHL0|GABRD_Danio_rerio                   SR-------SHR---ASDAAAAAAD----------------------PHEETETRSATSRRAQKESSEEKKCCKCKPIDADTIDIYARAVF-PFTFAVVN  429
A0A3Q1LQH4|GABRD_Bos_taurus                CRLPGNLMGSYSEGLSPKPSWGPSRWWFQWIHPHSAPCPHSRVFKLLTHSPGSPPSHPSFRELPQLYEVPPPSDGPPHSKVSTHINKLSPFQPYVHPAPI  483
F1QZB5|GABRZ_Danio_rerio                   GSVNNPTPSESK---EDVSSSQPQT----------------------RCTKCMSRARRIIRCL---------YCCKVENPHYIDNYSRLTF-PLSFVIIN  437
                                           .......410.......420.......430.......440.......450.......460.......470.......480.......490.......500
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                                                 *
O14764|GABRD_Homo_sapiens                      VIYWAAYAM-------------------  452
A0A2R9B726|GABRD_Pan_paniscus                  VIYWAAYAM-------------------  465
A0A2J8K8J4|GABRD_Pan_troglodytes               VIYWAAYAM-------------------  452
F6QDC4|GABRD_Macaca_mulatta                    VIYWAAYAM-------------------  430
XP_004768807.1|GABRD_Mustela_putorius_furo     VLYWAAYAM-------------------  452
P18506|GABRD_Rattus_norvegicus                 IIYWAAYTM-------------------  449
P22933|GABRD_Mus_musculus                      IIYWAAYTM-------------------  449
E2R3M6|GABRD_Canis_lupus_familiaris            VLYWAAYAM-------------------  456
E9QHL0|GABRD_Danio_rerio                       VIYWVAYTM-------------------  438
A0A3Q1LQH4|GABRD_Bos_taurus                    PTSWRATPTRPSMGKPGVVGSPRPCSGH  511
F1QZB5|GABRZ_Danio_rerio                       LLYWTYYLYF------------------  447
                                               .......510.......520........
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                                          *  .***  :*:.   :  *       *:  :  :.  ...  *   *:*   **         .   ::*  :.*     *         :    *::    *:.**:  *:  ***:****.*
P78334|GABRE_Homo_sapiens                 MLSKVLPVLLGILLILQSRVEGPQTESKNEASSRDV-VYGPQPQ--------PLENQLLSEETKSTETETGSRV-GKLPEASRILNTILSNYDHKLRPGI   90
A0A2R8ZSJ9|GABRE_Pan_paniscus             MLSKVLPVLLGILLILQSRVEGPQTESKNEASSRDV-VYGPQPQ--------PLENQLLSEETKSTETETGRRV-GKLPEASRILNTILSNYDHKLRPGI   90
H2QZ86|GABRE_Pan_troglodytes              MLSKVLPVLLGILLILQSRVEGPQTESKNEASSRDV-VYGPQPQ--------PLENQLLSEETKSTETETGRRV-GKLPEASRILNTILSNYDHKLRPGI   90
F7GJ80|GABRE_Macaca_mulatta               MLSKVLPVLLGILLILQSRVEGPQTESKNEASAHDV-VYGPPPQ--------PLENKLFSEETKSTKTETGRRA-GKLPEASRILNTILSNYDHKLRPGI   90
Q9ES14|GABRE_Rattus_norvegicus            MLPKVLLMLLNMFLALQWRV-GPHIKLENKPPAQDKVVFGPQPQ--------PSGKKLPARETELTADHTTERPRGKLTRASQILNTILSNYDHKLRPSI   91
Q9JLE8|GABRE_Mus_musculus                 MLPKVLLMLLNMFLALQWRV-GPHVNLENKPPAQDKVVFGPQPQPVPQPLTPPSGKKLPAKEPELTVDCMTERPGGKLTRASRILNTILSNYDHKLRPGI   99
E2QVM5|GABRE_Canis_lupus_familiaris       MLAKVLLILLGTSIILPSRSEGPHVESKSAPSARDD-VYGPQPQ--------APEEKLLSEETKSTM-----RL-GNLPAATHILNSILNNYDHKLRPGI   85
M3YJW6|GABRE_Mustela_putorius_furo        MLTKVLLILLGMSIILPSRSEEPRVELENGPSAHDD-VYGHQPQ--------APENKLLSEETVPTFTGPYHRL-GNLQAATHILNSILNNYDHKLRPGI   90
G3MWU9|GABRE_Bos_taurus                   MSTKVLLILMGTLVILPC-IEGPQVESEEGSSGSDDKVYGPKPQ--------PPEKELSSEEIKPTAVGTHKKL-GKMPTATEILDGIFHNYDYKLRPGI   90
                                          1.......10.......20.......30.......40.......50.......60.......70.......80.......90.......100

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P78334|GABRE_Homo_sapiens                 GEKPTVVTVEISVNSLGPLSILDMEYTIDIIFSQTWYDERLCYNDTFESLVLNGNVVSQLWIPDTFFRNSKRTHEHEITMPNQMVRIYKDGKVLYTIRMT   190
A0A2R8ZSJ9|GABRE_Pan_paniscus             GEKPTVVTVEISVNSLGPLSILDMEYTIDIIFSQTWYDERLSYNDTFESLVLNGNVVSQLWIPDTFFRNSKRTHEHEITMPNQMVRIYKDGKVLYTIRMT   190
H2QZ86|GABRE_Pan_troglodytes              GEKPTVVTVEISVNSLGPLSILDMEYTIDIIFSQTWYDERLSYNDTFESLVLNGNVVSQLWIPDTFFRNSKRTHEHEITMPNQMVRIYKDGKVLYTIRMT   190
F7GJ80|GABRE_Macaca_mulatta               GEKPTVVTVEISVNSLGPLSILDMEYTIDIIFSQTWYDERLCYNDTFESLVLNGNVVSQLWIPDTFFRNSKRTHEHEITMPNQMVRIYKDGKVLYTIRMT   190
Q9ES14|GABRE_Rattus_norvegicus            GEKPTVVTVKVFVNSLGPISILDMEYSIDIIFYQTWYDERLRYNDTFETLILHGNVVSQLWIPDTFFRNSKRTQEYDITIPNQMALIHKDGKVLYTVRMT   191
Q9JLE8|GABRE_Mus_musculus                 GEKPTVVTVEVIVNSLGPISTPDMEYSIDIIFHQTWYDERLRYNDTFETLILHGNVVSQLWIPDTFFRNSKRTQEYDITIPNQMALIHKDGKVLYSVRMT   199
E2QVM5|GABRE_Canis_lupus_familiaris       GERPTVVTVELSVNTLGPISIMDMEYTIDITFCQTWYDERLRYNGSFESFVLNGNMVSQLWIPDTVFRNSKRTQEHVITMPNQMVRIHKDGKVLYTIRMT   185
M3YJW6|GABRE_Mustela_putorius_furo        GEKPTVVTVELSVNTLGPISILDMEYTIDITFCQTWYDERLRYNGSFESFVLNGDMVSQLWIPDTFFRNSKRTQEHVITMPNQMARIHKDGKVLYTIRMT   190
G3MWU9|GABRE_Bos_taurus                   GERPTVVTVELSVNTLGPLSIMDMEYTIDITFCQTWYDERLRFNGSFESFVLSGNLVSLLWVPDTFFRNSKRTYEHSITMPNQMVRIHKDGKVLYTIRMT   190
                                          .......110.......120.......130.......140.......150.......160.......170.......180.......190.......200

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P78334|GABRE_Homo_sapiens                 IDAGCSLHMLRFPMDSHSCPLSFSSFSYPENEMIYKWENFKLEINEKNSWKLFQFDFTGVSNKTEIITTPVGDFMVMTIFFNVSRRFGYVAFQNYVPSSV   290
A0A2R8ZSJ9|GABRE_Pan_paniscus             IDAGCSLHMLRFPMDSHSCPLSFSSFSYPENEMIYKWENFKLEINEKNSWKLFQFDFTGVSNKTEIITTPVGDFMVMTIFFNVSRRFGYVAFQNYVPSSV   290
H2QZ86|GABRE_Pan_troglodytes              IDAGCSLHMLRFPMDSHSCPLSFSSFSYPENEMIYKWENFKLEINEKNSWKLFQFDFTGVSNKTEIITTPVGDFMVMTIFFNVSRRFGYVAFQNYVPSSV   290
F7GJ80|GABRE_Macaca_mulatta               IDAGCSLHMLRFPMDSHSCPLSFSSFSYPENEMIYKWENFKLEINEKNSWKLFQFDFTGVSNKTEIITTPVGDFMVMTIFFNVSRRFGYVAFQNYVPSSV   290
Q9ES14|GABRE_Rattus_norvegicus            IDARCSLHMLNFPMDSHSCPLSFSSFSYDEHEMIYKWENFKLKIDAKNTWKLLEFDFTGVNNKTEIISTPVGDFMVMTFFFNVSRRFGFIVFQNYIPSSV   291
Q9JLE8|GABRE_Mus_musculus                 IDARCSLHMLNFPMDSHSCPLSFSSFSYDDSEMIYKWENFKLEINAKNTWKLLEFDFTGVNNKTEIISTPFGDFMVMTFFFNVSRRFGFIVFQNYVPSSV   299
E2QVM5|GABRE_Canis_lupus_familiaris       IDAGCLLYMLKFPMDSHSCPLSFSSFSYPESEMIYKWENFKLQIDERNTWKLFQFDFTGVTNTTETISTPAGDFTVMTLYFNVSRRFGFVAFQNYVPSSV   285
M3YJW6|GABRE_Mustela_putorius_furo        IDAGCLLHMLKFPMDSHSCPLSFSSFSYPESEMIYKWEDFKLEINEENSWKLFQFDFTGVTNTTETISTPAGDFMVMTLFFNVSRRFGFVAFQNYVPSSV   290
G3MWU9|GABRE_Bos_taurus                   IEAGCSLHMLKFPMDSHSCPLSFSSFSYPENDLIYKWENFMLKINESNSWKLFQFDFTGVSNTTETVTTLAGDFIVMTLYFNVSRRFGFVAFHNYVPSSV   290
                                          .......210.......220.......230.......240.......250.......260.......270.......280.......290.......300

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P78334|GABRE_Homo_sapiens                 TTMLSWVSFWIKTESAPARTSLGITSVLTMTTLGTFSRKNFPRVSYITALDFYIAICFVFCFCALLEFAVLNFLIYNQTKAHASPKL------RHPRINS   384
A0A2R8ZSJ9|GABRE_Pan_paniscus             TTMLSWVSFWIKTESAPARTSLGITSVLTMTTLGTFSRKNFPRVSYITALDFYIAICFVFCFCALLEFAVLNFLIYNQTKAHASPKL------RHPRINS   384
H2QZ86|GABRE_Pan_troglodytes              TTMLSWVSFWIKTESAPARTSLGITSVLTMTTLGTFSRKNFPRVSYITALDFYIAICFVFCFCALLEFAVLNFLIYNQTKAHASPKL------RHPRINS   384
F7GJ80|GABRE_Macaca_mulatta               TTMLSWVSFWIKTESAPARTSLGITSVLTMTTLGTFSRKNFPRVSYITALDFYIAICFVFCFCALLEFAVLNFLIYNQTKAHASPKL------RHPRIDS   384
Q9ES14|GABRE_Rattus_norvegicus            TTMLSWVSFWIKIEAAAARASVGVSSVLTMATLGTFSRKNFPRVSYLTALDFYIAICFVLCFCTLLEFTVLNFLTYNNIERQASPKFYQFPTNSRANART   391
Q9JLE8|GABRE_Mus_musculus                 TTMLSWVSFWIKIEAAAARASVGVSSVLTMATLGTLSCKNFPRVSYLTALDFYIAICFVLCFCTLLEFAVLNFLTYNNIKRQASLKVYQLPTNSRANAHT   399
E2QVM5|GABRE_Canis_lupus_familiaris       TTMLSWVSFWIKRDSAPARTSLGITSVLTMTTLGTFSRKNFPRVSYITALDFYIAICFVFCFCALMEFAVLNFLTYNRTNPRASPKL------RHP----   375
M3YJW6|GABRE_Mustela_putorius_furo        TTMLSWVSFWIKKDSAPARTSLGITSVLTMTTLGTFSRKNFPRVSYVTALDFYIAICFVFCFCALMEFAVLNFLTYNWTKPRASPRF------RHP----   380
G3MWU9|GABRE_Bos_taurus                   TTMVSWISFWIKKESAPARTSLGITSVLTMTTLGTFSRKNFPRVSYITALDFYIAICFVFCFCALMEFAVLNFLTYNRTAPRGSPIL------RHP----   380
                                          .......310.......320.......330.......340.......350.......360.......370.......380.......390.......400

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P78334|GABRE_Homo_sapiens                 RAHARTRARSRACARQHQEAFVCQIVTTE-----GSDGEERPSCSAQQPPSPGSPEGPRSLCSKLACCEWCKRFKKYFCMVPDCEGSTWQQGRLCIHVYR   479
A0A2R8ZSJ9|GABRE_Pan_paniscus             RAHARTRARSRACARQHQEAFVCQIVTTE-----GSDGEEHPSCSAQQPPSPGSPEGPRSLCSKLACCEWCKGFKKYFCMVPDCEGSTWQQGRLCIHVYR   479
H2QZ86|GABRE_Pan_troglodytes              RAHARTRARSRACARQHQEAFVCQIVTTE-----GSDGEERPSCSAQQPPSPGSPEGPRSLCSKLACCEWCKGFKKYFCMVPDCEGSTWQQGRLCIHVYR   479
F7GJ80|GABRE_Macaca_mulatta               RAHARTRARSRACARQHQEAFVCQIVTTE-----GSDGEDRPSCSAQQPPSPGSPEGPHSLCSKLACCEWCKGFKKYFCMVPDCEGSTWQQGRLCIHVYR   479
Q9ES14|GABRE_Rattus_norvegicus            RARARTRARARARARQQQEVFVCEIVTYE------ENAEEG----YQWSPRSRRPQCPWRRCGRSYVCFRV--LRKYFCMVPGCEGNNWQRGRICIHVYR   479
Q9JLE8|GABRE_Mus_musculus                 RARARTRARARARARQQQEVFVCEIVTYE------ENAEEG----YQWSPRSRRPQCPWRRCGRSYVCFRV--LRKYFCMAPGCEGGSWQRGRICIHVYR   487
E2QVM5|GABRE_Canis_lupus_familiaris       QIRARSRTPPRAHVHQHPEAFVCQIEDSE-----ESDEEEGPSCPAQQSLSLDTTQRPG-GC-----CKWC---KKYFCMVPSCEGSIWQKGRLFIHIYR   461
M3YJW6|GABRE_Mustela_putorius_furo        HIRARALTPPRAHVHQHPEAFVCQIEDSE-----ESDEEEGPFCPAQQSFRRRTIQKAGNGC-----CKWC---KKYFCVVPSCEGSIWQQGRLFIHIYR   467
G3MWU9|GABRE_Bos_taurus                   RATIRVRIP----VVEHPEAFVCDIEDSEEEELGESEEDEGPSCPARQATRPSRLQRSA-GC-----SRWC---DKYCCMVPTCDASSWQQGRLFIHVYR   467
                                          .......410.......420.......430.......440.......450.......460.......470.......480.......490.......500
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                                     *******:**::******:**::****
P78334|GABRE_Homo_sapiens            LDNYSRVVFPVTFFFFNVLYWLVCLNL   506
A0A2R8ZSJ9|GABRE_Pan_paniscus        LDNYSRVVFPVSFFFFNVLYWLVCLNL   506
H2QZ86|GABRE_Pan_troglodytes         LDNYSRVVFPVTFFFFNVLYWLVCLNL   506
F7GJ80|GABRE_Macaca_mulatta          LDNYSRVVFPVTFFFFNVLYWLVCLNL   506
Q9ES14|GABRE_Rattus_norvegicus       LDNYSRVLFPITFFFFNVVYWVICLNL   506
Q9JLE8|GABRE_Mus_musculus            LDNYSRVLFPITFFFFNVLYWLICLNL   514
E2QVM5|GABRE_Canis_lupus_familiaris  LDNYSRVIFPVTFFFFNVLYWLVCLNL   488
M3YJW6|GABRE_Mustela_putorius_furo   LDNYSRVVFPVTFFFFNVLYWLVCLNL   494
G3MWU9|GABRE_Bos_taurus              LDNYSRVIFPVTFFFFNVIYWLVCLNL   494
                                     .......510.......520.......
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Q8N1C3|GABRG1_Homo_sapiens          ---MGPLKAFLFSPFLLRSQSRGVRLVFLLLT--LHLGNCVDKADDEDDEDLTVNKTWVLAPKIHEGDITQILNSLLQGYDNKLRPDIGVRPTVIETDVY    95
A0A2R9A1L5|GABRG1_Pan_paniscus      ---MGPLKAFLFSPFLLRSQSRGVRLVFLLLT--LHLGNCVDKADDEDDEDLTVNKTWVLAPKIHEGDITQILNSLLQGYDNKLRPDIGVRPTVIETDVY    95
H2QPE4|GABRG1_Pan_troglodytes       ---MGPLKAFLFSPFLLRSQSRGVRLVFLLLT--LHLGNCVDKADDEDDEDLTVNKTWVLAPKIHEGDITQILNSLLQGYDNKLRPDIGVRPTVIETDVY    95
A0A5F8A0N7|GABRG1_Macaca_mulatta    ---MGPLKAFLFSPFLLRSQSRGVRLVFLLLT--LHLGNCVDKADDEDDEDLTVNKTWVLAPKIHEGDITQILNSLLQGYDNKLRPDIGVRPTVIETDVY    95
M3Y0V1|GABRG1_Mustela_putorius_furo ---MGSWKAFLFSPFLLWSQSRVVRLMFLLLT--LHLGNCVDKVDDEDDEDLTVNKTWVLAPRIHEGDITQILNSLLQGYDNKLRPDIGVRPTVIETDVY    95
P23574|GABRG1_Rattus_norvegicus     ---MGSGKVFLFSPSLLWSQTRGVRLIFLLLT--LHLGNCIDKADDEDDEDLTMNKTWVLAPKIHEGDITQILNSLLQGYDNKLRPDIGVRPTVIETDVY    95
Q9R0Y8|GABRG1_Mus_musculus          ---MGSGKAFLFSPSLLWSQTRGVRLIFLLLT--LHLGNCVDKADDEDDEDLTMNKTWVLAPKIHEGDITQILNSLLQGYDNKLRPDIGVRPTVIETDVY    95
E2RH22|GABRG1_Canis_lupus_familiaris MPPMGPWKAPLCSPFLLGSQSRVVRLLFLLLT--LHLGNCVDKVDDEDDEDLTVNKTWVLAPKIHEGDITQILNSLLQGYDNKLRPDIGVRPTVIETDVY    98
A0A0R4IPF9|GABRG1_Danio_rerio       YMLNSPLQE---AP----SHAYGYFVLPLLNNQIMAFGPS--KAEEEDYEDVPINKTWVLSPKVYESDVTLILNKLLQGYDNKLRPDIGVRPTVIETAVY    91
F6Q4V7|GABRG1_Bos_taurus            ---MGSWKALLFSPFLLWKQRRGVRLMFLLLT--VHLGNCVDKADDEDDEDLTVNKTWVLAPKIHEGDITQILNSLLQGYDNKLRPDIGVRPTVIETDVY    95
                                    1.......10........20........30........40........50........60........70........80........90.......100

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Q8N1C3|GABRG1_Homo_sapiens          VNSIGPVDPINMEYTIDIIFAQTWFDSRLKFNSTMKVLMLNSNMVGKIWIPDTFFRNSRKSDAHWITTPNRLLRIWNDGRVLYTLRLTINAECYLQLHNF   195
A0A2R9A1L5|GABRG1_Pan_paniscus      VNSIGPVDPINMEYTIDIIFAQTWFDSRLKFNSTMKVLMLNSNMVGKIWIPDTFFRNSRKSDAHWITTPNRLLRIWNDGRVLYTLRLTINAECYLQLHNF   195
H2QPE4|GABRG1_Pan_troglodytes       VNSIGPVDPINMEYTIDIIFAQTWFDSRLKFNSTMKVLMLNSNMVGKIWIPDTFFRNSRKSDAHWITTPNRLLRIWNDGRVLYTLRLTINAECYLQLHNF   195
A0A5F8A0N7|GABRG1_Macaca_mulatta    VNSIGPVDPINMEYTIDIIFAQTWFDSRLKFNSTMKVLMLNSNMVGKIWIPDTFFRNSRKSDAHWITTPNRLLRIWNDGRVLYTLRLTINAECYLQLHNF   195
M3Y0V1|GABRG1_Mustela_putorius_furo VNSIGPVDPINMEYTIDIIFAQTWFDSRLKFNSTMKVLMLNSNMVGKIWIPDTFFRNSRKSDAHWITTPNRLLRIWNDGRVLYTLRLTINAECYLQLHNF   195
P23574|GABRG1_Rattus_norvegicus     VNSIGPVDPINMEYTIDIIFAQTWFDSRLKFNSTMKVLMLNSNMVGKIWIPDTFFRNSRKSDAHWITTPNRLLRIWSDGRVLYTLRLTINAECYLQLHNF   195
Q9R0Y8|GABRG1_Mus_musculus          VNSIGPVDPINMEYTIDIIFAQTWFDSRLKFNSTMKVLMLNSNMVGKIWIPDTFFRNSRKSDAHWITTPNRLLRIWSDGRVLYTLRLTINAECYLQLHNF   195
E2RH22|GABRG1_Canis_lupus_familiaris VNSIGPVDPINMEYTIDIIFAQTWFDSRLKFNSTMKVLMLNSNMVGKIWIPDTFFRNSRKSDAHWITTPNRLLRIWNDGRVLYTLRLTINAECYLQLHNF   198
A0A0R4IPF9|GABRG1_Danio_rerio       VNSIGPVDPINMEYTIDIFFAQTWYDSRLKFNSSMKLLMLNSNMVGKIWIPDTFFRNSRKSDAHWITTPNRLLRLWSNGRVMYTLRLTINAECYLKLHNF   191
F6Q4V7|GABRG1_Bos_taurus            VNSIGPVDPINMEYTIDIIFAQTWFDSRLKFNSTMKVLMLNSNMVGKIWIPDTFFRNSRKSDAHWITTPNRLLRIWNDGRVLYTLRLTINAECYLQLHNF   195
                                    .......110.......120.......130.......140.......150.......160.......170.......180.......190.......200

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Q8N1C3|GABRG1_Homo_sapiens          PMDEHSCPLEFSSYGYPKNEIEYKWKKPSVEVADPKYWRLYQFAFVGLRNSTEITHTISGDYVIMTIFFDLSRRMGYFTIQTYIPCILTVVLSWVSFWIN   295
A0A2R9A1L5|GABRG1_Pan_paniscus      PMDEHSCPLEFSSYGYPKNEIEYKWKKPSVEVADPKYWRLYQFAFVGLRNSTEITHTISGDYVIMTIFFDLSRRMGYFTIQTYIPCILTVVLSWVSFWIN   295
H2QPE4|GABRG1_Pan_troglodytes       PMDEHSCPLEFSSYGYPKNEIEYKWKKPSVEVADPKYWRLYQFAFVGLRNSTEITHTISGDYVIMTIFFDLSRRMGYFTIQTYIPCILTVVLSWVSFWIN   295
A0A5F8A0N7|GABRG1_Macaca_mulatta    PMDEHSCPLEFSSYGYPKNEIEYKWKKPSVEVADPKYWRLYQFAFVGLRNSTEISHTISGDYVIMTIFFDLSRRMGYFTIQTYIPCILTVVLSWVSFWIN   295
M3Y0V1|GABRG1_Mustela_putorius_furo PMDEHSCPLEFSSYGYPKNEIEYKWKKPSVEVADPKYWRLYQFAFVGLRNSTEISHTISGDYVIMTIFFDLSRRMGYFTIQTYIPCILTVVLSWVSFWIN   295
P23574|GABRG1_Rattus_norvegicus     PMDEHSCPLEFSSYGYPKNEIEYKWKKPSVEVADPKYWRLYQFAFVGLRNSTEISHTISGDYIIMTIFFDLSRRMGYFTIQTYIPCILTVVLSWVSFWIN   295
Q9R0Y8|GABRG1_Mus_musculus          PMDEHSCPLEFSSYGYPKNEIEYKWKKPSVEVADPKYWRLYQFAFVGLRNSTEISHTISGDYIIMTIFFDLSRRMGYFTIQTYIPCILTVVLSWVSFWIN   295
E2RH22|GABRG1_Canis_lupus_familiaris PMDEHSCPLEFSSYGYPKNEIEYKWKKPSVEVADPKYWRLYQFAFVGLRNSTEISHTISGDYIIMTIFFDLSRRMGYFTIQTYIPCILTVVLSWVSFWIN   298
A0A0R4IPF9|GABRG1_Danio_rerio       PMDEHSCPLEFSSYGYPKNEIQYRWQRRSVEVADQRYWRLYQFAFVGLRNSSDVANTQSGEYVVMTIFFDLSRRMGYFTIQTYIPCSMIVVLSWVSFWIN   291
F6Q4V7|GABRG1_Bos_taurus            PMDEHSCPLEFSSYGYPKNEIEYKWKKPSVEVADPKYWRLYQFAFVGLRNTTEISHTISGDYIIMTIFFDLSRRMGYFTIQTYIPCILTVVLSWVSFWIN   295
                                    .......210.......220.......230.......240.......250.......260.......270.......280.......290.......300

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Q8N1C3|GABRG1_Homo_sapiens          KDAVPARTSLGITTVLTMTTLSTIARKSLPKVSYVTAMDLFVSVCFIFVFAALMEYGTLHYFTSNQKGKTATKDRKLKNKASMTPGLHPGSTLIPMNNIS   395
A0A2R9A1L5|GABRG1_Pan_paniscus      KDAVPARTSLGITTVLTMTTLSTIARKSLPKVSYVTAMDLFVSVCFIFVFAALMEYGTLHYFTSNQKGKTATKDRKLKNKASMTPGLHPGSTLIPMNNIS   395
H2QPE4|GABRG1_Pan_troglodytes       KDAVPARTSLGITTVLTMTTLSTIARKSLPKVSYVTAMDLFVSVCFIFVFAALMEYGTLHYFTSNQKGKTATKDRKLKNKASMTPGLHPGSTLIPMNNIS   395
A0A5F8A0N7|GABRG1_Macaca_mulatta    KDAVPARTSLGITTVLTMTTLSTIARKSLPKVSYVTAMDLFVSVCFIFVFAALMEYGTLHYFTSNKKGKTATKDRKLKNKASMTPGLHPGSTLIPMNSVS   395
M3Y0V1|GABRG1_Mustela_putorius_furo KDAVPARTSLGITTVLTMTTLSTIARKSLPKVSYVTAMDLFVSVCFIFVFAALMEYGTLHYFTSNKKVKT-TRDKKQKNKSSMSPGLHPGSTLIPMNNIS   394
P23574|GABRG1_Rattus_norvegicus     KDAVPARTSLGITTVLTMTTLSTIARKSLPKVSYVTAMDLFVSVCFIFVFAALMEYGTLHYFTSNNKGKT-TRDRKLKSKTSVSPGLHAGSTLIPMNNIS   394
Q9R0Y8|GABRG1_Mus_musculus          KDAVPARTSLGITTVLTMTTLSTIARKSLPKVSYVTAMDLFVSVCFIFVFAALMEYGTLHYFTSNNKGKT-TRGRKLKNKTSASPGLHAGSTLIPMNSIS   394
E2RH22|GABRG1_Canis_lupus_familiaris KDAVPARTSLGITTVLTMTTLSTIARKSLPKVSYVTAMDLFVSVCFIFVFAALMEYGTLHYFTSNKKGKT-TRNRKLKNKSSMSPGLHPGSTLIPMNSIS   397
A0A0R4IPF9|GABRG1_Danio_rerio       KDAVPARTSLGITTVLTMTTLSTISRKSLPKVSYVTAMDLFVSVCFIFTFAALMEYGTLHYFTSNRQNKK-TKSSHAQKPSMVN--IRPGTSLLQMNNIA   388
F6Q4V7|GABRG1_Bos_taurus            KDAVPARTSLGITTVLTMTTLSTIARKSLPKVSYVTAMDLFVSVCFIFVFAALMEYGTLHYFTSNKKVKT-TREKKPKLKTSMSSGLHPGSTLIPMNSLS   394
                                    .......310.......320.......330.......340.......350.......360.......370.......380.......390.......400

                                      : ::**.*:**:****::*****:***:*:***.*:*::::*********      * :  ::.*. :    :*:
Q8N1C3|GABRG1_Homo_sapiens          VPQ-EDDYGYQCLEGKDCASFFCCFEDCRTGSWREGRIHIRIAKIDSYSRIFF----PTAFALFNLVYWVGYLYL---------   465
A0A2R9A1L5|GABRG1_Pan_paniscus      VPQ-EDDYGYQCLEGKDCASFFCCFEDCRTGSWREGRIHIRIAKIDSYSRIFF----PTAFALFNLVYWVGYLYL---------   465
H2QPE4|GABRG1_Pan_troglodytes       VPQ-EDDYGYQCLEGKDCASFFCCFEDCRTGSWREGRIHIRIAKIDSYSRIFF----PTAFALFNLVYWVGYLYL---------   465
A0A5F8A0N7|GABRG1_Macaca_mulatta    VPQ-EEDYGYQCLEGKDCASFFCCFEDCRTGSWREGRIHIRIAKIDSYSRIFSQQLLPCSTWFIGLAIFTYKIYFISKSQKKFD   478
M3Y0V1|GABRG1_Mustela_putorius_furo LPHGEEDYGYQCLEGKDCASFFCCFEDCRTGSWREGRIHIRIAKIDSYSRIFF----PTAFALFNLVYWVGYLYL---------   465
P23574|GABRG1_Rattus_norvegicus     MPQGEDDYGYQCLEGKDCATFFCCFEDCRTGSWREGRIHIRIAKIDSYSRIFF----PTAFALFNLVYWVGYLYL---------   465
Q9R0Y8|GABRG1_Mus_musculus          LPQGEDDYGYQCLEGKDCTSFFCCFDDCRTGSWREGRIHIRIAKIDSYSRIFF----PTAFALFNLVYWVGYLYL---------   465
E2RH22|GABRG1_Canis_lupus_familiaris LPHGEEDYGYQCLEGKDCASFFCCFEDCRTGSWREGRIHIRIAKIDSYSRIFF----PTAFALFNLVYWVGYLYL---------   468
A0A0R4IPF9|GABRG1_Danio_rerio       PYHEDDDYAYECLDGKDCTSFFCCFDDCRSGAWRENRMHVHVSKIDSYSRIFF----PTAFGLFNLVYWIGYLYL---------   459
F6Q4V7|GABRG1_Bos_taurus            LPQGEDDYGYQCLEGKDCASFFCCFEDCRTGSWREGRIHIRIAKIDSYSRIFF----PTAFALFNLVYWVGYLYL---------   465
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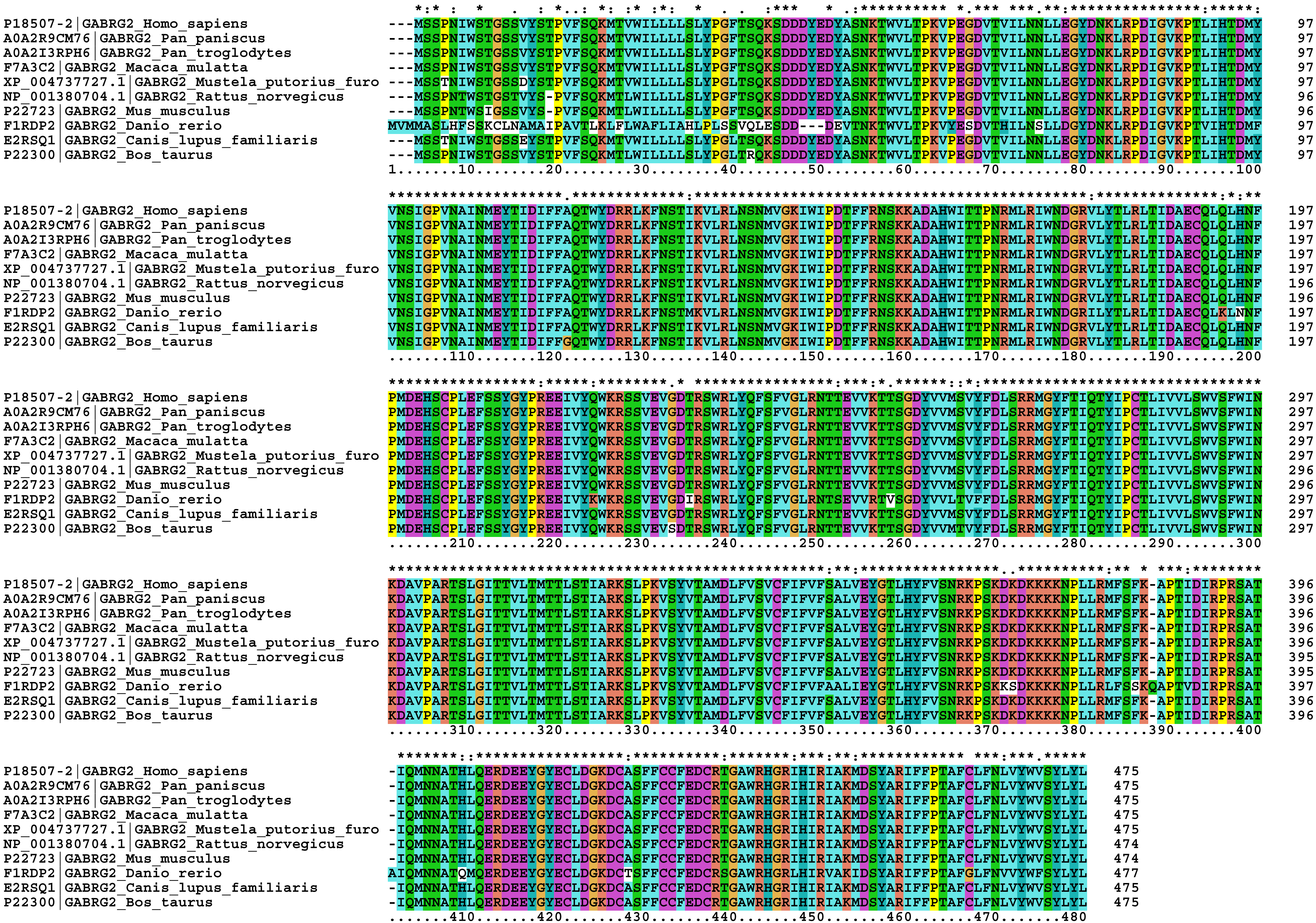
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A0A2R9CM76|GABRG2_Pan_paniscus ---MSSPNIWSTGSSVYSTPVFSQKMTVWILLLLSLYPGFTSQKSDDDYEDYASNKTWVLTPKVPEGDVTVILNNLLEGYDNKLRPDIGVKPTLIHTDMY 97
A0A2I3RPH6|GABRG2_Pan_troglodytes ---MSSPNIWSTGSSVYSTPVFSQKMTVWILLLLSLYPGFTSQKSDDDYEDYASNKTWVLTPKVPEGDVTVILNNLLEGYDNKLRPDIGVKPTLIHTDMY 97
F7A3C2|GABRG2_Macaca_mulatta ---MSSPNIWSTGSSVYSTPVFSQKMTVWILLLLSLYPGFTSQKSDDDYEDYASNKTWVLTPKVPEGDVTVILNNLLEGYDNKLRPDIGVKPTLIHTDMY 97
XP_004737727.1|GABRG2_Mustela_putorius_furo ---MSSTNIWSTGSSDYSTPVFSQKMTVWILLLLSLYPGLTSQKSDDDYEDYASNKTWVLTPKVPEGDVTVILNNLLEGYDNKLRPDIGVKPTLIHTDMY 97
NP_001380704.1|GABRG2_Rattus_norvegicus ---MSSPNTWSTGSTVYS-PVFSQKMTLWILLLLSLYPGFTSQKSDDDYEDYASNKTWVLTPKVPEGDVTVILNNLLEGYDNKLRPDIGVKPTLIHTDMY 96
P22723|GABRG2_Mus_musculus ---MSSPNTWSIGSSVYS-PVFSQKMTLWILLLLSLYPGFTSQKSDDDYEDYASNKTWVLTPKVPEGDVTVILNNLLEGYDNKLRPDIGVKPTLIHTDMY 96
F1RDP2|GABRG2_Danio_rerio MVMMASLHFSSKCLNAMAIPAVTLKLFLWAFLIAHLPLSSVQLESDD---DEVTNKTWVLTPKVYESDVTHILNSLLDGYDNKLRPDIGVKPTVIHTDMF 97
E2RSQ1|GABRG2_Canis_lupus_familiaris ---MSSTNIWSTGSSEYSTPVFSQKMTVWILLLLSLYPGLTSQKSDDDYEDYASNKTWVLTPKVPEGDVTVILNNLLEGYDNKLRPDIGVKPTLIHTDMY 97
P22300|GABRG2_Bos_taurus ---MSSPNIWSTGSSVYSTPVFSQKMTLWILLLLSLYPGLTRQKSDDDYEDYASNKTWVLTPKVPEGDVTVILNNLLEGYDNKLRPDIGVKPTLIHTDMY 97
P18507-2|GABRG2_Homo_sapiens VNSIGPVNAINMEYTIDIFFAQTWYDRRLKFNSTIKVLRLNSNMVGKIWIPDTFFRNSKKADAHWITTPNRMLRIWNDGRVLYTLRLTIDAECQLQLHNF 197
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A0A2I3RPH6|GABRG2_Pan_troglodytes VNSIGPVNAINMEYTIDIFFAQTWYDRRLKFNSTIKVLRLNSNMVGKIWIPDTFFRNSKKADAHWITTPNRMLRIWNDGRVLYTLRLTIDAECQLQLHNF 197
F7A3C2|GABRG2_Macaca_mulatta VNSIGPVNAINMEYTIDIFFAQTWYDRRLKFNSTIKVLRLNSNMVGKIWIPDTFFRNSKKADAHWITTPNRMLRIWNDGRVLYTLRLTIDAECQLQLHNF 197
XP_004737727.1|GABRG2_Mustela_putorius_furo VNSIGPVNAINMEYTIDIFFAQTWYDRRLKFNSTIKVLRLNSNMVGKIWIPDTFFRNSKKADAHWITTPNRMLRIWNDGRVLYTLRLTIDAECQLQLHNF 197
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P22723|GABRG2_Mus_musculus VNSIGPVNAINMEYTIDIFFAQTWYDRRLKFNSTIKVLRLNSNMVGKIWIPDTFFRNSKKADAHWITTPNRMLRIWNDGRVLYTLRLTIDAECQLQLHNF 196
F1RDP2|GABRG2_Danio_rerio VNSIGPVNAINMEYTIDIFFAQTWYDRRLKFNSTMKVLRLNSNMVGKIWIPDTFFRNSKKADAHWITTPNRMLRIWNDGRILYTLRLTIDAECQLKLNNF 197
E2RSQ1|GABRG2_Canis_lupus_familiaris VNSIGPVNAINMEYTIDIFFAQTWYDRRLKFNSTIKVLRLNSNMVGKIWIPDTFFRNSKKADAHWITTPNRMLRIWNDGRVLYTLRLTIDAECQLQLHNF 197
P22300|GABRG2_Bos_taurus VNSIGPVNAINMEYTIDIFFGQTWYDRRLKFNSTIKVLRLNSNMVGKIWIPDTFFRNSKKADAHWITTPNRMLRIWNDGRVLYTLRLTIDAECQLQLHNF 197
P18507-2|GABRG2_Homo_sapiens PMDEHSCPLEFSSYGYPREEIVYQWKRSSVEVGDTRSWRLYQFSFVGLRNTTEVVKTTSGDYVVMSVYFDLSRRMGYFTIQTYIPCTLIVVLSWVSFWIN 297
A0A2R9CM76|GABRG2_Pan_paniscus PMDEHSCPLEFSSYGYPREEIVYQWKRSSVEVGDTRSWRLYQFSFVGLRNTTEVVKTTSGDYVVMSVYFDLSRRMGYFTIQTYIPCTLIVVLSWVSFWIN 297
A0A2I3RPH6|GABRG2_Pan_troglodytes PMDEHSCPLEFSSYGYPREEIVYQWKRSSVEVGDTRSWRLYQFSFVGLRNTTEVVKTTSGDYVVMSVYFDLSRRMGYFTIQTYIPCTLIVVLSWVSFWIN 297
F7A3C2|GABRG2_Macaca_mulatta PMDEHSCPLEFSSYGYPREEIVYQWKRSSVEVGDTRSWRLYQFSFVGLRNTTEVVKTTSGDYVVMSVYFDLSRRMGYFTIQTYIPCTLIVVLSWVSFWIN 297
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P22723|GABRG2_Mus_musculus PMDEHSCPLEFSSYGYPREEIVYQWKRSSVEVGDTRSWRLYQFSFVGLRNTTEVVKTTSGDYVVMSVYFDLSRRMGYFTIQTYIPCTLIVVLSWVSFWIN 296
F1RDP2|GABRG2_Danio_rerio PMDEHSCPLEFSSYGYPKEEIVYKWKRSSVEVGDIRSWRLYQFSFVGLRNTSEVVRTVSGDYVVLTVFFDLSRRMGYFTIQTYIPCTLIVVLSWVSFWIN 297
E2RSQ1|GABRG2_Canis_lupus_familiaris PMDEHSCPLEFSSYGYPREEIVYQWKRSSVEVGDTRSWRLYQFSFVGLRNTTEVVKTTSGDYVVMSVYFDLSRRMGYFTIQTYIPCTLIVVLSWVSFWIN 297
P22300|GABRG2_Bos_taurus PMDEHSCPLEFSSYGYPREEIVYQWKRSSVEVSDTRSWRLYQFSFVGLRNTTEVVKTTSGDYVVMTVYFDLSRRMGYFTIQTYIPCTLIVVLSWVSFWIN 297
P18507-2|GABRG2_Homo_sapiens KDAVPARTSLGITTVLTMTTLSTIARKSLPKVSYVTAMDLFVSVCFIFVFSALVEYGTLHYFVSNRKPSKDKDKKKKNPLLRMFSFK-APTIDIRPRSAT 396
A0A2R9CM76|GABRG2_Pan_paniscus KDAVPARTSLGITTVLTMTTLSTIARKSLPKVSYVTAMDLFVSVCFIFVFSALVEYGTLHYFVSNRKPSKDKDKKKKNPLLRMFSFK-APTIDIRPRSAT 396
A0A2I3RPH6|GABRG2_Pan_troglodytes KDAVPARTSLGITTVLTMTTLSTIARKSLPKVSYVTAMDLFVSVCFIFVFSALVEYGTLHYFVSNRKPSKDKDKKKKNPLLRMFSFK-APTIDIRPRSAT 396
F7A3C2|GABRG2_Macaca_mulatta KDAVPARTSLGITTVLTMTTLSTIARKSLPKVSYVTAMDLFVSVCFIFVFSALVEYGTLHYFVSNRKPSKDKDKKKKNPLLRMFSFK-APTIDIRPRSAT 396
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P22723|GABRG2_Mus_musculus KDAVPARTSLGITTVLTMTTLSTIARKSLPKVSYVTAMDLFVSVCFIFVFSALVEYGTLHYFVSNRKPSKDKDKKKKNPLLRMFSFK-APTIDIRPRSAT 395
F1RDP2|GABRG2_Danio_rerio KDAVPARTSLGITTVLTMTTLSTIARKSLPKVSYVTAMDLFVSVCFIFVFAALIEYGTLHYFVSNRKPSKKSDKKKKNPLLRLFSSKQAPTVDIRPRSAT 397
E2RSQ1|GABRG2_Canis_lupus_familiaris KDAVPARTSLGITTVLTMTTLSTIARKSLPKVSYVTAMDLFVSVCFIFVFSALVEYGTLHYFVSNRKPSKDKDKKKKNPLLRMFSFK-APTIDIRPRSAT 396
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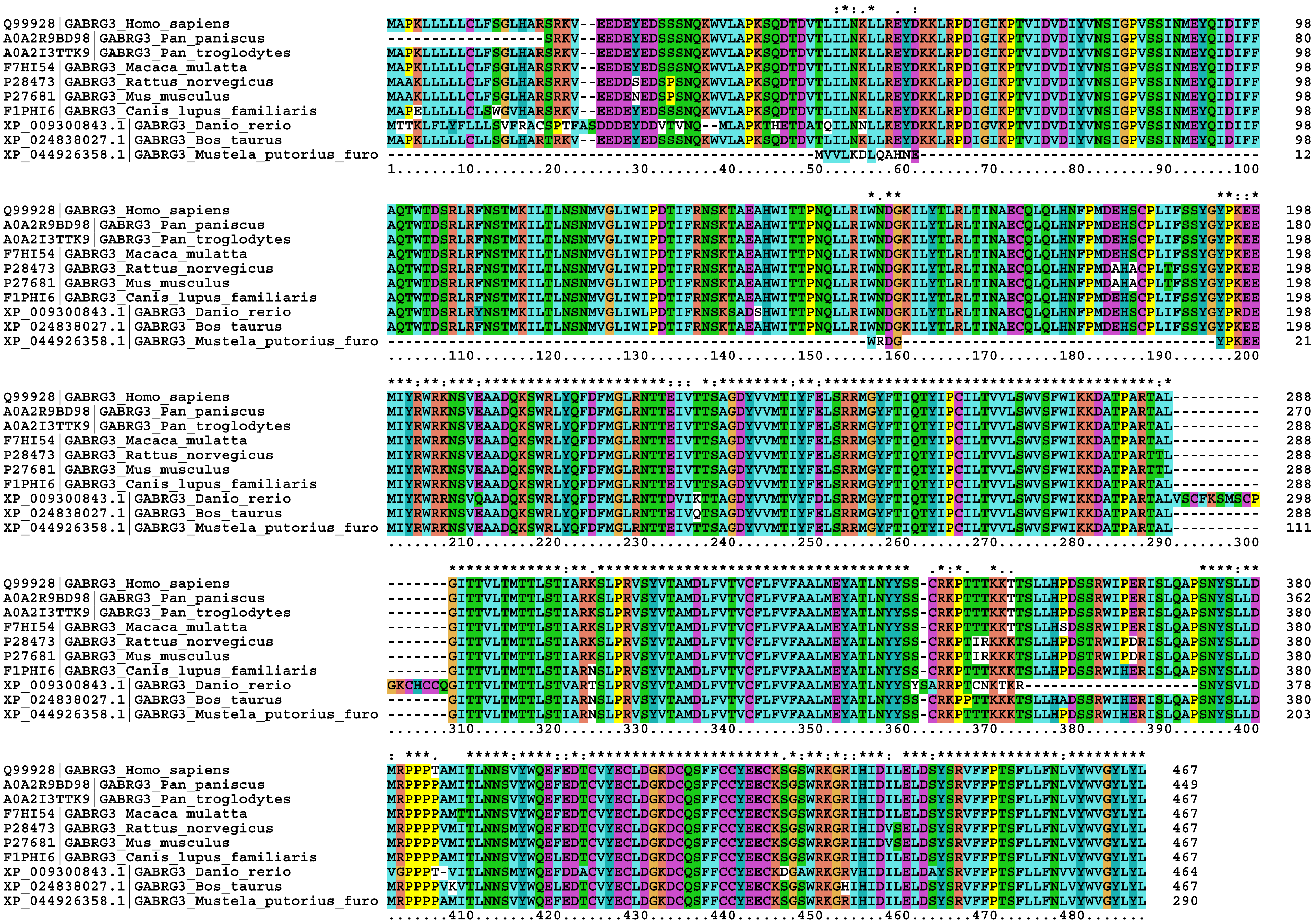
P18507-2|GABRG2_Homo_sapiens -IQMNNATHLQERDEEYGYECLDGKDCASFFCCFEDCRTGAWRHGRIHIRIAKMDSYARIFFPTAFCLFNLVYWVSYLYL 475
A0A2R9CM76|GABRG2_Pan_paniscus -IQMNNATHLQERDEEYGYECLDGKDCASFFCCFEDCRTGAWRHGRIHIRIAKMDSYARIFFPTAFCLFNLVYWVSYLYL 475
A0A2I3RPH6|GABRG2_Pan_troglodytes -IQMNNATHLQERDEEYGYECLDGKDCASFFCCFEDCRTGAWRHGRIHIRIAKMDSYARIFFPTAFCLFNLVYWVSYLYL 475
F7A3C2|GABRG2_Macaca_mulatta -IQMNNATHLQERDEEYGYECLDGKDCASFFCCFEDCRTGAWRHGRIHIRIAKMDSYARIFFPTAFCLFNLVYWVSYLYL 475
XP_004737727.1|GABRG2_Mustela_putorius_furo -IQMNNATHLQERDEEYGYECLDGKDCASFFCCFEDCRTGAWRHGRIHIRIAKMDSYARIFFPTAFCLFNLVYWVSYLYL 475
NP_001380704.1|GABRG2_Rattus_norvegicus -IQMNNATHLQERDEEYGYECLDGKDCASFFCCFEDCRTGAWRHGRIHIRIAKMDSYARIFFPTAFCLFNLVYWVSYLYL 474
P22723|GABRG2_Mus_musculus -IQMNNATHLQERDEEYGYECLDGKDCASFFCCFEDCRTGAWRHGRIHIRIAKMDSYARIFFPTAFCLFNLVYWVSYLYL 474
F1RDP2|GABRG2_Danio_rerio AIQMNNATQMQERDEEYGYECLDGKDCTSFFCCFEDCRSGAWRHGRLHIRVAKIDSYARIFFPTAFGLFNVVYWFSYLYL 477
E2RSQ1|GABRG2_Canis_lupus_familiaris -IQMNNATHLQERDEEYGYECLDGKDCASFFCCFEDCRTGAWRHGRIHIRIAKMDSYARIFFPTAFCLFNLVYWVSYLYL 475
P22300|GABRG2_Bos_taurus -IQMNNATHLQERDEEYGYECLDGKDCASFFCCFEDCRTGAWRHGRIHIRIAKMDSYARIFFPTAFCLFNLVYWVSYLYL 475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Q9UN88|GABRQ_Homo_sapiens              MGIRGMLRAAVILLLIRTWLAEGNYPSPIPKFHFEFSSAVPEVVLNLFNCKNCANEAVVQKILDRVLSRYDVRLRPNFGGAPVPVRISIYVTSIEQISEM  100
A0A2R9ALH2|GABRQ_Pan_paniscus          MGIRGMLRAAVILLLIRTWLAEGNYPSPIPKFHFEFSSAVPEVVLNLFNCKNCANEAVVQKILDRVLSRYDVRLRPNFGGAPVPVRISIYVTSIEQISEM  100
A0A6D2XHJ7|GABRQ_Pan_troglodytes       MGIRGMLRAAVILLLIRTWLAEGNYPSPIPKFHFEFSSAVPEVVLNLFNCKNCANEAVVQKILDRVLSRYDVRLRPNFGGAPVPVRISIYVTSIEQISEM  100
F7F4H9|GABRQ_Macaca_mulatta            MGIRGMLRAAVILLLIRTWLAEGNYPSPIPKFHFEFSSAVPEVVLNLFNCKNCANEAVVQKILDRVLSRYDVRLRPNFGGAPVPVRISIYVTSIEQISEM  100
M3YJY6|GABRQ_Mustela_putorius_furo     MGIRGMLRAAVLLLLIRTWLAESHDPSPTPEFHFELSPTVPEVVLSLFNCKNCANEAVVHKILDRVLSKYDVRLRPNFGGAPVPVGISMYVSSIEQISEM  100
G3V875|GABRQ_Rattus_norvegicus         -----MLRAAALLLLIRTWLAESNDPSPTPKFHFELSSSMPEVILDLFNCKNCANEAVVQKILDRVLSTYDVRLRPNFGGAPVPVSISIYVSSIEQISEI   95
Q9JLF1|GABRQ_Mus_musculus              MGIRGMLRAAALLLLIRTWLAESNGPSPTPKFHFELSSSTPEVILDLFNCKNCANEAVVQKILDRVLSTYDVRLRPNFGGAPVPVSVSIYVSSIEQISEI  100
F6XRX3|GABRQ_Canis_lupus_familiaris    MGIRGILRAAVLLLLIRTWLAEGNEPSPTPKFHFELSPTVPEVILNLFNCKKCANEGVVHKILDRVLSNYDVRLRPNFGGAPVPVGVSMYVSSIEQISEM  100
E1BJH4|GABRQ_Bos_taurus                MSIRGMLRAAVFLLLIRTWLAEGSDLSSTPKFHVELSSTVPEVVLNLFDCKNCANEATVHKILDRVLSNYDSRLRPNFGGTPVPVGVSIYVSSIEQISEV  100
F1Q4Y6|GABRB4_Danio_rerio              --------------------MLGPQEDKLCGIVSALAAL-SFVCFAQSSSTGSTGISVAKTTVDKLLKGYDIRLRPDFGGPPVIVGMSINIASIDSISEV   79
                                       1.......10........20........30........40........50........60........70........80........90.......100

                                       .*******:::*:*:* **** *  ****** *: ::**:** **:*.*.:*:*. *.:**:::******* **:*:****** ::*.::*:*:* *.*
Q9UN88|GABRQ_Homo_sapiens              NMDYTITMFFHQTWKDSRLAYYETTLNLTLDYRMHEKLWVPDCYFLNSKDAFVHDVTVENRVFQLHPDGTVRYGIRLTTTAACSLDLHKFPMDKQACNLV  200
A0A2R9ALH2|GABRQ_Pan_paniscus          NMDYTITMFFHQTWKDSRLAYYETTLNLTLDYRMHEKLWVPDCYFLNSKDAFVHDVTVENRVFQLHPDGTVRYGIRLTTTAACSLDLHKFPMDKQACNLV  200
A0A6D2XHJ7|GABRQ_Pan_troglodytes       NMDYTITMFFHQTWKDSRLAYYETTLNLTLDYRMHEKLWVPDCYFLNSKDAFVHDVTVENRVFQLHPDGTVRYGIRLTTTAACSLDLHKFPMDKQACNLV  200
F7F4H9|GABRQ_Macaca_mulatta            NMDYTITMFFHQTWKDSRLAYYETTLNLTLDYRMHEKLWVPDCYFLNSKDAFVHDVTVENRVFQLHPDGTVRYGIRLTTTAACSLDLHKFPMDKQACNLV  200
M3YJY6|GABRQ_Mustela_putorius_furo     TMDYTITMFFHQTWKDPRLAYHETNLNLTLDYRMLEKLWVPDCYFLNSKDAFVHDITVENRVFQLHPDGTVRYGMRLTTTAACSLDLQKFPLDKQACKLE  200
G3V875|GABRQ_Rattus_norvegicus         NMDYTITMFLHQTWKDTRLAYYETNLNLTLDYRMHEKLWVPDCYFVNSKDAFVHDVTAENRVFQLHPDGTVQYGIRLTTTAACSLDLQKFPMDKQSCKLE  195
Q9JLF1|GABRQ_Mus_musculus              NMDYTITMFLHQTWKDTRLAYYETNLNLTLDYRMHEKLWVPDCYFVNSKDAFVHDVTVENRVFQLHPDGTVRYGIRLTTTAACSLDLQKFPMDKQSCKLE  200
F6XRX3|GABRQ_Canis_lupus_familiaris    TMDYTITMFFHQTWKDPRLAYHETDLNLTLDYRMLEKLWVPDCYFLNSKDAFVHDVTVENRVFQLHPDGTVRYGMRLTTTAACSLDLQKFPLDKQTCKLE  200
E1BJH4|GABRQ_Bos_taurus                TMDYTITMFFHQTWKDPRLAYHETNLNLTLDYRLFEKLWVPDCYFLNSKEAFVHDATVENRVFQLHPDGTVRYGIRLTTTAACSMNLEKFPLDKQTCKLE  200
F1Q4Y6|GABRB4_Danio_rerio              NMDYTITMYFQQSWRDKRLAYAEMALNLTLDNRVADQLWLPDTYFLNDKKSFLHGVTVKNRMIRLHPDGTVLYGLRITTTAACMMDLRRYPLDEQNCTLE  179
                                       .......110.......120.......130.......140.......150.......160.......170.......180.......190.......200

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Q9UN88|GABRQ_Homo_sapiens              VESYGYTVEDIILFWDDNGNAIHMTEELHIPQFTFLGRTITSKEVYFYTGSYIRLILKFQVQREVNSYLVQVYWPTVLTTITSWISFWMNYDSSAARVTI  300
A0A2R9ALH2|GABRQ_Pan_paniscus          VESYGYTVEDIILFWDDNGNAIHMTEELHIPQFTFLGRTITSKEVYFYTGSYIRLILKFQVQREVNSYLVQVYWPTVLTTITSWISFWMNYDSSAARVTI  300
A0A6D2XHJ7|GABRQ_Pan_troglodytes       VESYGYTVEDIILFWDDNGNAIHMTEELHIPQFTFLGRTITSKEVYFYTGSYIRLILKFQVQREVNSYLVQVYWPTVLTTITSWISFWMNYDSSAARVTI  300
F7F4H9|GABRQ_Macaca_mulatta            VESYGYTVEDIILFWDDNGNAIHMTEELHIPQFTFLGRTITSKEVYFYTGSYIRLILKFQVQREVNSYLVQVYWPTVLTTITSWISFWMNYDSSAARVTI  300
M3YJY6|GABRQ_Mustela_putorius_furo     VESYGYTVEDIVLYWEGDGNAIQGTEQLHIPQFSFLGKTMSSKEMAFYTGSYVRLILRFLVQREITSYLVQIYWPTVLTTVVSWISFWMNYESSAARVTV  300
G3V875|GABRQ_Rattus_norvegicus         VESYGYTVEDIVLSWE-DDNAIHITDELHIPQYTYLGRTITSKEVYFYTGSYMRLIVKFQVQREVRSYLVQVYWPTVLTTILSWISFWMNYGSSAARVTI  294
Q9JLF1|GABRQ_Mus_musculus              VESYGYTVEDIVLSWE-DDNAIHITDGLHIPQYTYLGRTITSKEVYFYTGSYMRLIVKFQVQREVRSYLVQVYWPTVLTTILSWISFWMNYDSSAARVTI  299
F6XRX3|GABRQ_Canis_lupus_familiaris    VESYGYTVEDIVLYWEGNGNAIQGTEKLHIPQFSFLGKTMSSKEVFFYTGSYVRLILRFLVQREVTSYLVQIYWPTVLTTVVSWISFWMNYESSAARVTV  300
E1BJH4|GABRQ_Bos_taurus                VESYGYTVDDILLYWEGSGNAVQGTEKLHIPQFSFLGKTISTKEVFFYTGSYVRLILRFLVKREVTSYLVQIYWPSVLTTVVSWISFWMNYESSAARVTV  300
F1Q4Y6|GABRB4_Danio_rerio              IESYGYTTDDIVFFWQGGDNAVTGVDKLELPQFSIVELRLVSREVRFTTGSYPRLSLSFRIKRNIGYFILQTYMPSILITILSWVSFWINYDASAARVAL  279
                                       .......210.......220.......230.......240.......250.......260.......270.......280.......290.......300

                                       *:*::* :.:*::*: : :*:.. :****:*:: *:.****:*:**. :**:::.:   .::  :   .:  :.    : **.:  :
Q9UN88|GABRQ_Homo_sapiens              GLTSMLILTTIDSHLRDKLPNISCIKAIDIYILVCLFFVFLSLLEYVYINYLFYSR-GPRRQPRRHRRPRRV-IARYRYQQVVVG--------------  383
A0A2R9ALH2|GABRQ_Pan_paniscus          GLTSMLILTTIDSHLRDKLPNISCIKAIDIYILVCLFFVFLSLLEYVYINYLFYSR-GPRRQPRRRRRPRRV-IAHYRYQQVVVG--------------  383
A0A6D2XHJ7|GABRQ_Pan_troglodytes       GLTSMLILTTIDSHLRDKLPNISCIKAIDIYILVCLFFVFLSLLEYVYINYLFYSR-GPRRQPRRRRRPRRV-IAHYRYQQVVVG--------------  383
F7F4H9|GABRQ_Macaca_mulatta            GLTSMLILTTIDSHLRDKLPNISCIKAIDIYILVCLFFVFLSLLEYVYINYLFYSR-GPRRQPRRRRRPRRV-IAHYRYQQVVVE--------------  383
M3YJY6|GABRQ_Mustela_putorius_furo     GLTSMLILNAINSHLWNKLPQVACIKAIDIYMVVCFFFVFLSLLEYVSINYLFYSRGRSRRTQRRQRRARRV-MARYRYQEVTVQ--------------  384
G3V875|GABRQ_Rattus_norvegicus         GLTSILVLTTIDSHMRDKLPHISCIKAIDIYILVCLFFVFLSLLEYVYINYLFFSQ-APRRNHRRCRKPRRV-VARYRYQEVVVA--------------  377
Q9JLF1|GABRQ_Mus_musculus              GLTSILVLTTIDSHMRDKLPHISCIKAIDIYILVCLFFVFLSLLEYVYINYLFFSQ-VPRRNHRRCRKPRRV-VARYRYQEVVVA--------------  382
F6XRX3|GABRQ_Canis_lupus_familiaris    GLTSMLILNAINSHLRDKLPQVACIKAIDIYMVVCFLFVFLSLLEYVYINYLYYSRGGSRRNQRRRRARRV-MARYRYREVMLQPFANRRSQEEMILSV  399
E1BJH4|GABRQ_Bos_taurus                GLTSMLILNSINSHLRNQFPQFSCIKAIDIYMVVCFFFVFLSLVEYVYINYLFYNRRGSRHSQRQLRRALRV-MERYRYREVIVH--------------  384
F1Q4Y6|GABRB4_Danio_rerio              GVTTVLTMTTINTHLRETLPKIPYVKAIDVYLMGCFVFVFLALLEYAFVNYVFFGR-GPQQQKKINERLNKANNERPRYEEKRLR---------------  363
                                       .......310.......320.......330.......340.......350.......360.......370.......380.......390.......400

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Q9UN88|GABRQ_Homo_sapiens              ----------------------------------------NVQDGLINVEDGV----SSLPITPAQAPLASPESLGSLTSTSEQAQLATSESLSPLTS  437
A0A2R9ALH2|GABRQ_Pan_paniscus          ----------------------------------------NVQDGLINVEDGV----SSLPITPAQAPLASPESLGSLTSTSEQAQLATSESLSPLTS  437
A0A6D2XHJ7|GABRQ_Pan_troglodytes       ----------------------------------------NVQDGLINVEDGV----SSLPITPAQAPLASPESLGSLTSTSEQAQLATSESLSPLTS  437
F7F4H9|GABRQ_Macaca_mulatta            ----------------------------------------NVQDGLINVEDGV----SSLPITPAQAPLASPESLGSLTSTSEQAQLATSESLSPLTS  437
M3YJY6|GABRQ_Mustela_putorius_furo     -------------------------------------SRAGDGTEGDQISIEDES----GSLSSRPVQACLASLESLSSLISIPGQAPLATSESLSSLSS  443
G3V875|GABRQ_Rattus_norvegicus         ----------------------------------------NVQDGLINVEDRVEDRAGPLPDSPMQAHLASQESLGSLMFTSEEAQLATSESLSLLSS  435
Q9JLF1|GABRQ_Mus_musculus              ----------------------------------------NVQDGLINVEDRVEDRAGPLPDSPMQAHLASQESLGSLVFTSEQAQLATSESLSLLSS  440
F6XRX3|GABRQ_Canis_lupus_familiaris    EEIDTHIATVDEESPLSPLPSPPPVLIHVGLANPDAINVSTISLREDQISIEDES----GSLSSRPARACLASLESLSSLTSIPGQAPLATAESLSSLSS  495
E1BJH4|GABRQ_Bos_taurus                ----------------------------------------TAFNDQVHLEDET----DSPPSSPWWAGLASPESLGSLASLTKEAPLASSESLSSLSS  438
F1Q4Y6|GABRB4_Danio_rerio              ------------------------------------------EQDSISA----------------PFQS--------------------------NTLRS  379
                                       .......410.......420.......430.......440.......450.......460.......470.......480.......490.......500
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                                     :               .  :        . .     :: :   .: .             :.  .  . :.   :
Q9UN88|GABRQ_Homo_sapiens           LSGQAPLATGESLSDLPSTSEQARHSYG-VRFNGFQADDSIFPTEIRNRVEAHGHGVTHDHEDSNESLSSDERHGHG--PSGKPMLHHGEKGVQEAGWD-  533
A0A2R9ALH2|GABRQ_Pan_paniscus       LSGQAPLATGESLSDLPSTSEQARHSYG-VRFNGFQADDSIIPTEIRNRVEAHGHGVTHDHEDSNESLSSDERHGHG--PSGKPMLHHGEKGVQEAGWD-  533
A0A6D2XHJ7|GABRQ_Pan_troglodytes    LSGQAPLATGESLSDLPSTSEQARHSYG-VRFNGFQADDSIIPTEIRNRVEAHGHGVTHDHEDSNESLSSDERHGHG--PSGKPMLHHGEKGVQEAGWD-  533
F7F4H9|GABRQ_Macaca_mulatta         LSGQAPLATGESLSDLPSTSEQARHSYG-VRFNGFQADDSIIPTEIRNRAEAHGRGVTHDHEDSDESLSSDEHHGHG--PSGKPMLHDGKKGVQEAGWD-  533
M3YJY6|GABRQ_Mustela_putorius_furo  LSEQAWTGSRESLSDRPSTSEQALHRNG-FHFSGSETDDGVICPEIHNRADAHGHADTRDPEDPKENLSSDESHGHG--PRGRYLLVYGHRCVQEASYS-  539
G3V875|GABRQ_Rattus_norvegicus      ASSQVQLATGESLGDLPSTSEQTLPDYT-IHFHGFLTNDSILPIKIRSHSDALG------DEDSEESLSSEESYGHGGSPTGRLKLQISQRCVQEASWD-  527
Q9JLF1|GABRQ_Mus_musculus           ASSQTQLATGESLSDLPSTSEQTVPECT-IHFHGFLTNDSIIPIKIHSRSDACD------DEDSEESLSSEESHGHGSSHTGRLKLQISQRCVQEASWD-  532
F6XRX3|GABRQ_Canis_lupus_familiaris LSELASLASRESLSDLPSTPEQAPHRYG-IHVNGFENDDSVTPTEMRNLAEAHDHAEISDPEDPEENFSLDESPVHI--PSGRPLLLYGHRHVREASCSL  592
E1BJH4|GABRQ_Bos_taurus             LSDGASLATTESLSDLPSTSEQVVHDDG-IRVNGIDVDNSVVPTEIRNLAETHD-----PEEDPGESSDSDESEDNG--PSKKRLLVHGQRHVQEATYEL  530
F1Q4Y6|GABRB4_Danio_rerio           YTQRRNLY-------LEEQRKVGVDAYGNILLTTLEMNNEVMPSDVGSSV----------SDSRNSVMSFDSSGVQFRKPMGSR----------------  446
                                    .......510.......520.......530.......540.......550.......560.......570.......580.......590.......600

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Q9UN88|GABRQ_Homo_sapiens           -------------L-------DDNNDKSDCLAIKEQFKCDTNSTWGLNDDELMAHGQEKD--SSSESEDSCPPSPGCSFTEGFSFDLFNPDYVPKVDKWS  611
A0A2R9ALH2|GABRQ_Pan_paniscus       -------------L-------DDNNDKSDCLAIKEQFKCDTNSTWGLNDDGLMAHGQEKD--SSSESEDSCPPSPGCSFTEGFSFDLFNPDYVPKVDKWS  611
A0A6D2XHJ7|GABRQ_Pan_troglodytes    -------------L-------DDNNDKSDCLAIKEQFKCDTNSTWGLNDDGLMAHGQEKD--SSSESEDSCPPSPGCSFTEGFSFDLFNPDYVPKVDKWS  611
F7F4H9|GABRQ_Macaca_mulatta         -------------L-------DDNSVKSGCLAVEEQLKCDTNSTWGLNEDELMAHGHEKD--NSSESEDSCPPSPGCSFTEGFSFDLFNPDYVPKVDKWS  611
M3YJY6|GABRQ_Mustela_putorius_furo  -------------LEEIRSLPDDIRVESGYLDLEKQLRRDLDSTWSLNVDGLPGFDQDKD--SNSESDDSRPPSPRCSFSEVFSSKLFDPDYVPQVDKWS  624
G3V875|GABRQ_Rattus_norvegicus      -------------IDKIESLQDDISIKSSWLGLDEQRKGDADSIWSLTDEELMACDQEKDSSSSSESEESCSPSPGCSFNEGFSFQLFNPNRVPKVDRWS  614
Q9JLF1|GABRQ_Mus_musculus           -------------LDKIEILQDDISITSSWLGLDEQCKGDADSIWSLTDEELMACDQEKD--SSSESEENCSPSPGCSFNEGFSFQLFKPNRVPKVDRWS  617
F6XRX3|GABRQ_Canis_lupus_familiaris DEIRSNLNEIRSNLDEIRSLPDDIRVESGYLDLEKQLRYDL-YNWRPNAKKFMRLNRRKD--SNSESDDSCPPSPGCSFTEGFSSKLFDPDYVPKVDKWS  689
E1BJH4|GABRQ_Bos_taurus             QEICNT-------LRDIHSLPDDVIVESGYPDLEEQLKRKVDSTGSLHSDDFMDFDGDKE--SNSESDNSFPPSPGCSFSKGFSSDLFHPDYIPKVDQCC  621
F1Q4Y6|GABRB4_Danio_rerio           -------------------------------------------------DGFSHHSLDR---SAMRSRANCRLRRRSS---KLKLKIPNLTDVSTIDKWS  491
                                    .......610.......620.......630.......640.......650.......660.......670.......680.......690.......700

                                    *::**::* .**::** *::
Q9UN88|GABRQ_Homo_sapiens           RFLFPLAFGLFNIVYWVYHMY  632
A0A2R9ALH2|GABRQ_Pan_paniscus       RFLFPLAFGLFNIVYWVYHMY  632
A0A6D2XHJ7|GABRQ_Pan_troglodytes    RFLFPLAFGLFNIVYWVYHMY  632
F7F4H9|GABRQ_Macaca_mulatta         RFLFPLAFGLFNIVYWVYHMY  632
M3YJY6|GABRQ_Mustela_putorius_furo  RILFPLAFVVFNIVYWAYHLY  645
G3V875|GABRQ_Rattus_norvegicus      RFLFPLSFGLFNVVYWLYHIY  635
Q9JLF1|GABRQ_Mus_musculus           RFLFPLSFGLFNVVYWLYHVY  638
F6XRX3|GABRQ_Canis_lupus_familiaris RILFPLAFVVFNIVYWAYHLN  710
E1BJH4|GABRQ_Bos_taurus             RLLFPLAFVVFNIVYWVYHIY  642
F1Q4Y6|GABRB4_Danio_rerio           RIIFPITFGFFNLIYWLYYVN  512
                                    .......710.......720.
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                                                                                    :::    :                                       * :  :..*:**    *.: **::
P24046|GABRR1_Homo_sapiens                     ------------------MLAVPNMRFGIFLLWWGWVLATESRMHWPGREVHEMSKKG-RPQRQRREVHEDAHKQVSPILRRSPDITKSPLTKSEQLLRI    81
A0A2R9BL98|GABRR1_Pan_paniscus                 ------------------MLAVPNMRFGIFLLWWGWVLATESRVHWPGREVHEMSKKG-RPQRQRREVHEDAHKQVSPILRRSPDITKSPLTKSEQLLRI    81
H2QTE3|GABRR1_Pan_troglodytes                  ------------------MLAVPNMRFGIFLLWWGWVLATESRVHWPGREVHEMSKKG-RPQRQRREVHEDAHKQVSPILRRSPDITKSPLTKSEQLLRI    81
F6W0N4|GABRR1_Macaca_mulatta                   ------------------MLTVPNMKFGIFLLWWGWVLATESRVHWPGREVHEMSKKGSRPQRQRREVHEDAHKQISPILRRSPDITKSPLTKSEQLLRI    82
P50572|GABRR1_Rattus_norvegicus                ------------------MLAVRNMKFGIFLLWWGWVLAAESTVHWPGREVHEPSKKGSRPQRQRRGAHDDAHKQGSPILKRSSDITKSPLTKSEQLLRI    82
P56475|GABRR1_Mus_musculus                     ------------------MLAVQNMKFGIFLLWWGWVLAAESTAHWPGREVHEPSRKGSRPQRQRRGAHDDAHKQGSPILRRSSDITKSPLTKSEQLLRI    82
M3YD49|GABRR1_Mustela_putorius_furo            ------------------MLAVQNMKVGVFILWWGWVWATESKVH-GRREVHEMSKKG------------------STILKRSPDITKSPLTKSEQLLRI    63
XP_038538749.1|GABRR1_Canis_lupus_familiaris   MGKYNRTRDMRTIWIWKPMLAVQNMKVGVFLLWWGWVLTTEGRQHWQGREAHEMSKKG------------------SPILKRSPDITKSPLTKSEQLLRI    82
Q5TZ16|GABRR1_Danio_rerio                      ------------------MHADV-----CFLLCACFLCT----------GALQSCSKS-RSKRETTRAERDLHKPGSTLLMRSPDVTKAPVTKSEQLLKL    66
Q0II76|GABRR1_Bos_taurus                       ---------------------MPYFS-RLILFLFCLVVLVESR----------------KPKKRRWTGQLETSKP-SHLYKKNPDMTKIRHGKPQPLLRV    61
                                               1.......10........20........30........40........50........60........70........80........90.......100

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P24046|GABRR1_Homo_sapiens                     DDHDFSMRPGFGGPAIPVGVDVQVESLDSISEVDMDFTMTLYLRHYWKDERLSFPSTNNLSMTFDGRLVKKIWVPDMFFVHSKRSFIHDTTTDNVMLRVQ   181
A0A2R9BL98|GABRR1_Pan_paniscus                 DDHDFSMRPGFGGPAIPVGVDVQVESLDSISEVDMDFTMTLYLRHYWKDERLSFPSTNNLSMTFDGRLVKKIWVPDMFFVHSKRSFIHDTTTDNVMLRVQ   181
H2QTE3|GABRR1_Pan_troglodytes                  DDHDFSMRPGFGGPAIPVGVDVQVESLDSISEVDMDFTMTLYLRHYWKDERLSFPSTNNLSMTFDGRLVKKIWVPDMFFVHSKRSFIHDTTTDNVMLRVQ   181
F6W0N4|GABRR1_Macaca_mulatta                   DDHDFSMRPGFGGPAIPVGVDVQVESLDSISEVDMDFTMTLYLRHYWKDERLSFPSTNNLSMTFDGRLVKKIWVPDMFFVHSKRSFIHDTTTDNVMLRVQ   182
P50572|GABRR1_Rattus_norvegicus                DDHDFSMRPGFGGPAIPVGVDVQVESLDSISEVDMDFTMTLYLRHYWKDERLSFPSTNNLSMTFDGRLVKKIWVPDMFFVHSKRSFIHDTTTDNVMLRVQ   182
P56475|GABRR1_Mus_musculus                     DDHDFSMRPGFGGPAIPVGVDVQVESLDSISEVDMDFTMTLYLRHYWKDERLSFPSSNNLSMTFDGRLVKKIWVPDMFFVHSKRSFIHDTTTDNVMLRVQ   182
M3YD49|GABRR1_Mustela_putorius_furo            DDHDFSMRPGFGGPAIPVGVDVQVESLDSISEVDMDFTMTLYLRHYWKDERLSFPSTNNLSMTFDGRLVKKIWVPDMFFVHSKRSFIHDTTTDNVMLRVQ   163
XP_038538749.1|GABRR1_Canis_lupus_familiaris   DDHDFSMRPGFGGPAIPVGVDVQVESLDSISEVDMDFTMTLYLRHYWKDERLSFPSTNNLSMTFDGRLVKKIWVPDMFFVHSKRSFIHDTTTDNVMLRVQ   182
Q5TZ16|GABRR1_Danio_rerio                      DDHDFTMRPGFGGPAIPVGVDVQLESLDTISEVDMDFTMTLYLRHYWKDARLSFPSNTNQSMTFDGRLVKKIWVPDIFFVHSKRSFIHDTTTENVMLRVH   166
Q0II76|GABRR1_Bos_taurus                       DDHDFTMRPAFGGPAIPVGVDVQVESLDSISEVDMDFTMTLYLRHYWKDERLAFPSASNKSMTFDGRLVKKIWVPDVFFVHSKRSFIHDTTTDNIMLRVF   161
                                               .......110.......120.......130.......140.......150.......160.......170.......180.......190.......200

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P24046|GABRR1_Homo_sapiens                     PDGKVLYSLRVTVTAMCNMDFSRFPLDTQTCSLEIESYAYTEDDLMLYWKKGNDSLKTDERISLSQFLIQEFHTTTKLAFYSSTGWYNRLYINFTLRRHI   281
A0A2R9BL98|GABRR1_Pan_paniscus                 PDGKVLYSLRVTVTAMCNMDFSRFPLDTQTCSLEIESYAYTEDDLMLYWKKGNDSLKTDERISLSQFLIQEFHTTTKLAFYSSTGWYNRLYINFTLRRHI   281
H2QTE3|GABRR1_Pan_troglodytes                  PDGKVLYSLRVTVTAMCNMDFSRFPLDTQTCSLEIESYAYTEDDLMLYWKKGNDSLKTDERISLSQFLIQEFHTTTKLAFYSSTGWYNRLYINFTLRRHI   281
F6W0N4|GABRR1_Macaca_mulatta                   PDGKVLYSLRVTVTAMCNMDFSRFPLDTQTCSLEIESYAYTEDDLMLYWKKGNDSLKTDERISLSQFLIQEFHTTTKLAFYSSTGWYNRLYINFTLRRHI   282
P50572|GABRR1_Rattus_norvegicus                PDGKVLYSLRVTVTAMCNMDFSRFPLDTQTCSLEIESYAYTEDDLMLYWKKGNDSLKTDERISLSQFLIQEFHTTTKLAFYSSTGWYNRLYINFTLRRHI   282
P56475|GABRR1_Mus_musculus                     PDGKVLYSLRVTVTAMCNMDFSRFPLDTQTCSLEIESYAYTEDDLMLYWKKGNDSLKTDERISLSQFLIQEFHTTTKLAFYSSTGWYNRLYINFTLRRHI   282
M3YD49|GABRR1_Mustela_putorius_furo            PDGTVLYSLRVTVTAMCNMDFSRFPLDTQTCSLEIESYAYTEDDLMLYWKKGNDSLKTDERISLSQFLIQEFHTTTKLAFYSSTGWYNRLYINFTLRRHI   263
XP_038538749.1|GABRR1_Canis_lupus_familiaris   PDGKVLYSLRVTVTAMCNMDFSRFPLDTQTCSLEIESYAYTEDDLMLYWKKGNDSLKTDERISLSQFLIQEFHTTTKLAFYSSTGWYNRLYINFTLRRHI   282
Q5TZ16|GABRR1_Danio_rerio                      PDGKVLYSLRVTVTSMCNMDLSHFPLDTQTCSLEIESYAYTDDDLMLYWKRGNKSLQTDDKISLSQFLIQEFHTTTKLAFYSSTGWYNRLYINFTLRRHI   266
Q0II76|GABRR1_Bos_taurus                       PDGQVLYSMRITVTAMCNMDFSHFPLDSQTCSLELESYAYTDEDLMLYWKNGDESLKTDEKISLSQFLIQKFHTTSRLAFYSSTGWYNRLYINFTLRRHI   261
                                               .......210.......220.......230.......240.......250.......260.......270.......280.......290.......300

                                               *********************************.*******************************:************:*******:*:*****:. ..
P24046|GABRR1_Homo_sapiens                     FFFLLQTYFPATLMVMLSWVSFWIDRRAVPARVPLGITTVLTMSTIITGVNASMPRVSYIKAVDIYLWVSFVFVFLSVLEYAAVNYLTTVQERKEQKLRE   381
A0A2R9BL98|GABRR1_Pan_paniscus                 FFFLLQTYFPATLMVMLSWVSFWIDRRAVPARVPLGITTVLTMSTIITGVNASMPRVSYIKAVDVYLWVSFVFVFLSVLEYAAVNYLTTVQERKEQKLRE   381
H2QTE3|GABRR1_Pan_troglodytes                  FFFLLQTYFPATLMVMLSWVSFWIDRRAVPARVPLGITTVLTMSTIITGVNASMPRVSYIKAVDIYLWVSFVFVFLSVLEYAAVNYLTTVQERKEQKLRE   381
F6W0N4|GABRR1_Macaca_mulatta                   FFFLLQTYFPATLMVMLSWVSFWIDRRAVPARVPLGITTVLTMSTIITGVNASMPRVSYIKAVDIYLWVSFVFVFLSVLEYAAVNYLTTVQERKEQKLRE   382
P50572|GABRR1_Rattus_norvegicus                FFFLLQTYFPATLMVMLSWVSFWIDRRAVPARVPLGITTVLTMSTIITGVNASMPRVSYIKAVDIYLWVSFVFVFLSVLEYAAVNYLTTVQERKERKLRE   382
P56475|GABRR1_Mus_musculus                     FFFLLQTYFPATLMVMLSWVSFWIDRRAVPARVPLGITTVLTMSTIITGVNASMPRVSYIKAVDIYLWVSFVFVFLSVLEYAAVNYLTTVQERKERKLRE   382
M3YD49|GABRR1_Mustela_putorius_furo            FFFLLQTYFPATLMVMLSWVSFWIDRRAVPARVPLGITTVLTMSTIITGVNASMPRVSYIKAVDIYLWVSFVFVFLSVLEYAAVNYLTTVQERKERKLRE   363
XP_038538749.1|GABRR1_Canis_lupus_familiaris   FFFLLQTYFPATLMVMLSWVSFWIDRRAVPARVPLGITTVLTMSTIITGVNASMPRVSYIKAVDIYLWVSFVFVFLSVLEYAAVNYLTTVQERKERKLRD   382
Q5TZ16|GABRR1_Danio_rerio                      FFFLLQTYFPATLMVMLSWVSFWIDRRAVPARVPLGITTVLTMSTIITGVNASMPRVSYIKAVDIYLWVSFVFVFLSVIEYAAVNYLSTLQERKERSQNS   366
Q0II76|GABRR1_Bos_taurus                       FFFLLQTYFPATLMVMLSWVSFWIDRRAVPARVSLGITTVLTMSTIITGVNASMPRVSYIKAVDIYLWVSFVFVFLSVLEYAAVNYLTTVQERKERKLQE   361
                                               .......310.......320.......330.......340.......350.......360.......370.......380.......390.......400

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P24046|GABRR1_Homo_sapiens                     K-LPCTSGLPPPRTAMLDGNYSDGEVNDLDNY-----MPE-NGEKPDRMMVQLTLASERSSPQRKSQRSSYVSMRI--DTHAIDKYSRIIFPAAYILFNL   472
A0A2R9BL98|GABRR1_Pan_paniscus                 K-LPCTSGLPPPRTVMLDGNYSDGEVNDLDNY-----MPE-NGEKPDRMMVQLTLASERSSPRRKSQRSSYVSMRI--DTHAIDKYSRIIFPAAYILFNL   472
H2QTE3|GABRR1_Pan_troglodytes                  K-LPCTSGLPPPRTVMLDGNYSDGEVNDLDNY-----VPE-NGEKPDRMMVQLTLASERSSPRRKSQRSSYVSMRI--DTHAIDKYSRIIFPAAYILFNL   472
F6W0N4|GABRR1_Macaca_mulatta                   K-LPCTSGLPPPRTVMLDGNYSDGEVNDLDNY-----MPE-NGEKPDRMMVQLTLASERSSPQRKSQRSSYVSMRI--DTHAIDKYSRIIFPAAYILFNL   473
P50572|GABRR1_Rattus_norvegicus                K-ISCTCGLPQPRGVMLDSSYSDGEVNDLGGY-----MPE-NGEKPDRMMVQLTLASERGSPQRKSQRGSYVSMRI--NTHAIDKYSRIIFPAAYILFNL   473
P56475|GABRR1_Mus_musculus                     K-ISCTCGLPQPRGVMLDSSYSDGEVNDLGGY-----MPE-NGEKPDRMMVQLTLASERGSPQRKGQRGSYVSMRI--NTHAIDKYSRIIFPAAYILFNL   473
M3YD49|GABRR1_Mustela_putorius_furo            K-LPCTCGIPQPRGVMLEGSYSDGEVNDLGSY-----IPE-NGEKPDKMMVQLTLASERSSPQRKSQRSSYVSMRI--DTHAIDKYSRIIFPAAYILFNL   454
XP_038538749.1|GABRR1_Canis_lupus_familiaris   K-LPCTCGMPQPRGVMLDGSYSDGEVNDLGNY-----TPE-NGEKPDKMMVQLTLASERSSPQRKSQRSSYVSMRI--DTHAIDKYSRIIFPAAYILFNL   473
Q5TZ16|GABRR1_Danio_rerio                      RQLPCTCGMTHPGQMMMSSSYSEMDMMTTGNYG----MSELSDTKQEQFLVHLVMDNEQGAQRPAVTTSS--TLNI--DTHAIDKYSRVIFPGAYTLFNI   458
Q0II76|GABRR1_Bos_taurus                       K-FPCMCGMLHSRTMMLDGSYSESEANSLAGYPRSHILPE--EERQDKIVVHLALSNESSSSRKKGLLKGQVGLRIFQNTHAIDKYSRLIFPASYIFFNL   458
                                               .......410.......420.......430.......440.......450.......460.......470.......480.......490.......500
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                                             ****:::
P24046|GABRR1_Homo_sapiens                   IYWSIFS-    479
A0A2R9BL98|GABRR1_Pan_paniscus               IYWSIFS-    479
H2QTE3|GABRR1_Pan_troglodytes                IYWSIFS-    479
F6W0N4|GABRR1_Macaca_mulatta                 IYWSIFS-    480
P50572|GABRR1_Rattus_norvegicus              IYWSIFS-    480
P56475|GABRR1_Mus_musculus                   IYWSIFS-    480
M3YD49|GABRR1_Mustela_putorius_furo          IYWSIFS-    461
XP_038538749.1|GABRR1_Canis_lupus_familiaris IYWSIFS-    480
Q5TZ16|GABRR1_Danio_rerio                    IYWSIYSQ    466
Q0II76|GABRR1_Bos_taurus                     IYWSVFA-    465
                                             ........
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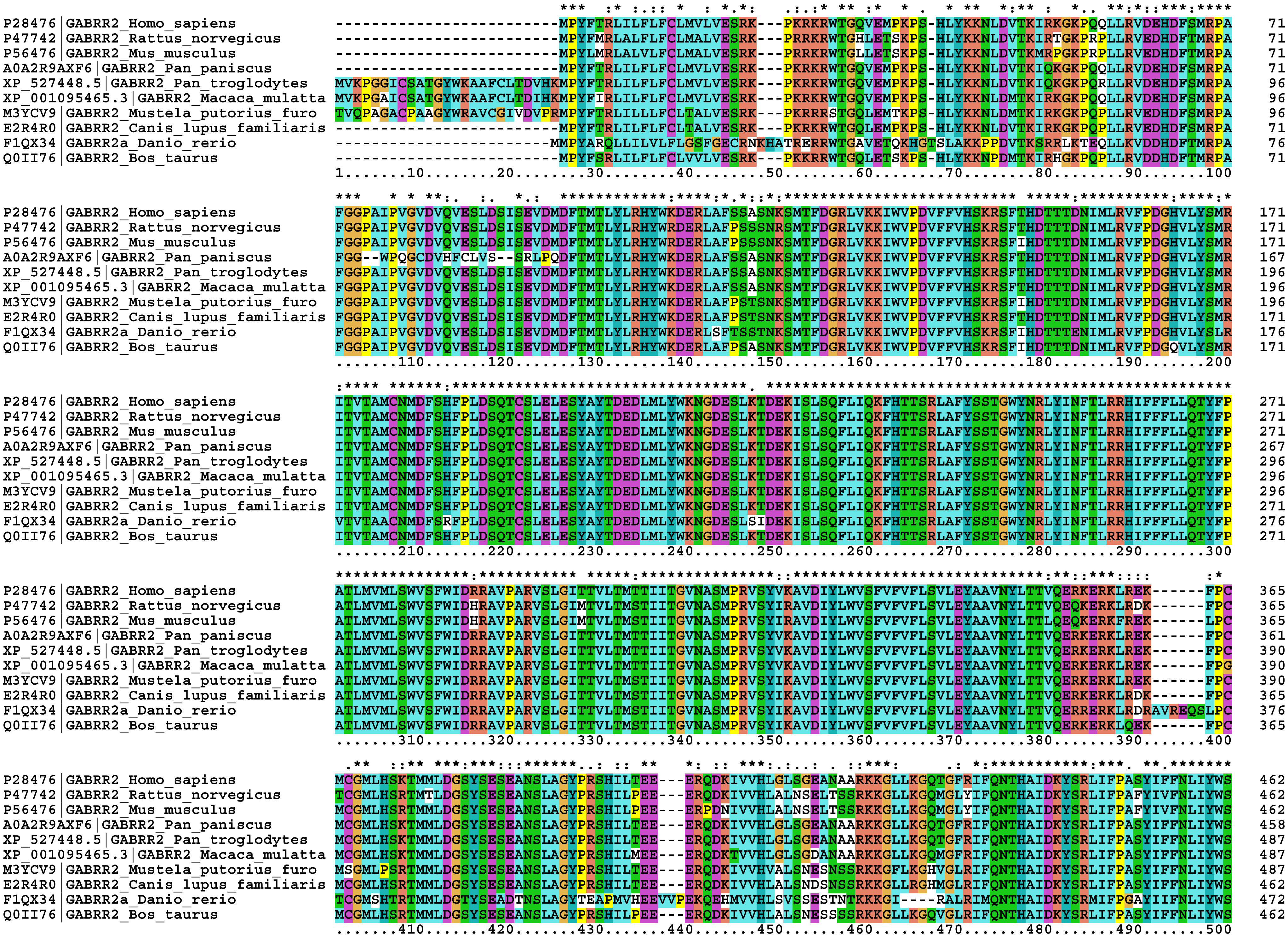
P28476|GABRR2_Homo_sapiens
P47742|GABRR2_Rattus_norvegicus
P56476|GABRR2_Mus_musculus
A0A2R9AXF6|GABRR2_Pan_paniscus
XP_527448.5|GABRR2_Pan_troglodytes
XP_001095465.3|GABRR2_Macaca_mulatta
M3YCV9|GABRR2_Mustela_putorius_furo
E2R4R0|GABRR2_Canis_lupus_familiaris
F1QX34|GABRR2a_Danio_rerio
Q0II76|GABRR2_Bos_taurus

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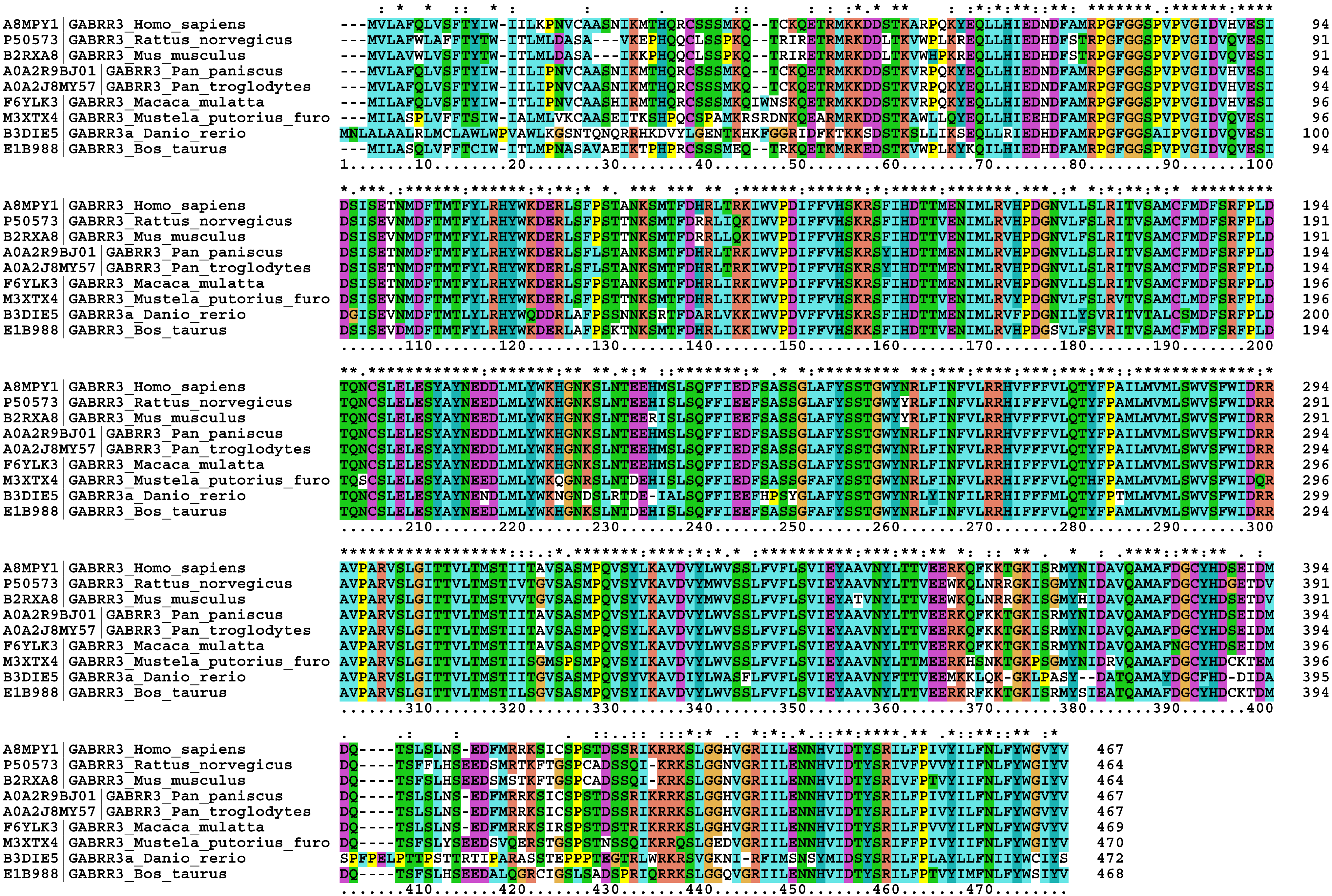
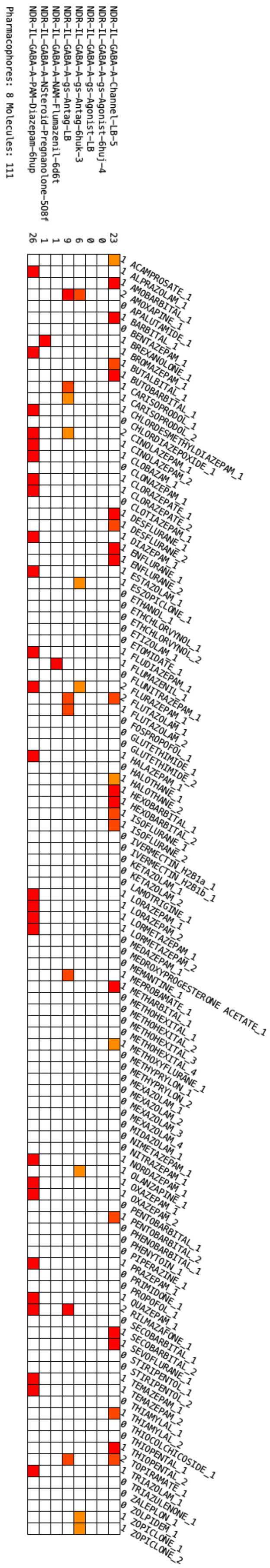
                                      *:.
P28476|GABRR2_Homo_sapiens            VFS   465
P47742|GABRR2_Rattus_norvegicus       VFS   465
P56476|GABRR2_Mus_musculus            VFS   465
A0A2R9AXF6|GABRR2_Pan_paniscus        VFS   461
XP_527448.5|GABRR2_Pan_troglodytes    VFS   490
XP_001095465.3|GABRR2_Macaca_mulatta  VFS   490
M3YCV9|GABRR2_Mustela_putorius_furo   VFS   490
E2R4R0|GABRR2_Canis_lupus_familiaris  VFS   465
F1QX34|GABRR2a_Danio_rerio            VYC   475
Q0II76|GABRR2_Bos_taurus              VFA   465
                                      ...
```


A8MPY1|GABRR3_Homo_sapiens 467
P50573|GABRR3_Rattus_norvegicus 464
B2RXA8|GABRR3_Mus_musculus 464
A0A2R9BJ01|GABRR3_Pan_paniscus 467
A0A2J8MY57|GABRR3_Pan_troglodytes 467
F6YLK3|GABRR3_Macaca_mulatta 469
M3XTX4|GABRR3_Mustela_putorius_furo 470
B3DIE5|GABRR3a_Danio_rerio 472
E1B988|GABRR3_Bos_taurus 468

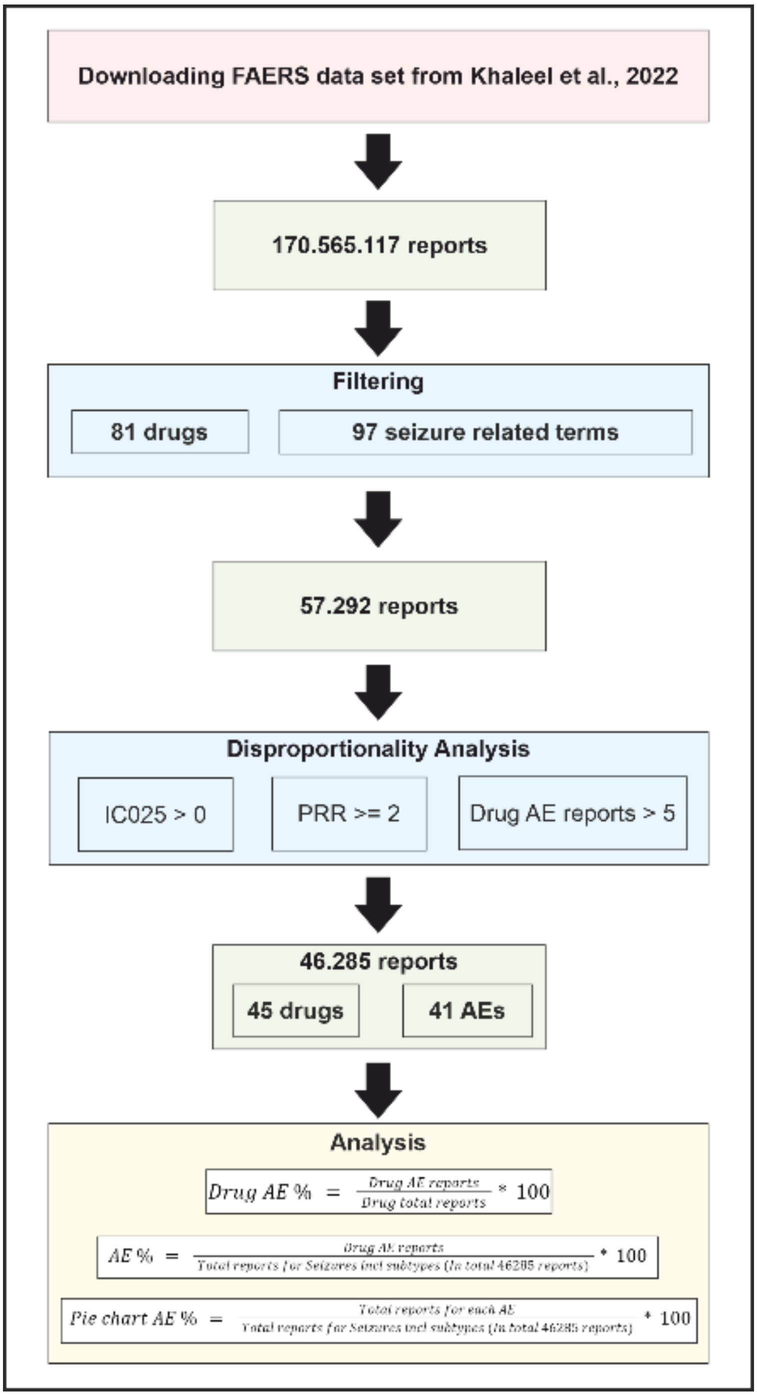
#	Name	SMILES
1	ACAMPROSATE	S(=O)(=O)(O)CCCNC(=O)C
2	ALPRAZOLAM	Clc4cc2c(n1c(nnc1C)CN=C2c3ccccc3)cc4
3	AMOBARBITAL	O=C1NC(=O)C(C(=O)N1)(CCC(C)C)CC
4	AMOXAPINE	Clc4cc2c(Oc3c(N=C2N1CCNCC1)cccc3)cc4
5	APALUTAMIDE	S=C3N(c1cnc(c(c1)C(F)(F)F)C#N)C(=O)C4(N3c2cc(F)c(cc2)C(=O)NC)CCC4
6	BARBITAL	O=C1NC(=O)C(C(=O)N1)(CC)CC
7	BENTAZEPAM	s2c1NC(=O)CN=C(c1c3c2CCCC3)c4ccccc4
8	BREXANOLONE	O=C([C@@H]4[C@@]3([C@H]([C@H]2[C@@H]([C@@]1([C@H](C[C@H](O)CC1)CC2)C)CC3)CC4)C)C
9	BROMAZEPAM	Brc3cc2c(NC(=O)CN=C2c1ncccc1)cc3
10	BUTALBITAL	O=C1NC(=O)C(C(=O)N1)(CC=C)CC(C)C
11	BUTOBARBITAL	O=C1NC(=O)C(C(=O)N1)(CCCC)CC
12	CARISOPRODOL	O=C(OCC(COC(=O)N)(CCC)C)NC(C)C
13	CHLORDESMETHYLDIAZEPAM	Clc1c(cccc1)C3=NCC(=O)Nc2c3cc(Cl)cc2
14	CHLORDIAZEPOXIDE	Clc3cc1c(N=C(NC)CN(=O)=C1c2ccccc2)cc3
15	CINOLAZEPAM	Clc3cc2c(N(C(=O)C(O)N=C2c1c(F)cccc1)CCC#N)cc3
16	CLOBAZAM	Clc3cc2N(c1ccccc1)C(=O)CC(=O)N(c2cc3)C
17	CLONAZEPAM	Clc1c(cccc1)C3=NCC(=O)Nc2c3cc([N+](=O)[O-])cc2
18	CLORAZEPATE	Clc3cc1c(NC(=O)C(N=C1c2ccccc2)C(=O)O)cc3
19	CLOTIAZEPAM	Clc1c(cccc1)C3=NCC(=O)N(c2sc(cc23)CC)C
20	DESFLURANE	FC(F)(F)C(F)OC(F)F
21	DIAZEPAM	Clc3cc1c(N(C(=O)CN=C1c2ccccc2)C)cc3
22	ENFLURANE	ClC(F)C(F)(F)OC(F)F
23	ESTAZOLAM	Clc4cc2c(n1c(nnc1)CN=C2c3ccccc3)cc4
24	ESZOPICLONE	Clc4cnc(N3C(=O)c1nccnc1[C@@H]3OC(=O)N2CCN(CC2)C)cc4
25	ETHANOL	OCC
26	ETHCHLORVYNOL	Cl¥C=C¥C(O)(C#C)CC
27	ETIZOLAM	Clc1c(cccc1)C3=NCc4nnc(n4c2sc(cc23)CC)C
28	ETOMIDATE	O=C(OCC)c1n(cnc1)[C@@H](c2ccccc2)C
29	FLUDIAZEPAM	Clc3cc2c(N(C(=O)CN=C2c1c(F)cccc1)C)cc3
30	FLUMAZENIL	Fc3cc2c(n1cnc(c1CN(C2=O)C)C(=O)OCC)cc3
31	FLUNITRAZEPAM	Fc1c(cccc1)C3=NCC(=O)N(c2c3cc([N+](=O)[O-])cc2)C
32	FLURAZEPAM	Clc3cc2c(N(C(=O)CN=C2c1c(F)cccc1)CCN(CC)CC)cc3
33	FLUTAZOLAM	Clc4cc3c(N(C(=O)CN1CCOC13c2c(F)cccc2)CCO)cc4
34	FOSPROPOFOL	P(=O)(OCOc1c(cccc1C(C)C)C(C)C)(O)O
35	GLUTETHIMIDE	O=C1NC(=O)CCC1(c2ccccc2)CC
36	HALAZEPAM	Clc3cc1c(N(C(=O)CN=C1c2ccccc2)CC(F)(F)F)cc3
37	HALOTHANE	BrC(Cl)C(F)(F)F
38	HEXOBARBITAL	O=C1N(C(=O)C(C(=O)N1)(C2=CCCCC2)C)C
39	ISOFLURANE	ClC(OC(F)F)C(F)(F)F
40a	IVERMECTIN H2B1a	O=C6O[C@@H]2C[C@@]1(O[C@@H]([C@H](CC1)C)[C@@H](CC)C)O[C@@H](C2)CC=C([C@@H](O[C@@H]4O[C@H]([C@H](O[C@@H]3O[C@H]([C@H](O)[C@@H](OC)C3)C)[C@@H](OC)C4)C)[C@H](C=CC=C5[C@]7(O)[C@H](OC5)[C@H](O)C(=C[C@@H]67)C)C)C
40b	IVERMECTIN H2B1b	O=C6O[C@@H]2C[C@@]1(O[C@@H]([C@H](CC1)C)C(C)C)O[C@@H](C2)CC=C([C@@H](O[C@@H]4O[C@H]([C@H](O[C@@H]3O[C@H]([C@H](O)[C@@H](OC)C3)C)[C@@H](OC)C4)C)[C@H](C=CC=C5[C@]7(O)[C@H](OC5)[C@H](O)C(=C[C@@H]67)C)C)C
41	KETAZOLAM	Clc4cc2c(N(C(=O)CN1C(=O)C=C(OC12c3ccccc3)C)C)cc4
42	LAMOTRIGINE	Clc1c(Cl)cccc1c2nnc(nc2N)N
43	LORAZEPAM	Clc1c(cccc1)C3=NC(O)C(=O)Nc2c3cc(Cl)cc2
44	LORMETAZEPAM	Clc1c(cccc1)C3=NC(O)C(=O)Nc2c3cc(Cl)cc2
45	MEDAZEPAM	Clc3cc1c(N(CCN=C1c2ccccc2)C)cc3
46	MEDROXYPROGESTERONE ACETATE	O=C(O[C@@]4(C(=O)C)[C@@]3([C@H]([C@H]2[C@@H]([C@@]1(C(=CC(=O)CC1)[C@H](C2)C)C)CC3)CC4)C)C
47	MEMANTINE	NC12C[C@]3(C[C@@](C1)(CC(C2)C3)C)C

48	MEPROBAMATE	O=C(OCC(COC(=O)N)(CCC)C)N
49	METHARBITAL	O=C1N(C(=O)C(C(=O)N1)(CC)CC)C
50	METHOHEXITAL	O=C1N(C(=O)C(C(=O)N1)(C(C#CCC)C)CC=C)C
51	METHOXYFLURANE	ClC(Cl)C(F)(F)OC
52	METHYPRYLON	O=C1NCC(C(=O)C1(CC)CC)C
53	MEXAZOLAM	Clc1c(cccc1)C34OCC(N4CC(O)=Nc2c3cc(Cl)cc2)C
54	MIDAZOLAM	Clc4cc3c(n1c(ncc1CN=C3c2c(F)cccc2)C)cc4
55	NIMETAZEPAM	O=[N+]([O-])c3cc1c(N(C(=O)CN=C1c2ccccc2)C)cc3
56	NITRAZEPAM	O=[N+]([O-])c3cc1c(NC(=O)CN=C1c2ccccc2)cc3
57	NORDAZEPAM	Clc3cc1c(NC(=O)CN=C1c2ccccc2)cc3
58	OLANZAPINE	s3c2Nc4c(N=C(N1CCN(CC1)C)c2cc3C)cccc4
59	OXAZEPAM	Clc3cc1c(NC(=O)C(O)N=C1c2ccccc2)cc3
60	PENTOBARBITAL	O=C1NC(=O)C(C(=O)N1)(C(CCC)C)CC
61	PHENOBARBITAL	O=C2NC(=O)C(c1ccccc1)(C(=O)N2)CC
62	PHENYTOIN	O=C1NC(=O)C(N1)(c2ccccc2)c3ccccc3
63	PIPERAZINE	N1CCNCC1
64	PRAZEPAM	Clc4cc1c(N(C(=O)CN=C1c2ccccc2)CC3CC3)cc4
65	PRIMIDONE	O=C2NCNC(=O)C2(c1ccccc1)CC
66	PROPOFOL	Oc1c(cccc1C(C)C)C(C)C
67	QUAZEPAM	Clc3cc2c(N(C(=S)CN=C2c1c(F)cccc1)CC(F)(F)F)cc3
68	RILMAZAFONE	Clc1c(cccc1)C(=O)c3c(n2nc(nc2CNC(=O)CN)C(=O)N(C)C)ccc(Cl)c3
69	SECOBARBITAL	O=C1NC(=O)C(C(=O)N1)(C(CCC)C)CC=C
70	SEVOFLURANE	FC(F)(F)C(OCF)C(F)(F)F
71	STIRIPENTOL	O1c2c(OC1)ccc(c2)¥C=C¥C(O)C(C)(C)C
72	TEMAZEPAM	Clc3cc1c(N(C(=O)C(O)N=C1c2ccccc2)C)cc3
73	THIAMYLAL	S=C1NC(=O)C(C(=O)N1)(C(CCC)C)CC=C
74	THIOCOLCHICOSIDE	S(c4c(=O)cc3c(c2c(OC)c(OC)c(O[C@@H]1O[C@@H]([C@@H](O)[C@H](O)[C@H]1O)CO)cc2CC[C@@H]3NC(=O)C)cc4)C
75	THIOPENTAL	S=C1NC(=O)C(C(=O)N1)(C(CCC)C)CC
76	TOPIRAMATE	S(=O)(=O)(OC[C@]13OC(O[C@H]1[C@@H]2OC(O[C@@H]2CO3)(C)C)(C)C)N
77	TRIAZOLAM	Clc1c(cccc1)C4=NCc2nnc(n2c3c4cc(Cl)cc3)C
78	TRIAZULENONE	Clc1c(cccc1)C3=NCC4N(c2c3cc(N(=O)=O)cc2)C(=O)/C(/N=4)=C/N5CCN(CC5)C
79	ZALEPLON	O=C(N(c3cc(c2n1ncc(c1ncc2)C#N)ccc3)CC)C
80	ZOLPIDEM	O=C(N(C)C)Cc1n3c(nc1c2ccc(cc2)C)ccc(c3)C
81	ZOPICLONE	Clc4cnc(N3C(=O)c1nccnc1C3OC(=O)N2CCN(CC2)C)cc4

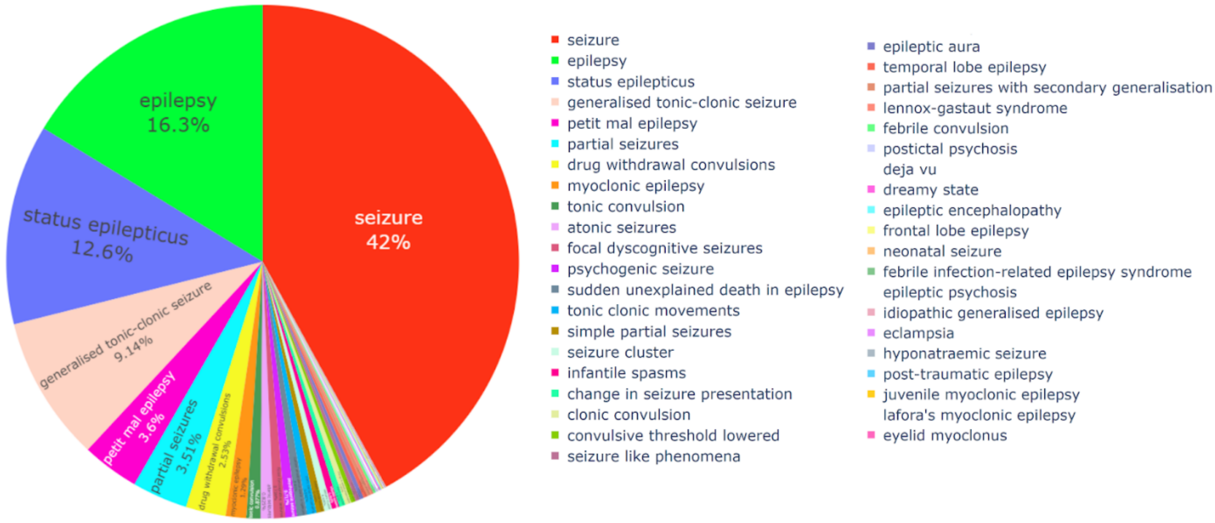
Supplementary Table S3: All drugs selected in this study and their respective SMILES codes.



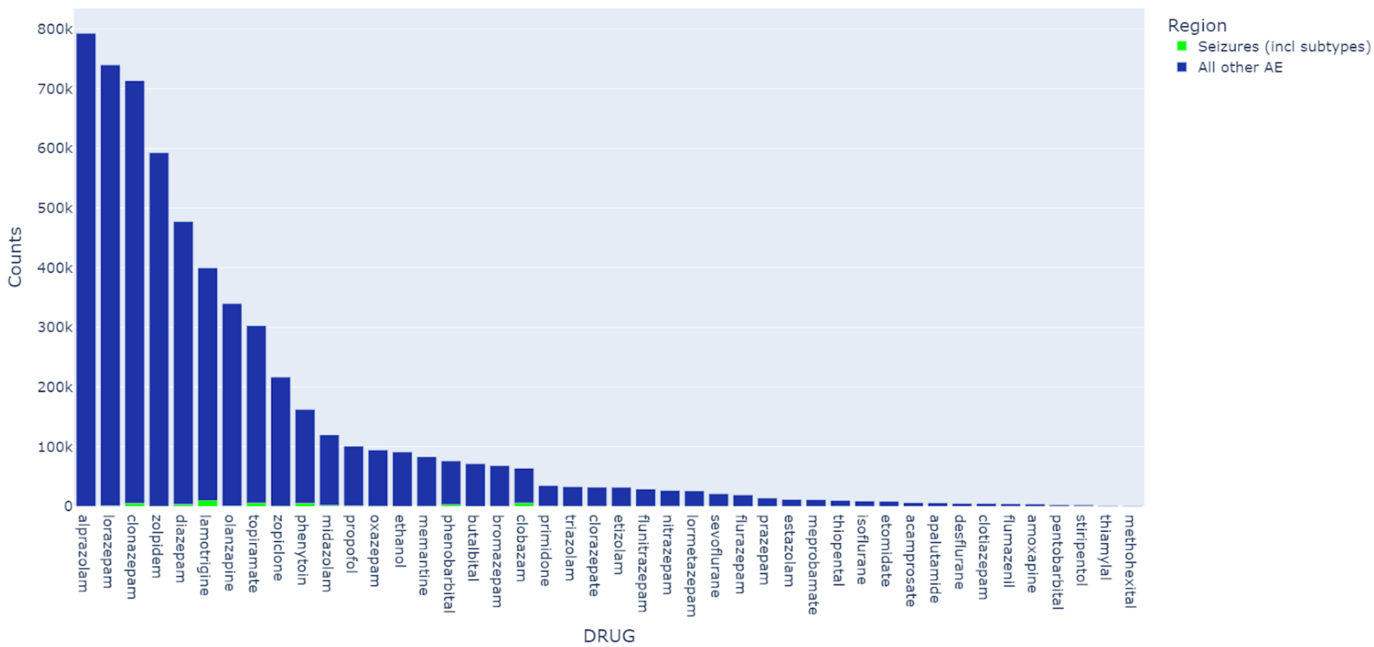
Supplementary Figure S9: Results from *in silico* profiling of 81 Drugs using selected GABAA models in the NeuroDeRisk IL Profiler V1.0.



Supplementary Figure S10. Workflow of the FAERS analysis. Greenish boxes contain the results of the filtering processes, blue boxes describe the filtering process which was used for the FAERS data set obtained from Khaleel et al., 2022 (Khaleel *et al.*, 2022).



Supplementary Figure S11: Distribution of all seizure related reports from our analysis which are in total 46285 reports.



Supplementary Figure S12. Total report number for all drugs used, which are displayed in Figure 4. All seizure-category reports are displayed in green, all other adverse event terms as a lump sum in blue.

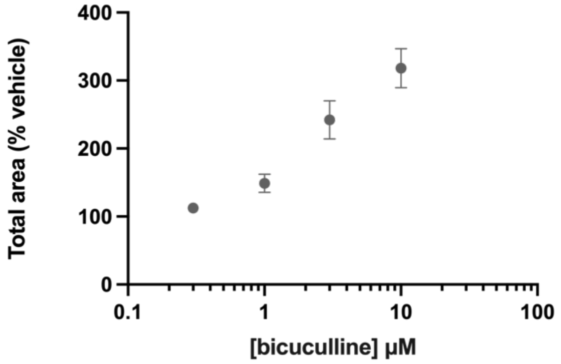
Drug	Total reports	AE	AE reports
ACAMPROSATE	587	SEIZURE	17
ALPRAZOLAM	30031	SEIZURE	410
		EPILEPSY	71
		STATUS EPILEPTICUS	44
		DRUG WITHDRAWAL CONVULSIONS	29
AMOXAPINE	134	SEIZURE	9
BROMAZEPAM	3047	SEIZURE	26
		STATUS EPILEPTICUS	14
		EPILEPSY	13
CARISOPRODOL	3493	SEIZURE	56
CHLORDIAZEPOXIDE	756	SEIZURE	14
CLOBAZAM	2070	SEIZURE	325
		DRUG WITHDRAWAL CONVULSIONS	76
		EPILEPSY	55
		STATUS EPILEPTICUS	51
		PARTIAL SEIZURES	19
		FOCAL DYSCOGNITIVE SEIZURES	12
		PETIT MAL EPILEPSY	11
CLONAZEPAM	18338	SEIZURE	498
		STATUS EPILEPTICUS	66
		EPILEPSY	59
		DRUG WITHDRAWAL CONVULSIONS	29
		PARTIAL SEIZURES	18
		PETIT MAL EPILEPSY	18
CLORAZEPATE DIPOTASSIUM	1205	SEIZURE	16
DESFLURANE	509	SEIZURE LIKE PHENOMENA	12
DIAZEPAM	20293	SEIZURE	345
		STATUS EPILEPTICUS	88
		DRUG WITHDRAWAL CONVULSIONS	69
		EPILEPSY	58
		PARTIAL SEIZURES	12
		PETIT MAL EPILEPSY	9
ESTAZOLAM	263	SEIZURE	16

ESZOPICLONE	7811	SEIZURE	22
ETHANOL	13742	SEIZURE	164
		STATUS EPILEPTICUS	14
		EPILEPSY	14
		PARTIAL SEIZURES	12
ETIZOLAM	630	SEIZURE	13
FLUMAZENIL	248	SEIZURE	51
		DRUG WITHDRAWAL CONVULSIONS	42
FLUNITRAZEPAM	875	SEIZURE	28
ISOFLURANE	502	STATUS EPILEPTICUS	33
IVERMECTIN	1764	SEIZURE	21
LAMOTRIGINE	26598	SEIZURE	1723
		EPILEPSY	228
		STATUS EPILEPTICUS	189
		PETIT MAL EPILEPSY	140
		PARTIAL SEIZURES	76
		FOCAL DYSCOGNITIVE SEIZURES	54
		SUDDEN UNEXPLAINED DEATH IN EPILEPSY	41
		MYOCLONIC EPILEPSY	33
LORAZEPAM	16200	SEIZURE	291
		STATUS EPILEPTICUS	79
		EPILEPSY	76
		DRUG WITHDRAWAL CONVULSIONS	27
		PETIT MAL EPILEPSY	10
		PARTIAL SEIZURES	9
LORMETAZEPAM	1347	EPILEPSY	14
		SEIZURE	8
MEDROXYPROGESTERONE ACETATE	10874	SEIZURE	34
		EPILEPSY	12
MEMANTINE	6510	SEIZURE	190
		EPILEPSY	61
		PETIT MAL EPILEPSY	16
		PARTIAL SEIZURES	12
		STATUS EPILEPTICUS	8
MEPROBAMATE	781	SEIZURE	10

MIDAZOLAM	4298	SEIZURE	123
		STATUS EPILEPTICUS	57
		EPILEPSY	21
		SEIZURE LIKE PHENOMENA	15
		FOCAL DYSCOGNITIVE SEIZURES	13
		PETIT MAL EPILEPSY	13
		PARTIAL SEIZURES	12
NITRAZEPAM	627	SEIZURE	13
OLANZAPINE	28785	SEIZURE	554
		EPILEPSY	137
		STATUS EPILEPTICUS	42
		PARTIAL SEIZURES	18
		PETIT MAL EPILEPSY	17
OXAZEPAM	4627	SEIZURE	53
		STATUS EPILEPTICUS	17
		EPILEPSY	14
PHENOBARBITAL	2287	SEIZURE	143
		STATUS EPILEPTICUS	80
		EPILEPSY	35
		PARTIAL SEIZURES	15
		MYOCLONIC EPILEPSY	8
PHENYTOIN	10126	SEIZURE	972
		STATUS EPILEPTICUS	204
		EPILEPSY	100
		PARTIAL SEIZURES	59
		PETIT MAL EPILEPSY	42
		FOCAL DYSCOGNITIVE SEIZURES	23
		SUDDEN UNEXPLAINED DEATH IN EPILEPSY	13
PRIMIDONE	711	SEIZURE	51
		EPILEPSY	17
		PETIT MAL EPILEPSY	14
		MYOCLONIC EPILEPSY	11
PROPOFOL	7869	SEIZURE	165
		STATUS EPILEPTICUS	70
		EPILEPSY	27
		SEIZURE LIKE PHENOMENA	25

		MYOCLONIC EPILEPSY	12
SEVOFLURANE	1650	SEIZURE	23
	1650	SEIZURE LIKE PHENOMENA	11
TEMAZEPAM	3519	SEIZURE	46
THIOPENTAL	530	STATUS EPILEPTICUS	30
	530	SEIZURE	20
TOPIRAMATE	11307	SEIZURE	562
	11307	EPILEPSY	85
	11307	STATUS EPILEPTICUS	82
	11307	PETIT MAL EPILEPSY	59
	11307	FOCAL DYSCOGNITIVE SEIZURES	31
	11307	PARTIAL SEIZURES	30
	11307	DRUG WITHDRAWAL CONVULSIONS	21
	11307	MYOCLONIC EPILEPSY	21
	11307	SUDDEN UNEXPLAINED DEATH IN EPILEPSY	16
	11307	SEIZURE LIKE PHENOMENA	9
TRIAZOLAM	1431	SEIZURE	12
ZOLPIDEM	18040	SEIZURE	248
		DRUG WITHDRAWAL CONVULSIONS	39
		EPILEPSY	38
		STATUS EPILEPTICUS	22
ZOPICLONE	5399	SEIZURE	70
		EPILEPSY	28
		STATUS EPILEPTICUS	12

Supplementary Table S4: Showing all total reports and the reports per AE per drug for the selected drugs and AEs.



Supplementary Figure S13: Mean (± SEM) concentration-response data summary of bicuculline (0.3, 1, 3, and 10 µM) effects on CA1 population spike area in rat hippocampal brain slices. Statistical testing was run on raw concentration data (dose: $F(4, 27)=57.10$ $p<.001$; Dunnett's posthoc: bicuculline 1, 3, and 10 µM vs veh, $p<.001$).

BEHAVIORAL ALTERATIONS		
Amoxapine (mM)	**Occurrence**	**%**
0	0/8	0%
0.003	0/8	0%
0.01	0/8	0%
0.03	3/8	38%
0.1	8/8	100%
0.3	8/8	100%

MORTALITY		
Amoxapine (mM)	**Occurrence**	**%**
0	0/8	0%
0.003	0/8	0%
0.01	0/8	0%
0.03	0/8	0%
0.1	0/8	0%
0.3	2/8	25%

Supplementary Table S5: Behavioral alterations (abnormal swimming behaviour, like circling or erratic movements) and mortality of zebrafish larvae.

	1.58µM	**5µM**	**15.8µM**	**50µM**
Firing Rate	**146**	**173**	**145**	76
Median Burst Rate	104	127	**160**	**158**
Median Num of Spikes in Burst	**296**	**303**	183	92
Percent Isolated Spikes	72	**14**	**9**	**35**
ISI CV	135	**170**	129	92
Normalized IQR Burst Duration	73	**27**	**42**	**49**
Median Burst Duration	**145**	**124**	71	57
Mean Interburst Interval	97	79	**65**	**66**
Mean of ISI-distance	75	**41**	**36**	**39**
Normalized MAD Burst Spike	**59**	**22**	**25**	**38**
Median/Mean ISI	101	76	70	89
Median ISI	**67**	**32**	**37**	76

Supplementary Table S6: Heatmap of bicuculline from MEA recordings from rat cortical neurons showing % changes versus vehicle in each parameter. Green color indicates an increase in a parameter, red color indicates a decrease. A value of 100% corresponds to no change from vehicle (pure white color). The intensity of the color in a box increases with the magnitude of the change. Statistically significant values compared to vehicle are indicated in bold.

Subunit	Species	Sequence ID	Database	Date accessed
GABRA1	Homo sapiens	P14867	UniProt	11.04.2022
GABRA1	Pan paniscus	A0A2R9AU65	UniProt	11.04.2022
GABRA1	Mustela putorius furo	M3YP62	UniProt	11.04.2022
GABRA1	Rattus norvegicus	P62813	UniProt	11.04.2022
GABRA1	Mus musculus	P62812	UniProt	11.04.2022
GABRA1	Canis lupus familiaris	E2RSP8	UniProt	11.04.2022
GABRA1	Pan troglodytes	A0A2I3SDV7	UniProt	11.04.2022
GABRA1	Danio rerio	Q08BJ3	UniProt	11.04.2022
GABRA1	Macaca mulatta	A0A1D5Q406	UniProt	11.04.2022
GABRA1	Bos taurus	P08219	UniProt	11.04.2022
GABRA2	Homo sapiens	P47869	UniProt	11.04.2022
GABRA2	Pan paniscus	XP_003816076.1	NCBI Protein	11.04.2022
GABRA2	Mustela putorius furo	M3Y0V9	UniProt	11.04.2022
GABRA2	Rattus norvegicus	P23576	UniProt	11.04.2022
GABRA2	Mus musculus	P26048	UniProt	11.04.2022

GABRA2	**Canis lupus familiaris**	**A0A5F4CYJ7**	**UniProt**	**11.04.2022**
GABRA2	**Pan troglodytes**	**H2QPE5**	**UniProt**	**11.04.2022**
GABRA2a	**Danio rerio**	**E7F635**	**UniProt**	**11.04.2022**
GABRA2	**Macaca mulatta**	**XP_028703913.1**	**NCBI Protein**	**11.04.2022**
GABRA2	**Bos taurus**	**P10063**	**UniProt**	**11.04.2022**

GABRA3	**Homo sapiens**	**P34903**	**UniProt**	**11.04.2022**
GABRA3	**Pan paniscus**	**A0A2R9AHX2**	**UniProt**	**11.04.2022**
GABRA3	**Mustela putorius furo**	**M3YJX4**	**UniProt**	**11.04.2022**
GABRA3	**Rattus norvegicus**	**P20236**	**UniProt**	**11.04.2022**
GABRA3	**Mus musculus**	**P26049**	**UniProt**	**11.04.2022**
GABRA3	**Canis lupus familiaris**	**E2QVJ7**	**UniProt**	**11.04.2022**
GABRA3	**Pan troglodytes**	**H2QZ88**	**UniProt**	**11.04.2022**
GABRA3	**Danio rerio**	**A0A2R8QC15**	**UniProt**	**11.04.2022**
GABRA3	**Macaca mulatta**	**F7G8R5**	**UniProt**	**11.04.2022**
GABRA3	**Bos taurus**	**P10064**	**UniProt**	**11.04.2022**

GABRA4	**Homo sapiens**	**P48169**	**UniProt**	**11.04.2022**
GABRA4	**Pan paniscus**	**A0A2R9C6M3**	**UniProt**	**11.04.2022**
GABRA4	**Mustela putorius furo**	**M3Y0X5**	**UniProt**	**11.04.2022**
GABRA4	**Rattus norvegicus**	**P28471**	**UniProt**	**11.04.2022**
GABRA4	**Mus musculus**	**Q9D6F4**	**UniProt**	**11.04.2022**
GABRA4	**Canis lupus familiaris**	**F1P9P9**	**UniProt**	**11.04.2022**
GABRA4	**Pan troglodytes**	**H2QPE7**	**UniProt**	**11.04.2022**
GABRA4	**Danio rerio**	**Q568M9**	**UniProt**	**11.04.2022**
GABRA4	**Macaca mulatta**	**F6YLN4**	**UniProt**	**11.04.2022**
GABRA4	**Bos taurus**	**P20237**	**UniProt**	**11.04.2022**

GABRA5	**Homo sapiens**	**P31644**	**UniProt**	**11.04.2022**
GABRA5	**Pan paniscus**	**A0A2R9BFW8**	**UniProt**	**11.04.2022**
GABRA5	**Mustela putorius furo**	**M3XNJ0**	**UniProt**	**11.04.2022**
GABRA5	**Rattus norvegicus**	**P19969**	**UniProt**	**11.04.2022**
GABRA5	**Mus musculus**	**Q8BHJ7**	**UniProt**	**11.04.2022**
GABRA5	**Canis lupus familiaris**	**E2RG38**	**UniProt**	**11.04.2022**

GABRA5	**Pan troglodytes**	**H2Q919**	**UniProt**	**11.04.2022**
GABRA5	**Danio rerio**	**E9QE70**	**UniProt**	**11.04.2022**
GABRA5	**Macaca mulatta**	**G7MW76**	**UniProt**	**11.04.2022**
GABRA5	**Bos taurus**	**Q08E50**	**UniProt**	**11.04.2022**
GABRA6	**Homo sapiens**	**Q16445**	**UniProt**	**11.04.2022**
GABRA6	**Pan paniscus**	**A0A2R8ZQ75**	**UniProt**	**11.04.2022**
GABRA6	**Mustela putorius furo**	**M3YP65**	**UniProt**	**11.04.2022**
GABRA6	**Rattus norvegicus**	**P30191**	**UniProt**	**11.04.2022**
GABRA6	**Mus musculus**	**P16305**	**UniProt**	**11.04.2022**
GABRA6	**Canis lupus familiaris**	**E2RSN3**	**UniProt**	**11.04.2022**
GABRA6	**Pan troglodytes**	**H2QRY3**	**UniProt**	**11.04.2022**
GABRA6b	**Danio rerio**	**F1QGW0**	**UniProt**	**11.04.2022**
GABRA6	**Macaca mulatta**	**A0A1D5RL15**	**UniProt**	**11.04.2022**
GABRA6	**Bos taurus**	**E1BE96**	**UniProt**	**11.04.2022**
GABRB1	**Homo sapiens**	**P18505**	**UniProt**	**11.04.2022**

GABRB1	**Pan paniscus**	**A0A2R9C4H8**	**UniProt**	**11.04.2022**
GABRB1	**Mustela putorius furo**	**M3Y0Y5**	**UniProt**	**11.04.2022**
GABRB1	**Rattus norvegicus**	**P15431**	**UniProt**	**11.04.2022**
GABRB1	**Mus musculus**	**P50571**	**UniProt**	**11.04.2022**
GABRB1	**Canis lupus familiaris**	**F1PEG2**	**UniProt**	**11.04.2022**
GABRB1	**Pan troglodytes**	**H2QPE8**	**UniProt**	**11.04.2022**
GABRB1	**Danio rerio**	**F1QPW7**	**UniProt**	**11.04.2022**
GABRB1	**Macaca mulatta**	**G7MSU9**	**UniProt**	**11.04.2022**
GABRB1	**Bos taurus**	**P08220**	**UniProt**	**11.04.2022**

GABRB2	**Homo sapiens**	**P47870**	**UniProt**	**11.04.2022**
GABRB2	**Pan paniscus**	**A0A2R9C8C8**	**UniProt**	**11.04.2022**
GABRB2	**Mustela putorius furo**	**M3YP67**	**UniProt**	**11.04.2022**
GABRB2	**Rattus norvegicus**	**P63138**	**UniProt**	**11.04.2022**
GABRB2	**Mus musculus**	**P63137**	**UniProt**	**11.04.2022**
GABRB2	**Canis lupus familiaris**	**A0A5F4CKV8**	**UniProt**	**11.04.2022**
GABRB2	**Pan troglodytes**	**A0A2J8NKG3**	**UniProt**	**11.04.2022**

GABRB2	Danio rerio	Q9DDD9	UniProt	11.04.2022
GABRB2	Macaca mulatta	D1LYT2	UniProt	11.04.2022
GABRB2	Bos taurus	P0C2W5	UniProt	11.04.2022
GABRB3	Homo sapiens	P28472	UniProt	11.04.2022
GABRB3	Pan paniscus	A0A2R9BWE5	UniProt	11.04.2022
GABRB3	Mustela putorius furo	M3XNI6	UniProt	11.04.2022
GABRB3	Rattus norvegicus	P63079	UniProt	11.04.2022
GABRB3	Mus musculus	P63080	UniProt	11.04.2022
GABRB3	Canis lupus familiaris	J9P3X1	UniProt	11.04.2022
GABRB3	Pan troglodytes	A0A2I3THQ1	UniProt	11.04.2022
GABRB3	Danio rerio	A0A0R4ILP2	UniProt	11.04.2022
GABRB3	Macaca mulatta	F6ZKJ4	UniProt	11.04.2022
GABRB3	Bos taurus	A5D7U6	UniProt	11.04.2022
GABRD	Homo sapiens	O14764	UniProt	11.04.2022
GABRD	Pan paniscus	A0A2R9B726	UniProt	11.04.2022

GABRD	**Mustela putorius furo**	**XP_004768807.1**	**NCBI Protein**	**11.04.2022**
GABRD	**Rattus norvegicus**	**P18506**	**UniProt**	**11.04.2022**
GABRD	**Mus musculus**	**P22933**	**UniProt**	**11.04.2022**
GABRD	**Canis lupus familiaris**	**E2R3M6**	**UniProt**	**11.04.2022**
GABRD	**Pan troglodytes**	**A0A2J8K8J4**	**UniProt**	**11.04.2022**
GABRD	**Danio rerio**	**E9QHL0**	**UniProt**	**11.04.2022**
GABRD	**Macaca mulatta**	**F6QDC4**	**UniProt**	**11.04.2022**
GABRD	**Bos taurus**	**A0A3Q1LQH4**	**UniProt**	**11.04.2022**

GABRE	**Homo sapiens**	**P78334**	**UniProt**	**11.04.2022**
GABRE	**Pan paniscus**	**A0A2R8ZSJ9**	**UniProt**	**11.04.2022**
GABRE	**Mustela putorius furo**	**M3YJW6**	**UniProt**	**11.04.2022**
GABRE	**Rattus norvegicus**	**Q9ES14**	**UniProt**	**11.04.2022**
GABRE	**Mus musculus**	**Q9JLE8**	**UniProt**	**11.04.2022**
GABRE	**Canis lupus familiaris**	**E2QVM5**	**UniProt**	**11.04.2022**
GABRE	**Pan troglodytes**	**H2QZ86**	**UniProt**	**11.04.2022**
-	**Danio rerio**	**-**	**-**	**-**

GABRE	**Macaca mulatta**	**F7GJ80**	**UniProt**	**11.04.2022**
GABRE	**Bos taurus**	**G3MWU9**	**UniProt**	**11.04.2022**
GABRG1	**Homo sapiens**	**Q8N1C3**	**UniProt**	**11.04.2022**
GABRG1	**Pan paniscus**	**A0A2R9A1L5**	**UniProt**	**11.04.2022**
GABRG1	**Mustela putorius furo**	**M3Y0V1**	**UniProt**	**11.04.2022**
GABRG1	**Rattus norvegicus**	**P23574**	**UniProt**	**11.04.2022**
GABRG1	**Mus musculus**	**Q9R0Y8**	**UniProt**	**11.04.2022**
GABRG1	**Canis lupus familiaris**	**E2RH22**	**UniProt**	**11.04.2022**
GABRG1	**Pan troglodytes**	**H2QPE4**	**UniProt**	**11.04.2022**
GABRG1	**Danio rerio**	**A0A0R4IPF9**	**UniProt**	**11.04.2022**
GABRG1	**Macaca mulatta**	**A0A5F8A0N7**	**UniProt**	**11.04.2022**
GABRG1	**Bos taurus**	**F6Q4V7**	**UniProt**	**11.04.2022**
GABRG2	**Homo sapiens**	**P18507-2**	**UniProt**	**11.04.2022**
GABRG2	**Pan paniscus**	**A0A2R9CM76**	**UniProt**	**11.04.2022**
GABRG2	**Mustela putorius furo**	**XP_004737727.1**	**NCBI Protein**	**11.04.2022**

GABRG2	**Rattus norvegicus**	**NP_001380704.1**	**NCBI Protein**	**11.04.2022**
GABRG2	**Mus musculus**	**P22723**	**UniProt**	**11.04.2022**
GABRG2	**Canis lupus familiaris**	**E2RSQ1**	**UniProt**	**11.04.2022**
GABRG2	**Pan troglodytes**	**A0A2I3RPH6**	**UniProt**	**11.04.2022**
GABRG2	**Danio rerio**	**F1RDP2**	**UniProt**	**11.04.2022**
GABRG2	**Macaca mulatta**	**F7A3C2**	**UniProt**	**11.04.2022**
GABRG2	**Bos taurus**	**P22300**	**UniProt**	**11.04.2022**

GABRG3	**Homo sapiens**	**Q99928**	**UniProt**	**11.04.2022**
GABRG3	**Pan paniscus**	**A0A2R9BD98**	**UniProt**	**11.04.2022**
GABRG3	**Mustela putorius furo**	**XP_044926358.1**	**NCBI Protein**	**11.04.2022**
GABRG3	**Rattus norvegicus**	**P28473**	**UniProt**	**11.04.2022**
GABRG3	**Mus musculus**	**P27681**	**UniProt**	**11.04.2022**
GABRG3	**Canis lupus familiaris**	**F1PHI6**	**UniProt**	**11.04.2022**
GABRG3	**Pan troglodytes**	**A0A2I3TTK9**	**UniProt**	**11.04.2022**
GABRG3	**Danio rerio**	**XP_009300843.1**	**NCBI Protein**	**11.04.2022**

GABRG3	Macaca mulatta	F7HI54	UniProt	11.04.2022
GABRG3	Bos taurus	XP_024838027.1	NCBI Protein	11.04.2022
GABRP	Homo sapiens	O00591	UniProt	11.04.2022
GABRP	Pan paniscus	A0A2R8ZJ77	UniProt	11.04.2022
GABRP	Mustela putorius furo	M3YN80	UniProt	11.04.2022
GABRP	Rattus norvegicus	O09028	UniProt	11.04.2022
GABRP	Mus musculus	Q8QZW7	UniProt	11.04.2022
GABRP	Canis lupus familiaris	E2RS87	UniProt	11.04.2022
GABRP	Pan troglodytes	A0A2J8LXV6	UniProt	11.04.2022
GABRP	Danio rerio	XP_021330000.1	NCBI Protein	11.04.2022
GABRP	Macaca mulatta	A0A1D5QJZ6	UniProt	11.04.2022
GABRP	Bos taurus	Q5EA06	UniProt	11.04.2022
GABRQ	Homo sapiens	Q9UN88	UniProt	11.04.2022
GABRQ	Pan paniscus	A0A2R9ALH2	UniProt	11.04.2022
GABRQ	Mustela putorius furo	M3YJY6	UniProt	11.04.2022

GABRQ	Rattus norvegicus	G3V875	UniProt	11.04.2022
GABRQ	Mus musculus	Q9JLF1	UniProt	11.04.2022
GABRQ	Canis lupus familiaris	F6XRX3	UniProt	11.04.2022
GABRQ	Pan troglodytes	A0A6D2XHJ7	UniProt	11.04.2022
GABRB4	Danio rerio	F1Q4Y6	UniProt	11.04.2022
GABRQ	Macaca mulatta	F7F4H9	UniProt	11.04.2022
GABRQ	Bos taurus	E1BJH4	UniProt	11.04.2022

GABRR1	Homo sapiens	P24046	UniProt	11.04.2022
GABRR1	Pan paniscus	A0A2R9BL98	UniProt	11.04.2022
GABRR1	Mustela putorius furo	M3YD49	UniProt	11.04.2022
GABRR1	Rattus norvegicus	P50572	UniProt	11.04.2022
GABRR1	Mus musculus	P56475	UniProt	11.04.2022
GABRR1	Canis lupus familiaris	XP_038538749.1	NCBI Protein	11.04.2022
GABRR1	Pan troglodytes	H2QTE3	UniProt	11.04.2022
GABRR1	Danio rerio	Q5TZ16	UniProt	11.04.2022

GABRR1	**Macaca mulatta**	**F6W0N4**	**UniProt**	**11.04.2022**
GABRR1	**Bos taurus**	**Q0II76**	**UniProt**	**11.04.2022**

GABRR2	**Homo sapiens**	**P28476**	**UniProt**	**11.04.2022**
GABRR2	**Pan paniscus**	**A0A2R9AXF6**	**UniProt**	**11.04.2022**
GABRR2	**Mustela putorius furo**	**M3YCV9**	**UniProt**	**11.04.2022**
GABRR2	**Rattus norvegicus**	**P47742**	**UniProt**	**11.04.2022**
GABRR2	**Mus musculus**	**P56476**	**UniProt**	**11.04.2022**
GABRR2	**Canis lupus familiaris**	**E2R4R0**	**UniProt**	**11.04.2022**
GABRR2	**Pan troglodytes**	**XP_527448.5**	**NCBI Protein**	**11.04.2022**
GABRR2a	**Danio rerio**	**F1QX34**	**UniProt**	**11.04.2022**
GABRR2	**Macaca mulatta**	**XP_001095465.3**	**NCBI Protein**	**11.04.2022**
GABRR2	**Bos taurus**	**Q0II76**	**UniProt**	**11.04.2022**

GABRR3	**Homo sapiens**	**A8MPY1**	**UniProt**	**11.04.2022**
GABRR3	**Pan paniscus**	**A0A2R9BJ01**	**UniProt**	**11.04.2022**
GABRR3	**Mustela putorius furo**	**M3XTX4**	**UniProt**	**11.04.2022**

GABRR3	**Rattus norvegicus**	**P50573**	**UniProt**	**11.04.2022**
GABRR3	**Mus musculus**	**B2RXA8**	**UniProt**	**11.04.2022**
GABRR3	**Canis lupus familiaris**	**-**	**-**	**-**
GABRR3	**Pan troglodytes**	**A0A2J8MY57**	**UniProt**	**11.04.2022**
GABRR3a	**Danio rerio**	**B3DIE5**	**UniProt**	**11.04.2022**
GABRR3	**Macaca mulatta**	**F6YLK3**	**UniProt**	**11.04.2022**
GABRR3	**Bos taurus**	**E1B988**	**UniProt**	**11.04.2022**

Supplementary Table S7: All sources (subunit, sequence ID, database) of protein sequences used for the alignments of all species in different coloured groups (all alpha subunits, all beta subunits, etc).

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